

UNIVERSITÉ DU QUÉBEC EN ABITIBI-TÉMISCAMINGUE

DÉTERMINER LES EFFETS HORS SITE DES MINES ET LEUR EMPREINTE
EN UTILISANT LA VÉGÉTATION ET LA PHYLLOSPHÈRE : INTÉGRATION
DU TYPE D'ÉCOSYSTÈME ET DU CYCLE DE VIE DES MINES

THÈSE

PRÉSENTÉE

COMME EXIGENCE PARTIELLE

DU DOCTORAT EN SCIENCES DE L'ENVIRONNEMENT

PAR

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DÉCEMBRE 2022

REMERCIEMENTS

Je tiens à exprimer ma gratitude à tous ceux qui m'ont aidé durant la rédaction de cette thèse. Ma plus profonde gratitude va d'abord et avant tout à ma superviseure, Dre Nicole J. Fenton (Université du Québec en Abitibi-Témiscamingue) et à ma co-superviseure, Dre Christine Martineau (Ressources Naturelles Canada) pour leurs encouragements et leurs conseils constants. Ils m'ont accompagné à travers toutes les étapes de cette thèse, y compris la proposition, le travail sur le terrain, le travail en laboratoire, l'analyse statistique et la rédaction. Sans leurs instructions constantes et éclairantes, cette thèse n'aurait pu atteindre sa forme actuelle. Merci, Nicole, de m'avoir offert un environnement libre et une plateforme professionnelle pour me permettre de profiter de cette thèse et de la mener à bien. Je veux particulièrement souligner ma gratitude envers elle pour m'avoir soutenu dans la poursuite de mes idées, en ajoutant les micro-organismes à mon projet et en me mettant en contact avec la Dre Christine Martineau pour me superviser dans cette section. Vous n'êtes pas seulement une superviseure qui m'a fait découvrir le domaine de l'écologie forestière, mais aussi une guide qui m'a montré comment devenir un bon conseiller dans ma future carrière. Je tiens également à remercier ma co-superviseure, la Dre Christine Martineau, qui a été d'un grand soutien dans ce travail, en particulier dans la conception, le travail de terrain, les expériences et l'analyse bioinformatique pour l'étude du microbiome de la phyllosphère. Enfin, merci à mes superviseures, leurs conseils m'ont aidé tout au long de la recherche et de la rédaction de cette thèse.

En plus de mes superviseures, je voudrais remercier les autres membres de mon comité de thèse, Dre Isabelle Demers et Dr Nathan Basiliko, pour leurs commentaires perspicaces et leurs suggestions pour mon proposé de recherche de doctorat et la rédaction de l'article de synthèse, qui m'ont aidé à élargir ma recherche à partir de

diverses perspectives, en particulier dans les questions environnementales liées à l'exploitation minière. Merci au Dr Mebarek Lamara pour son aide dans l'extraction d'ADN, l'analyse bioinformatique et la rédaction de ma publication sur les nouvelles espèces. Je tiens également à remercier le Dr Abdul Samad pour son aide précieuse dans l'analyse bioinformatique et la révision pour ma rédaction du Chapitre III.

Je tiens à remercier profondément Julie Arseneault et Dre Mélanie Jean pour leur soutien technique dans la conception du travail de terrain, l'établissement des parcelles, l'échantillonnage et l'identification des espèces sur le terrain. Je tiens à remercier Elizabeth Riendeau, Andréane Garant, Cyrielle Bigonneau, Simon Bégin, Matilde Offroy, Pamela Parisien, Jeffrey Raymond, Juliette Larrivée, Marie-Anyse Dubuc, Emmanuelle Richard, Maxime Thomas, Terresa Kim, Xavier Saint-Amant, Jolianne Garand et Natalie Vuong pour leur aide précieuse dans la collecte des données et le travail de terrain. Je remercie également Josée Noël, Lucienne Anctil, Stéphanie Lafrenière et les autres membres du personnel des mines de Canadian Malartic, Casa Berardi, LaRonde, Lapa, Akasaba et Joutel pour leur aide dans l'accès aux sites miniers.

Je remercie tous les membres actuels et passés du laboratoire de bryophytes pour les discussions stimulantes, pour tout le plaisir que nous avons eu au cours des quatre dernières années. J'ai eu la chance de faire partie d'un groupe extraordinaire. Je suis entouré de collègues de laboratoire fantastiques et sympathiques. Bien que nous venions de pays et de milieux différents, nous nous comprenons et nous entraïdons au sein de la grande famille. Je remercie tout particulièrement mes collègues et amis les plus proches, Enrique Hernández-Rodríguez, Amira Fetouab, Andréane Garant et Marc-Frédéric Indorf, qui m'ont permis d'avoir une vie merveilleuse en dehors de mon projet. Merci pour votre amitié et votre soutien dans mon parcours de doctorat.

Enfin, un merci particulier à ma famille pour son soutien. Merci à ma mère, Fengjie, pour son soutien indéfectible et ses encouragements continus tout au long de mon

parcours de doctorat. Merci au partenaire de ma mère, Jinlong, qui a accompagné ma mère et s'est occupé de la famille afin que je puisse me concentrer sur le projet dans un pays étranger sans aucun souci. Merci à ma conjointe, Leeping, qui comprend mon emploi du temps chargé et les pressions que je subis. Cet accomplissement n'aurait pas été possible sans eux. Merci.

Cette recherche a été financée par la Chaire industrielle CRSNG-UQAT sur la biodiversité nordique dans un contexte minier.

J'apprécie vraiment chaque personne dans ma vie. Vous avez fait de moi ce que je suis aujourd'hui.

DÉDICACE

À la mémoire de mon père, Deguo Yin,

*Bien qu'il n'ait pas pu assister à la remise de mon diplôme, il a été ma source d'inspiration
pour poursuivre mon doctorat.*

À ma mère, Fengjie Shi,

Son soutien, ses encouragements et son amour infini m'ont soutenu tout au long de ma vie.

À mon mentor, la Dre Nicole J. Fenton.

Sa sagesse et ses encouragements m'ont guidé tout au long de mon parcours.

AVANT-PROPOS

Cette thèse de doctorat a été réalisée en co-tutelle entre l'Université du Québec en Abitibi-Témiscamingue (UQAT) et Ressources Naturelles Canada (RNCan) sous la direction de Nicole J. Fenton (UQAT) et Christine Martineau (RNCan). Cette recherche a été soutenue financièrement par la Chaire industrielle CRSNG-UQAT sur la biodiversité nordique en contexte minier. Cette thèse est composée de cinq chapitres. Le Chapitre I et le Chapitre V constituent respectivement l'introduction générale et la conclusion. Les trois autres chapitres constituent le contenu essentiel de cette thèse, ils ont été rédigés sous forme de manuscrits préparés pour être publiés comme articles dans des revues à comité de lecture. La thèse est ma propre composition et a nécessité des collaborations avec mes superviseuses et d'autres collaborateurs. Mes superviseuses m'ont aidé à établir l'orientation de la recherche, la conception et l'échantillonnage du travail de terrain, l'analyse des données et la rédaction académique. Dans le chapitre III, le Dr Abdul Samad de RNCan a contribué à l'analyse microbiologique et bioinformatique. J'ai la permission de tous les co-auteurs d'inclure les publications ou les manuscrits dans ma thèse. Les publications et les manuscrits préparés pour cette thèse ainsi que les présentations de conférences sont listés comme suit :

Articles de Revues avec comité de lecture :

Yin, X., Martineau, C. and Fenton, N.J., 2022. Synergistic effects in mine offsite landscapes: Predicted ecosystem shifts could exacerbate mining effects on bryophyte community structure. *Ecological Indicators*, 144: 109555. doi: 10.1016/j.ecolind.2022.109555 (CHAPTRE IV)

Yin, X. B., Martineau, C., Demers, I., Basiliko, Nathan & Fenton, N. 2021. The potential environmental risks associated with the development of rare earth element production in Canada. *Environmental Reviews*, 29(3): 354-377. doi: [10.1139/er-2020-0115](https://doi.org/10.1139/er-2020-0115). (ANNEXE A)

Boisvert, R., Yin, X., & Fenton, J. N. 2021. Offsite effects of mining on the frequency and abundance of five herbaceous species in western Québec (Canada). *Botany*, 99(7): 449-455. doi: [10.1139/cjb-2020-0158](https://doi.org/10.1139/cjb-2020-0158). (ANNEXE C)

Yin, X., Lamara, M., Liyanage, N.S., Zhu, R.L. & Fenton, J.N., 2021. *Riccardia vitrea* (Aneuraceae), a liverwort species new to China and North America. *Herzogia*, 34(2): 279-285. doi: [10.13158/heia.34.2.2021.279](https://doi.org/10.13158/heia.34.2.2021.279). (ANNEXE B)

Manuscrits préparés pour la publication:

Yin, X., Martineau, C. & Fenton, J. N. How big is the footprint? Quantifying offsite effects of mines on boreal plant communities. (en cours de révision, *Global ecology and conservation*, CHAPTRE II)

Yin, X., Martineau, C., Samad, A. & Fenton, J. N. Out of sight, out of mind: changes in feather moss phyllosphere structure in mine offsite boreal landscapes. (en cours de révision, *Environmental Microbiology*, CHAPTRE IV)

Présentation de posters et de conférences :

Présentations orales

Yin, X., Martineau, C., Samad, A., Boisvert, R. & Fenton, J. N., 2022. Offsite effects of mines on understory plants and moss phyllosphere in boreal forests: integrating mining stage and ecosystem type. Le 4e Colloque annuel de la Chaire industrielle

CRSNG-UQAT sur la biodiversité en contexte minier. April 2022, Rouyn-Noranda, Canada.

Yin, X., Martineau, C., Samad, A. & Fenton, J. N., 2021. Offsite impacts of mines on understory plants and moss phyllosphere bacteria in boreal ecosystems: integrating mine stages and habitats. Bryophytes, lichens, and northern ecosystems in a changing world (BL2021) Conference. July 2021, Quebec, Canada.

Yin, X., Martineau, C., Samad, A., Boisvert, R. & Fenton, J. N., 2021. Offsite impacts of mines on understory plants and moss phyllosphere in boreal ecosystems: Integrating mine stages and forest types. Le 3e Colloque annuel de la Chaire industrielle CRSNG-UQAT sur la biodiversité en contexte minier. September 2021, Rouyn-Noranda, Canada.

Yin, X., Martineau, C., Boisvert, R. & Fenton, J. N., 2020. The offsite impacts of mines on the diversity of vegetation and associated phyllosphere microbiome in boreal areas. The 2020 Canadian Botanical Association Virtual Meeting. July 2020, Quebec, Canada.

Yin, X., Martineau, C., Boisvert, R. & Fenton, J. N., 2020. Determining the footprint of mines for plant diversity in different ecosystem types: integrating offsite impacts and the mine life cycle. Le 2e Colloque annuel de la Chaire industrielle CRSNG-UQAT sur la biodiversité en contexte minier. April 2020, Rouyn-Noranda, Canada.

Yin, X., Martineau, C., Boisvert, R. & Fenton, J. N., 2019. Determining the footprint of mines on plant diversity: integrating enigmatic impacts and the mine life cycle. Le 1e Colloque annuel de la Chaire industrielle CRSNG-UQAT sur la biodiversité en contexte minier. April 2019, Rouyn-Noranda, Canada.

Présentation de posters

Yin, X., Martineau, C., Boisvert, R. & Fenton, J. N., 2021. How big is the footprint? Quantifying offsite effects of mines on boreal plant communities.” 23e colloque annuel de la Chaire AFD. December 2021, Amos, Canada.

Yin, X., Lamara, M., Liyanage, N.S., Zhu, R.L. & Fenton, J.N., 2021. *Riccardia virea* (Aneuraceae, Marchantiophyta), a liverworts species new to Canada and North America. Bryophytes, lichens, and northern ecosystems in a changing world (BL2021) Conference. July 2021, Quebec, Canada.

Yin, X., Martineau, C., Boisvert, R. & Fenton, J. N., 2021. The offsite impacts of mines on plant diversity in boreal areas.” 22e colloque annuel de la Chaire AFD. December 2020, Rouyn-Noranda, Canada.

Effets hors site des mines

comme la propagation du COVID-19 par les éternuements et la toux.

à l'abri des regards

mais le virus dans l'air ambiant

comme les poussières qui recouvrent les lunettes

petites, voire minuscules

mais qui deviennent grandes

quand on les néglige ou qu'on les laisse reposer

comme une pomme pourrie

intacte et belle à l'extérieur

mais qui commence à pourrir de l'intérieur

comme un diamant qui sort de terre

en enlevant seulement une particule

mais qui pourrait avoir plus de valeur qu'une tonne de roche.

"It's the little details that are vital. Little things make big things happen." — John Wooden



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RÉSUMÉ

Les plantes et les microorganismes de leur phyllosphère contribuent à la biodiversité et aux services écologiques dans différents biomes forestiers, dont la forêt boréale. Ils jouent un rôle important dans les réservoirs de carbone mondiaux et la fixation de l'azote. L'augmentation de l'activité minière dans le but de satisfaire les besoins en minéraux constitue une menace potentielle pour les services écosystémiques fournis par les plantes de sous-bois et le microbiote de la phyllosphère. Ces menaces ne concernent pas seulement les zones directement perturbées, mais pourraient également s'étendre à des paysages hors site relativement intacts. Les changements prévus dans le type d'écosystème dans le cadre des scénarios de changements globaux constituent une autre perturbation potentielle pour les communautés du sous-bois boréal. Les effets qui se produisent dans les paysages hors site (effets hors site) sont facilement et souvent négligés dans les évaluations d'impact car ils se situent en dehors de l'emplacement immédiat des activités minières. Par conséquent, on en sait peu sur la façon dont les effets hors site des mines sur les communautés du sous-étage et sur le rôle du type d'écosystème sur ces effets.

Afin d'améliorer notre compréhension des effets hors site des mines sur les plantes boréales et le microbiote de la phyllosphère, nous avons étudié la diversité des espèces du sous-étage et l'ADN microbien de la phyllosphère d'une espèce de mousse à plumes (*Pleurozium schreberi* (Brid.) Mitt. dans des paysages en mosaïque de 1 km contenant quatre types d'écosystèmes (conifères, feuillus, forêts mixtes et milieux ouverts) autour de six mines à différents stades d'exploitation (trois mines en exploitation et trois mines non exploitées) dans la zone boréale du Canada. Le métabarcodage du gène de l'ARNr 16S bactérien et la région ITS2 fongique ont été utilisés pour la caractérisation taxonomique du microbiome. Nos résultats ont confirmé la présence d'effets hors site des mines sur la diversité et la composition de la phyllosphère des plantes vasculaires de sous-bois, des bryophytes et des mousses à plumes. L'ampleur des effets hors site était associée au stade d'exploitation et au type d'écosystème. Les effets combinés du type d'écosystème et de l'exploitation minière dans les paysages hors site étaient généralement synergiques pour les plantes vasculaires de sous-bois et les bryophytes, mais pas pour la phyllosphère de la mousse à plumes. Nous avons également constaté que les microhabitats pouvaient atténuer les effets hors site des mines sur la diversité du sous-étage (c'est-à-dire la richesse en bryophytes). Les effets hors site des mines les plus forts se sont produits à une distance de 0,2 km des mines, en particulier près des mines en exploitation.

Compte tenu des changements prévus dans les écosystèmes de la forêt boréale, avec l'empiètement des espèces à feuilles caduques sur les forêts de conifères, et la sensibilité accrue des forêts mixtes et à feuilles caduques, la zone touchée par les effets hors site des mines pourrait augmenter dans le futur. La résistance des forêts pourrait être réduite dans les paysages hors site par la réduction de la diversité des plantes vasculaires et des bryophytes, l'augmentation des différences de composition et la diminution de l'abondance de groupes fonctionnels importants (c'est-à-dire les mousses à plumes et les sphaignes).

Les effets hors site sur la diversité et la composition du sous-étage observés étaient généralement faibles et semblaient négligeables dans la biodiversité globale de la forêt boréale. Cependant, la perte d'espèces clés, l'invasion d'espèces introduites et les impacts cumulatifs à grande échelle peuvent augmenter l'impact des effets hors site des mines sur les services écosystémiques. Par conséquent, les effets hors site des mines devraient être inclus dans les évaluations écologiques futures afin de maintenir la biodiversité et les services écologiques dans la forêt boréale. La création d'une zone tampon autour des mines pourrait permettre de compenser les effets hors site potentiels. D'autres études seront nécessaires pour évaluer comment ces effets hors site des mines sur la structure du sous-étage et de la phyllosphère affectent les services des écosystèmes boréaux et pour développer des stratégies d'atténuation appropriées.

Mots-clés : impacts indirects, végétation de sous-bois, surface foliaire, bactéries, champignons, écologie du paysage, indicateurs, taïga forestière.

CHAPITRE I

INTRODUCTION GÉNÉRALE

1.1 Mise en contexte

Les activités humaines sont devenues un moteur essentiel de la dynamique de la biodiversité mondiale dans l'Anthropocène (Ellis et al., 2012 ; Seddon et al., 2016 ; Storch et al., 2021). Les activités minières constituent un héritage à long terme de l'industrialisation propre à l'époque (McMahon et Moreira, 2014 ; Zhang et al., 2019) et ont un impact considérable sur la biodiversité par le biais de divers mécanismes (par exemple, la dégradation de l'habitat, les espèces envahissantes, le rejet de déchets et la pollution) et à travers les échelles spatiales (Sonter et al., 2018). En général, les mines et les gisements ne sont pas répartis de manière aléatoire, mais concentrés dans des zones présentant une biodiversité intermédiaire à élevée (Murguia et al., 2016). Les besoins croissants en matière de production d'énergie renouvelable vont également exacerber les menaces minières sur la biodiversité (Sonter et al., 2020). Par exemple, les éléments de terres rares sont de plus en plus nécessaires pour les technologies propres telles que les véhicules électriques et les éoliennes (Zhou et al., 2017), ce qui entraînera une augmentation des zones influencées directement et indirectement par l'exploitation minière (Goodenough et al., 2018).

Les impacts de l'exploitation minière sur la biodiversité sont susceptibles d'être modulés par des changements globaux tels que les changements prévus dans la composition des forêts dans les zones boréales. La prédominance relative des arbres à feuilles caduques devrait augmenter au cours des prochaines décennies, la composition des peuplements passant des conifères aux feuillus sous l'effet des changements climatiques et des incendies de forêt (Gray et He, 2009 ; Mack et al., 2021 ; Mekonnen et al., 2019). En outre, le passage des tourbières aux peuplements forestiers est également un scénario pour les changements de végétation prévus dans les forêts boréales (Ratcliffe et al., 2017 ; van der Velde et al., 2021), en particulier dans le sud des forêts boréales où le boisement des tourbières est causé par des perturbations naturelles et anthropiques, telles que les changements climatiques,

l'agriculture et l'exploitation minière (Howson et al., 2021 ; Lachance et al., 2005 ; Lavoie et al., 2005). Les effets de l'exploitation minière et des changements prévus dans le type d'écosystème sur la biodiversité ont été largement étudiés séparément, mais peu d'études sont disponibles sur les effets de leur interaction sur la biodiversité, en particulier pour les impacts miniers hors site potentiels. Par conséquent, on ne sait toujours pas si les changements prévus dans la composition des forêts exacerberont les effets de l'exploitation minière sur la biodiversité des paysages environnants. Les effets hors site des mines sur la biodiversité appartiennent à la catégorie des impacts énigmatiques (figure 1.1) car ils sont souvent négligés dans les évaluations d'impact (Raiter et al., 2014). Les effets hors site sont les impacts qui se produisent loin de l'emplacement immédiat de la perturbation ou de la zone d'étude et qui peuvent s'infiltrer loin dans des paysages apparemment intacts (Raiter et al., 2014). Des études sur les effets hors site ont été réalisées à proximité des sites de diverses activités humaines telles que les routes (Jakobsson et al., 2018), l'exploitation forestière (Brudvig et al., 2009) et l'exploitation minière (Weir et al., 2007), cependant, nos connaissances sur les effets hors site des mines sur la diversité végétale et microbienne sont encore limitées, en particulier dans les paysages boréaux. La recherche d'indicateurs appropriés pour surveiller l'étendue de la zone affectée par les mines (empreinte hors site) constitue également un défi dans l'évaluation des effets hors site. Nous prévoyons que les types d'écosystèmes et le cycle de vie des mines sont des facteurs déterminants pour la présence et l'ampleur des effets hors site des mines sur la biodiversité. Trois groupes de communautés d'organismes de sous-bois, les plantes vasculaires, les bryophytes et le microbiote de la phyllosphère (micro-organismes vivant à la surface des feuilles), ont été utilisés dans le projet pour confirmer cette prédiction et pour sélectionner des groupes d'indicateurs appropriés.

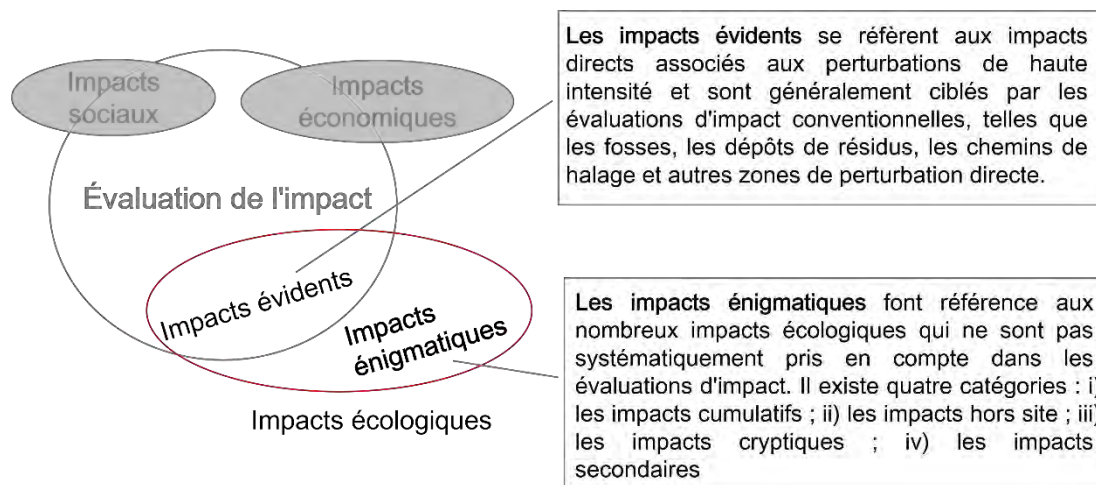


Figure 1.1 Le concept d'impact énigmatique

1.1.1 Effets des types d'écosystèmes sur la biodiversité du sous-bois

Les forêts de conifères, les forêts de feuillus, les forêts mixtes et les milieux ouverts (sans arbres de canopée, constitués principalement de tourbières) sont les quatre principaux types d'écosystèmes de la forêt boréale en fonction de la composition de leur canopée (Figures 1.2). Les types d'écosystèmes peuvent déterminer la structure du sous-étage en créant des conditions variées pour la croissance des espèces qui y vivent. En général, les arbres à feuilles caduques permettent une transmission plus élevée des nutriments foliaires et de la lumière que les conifères (Kembel et Dale, 2006 ; Qian et al., 2003), tandis que les peuplements de conifères créent un pH plus faible, des cations basiques plus bas et une humidité de l'air plus élevée (Augusto et al., 2002). Les forêts mixtes créent généralement des environnements plus hétérogènes que les forêts monospécifiques (Pollastrini et al., 2017). Les milieux ouverts, sans arbres de la canopée, reçoivent plus de lumière, de précipitations, de rayonnement solaire et de dépôts atmosphériques que les forêts à canopée fermée. Ces différences abiotiques influencent la composition du sous-étage dans chaque type d'écosystème. Les peuplements de conifères sont généralement considérés comme

moins favorables à la diversité et à la couverture des plantes vasculaires que les peuplements de feuillus (Kembel et Dale, 2006), et tendent à supporter une couverture plus élevée de bryophytes (Bartels et al., 2018b ; Légraré et al., 2005). Les forêts mixtes ne favorisent ni les strates vasculaires ni les bryophytes (Cavard et al., 2011) et présentent une diversité de sous-étage similaire à celle des peuplements de conifères et de feuillus (Brassard et al., 2008). Les zones ouvertes ont tendance à abriter des communautés de lichens ou de bryophytes en fonction du taux d'humidité (Horstkotte et Moen, 2019). En outre, certaines études ont révélé que les forêts de conifères et de feuillus soutiennent différents groupes microbiens spécifiques dans leur couche de sous-bois. Dans la phyllospère de la mousse à plumes, les cyanobactéries étaient davantage associées aux forêts de feuillus (*Populus tremuloides* Michx.) qu'aux forêts de conifères (*Picea mariana* (Mill.) Britton, Sterns & Poggenb., Rodríguez-Rodríguez et al., 2022), tandis que la forêt de conifères supportait une plus grande abondance relative du phylum bactérien WPS-2 (Jean et al., 2020b).

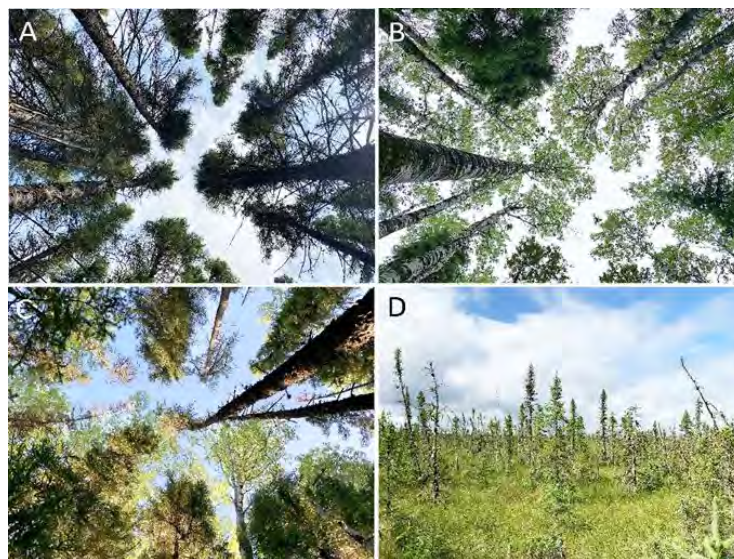


Figure 1.2 Principaux types d'écosystèmes dans les paysages boréaux. A: forêt de conifères. B: forêt de feuillus. C: forêt mixte. D: milieux ouverts (tourbières).

Les différences dans la structure de la canopée et la composition des espèces du sous-étage contribuent à la résistance variable de chaque type d'écosystème aux perturbations humaines (par exemple, le dépôt de la pollution atmosphérique provenant de l'exploitation minière) et aux changements climatiques. Les espèces végétales soutenues par les canopées de feuillus sont généralement intolérantes à l'ombre et exigeantes en nutriments, tandis que les conifères favorisent l'établissement d'espèces végétales tolérantes à l'ombre et à faible teneur en nutriments du sol, comme les bryophytes (Su et al., 2019), ce qui indique que l'écosystème de conifères pourrait être plus résistant aux changements climatiques naturels et aux perturbations anthropiques que l'écosystème de feuillus. En outre, les canopées de conifères peuvent tamponner les variations environnementales et éliminer la pollution atmosphérique grâce aux caractéristiques de la longueur de la période de défoliation et de la forte densité du couvert arboré (Barbé et al., 2020 ; De Jalón et al., 2019). Dans les milieux ouverts, malgré une exposition directe aux changements environnementaux avec l'absence d'arbres de la canopée, les données paléoécologiques montrent une résistance relativement élevée des tourbières aux changements climatiques du passé (Łuców et al., 2020) car elles sont protégées par un microclimat spécifique avec des conditions humides générées par la tourbe accumulée (Augusto et al., 2002). Cependant, la résistance des tourbières aux activités humaines pourrait être faible, et la dégradation des tourbières par l'exploitation minière, les routes, la foresterie et l'agriculture par le drainage et l'extraction de la tourbe (Glina et al., 2019 ; Page et Baird, 2016 ; Williams-Mounsey et al., 2021) a été signalée. Par conséquent, nous pensons que la structure du sous-étage dans chaque type d'écosystème pourrait présenter une résistance différente aux effets hors site des mines.

1.1.2 Dynamique des perturbations au cours du cycle de vie de l'exploitation minière

L'intensité et l'échelle des perturbations minières évoluent de manière dynamique avec le cycle de vie de l'exploitation. Le processus d'exploitation minière se compose de quatre grandes étapes de développement: 1) l'exploration; 2) la construction; 3) l'exploitation; 4) la fermeture et la réhabilitation (Gorman et Dzombak, 2018). La force et les échelles de perturbations ne sont pas les mêmes à chaque étape. La tâche principale de l'étape d'exploration est de sonder pour trouver des minéraux, y compris des relevés sur le terrain et par satellite, ainsi que la collecte d'échantillons de minéraux pour une évaluation économique. Les perturbations (par exemple, les camps et l'équipement sur le terrain) sur les paysages sont limitées à ce stade. L'intensité des perturbations augmente au cours de la phase de construction, lorsque les bâtiments et les infrastructures sont développés et que la végétation et le sol de surface sont défrichés, ce qui détruit directement la structure et la fonction des écosystèmes de surface. L'intensité des perturbations (c'est-à-dire les effets sur les milieux environnants) et les échelles (c'est-à-dire les zones influencées) continuent de s'étendre au stade de l'exploitation, car les opérations d'extraction et de traitement des minéraux sont les principales activités sur les sites miniers en exploitation, comme le forage, le dynamitage, la ventilation, l'assèchement, le chargement et le transport, le concassage et le broyage, et les séparations (Norgate et Haque, 2010). De nombreux polluants (par exemple, les métaux lourds, la poussière et le drainage minier acide) sont créés et libérés dans le milieu environnant au cours de ces activités (Punia, 2021), ce qui entraîne une pollution atmosphérique, une pollution de l'eau et une dégradation des sols dans les paysages environnants (Li et al., 2014 ; Tolvanen et al., 2019). Au stade de la fermeture et de la réhabilitation, les perturbations commencent à être réduites en raison de la fermeture de toutes les opérations minières, du retrait des équipements et de la revégétalisation (Limpitlaw, 2004).

En général, étant donné que l'étendue et l'intensité des perturbations minières changent avec le temps, le cycle de vie minier peut être classé en deux groupes principaux en fonction de l'intensité des perturbations : exploitation et non-exploitation (y compris l'exploration, la construction, ainsi que la fermeture et la réhabilitation). Les différences de perturbations minières à différentes étapes peuvent conduire à des effets distincts des mines sur la biodiversité environnante entre les étapes.

1.1.3 Rôles écologiques des communautés de sous-bois dans l'écosystème boréal

La forêt boréale est l'un des plus grands biomes sur Terre, représentant 11% de la surface terrestre globale, qui assure de multiples fonctions écologiques à l'échelle locale et globale (Gauthier et al., 2015). Les forêts boréales canadiennes, qui occupent environ 30 % des forêts boréales du monde entier, jouent des rôles essentiels dans le cycle global du carbone, la régulation du climat et la conservation de la biodiversité (Peng et al., 2011 ; Venier et al., 2014). Les communautés végétales du sous-bois (y compris les espèces vasculaires et bryophytes, Figure 1.3 présentant la biodiversité du sous-bois dans la forêt boréale) apportent une contribution indispensable à ces services écologiques, notamment la succession forestière (Royo et Carson, 2006) et le cycle des nutriments (Nilsson et Wardle, 2005), la nourriture et les habitats pour les espèces sauvages et même les médecines traditionnelles pour les communautés autochtones (Upreti et al., 2012). En outre, le microbiote résidant dans ou sur les tissus végétaux d'espèces de sous-bois (comme les bryophytes) soutient également la biodiversité et les services écologiques dans les paysages boréaux (Compant et al., 2019 ; Trivedi et al., 2020).



Figure 1.3 Biodiversité du sous-étage dans les paysages boréaux. A-D: la couche de sous-bois des forêts de conifères, de feuillus, mixtes et de milieux ouverts (de A à D); E: *Chamerion angustifolium* (L.) Holub; F: *Gaultheria hispidula* (L.) Muhl. ex Bigelow; G: *Pteridium aquilinum* (L.) Kuhn; H: *Vaccinium myrtilloides* Michx.; I: *Rubus idaeus* L.; J: *Ribes glandulosum* Grauer; K: *Sarracenia purpurea* L.; L: *Rhododendron groenlandicum* (Oeder) Kron & Judd; M: *Drosera rotundifolia* L.; N: *Cladonia cristatella* Tuck.; O: *Sphagnum fuscum* (Schimp.) Klinggr.; P: *Sphagnum capillifolium* (Ehrh.) Hedw.; Q: *Riccardia chamedryfolia* (With.) Grolle.; R: *Pleurozium schreberi* (Willd. ex Brid.) Mitt.; S: *Marchantia polymorpha* L. subsp. *ruderalis* Bischl. & Boissel.-Dub.; T: *Aneura pinguis* (L.) Dumort.; U-V: champignons.

La végétation de sous-bois est la composante la plus diversifiée des communautés végétales boréales (Hart et Chen, 2006). Les plantes de sous-bois représentent une faible proportion de la biomasse totale par rapport aux arbres, mais elles contribuent de manière significative aux cycles du carbone et des nutriments et à la succession des écosystèmes boréaux, avec des taux de renouvellement plus élevés que les arbres de la canopée (Kumar et al., 2018 ; Nilsson et Wardle, 2005). Par exemple, les mousses à plumes peuvent représenter jusqu'à 30 % de la productivité primaire nette

des forêts boréales (Wardle et al., 2012) et leur phyllosphère peut contribuer jusqu'à > 50 % des sources totales d'azote dans les forêts boréales de conifères (Jean et al., 2021). Les espèces du sous-étage sont aussi généralement utilisées comme indicateurs biologiques pour évaluer les perturbations humaines, car leur diversité et la composition des communautés sont sensibles aux activités anthropiques, notamment l'exploitation forestière, l'exploitation minière et le pâturage (Ekka et Behera, 2011; Kariuki et al., 2006 ; Metzger et al., 2005). Les espèces végétales envahissantes sont des indicateurs typiques qui sont généralement utilisés pour évaluer la gravité et la fréquence des perturbations (Dyderski et Jagodziński, 2018; Lavoie, 2010). En outre, les bryophytes ont également été largement utilisés comme indicateurs fiables de la pollution atmosphérique, des changements climatiques (température et humidité), des dépôts de métaux lourds et d'autres perturbations (Balabanova et al., 2017; Ceschin et al., 2012; Gignac, 2001; Mahapatra et al., 2019; Szczepaniak et Biziuk, 2003). En effet, elles sont dépourvues de véritables racines, de systèmes vasculaires et de cuticules épaisses, et sont donc très sensibles aux changements du milieu environnant, notamment aux microhabitats, aux types d'écosystèmes et aux propriétés physicochimiques.

La microflore (ou microbiote) de la phyllosphère (la surface des feuilles) est une autre composante importante de la couche de sous-bois qui maintient les services écologiques des écosystèmes boréaux. Des études ont révélé que la microflore de la phyllosphère joue un rôle critique dans la résistance aux agents pathogènes et aux herbivores (Ab Rahman et al., 2018 ; Saleem et al., 2017), la tolérance au stress (Arun et al., 2020), la promotion de la croissance et de la reproduction (Canto et Herrera, 2012 ; Stone et al., 2018), la dégradation du méthanol (Galbally et Kirstine, 2002 ; Iguchi et al., 2012) et la fixation de l'azote (Abril et al., 2005 ; Fürnkranz et al., 2008). Dans les écosystèmes boréaux, les micro-organismes de la phyllosphère des mousses jouent un rôle important dans les services écologiques. En particulier, les associations mousse à plumes-cyanobactéries (Rousk et al., 2013) et *Sphagnum*-

méthanotrophes (Putkinen et al., 2012) sont des moteurs importants du cycle du carbone et de l'azote dans les forêts de conifères et les tourbières. Une étude a montré que *Pleurozium schreberi* (une espèce commune de mousse à plumes) pouvait à elle seule, grâce à l'activité des bactéries fixatrices d'azote de la phyllosphère (par exemple, les genres *Nostoc* et *Scytonema*), fixer 1,5 à 2,0 kg d'azote ha⁻¹ an⁻¹ dans les forêts boréales (DeLuca et al., 2002). En outre, les communautés microbiennes (la plupart des études portent sur le microbiome du sol) sont sensibles aux changements environnementaux causés par les activités humaines, notamment la teneur en métaux lourds et les dépôts atmosphériques (Karimi et al., 2017 ; Li et al., 2017 ; Sharma et al., 2010). Les micro-organismes sensibles aux perturbations (par exemple, la toxicité des métaux lourds) vont diminuer en abondance, tandis que les espèces tolérantes deviennent prédominantes (Guo et al., 2017 ; Zhang et al., 2022). Les changements dans la structure de la communauté microbienne peuvent sérieusement affecter les services écologiques fournis par les micro-organismes (Giller et al., 1998). Les changements dans le microbiome de la phyllosphère pourraient également être un indicateur sensible du stress de l'hôte par les perturbations humaines, y compris l'exploitation minière (Xu et al., 2020). Cependant, peu d'études ont jusqu'à présent permis de comprendre comment l'exploitation minière affecte les communautés de la phyllosphère.

1.2 Objectifs et hypothèses de recherche

L'objectif général de cette thèse de doctorat est de comprendre et de caractériser les effets hors site des mines sur les communautés du sous-étage et de la phyllosphère dans les paysages boréaux environnants et d'identifier si la composition de la forêt et le cycle de vie de l'exploitation minière influencent ces effets. Pour cet objectif global de recherche, nous avons sélectionné trois groupes : les communautés végétales du sous-étage au sol, les bryophytes dans les microhabitats et les micro-organismes de la phyllosphère des mousses à plumes. Les objectifs spécifiques et les hypothèses pour chaque groupe de recherche sont les suivants :

1.2.1 Communautés générales du sous-bois

Objectifs :

- 1) Évaluer comment le type d'écosystème et le cycle de vie des mines affectent les effets hors site des mines ;
- 2) Evaluer la taille de l'empreinte hors site des mines sur le sous-étage.

Hypothèses:

- 1) Étant donné que plus d'effets environnementaux (par exemple, les émissions) se produisent pendant la phase d'exploitation, on s'attend à ce que la diversité globale du sous-étage soit plus faible à proximité des sites d'exploitation que des sites non exploités ;
- 2) On s'attend à ce que les communautés du sous-étage diffèrent dans leur sensibilité aux effets hors site de l'exploitation minière en fonction du type d'écosystème parce que les différents écosystèmes favorisent des groupes distincts qui pourraient avoir des niveaux variables de tolérance aux perturbations ; en outre, les différences dans la structure de la canopée entre les types d'écosystèmes pourraient fournir différents

niveaux de protection aux communautés du sous-étage contre les perturbations des mines;

3) Si l'on considère que les effets hors site se produisent dans des paysages relativement intacts qui ne sont pas visuellement distincts, les changements et la distance influencés par les mines devraient être relativement faibles, mais détectables si l'on considère la richesse et la composition du sous-étage au niveau de la communauté, et leur empreinte cumulative pourrait être significative.

1.2.2 Bryophytes sur les microhabitats

Objectifs:

- 1) Évaluer le rôle des microhabitats dans la médiation des effets d'interaction des types d'écosystèmes et de l'exploitation minière sur la structure des communautés de bryophytes.
- 2) Évaluer si les types d'écosystèmes et l'exploitation minière affectent les services écologiques des bryophytes en modifiant l'abondance de groupes fonctionnels importants (i.e. les mousses à plumes et les sphaignes).

Hypothèses:

- 1) On s'attend à ce que les types d'écosystèmes affectent la présence et l'ampleur des effets des mines sur la structure des bryophytes environnantes, étant donné les effets variables des différents types d'écosystèmes sur la diversité des bryophytes ;
- 2) Les interactions entre les effets hors site des mines et les types d'écosystèmes sur les bryophytes devraient également être influencées par les microhabitats, étant donné que les microhabitats présentent des capacités différentes pour amortir les changements environnementaux ;

3) Les changements de type de mines et d'écosystèmes pourraient affecter les services écologiques des paysages hors site en réduisant l'abondance de groupes fonctionnels importants (c'est-à-dire les mousses à plumes et les sphaignes), car les mousses à plumes et les sphaignes sont principalement soutenues par les forêts de conifères et les tourbières par rapport aux forêts de feuillus et mixtes.

1.2.3 Microbiote de la phyllosphère de la mousse de plumes

Objectifs :

- 1) Déterminer la présence d'effets hors site des mines sur le microbiote de la phyllosphère des mousses à plumes dans les paysages environnants des mines.
- 2) Évaluer les effets du type d'écosystème, du stade d'exploitation minière et de leur interaction sur les effets hors site de l'exploitation minière sur la structure de la communauté microbienne de la phyllosphère des mousses.
- 3) Déterminer la distance d'influence des effets hors site des mines sur le microbiote de la phyllosphère des mousses.

Hypothèses:

- 1) En se basant sur les résultats prédits selon lesquels la structure des communautés de bryophytes serait affectée par le stade de l'exploitation minière (Hypothèse en 1.2.2), on s'attend à ce que les effets sur la diversité microbienne, la structure des communautés et la composition de la phyllosphère soient plus importants près des mines en exploitation que près des sites non exploités.
- 2) Étant donné que le type d'écosystème est censé être un déterminant clé de la structure des communautés de bryophytes (hypothèse du point 1.2.2), les effets des mines sur le microbiote de la phyllosphère de la mousse à plumes devraient également être influencés par les types d'écosystèmes.

1.3. Sites étudiés

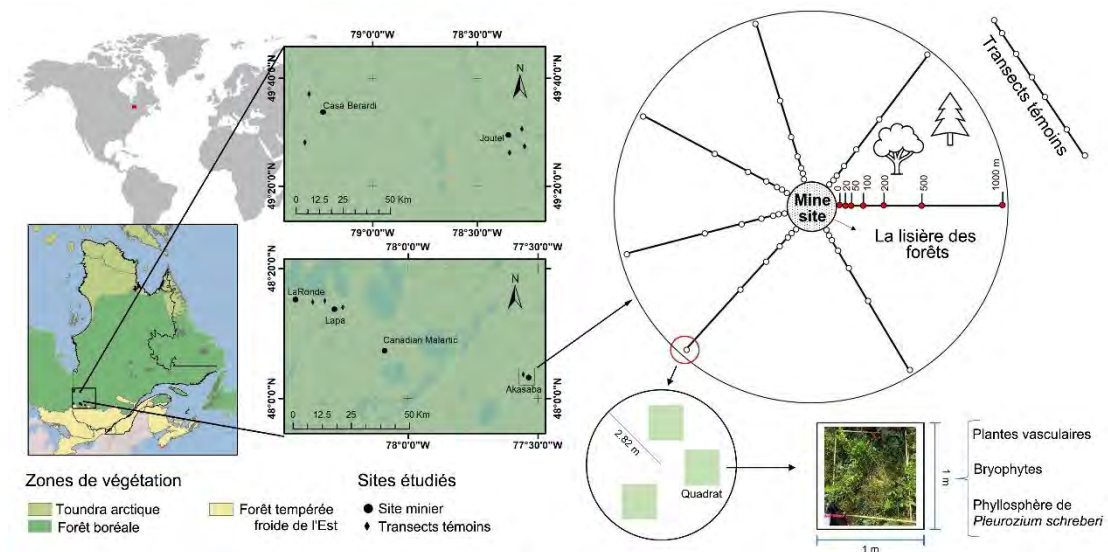


Figure 1.4 Sites étudiés. À gauche, carte du Canada et du Québec montrant les principales zones de végétation. À droite, carte de la région d'étude indiquant la position des parcelles d'échantillonnage sur six sites de mines d'or et 9 transects de contrôle

Six sites miniers ont été sélectionnés en Abitibi-Témiscamingue et dans le Nord-du-Québec, dans la province de Québec au Canada (Figure 1.4): Akasaba (48°3'12"N; 77°32'8"W, Agnico Eagle Mines Ltd), Canadian Malartic (48°7'21"N; 78°5'23"W, Canadian Malartic Corp.), Casa Berardi (49°33'43"N; 79°14'8"W, Hecla Québec Inc.), Joutel (49°29'28"N; 78°21'8"W, Agnico Eagle Mines Ltd), Lapa (48°13'45"N; 78°17'1"W, Agnico Eagle Mines Ltd), LaRonde (48°15'14"N; 78°25'59"W, Agnico Eagle Mines Ltd). Trois d'entre eux (Casa Berardi, en exploitation depuis 1988 ; LaRonde, depuis 1988 et Canadian Malartic, depuis 2005) sont en exploitation et les trois autres (Akasaba, en cours d'établissement ; Joutel, fermé depuis 1988 et Lapa, fermé depuis 2018) sont hors exploitation. Casa Berardi et Joutel sont situés dans le domaine bioclimatique *Picea mariana* - mousses, tandis que les quatre autres sites miniers sont tous situés dans le domaine bioclimatique *Abies balsamea* - *Betula*

papyrifera. Neuf transects non perturbés ont été sélectionnés comme sites témoins : cinq dans le domaine bioclimatique *Picea mariana* - mousses et quatre dans le domaine bioclimatique *Abies balsamea* - *Betula papyrifera*.

Les sites d'étude se trouvent dans une région qui est caractérisée par un climat continental froid et humide avec une température moyenne comprise entre -14 et -10°C en hiver et entre 16 et 18°C en été (Environnement Canada, 2021). Les précipitations annuelles moyennes vont de 790 à plus de 1 300 mm (Wang et al., 2013).

1.4. Plan de la thèse

Cette thèse est structurée en six chapitres, comme le montre la figure (Figure 1.5). Le contenu principal de ces six chapitres est le suivant :

Le chapitre I (ce chapitre) présente le contexte de la recherche, les objectifs, la méthodologie et le plan de la thèse.

Le chapitre II présente les effets hors site des mines sur la richesse, la couverture et la composition des espèces du sous-étage entourant les paysages boréaux (zones de 1 km près des sites miniers) au niveau des communautés (richesse, couverture et composition des espèces de plantes vasculaires herbacées et ligneuses ainsi que la couverture des bryophytes et des lichens) et évalue les rôles potentiels du cycle de vie minier (mines en phase d'exploitation et de non-exploitation) et des types d'écosystèmes comme facteur déterminant de ces paramètres. La distance influencée par les effets hors site est déterminée en utilisant différents groupes de sous-étage.

Le chapitre III présente les effets d'interaction de l'exploitation minière et de la composition de la forêt sur la structure des communautés de bryophytes dans les paysages boréaux entourant les mines en utilisant la richesse et la composition en

espèces de toutes les bryophytes, mousses et hépatiques. Ensuite, les effets tampons des microhabitats (c'est-à-dire le sol, les arbres, les billes, les souches, les chicots et les rochers) sur ces effets d'interaction ont été évalués. Enfin, l'abondance des mousses à plumes et des sphaignes a été utilisée pour estimer les impacts potentiels des effets d'interaction sur les services écologiques des forêts boréales. Dans ce chapitre, nous avons constaté que la couverture végétale de la mousse à plumes a des effets significatifs en fonction du stade d'exploitation et du type d'écosystème. Nous avons donc choisi d'étudier le microbiote de la mousse à plumes dans le chapitre IV pour évaluer comment les effets hors site des mines affectent la structure de la communauté microbienne de la phyllosphère.

Le chapitre IV présente les effets des mines sur la diversité et la composition du microbiote de la phyllosphère des mousses à plumes (indices de richesse et de diversité bactérienne et fongique ainsi que l'abondance relative des principaux phyla) dans les paysages boréaux et évalue l'influence potentielle du cycle de vie minier et des types d'écosystèmes. Les taxons indicateurs de chaque type d'écosystème et de chaque stade d'exploitation minière (exploitation, non-exploitation et contrôles) ont été évalués, respectivement. Ensuite, la distance influencée par les effets hors site des mines a été déterminée en utilisant la diversité et la structure de la communauté du microbiote de la phyllosphère ainsi que l'abondance relative des taxons individuels (principaux phyla bactériens et fongiques, genres et ASVs, amplicon sequence variant).

Le chapitre V résume brièvement les résultats obtenus dans la thèse, discute de leur signification, identifie les limites de la recherche et propose des orientations futures pour la recherche.

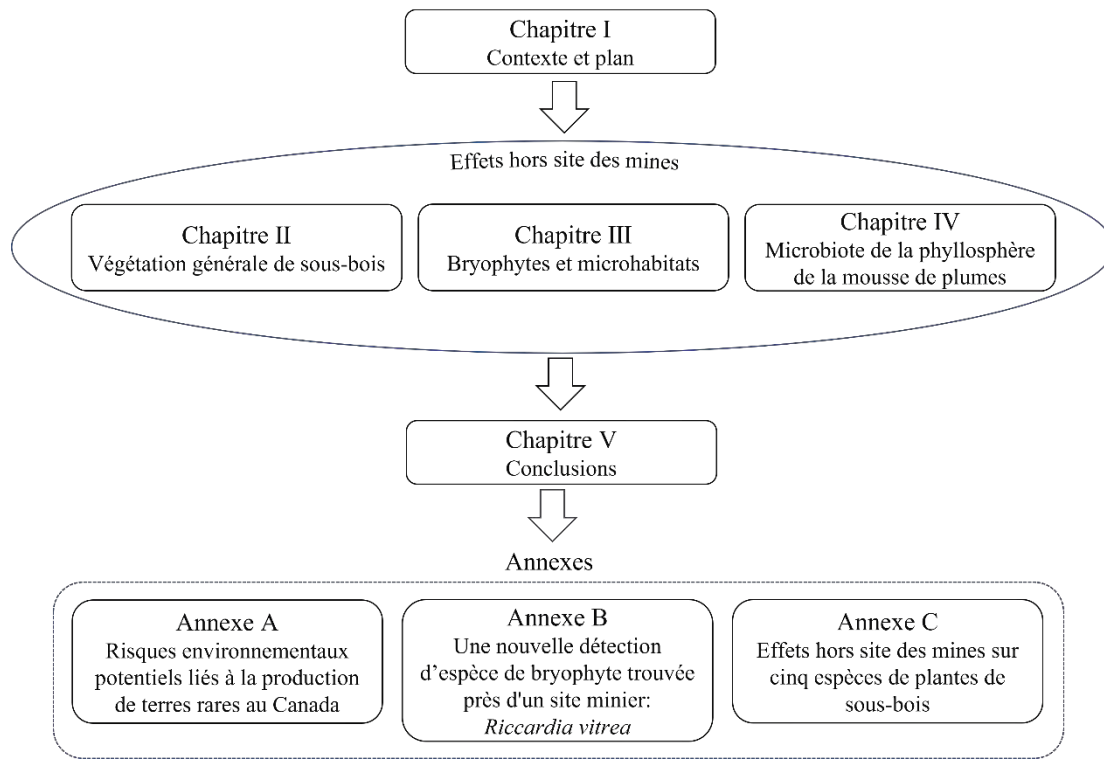


Figure 1.5 Le plan de la thèse

En outre, trois chapitres supplémentaires sont inclus dans la section ‘Annexe’ de cette thèse. Le premier chapitre annexé (annexe A) concerne les risques environnementaux potentiels associés à ‘activités minière en utilisant la production de terres rares comme cas. Il traite des impacts potentiels de l'exploitation des terres rares sur les services écologiques et la santé humaine au Canada, ce qui fournit une revue de la littérature en lien avec le contexte de la thèse. Le deuxième chapitre annexé (annexe B) décrit une nouvelle détection d'espèce (*Riccardia vitrea*) en Amérique du Nord. Cette espèce a été trouvée près d'un site minier, ce qui suggère que l'exploitation minière pourrait fournir un habitat assez spécifique pour certaines espèces. L'annexe B présente la contribution de cette thèse au nombre d'espèces de bryophytes présentes au Canada. Le troisième chapitre annexé (annexe C) porte sur les effets hors site des mines sur cinq espèces de sous-bois, ce qui constitue nos résultats préliminaires. Les

résultats de l'annexe C ont confirmé la présence d'effets hors site des mines au niveau des espèces, ce qui fournit des bases pour des études ultérieures dans les chapitres principaux.

CHAPITRE II

QUELLE EST LA TAILLE DE L'EMPREINTE ? QUANTIFIER LES EFFETS
HORS SITE DES MINES SUR LES COMMUNAUTÉS VÉGÉTALES BORÉALES

HOW BIG IS THE FOOTPRINT? QUANTIFYING OFFSITE EFFECTS OF
MINES ON BOREAL PLANT COMMUNITIES

Ce chapitre a été soumis à la revue *Global Ecology and Conservation*

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2.1 Résumé

Les menaces que l'exploitation minière fait peser sur la biodiversité et les services écologiques des forêts boréales augmentent à mesure que la demande de minéraux s'accroît à l'échelle mondiale. Cependant, on en sait beaucoup moins sur la façon dont les effets hors site des mines affectent les communautés du sous-étage, car ils se produisent en dehors de l'emplacement immédiat de la mine et sont souvent négligés lors des évaluations écologiques. Nous avons mené une étude approfondie sur le terrain le long de transects d'un kilomètre autour de six sites miniers à différents stades d'exploitation (en exploitation ou non), qui traversent quatre types d'écosystèmes (feuillus, conifères, forêts mixtes et canopée ouverte) dans la forêt boréale canadienne. Les effets hors site des mines sur le sous-étage ont été quantifiés en utilisant des plantes vasculaires (ligneuses et herbacées), des bryophytes et des lichens. Les effets hors site des mines ont eu un impact sur la diversité et la composition du sous-étage. La richesse et la couverture du sous-étage étaient plus réduites à proximité des mines en exploitation que des mines non exploitées. Le stade d'exploitation minière a principalement altéré de manière significative la diversité du sous-étage et la structure de la communauté dans les forêts à feuilles caduques et mixtes, tandis que les communautés du sous-étage étaient plus résistantes aux effets hors site dans les forêts de conifères. L'empreinte a été quantifiée en utilisant la distance influencée et les effets les plus forts se situaient généralement à moins de 0,2 km des mines. Étant donné les changements prévus dans les écosystèmes de la forêt boréale avec l'empiètement des espèces à feuilles caduques sur les forêts de conifères et la sensibilité accrue des forêts mixtes et à feuilles caduques, la zone affectée par les effets hors site des mines pourrait augmenter à l'avenir. Nous suggérons que les effets hors site soient inclus dans les évaluations écologiques.

Mots-clés: impacts énigmatiques, empreinte minière, végétation de sous-bois, activités humaines.

2.2 Abstract

Threats from mining to the biodiversity and ecological services of boreal forests are increasing as demand for minerals increases globally. However, much less is known about how offsite effects of mines affect understory communities as they occur outside the immediate location of mining and are often overlooked during ecological evaluations. We conducted an extensive field survey along 1-km transects surrounding six mine sites of different mining stages (operating vs non-operating), that crossed four ecosystem types (deciduous, coniferous, mixed forests and open canopy) in Canada's boreal forest. The offsite effects of mines on the understory were quantified using vascular plants (woody and herbaceous), bryophytes and lichens. Mine offsite effects impacted understory diversity and composition. Understory richness and cover was more reduced near operating than non-operating mines. Mining stage mainly significantly altered understory diversity and community structure in deciduous and mixed forests while understory communities were more resistant to the offsite effects in coniferous forest ($P > 0.05$). The footprint was quantified using the influenced distance and the strongest effects were generally within 0.2 km from mines. Given the predicted changes in boreal forest ecosystems with encroachment of deciduous species into coniferous forests and the increased sensitivity of mixed and deciduous forests, the area affected by offsite effects of mines could grow in the future. We suggest that offsite effects should be included in ecological evaluations.

Keywords: enigmatic impacts, mine footprint, understory vegetation, human activities

2.3 Introduction

Ecological degradation, particularly biodiversity loss, is a global process with mining activities as one of the main drivers. This is due to habitat degradation, invasive species propagation, and pollution (Sonter et al., 2018; Tolvanen et al., 2019; Žibret et al., 2018). Currently, studies focus on impacts within industrial mine sites, however the effect of mine sites on surrounding landscape has not been studied. Offsite effects are impacts that extend beyond the edge of the immediate mine location reaching into undisturbed areas (Raiter et al. 2014). They are often overlooked in ecological evaluations and decision-making processes as they are outside of the designated project area and their impacts are often less visually distinct. The offsite landscapes surrounding mines are exposed to various disturbances including deposition of heavy metals, salt, and excess nutrients that are generated by mining operations, transportation, forest fragmentation and edge effects (Raiter et al., 2014; Sonter et al., 2018). A few studies report offsite effects on plants in the areas surrounding mines (Bech et al., 2012; Boojar and Tavakkoli, 2011; Christou et al., 2017). However, the potential for characteristics of mine sites (e.g., mining stages) and surrounding landscapes (e.g., ecosystem types over hundreds of meters) to modify the strength of offsite effects remains unstudied to our knowledge. This is particularly true in the boreal forest where increasing mining activity is becoming a highly visible threat to biodiversity (Venier et al. 2014; Musetta-Lambert et al. 2019).

Mine sites at different stages of the mining lifecycle generate different environmental disturbances. Generally, the mining lifecycle includes establishment (including surface vegetation cleared, building roads and processing facilities), operation (including extracting, processing, grinding and smelting), closure and rehabilitation (including removing waste and hazardous material and establishing new landforms and vegetation), which could be grouped into two stages of operating (operation) and non-operating (i.e., establishment, closure and rehabilitation) basing on the strength

of mining activities. At operating mine sites, wind, water and moving vehicles transport large amounts of particulate matter that are generated by excavation, roads, and residues (Csavina et al., 2012; Ochieng et al., 2010; Patra et al., 2016). Non-operating mine sites (i.e., mine sites at the stage of establishment, closure, and rehabilitation) have no intensive mining activities; however, clearing of surface vegetation and soil and residual pollution are their main sources of pollutants. Therefore, mine sites at different stages could lead to different effects on surrounding understory plant communities.

Deciduous, mixed (deciduous and coniferous trees), coniferous forests, and open canopy areas (i.e., peatlands without canopy trees) are four ecosystem types frequently found in the Canadian boreal forest, and their different environmental characteristics result in a varying diversity and composition of understory plant communities (Barbier et al., 2008; Kembel and Dale, 2006; Qian et al., 2003). Broadleaf overstories usually support higher understory cover and richness than conifer overstories (Chen et al., 2018a). Thus, factors that drive changes in ecosystem types will likely cause concomitant changes in understory communities and canopy structure. Coniferous forest ecosystems currently dominate many boreal forest landscapes, but global change is predicted to result in an increased abundance of deciduous and mixed forests (Boisvert-Marsh and de Blois, 2021; Mack et al., 2021). A shift towards deciduous and mixed forests is predicted to inhibit nonvascular species growth, increase vascular species establishment and lower canopy closure (Hart and Chen, 2008; Turetsky et al., 2010). Nonvascular groups (bryophytes and lichens) are typically more resistant to mining disturbances (e.g., heavy metals) than vascular species (Salemaa et al., 2004), whereas lower canopy closure can result in more dust falling on the ground (Amezaga et al., 1997). However, whether ecosystem type has the potential to modulate offsite effects of mines on understory diversity and composition remains unknown.

Our objective is to determine the offsite effects of mines on understory plant communities. To do this, we quantified the diversity and composition of understory vegetation (the presence and abundance of wood and herbaceous species as well as bryophyte and lichen abundance) along 1-km transects near six gold mines in Abitibi-Témiscamingue and Nord-du-Québec (Quebec, Canada) to examine whether community diversity and composition is associated with mining stage and ecosystem type. We also quantified the distance up to which impacts from the mine sites were detected. We tested three hypotheses: 1) more elements of the understory in terms of diversity and composition are expected to be affected surrounding operating than non-operating mine sites (H1); 2) understory communities differ in their sensitivity to offsite effects of mines among different ecosystem types (H2); 3) the influenced distance and understory community changes are expected to be small (H3) as previous studies found that the mean distance of edge influence on forest variables was less than 50 m (Franklin et al., 2021). By examining these three hypotheses, we will characterize the offsite effects of mines in the boreal forest and highlight their importance in future ecological evaluations and forest management planning.

2.4.1 Study area

Six mine sites were selected in the regions of Abitibi-Témiscamingue and Nord-du-Québec in the province of Québec, Canada (Fig. 1): Akasaba (48°3'12"N; 77°32'8"W, Agnico Eagle Mines Ltd), Canadian Malartic (48°7'21"N; 78°5'23"W, Canadian Malartic Corp.), Casa Berardi (49°33'43"N; 79°14'8"W, Hecla Québec Inc.), Joutel (49°29'28"N; 78°21'8"W, Agnico Eagle Mines Ltd), Lapa (48°13'45"N; 78°17'1"W, Agnico Eagle Mines Ltd), LaRonde (48°15'14"N; 78°25'59"W, Agnico Eagle Mines Ltd). The selected sites are different in size, mining method, life stage and bioclimatic domain, but together represent the main characteristics of mines in Canada's boreal forest. Lapa is a small underground mine, while Canadian Malartic is one of the largest open pit mines in North America. Three sites (Casa Berardi, LaRonde and

Canadian Malartic) are operating and the rest (Akasaba, Joutel and Lapa) are non-operating. Casa Berardi and Joutel are located in the *Picea mariana* –moss bioclimatic domain, while the other four are in the *Abies balsamea* – *Betula papyrifera* bioclimatic domain. Study sites are in the region that is characterized by a cold and humid continental climate with the mean temperature between -14 to -10°C in winter and between 16 to 18°C in summer (Environment Canada, 2021). Mean annual precipitation ranges from 790 to over 1,300 mm (Wang et al., 2013). More detailed information about individual mines is available in Table S1.1.

2.4.2 Fieldwork design and sampling

Six to eight transects (the number selected based on the size of each mine area, Fig. S1.1) were established perpendicular to each mine periphery through different ecosystem types in 2017-2019. Considering the environmental effects (e.g., dust) of mining are usually detected more than 1 km from mines (Gillings et al., 2022; Mattielli et al., 2009), the transects were established to 1 km from the edge of mine sites to make sure sampling plots were in the potential area influenced by the mine sites. Transects were positioned in different cardinal directions, but to avoid other non-mining disturbances (e.g., highways and towns), not all four cardinal directions are represented in some mine sites. This is particularly true at the Canadian Malartic and Lapa mine sites; to the north of the Canadian Malartic mine is the town of Malartic and to the south of Lapa mine is a highway, a natural gas pipeline and a hydroelectric corridor (Fig. S1.1). In addition, nine control transects with seven plots per transect were established in undisturbed natural sites (Fig. S1.1), five in the *Picea mariana* – moss bioclimatic domain and four in the *Abies balsamea* – *Betula papyrifera* bioclimatic domain.

Seven vegetation plots (3 m in radius) were established along each transect at distances from the edge of adjacent forest at 0, 20, 0.02, 0.1, 0.2, 0.5 and 1 km from

the mine. However, the plot in control transects did not follow the distance pattern of the transects around the mine and they were selected depending on ecosystem types to have balance the types in the controls. The forest stand of each plot was categorized into one of four ecosystem types based on tree species composition and size (diameter at breast height (DBH) $\geq 10\text{cm}$) within a prism plot (2M basal area factor) centered on the vegetation plot: coniferous ($> 70\%$ of coniferous tree stems, $N=173$), deciduous ($>70\%$ deciduous tree stems, $N=48$), mixed forests (coniferous tree stems between 69% and 31% , $n=38$), and open canopy ecosystem (DBH of all trees below 10 cm , primarily peatlands in our study area, $N=110$, Table S1.2). In total, 371 plots were established, but 369 plots were used because two control plots were destroyed by logging.

Understory composition in each vegetation plot was determined based on the Canadensys Database (www.canadensys.net) during the summer of 2019. In total, 172 vascular species (60 woody species and 112 herbaceous species) were identified (species list in Table S1.3). Cover (%) of each vascular species, total bryophyte and total lichen was determined and value in three random 1 m^2 quadrats were averaged by plot. The cover of individual species often overlapped and, therefore, total cover sometimes exceeded 100% .

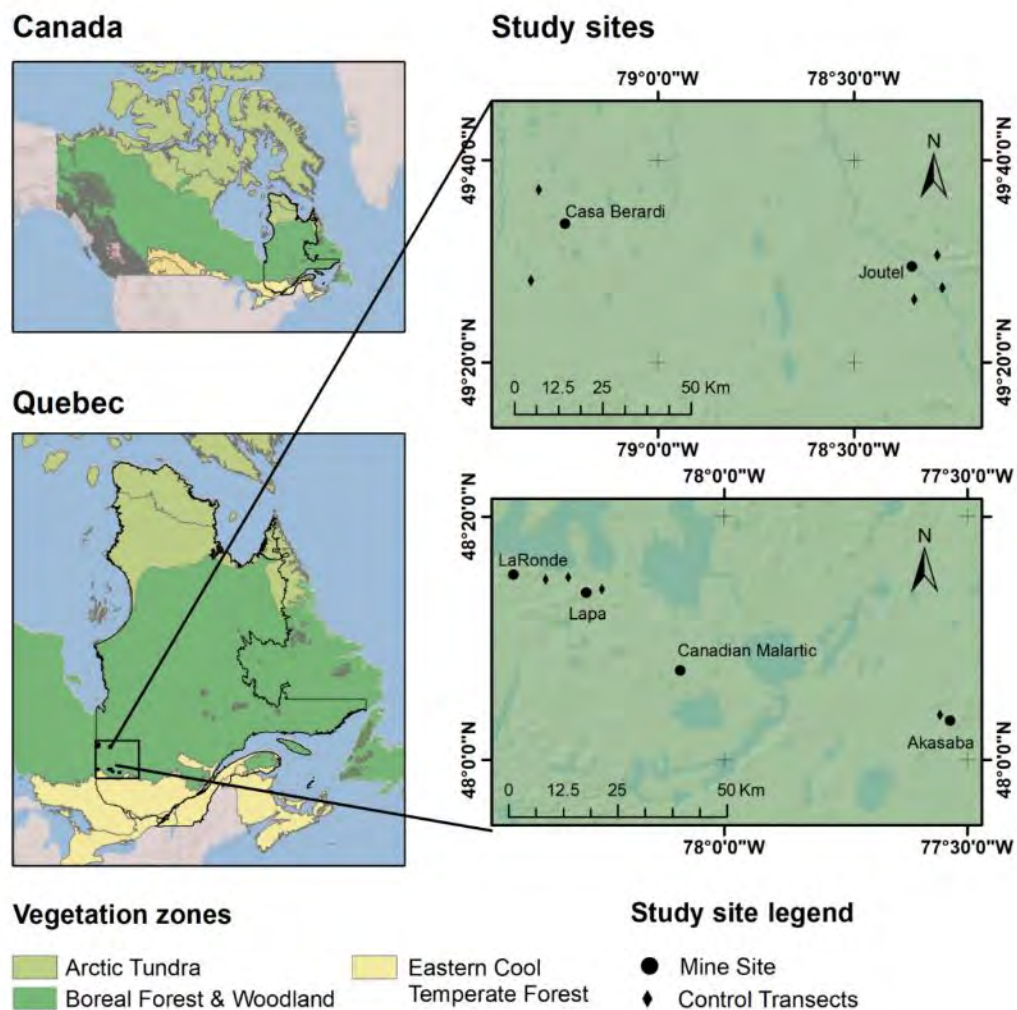


Figure 2.1 Study area. Left, map of Canada and Quebec showing the main vegetation zones. Right, map of study region indicating the position of sampling sites at six gold mine sites and 9 control transects (graph by Enrique Hernández-Rodríguez using GIS, August 9th, 2021)

Furthermore, 13 structural variables (i.e., ground cover of leaf litter, woody debris, organic matter, water and rock, soil humidity and organic soil depth, canopy openness, conifer proportion, live crown ratio and basal areas) were measured at each plot (listed in Table S1.4). A principal component analysis (PCA) for structural variables illustrates that the 13 variables differed significantly among ecosystem types

(ANOVA on the first two axis scores, $p < 0.001$, Fig. 2.2), therefore, structural variables were grouped into ecosystem types in the analysis. Furthermore, no differences in almost all structural variables were found between distances along the transects (Table S1.5), which supports our sampling design and confirms that the plots were established in relatively intact landscapes.

2.4.3 Statistical analysis

2.4.3.1 The impact of mining stage and ecosystem type on understory structure in the mine offsite landscapes

To describe patterns of understory diversity affected by mining stage and ecosystem type in mine offsite landscapes, we fitted a generalized linear mixed model (GLMM, “glmer” function from the *lme4* R package, Bates et al., 2018, and “glmmTMB” function from *glmmTMB* package, Brooks et al., 2017). Richness, cover and alpha diversity indices (“diversity” function from *vegan* package, Oksanen et al., 2013) of all vascular species, woody and herbaceous richness and cover and bryophyte and lichen cover were used as the response variables (Table S1.6). The predictor variables included mining stage, ecosystem type and their interaction as fixed effects. To avoid spatial autocorrelation, the nested terms representing the sampling design (site, transect and plot) were included to take into account the spatial location of the data. Different distribution functions were used, Conway-Maxwell-Poisson (“glmmTMB” function and family= “compois”) for underdispersion of woody richness counts, Negative Binomial (“glmer” function and family=MASS::negative.binomial) for overdispersed of all vascular and herbaceous richness counts, Gaussian (“glmmTMB” function and family=gaussian) for continuous number with normally distributed residuals of Shannon Index, bryophyte cover, all vascular, woody and herbaceous covers, Gamma (“glmer” function and family=Gamma) for continuous number with not normally distributed residuals of Simpson Index, and Zero-inflated beta distributions (“glmmTMB” function and family=beta_family) for Zero-Inflated data

of lichen cover. Marginal significance of the predictors was tested by type II analysis of variance with the “Anova” function in *car* package (Fox et al., 2012). Pairwise comparisons for all response variables among mine stages and ecosystem type were conducted with Tukey’s post hoc test (“emmeans” function in the *emmeans* package, Lenth et al., 2018). When an interaction term was significant, pairwise comparisons were performed between mine stages within each ecosystem type, whereas pairwise comparisons between mine stages and between ecosystem types were performed independently when the interaction term was not significant. All analyses were performed with R 3.4. Results were visualized with the *ggplot2* package (Wickham et al., 2015).

To assess variations in understory composition in mine offsite landscapes, the differences in species composition patterns of woody and herbaceous plants among mining stages and their interactions with ecosystem types were tested by permutational multivariate analysis of variance (PERMANOVA, “adonis” function from the *vegan* package, Anderson, 2005) on Bray–Curtis dissimilarity matrices with 999 permutations. Dissimilarity between plots was further visualized by distance-based Principal co-ordinates analysis (PCoA, “PCOA” function in the *vegan* package, Oksanen et al., 2013). We also ran pairwise PERMANOVAs (999 permutat, “pairwise.adonis” function in the package *RVAidemoire*, Hervé, 2017) to separately compare the mining stages with a multiple comparison correction based on Benjamini-Hochberg method (Benjamini and Hochberg, 1995). In addition, to further define communities specifically associated with each mining stage, indicator species analysis was performed with the “multipatt” function in the *indicspecies* package (De Caceres et al., 2016).

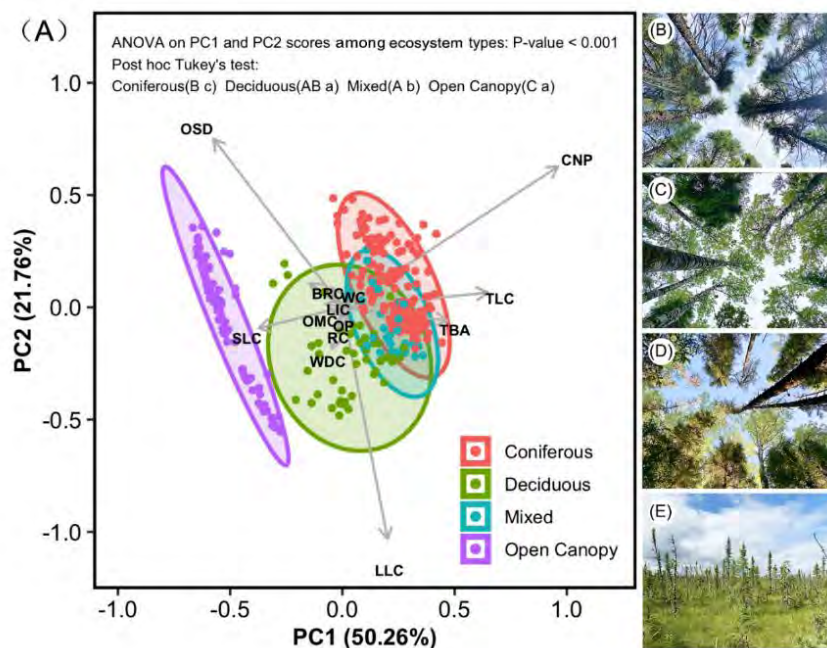


Figure 2.2 Biplot of principal component analysis (PCA) of structural variables across plots and forest types. Each ecosystem type (coniferous (B), deciduous (C), mixed forest (D) and open canopy ecosystem(E)) is represented with a specific color and ellipse. All plots are displayed (small circles). SLC: live crown ratio for sapling trees; TLC: live crown ratio for canopy trees; SBA: basal area for sapling trees; TBA: basal area for canopy trees; OP: Openness; LLC: leave litter ground cover; WDC: woody debris ground cover; WC: water ground cover; RC: rock ground cover; OMC: organic matter ground cover; OSD: depth of organic soil; CNP: conifer proportion; HM: humility; BRC: bryophyte ground cover; LIC: lichen ground cover.

2.4.3.2 Understory diversity patterns along distance from mine sites and footprint

To assess the patterns of understory diversity with distance from mine sites, generalized linear mixed models were performed. As the distance between plots in control transects and mine sites varied and to ensure the independence of predictors, control transects were not included in the models. Response variables (understory richness, cover and alpha diversity indexes) were the same as those used in 2.3.1 while nested terms (mine site, transect and plot) and ecosystem types were included as random factors. The predictor variables included the distance from mine sites and

its interaction with mining stage. When the graphs (“ggplot” function in *ggplot2* package) of responses present a non-linear relationship with distance, a second-order polynomial of distance (“poly” function in *stats* package) was included in the predictors to account for unimodal patterns (all vascular and herbaceous plant cover). As we expected the disturbance from mines to decrease with increased distance, only linear and second-order polynomial patterns were considered. The insignificant interactions and quadratic terms were removed from the final models. As the presence of ecosystem types with distance was random, generated by the structure of the landscape (Fig. S1.1, Table S1.2) and unbalanced, ecosystem type and cardinal direction were included as random factors.

To determine influenced distances, we calculated means and plotted all responses against the distance using the *ggplot2* package. Based on the general average influenced distance, the offsite footprint of the selected six mine sites were calculated with the Google Earth Engine Platform. As the selected mine sites in our project varied in size, shape, mining method, stage of mining lifecycle and bioclimatic domain, their average offsite footprint was considered representative and was further used to estimate total offsite footprint of operating mine sites in the boreal biome of Canada. To avoid overestimation of influence area, we used the distance that had the strongest effects in estimation of average offsite footprint.

2.5 Results

2.5.1 Effects of mine stage and ecosystem type

The effect of mining stage on understory diversity was not consistent among the ecosystem types in the offsite landscapes. The richness and cover of all vascular and herbaceous plants as well as the cover of bryophytes and lichens were affected by the interaction between mining stage and ecosystem type (the results of GLMMs, ANOVA type II sums, $P < 0.05$, Table S1.7 and S1.8). All vascular and herbaceous

plants had lower richness and cover in deciduous or mixed forests near operating mines than in controls (the results of Tukey HSD pairwise comparisons in Figure 2.3). Notably, herbaceous richness and cover were also lower near non-operating sites than near controls (Figure 2.3). Unlike vascular plants, bryophyte cover differed between mining stages in coniferous forests and lichen cover was affected in open canopy ecosystems. These trends were observed near both operating and non-operating sites (Figure 2.3).

The composition of offsite understory vascular communities was also significantly associated with mining stage, ecosystem type and their interaction for both woody and herbaceous plants (see results of PERMANOVAs in Figure 2.4), although the effects were small (the sum contribution of mining stage, ecosystem type and their interaction in PERMANOVA results, $R^2 < 0.11$, Table S1.9). This pattern was visible in the principal component analysis (PCoA), where the communities clustered by mining stage in each ecosystem type (Figure 2.4). Ecosystem type was the largest contributing variable to community composition of woody ($F=9.37$, $R^2 = 0.071$, $p < 0.001$) and herbaceous species patterns ($F=5.71$, $R^2 = 0.046$, $p < 0.001$), while mining stage (woody and herbaceous, $F=2.898$, $R^2 = 0.015$, $p < 0.001$ and $F=4.209$, $R^2 = 0.023$, $p < 0.001$, respectively) and their interaction (woody and herbaceous, $F=1.445$, $R^2 = 0.022$, $p = 0.007$ and $F=1.555$, $R^2 = 0.025$, $p = 0.002$, respectively) were also significant but contributed less (PERMANOVA results in Table S1.9). Pairwise PERMANOVA revealed that woody community composition differed among mining stages in coniferous and mixed forests, while herbaceous composition differed more among mining stages in deciduous forests and open canopy ecosystems than in coniferous and mixed forests (pairwise PERMANOVA results in Table S1.10).

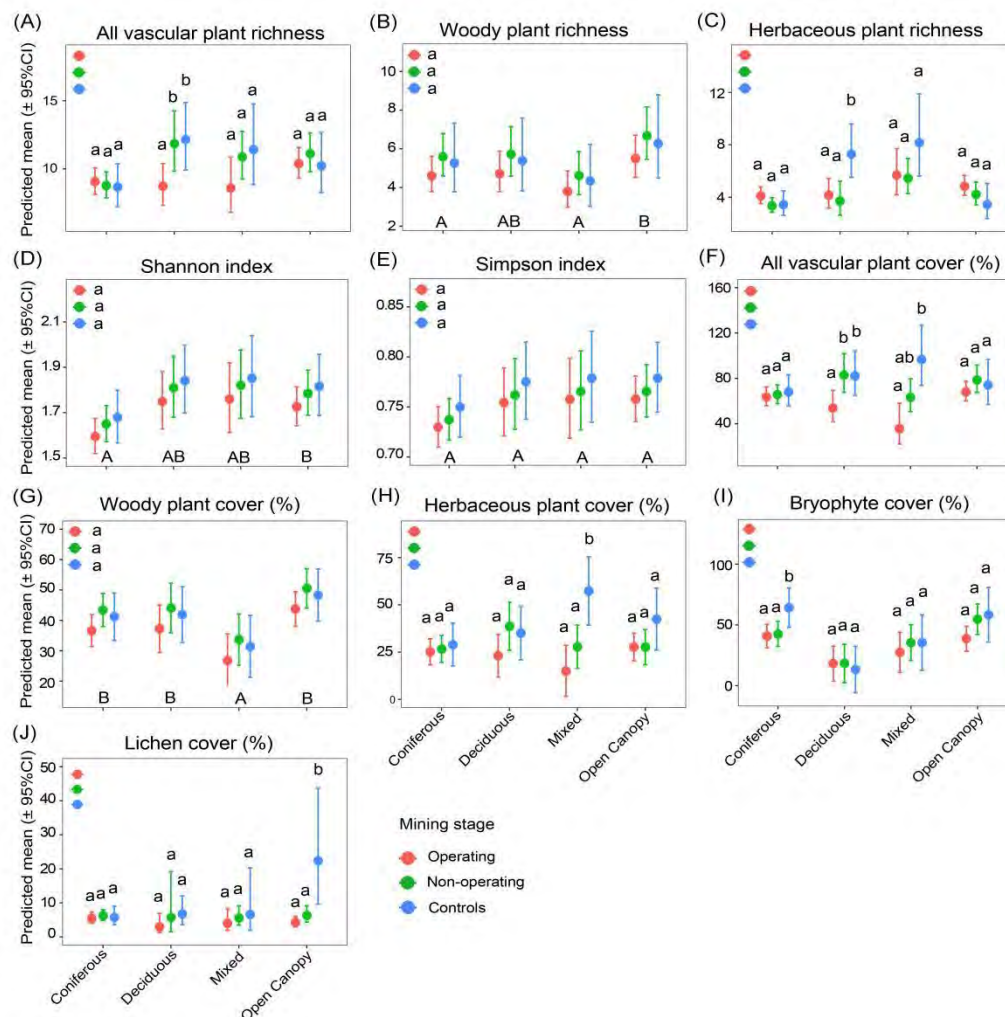


Figure 2.3 Comparison of understory diversity between mining stages for each ecosystem type. Points show means for all samples; bars show a 95% confidence interval around the mean. Different lowercase letters indicate significantly different means between groups based on Tukey HSD pairwise comparisons ($\alpha = 0.05$) from the results of generalized linear mixed model (Table S1.4 and Table S1.5). When interactions between ecosystem type and mining stage were significant, pairwise comparisons were used between mine stage in each ecosystem type, while when the interactions were not significant, the interaction terms were removed from the models and pairwise comparison were used between forest types and mining stages, respectively.

Indicator species analysis indicated that more species were associated with controls than non-operating and operating mine sites (Table 2.1). Two non-native species (*Pilosella aurantiaca* (L.) FW Schultz & Sch. Bip. and *P. caespitosa* (Dumort.) P. D. Sell & C. West.) and one non-native species (*Galium sylvaticum* L.) were closely associated with operating and non-operating sites, respectively. However, all indicator species for controls were native species (Table 2.1).

Table 2.1 List of indicator species for the landscape near sites at different mining stage (only significant species are shown, $P < 0.05$)

Species	Types	Operating (5)		Non-operating		Controls (15)	
		Stat	P value	stat	P value	stat	P value
<i>Amelanchier arborea</i>	Woody			0.172	0.003		
<i>Aralia nudicaulis</i>	Herbaceous					0.182	0.003
<i>Athyrium filix-femina</i>	Herbaceous					0.136	0.013
<i>Coptis trifolia</i>	Herbaceous	0.129	0.045				
<i>Cornus sericea</i>	Woody			0.134	0.02		
<i>Corylus cornuta</i>	Woody					0.137	0.028
<i>Dendrolycopodium</i>	Herbaceous					0.14	0.015
<i>Dryopteris carthusiana</i>	Herbaceous					0.217	0.001
<i>Epilobium angustifolium</i>	Herbaceous			0.147	0.01		
<i>Eriophorum vaginatum</i>	Herbaceous			0.125	0.047		
<i>Galium sylvaticum</i> *	Herbaceous			0.118	0.029		
<i>Gymnocarpium dryopteris</i>	Herbaceous					0.13	0.018
<i>Lonicera hirsute</i>	Woody					0.146	0.023
<i>Maianthemum canadense</i>	Herbaceous					0.137	0.029
<i>Moneses uniflora</i>	Herbaceous			0.141	0.018		
<i>Pilosella aurantiaca</i> *	Herbaceous	0.129	0.046				
<i>Pilosella caespitosa</i> *	Herbaceous	0.12	0.047				
<i>Rhododendron canadense</i>	Woody			0.142	0.013		
<i>Ribes hudsonianum</i>	Woody					0.166	0.003
<i>Rubus chamaemorus</i>	Herbaceous					0.133	0.041
<i>Salix bebbiana</i>	Woody	0.15	0.009				
<i>Sorbus Americana</i>	Woody	0.14	0.026				
<i>Sparganium americanum</i>	Herbaceous					0.169	0.005
<i>Spinulum annotinum</i>	Herbaceous					0.169	0.005
<i>Vaccinium myrtilloides</i>	Woody			0.151	0.017		
<i>Viburnum lentago</i>	Woody					0.188	0.001
<i>Viburnum nudum</i>	Woody					0.132	0.037

Notes: Stat is the association statistic and P value is the p -value of the permutational test. “*” indicates the non-native plant species.

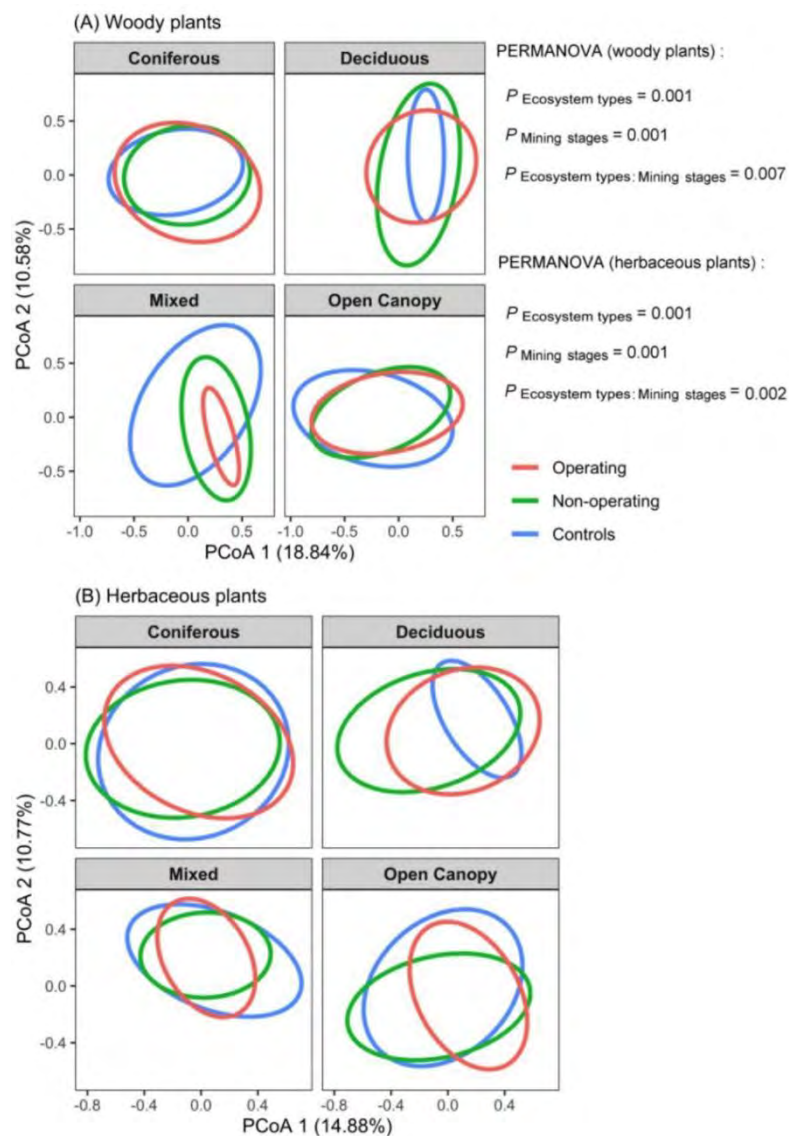


Figure 2.4 Ordination diagrams and PERMANOVA results of understory vascular plant composition at the community level. Principal co-ordinates analysis (PCoA) for woody plants (A) and herbaceous plants (B) based on Bray–Curtis dissimilarity. Ellipses represent 95% confidence intervals (individual data points not shown for clarity). Texts on the right side present the results of PERMANOVA.

2.5.2 Distance influenced by offsite effects of mines and offsite footprint

The impacts of mining activities on understory diversity generally decreased with increasing distance from operations, although the pattern was mediated by mining stage (the results of GLMM in Table S2.11 and Fig. 2.5). All six response variables (richness of all vascular, woody, and herbaceous plant, Shannon and Simpson indices and bryophyte cover) were associated with distance except for two response variables (all vascular, herbaceous richness) that varied not only with distance but also with by mining stage (operating vs. non-operating, Table S2.11).

The variations with distance occurred mostly in 0.2-0.5 km for Shannon and Simpson indices, 0.2 km for woody plant richness and 0.1 m for bryophyte cover near all mine sites, according to the pattern of mean values with the distance (Fig. 2.5). Moreover, the richness of all vascular and herbaceous plants had no significant patterns with distance near operating mines but there was a pattern near non-operating sites where richness of all vascular plants showed a slight threshold (turning point) at 0.5 km (Fig. 2.5A), and herbaceous richness gradually increased without a threshold (Fig. 2.5B).

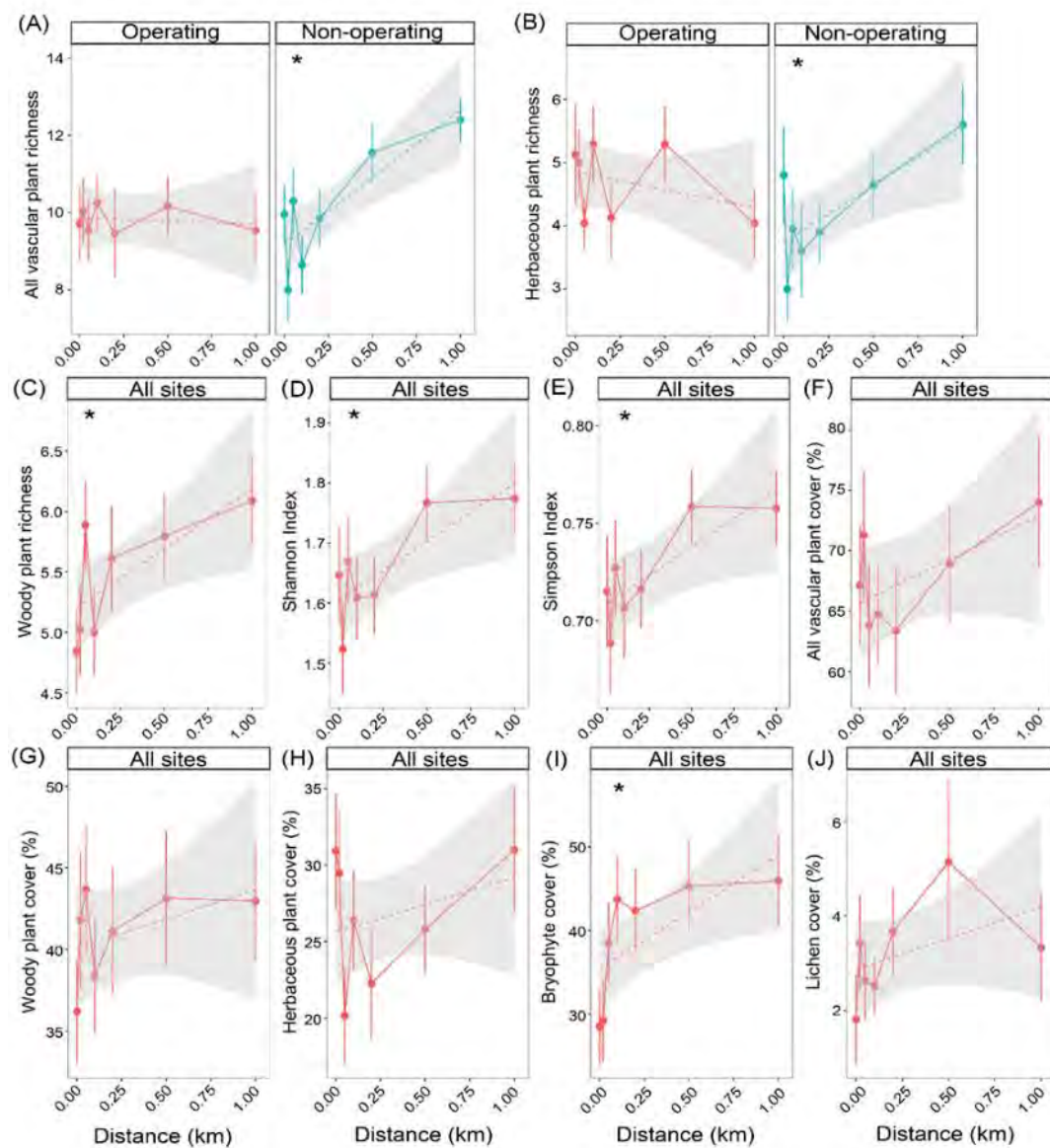


Figure 2.5 Effects of distance from mine site on understory diversity. Points show means for all samples; bars show standard error (SE). The significant associations were examined based on the results of (generalized) linear mixed model (Table S1.8). Dashed lines show the fitted trends and asterisks show significant correlations ($P < 0.05$).

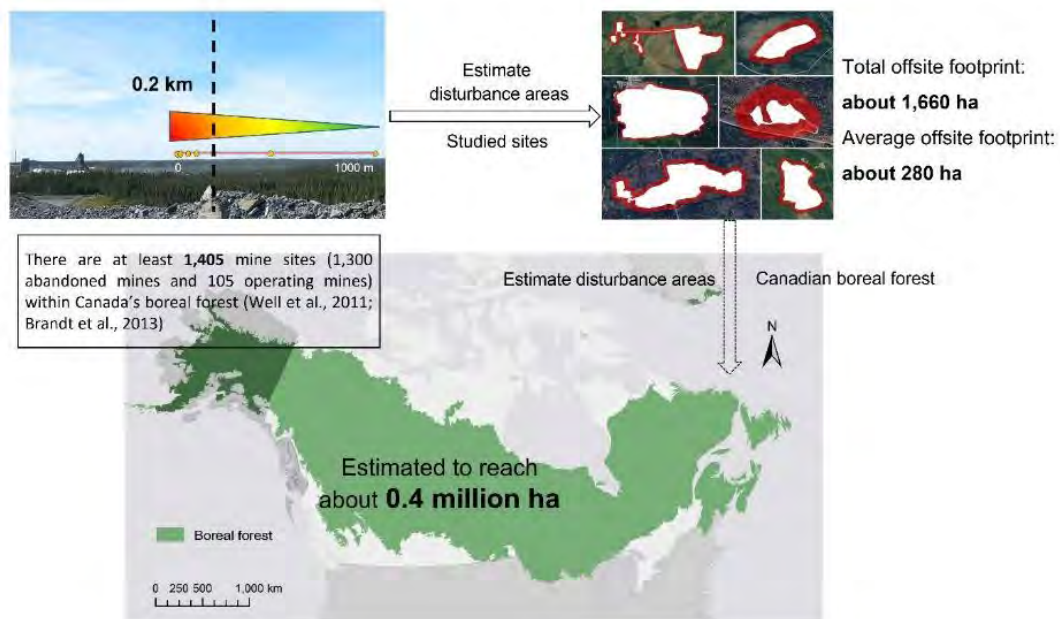


Figure 2.6 Total disturbance offsite footprint of mines in Canada's boreal forest. The offsite footprint was calculated within the Google Earth Engine Platform. The data of operating and abandoned mines is from previous reports (Brandt et al., 2013; Wells et al., 2010)

2.6. Discussion

2.6.1 Determinants of understory diversity and composition in surrounding landscapes of mine sites

The presence of offsite effects of mines on understory communities was confirmed. The offsite effects decreased understory diversity (richness and cover) in the surrounding landscapes, but the severity of the effects was modulated by mining stage and ecosystem type. As expected in H1, reduced richness and cover were mainly found near operating rather than near non-operating mines, which suggests potential recovery in plant diversity after mine closure. Alternatively, mining activities including digging, blasting and grinding mainly occurred during the operating and not the non-operating stage, which could lead to overall stronger effects

on understory communities near operating mines because of higher heavy metal contents and dust deposition (David et al. 2021). However, some offsite effects were still detected near non-operating sites (e.g., lower herbaceous, bryophyte and lichen cover, Fig. 2.3). This may be the result of residual pollution from the operating period that may persist after closure (Festin et al., 2019; Frouz et al., 2001). Some studies reported that the residuals effects on onsite surface vegetation diversity could persist over 35 years after mine closure (Gould, 2012; Holl, 2002; Norman et al., 2006) and our results confirmed that persistent residual effects of mining might be also remaining surrounding intact landscapes.

Notably, herbaceous plants, bryophytes and lichens were more sensitive to the offsite effects than woody plants. With high water absorption efficiency and a greater ability to store nutrients (Craine and Dybzinski, 2013), woody plants usually have a higher resilience to environmental changes than other understory groups (Kirk et al., 2019; Salemaa et al., 2001; Watkinson et al., 2021). Bryophytes and lichens do not have roots and conductive tissues and absorb mineral salts as well as heavy metal ions from precipitation and dry deposition on their surface (Kłos et al., 2018; Korzeniowska et al., 2021). Bryophytes and lichens have been widely reported as indicators of air and metal pollution (Gerdol et al., 2014; Onianwa, 2001; Szczepaniak and Biziuk, 2003). Therefore, they have the potential to be applied in future offsite effect evaluation as specific ecological indicators of mining, but more work is needed to explore which species are the most reliable.

Ecosystem type was also an important factor associated with the offsite effects as we expected in H2. Understory communities showed variable resistance to the offsite effects among ecosystem types. Vascular plant diversity and composition changed in deciduous and mixed but not in coniferous forests, which may be due to the possibility that coniferous trees intercept more dust, heavy metals and other pollutants than deciduous trees (Augusto et al., 2002; Barbier et al., 2008; Nguyen et al., 2015)

because of higher canopy cover, leaf area index and a more persistent foliage (Augusto et al., 2002). Mixed and deciduous forests had similar responses to the offsite effects, which could be due to their similar canopy structure with high seasonal variation in canopy gaps because deciduous trees drop their leaves in autumn (Brassard et al., 2008; Chávez and Macdonald, 2010). Furthermore, deciduous trees can translocate high amounts of heavy metals into the foliage which can then result in more heavy metal accumulation in topsoil (Van Nevel et al., 2011). Consequently, acidic deposition (e.g., sulfides and carbonates) from mining could lead to an increase in available heavy metals in the soil, resulting in greater effects on the understory species in deciduous and mixed forests.

Both woody and herbaceous plants were more resistant in open canopy ecosystems, potentially because the vegetation in these ecosystems (acidic and nutrient poor) develops unique adaptations to survive (Williams-Mounsey et al., 2021) and therefore is less vulnerable to the offsite effects. An example are the insectivorous species *Sarracenia purpurea* L. and *Drosera rotundifolia* L. (Table S1.3) are able to tolerate low oxygen, acidity and poor nutrient availability (Ellison and Gotelli, 2001; Hoyo and Tsuyuzaki, 2014).

In contrast to the findings on vascular plants, bryophyte cover was affected in coniferous forests while lichen cover was affected in open canopy ecosystems. Reduced bryophyte cover in coniferous forests may be due to canopy cover loss near mine sites given that forest-dwelling bryophytes are considered to be shade-tolerant (Tinya et al., 2009). Similarly, previous studies have shown that bryophyte abundance can be used to determine the depth of edge effects, which could be attributed to the removal of large trees and canopy disruption (Benítez et al., 2015, Stewart and Mallik, 2006). The negative offsite effects of mines on lichen cover in open canopy ecosystem is consistent with previous studies that showed a decrease in lichen cover near a diamond mine in the boreal forest (Watkinson et al., 2021). This

is probably a consequence of the high sensitivity of lichens to dust deposition (Farmer, 1993; Salemaa et al., 2004).

Three non-native species (*Pilosella aurantiaca*, *P. caespitosa* and *Galium sylvaticum*) were identified as indicators of offsite effects, supporting the finding of earlier studies that mining activities promote the establishment and expansion of non-native species (Boyce, 2002; Calinger et al., 2015; Lemke et al., 2013). Removal of native vegetation, soil, soil microbes, and seed banks from the surface of mine sites (Dhar et al., 2018) may have provided opportunities for non-native species. With small plant height and greater capacity to utilise soil resources than native species, non-native species usually can establish rapidly and grow under harsh conditions such as nutrient-poor soils (Jo et al., 2015; Mathakutha et al., 2019; Schumacher et al., 2009). Our results indicate that these non-native species growing on the surface of mines could reach into surrounding intact forests. Therefore, mine offsite effects would also be an important driver of the spread of non-native species.

Our results supported our first and second hypothesis. Mining stage and ecosystem type were important factors affecting the magnitude of mine offsite effects. The combination of canopy cover loss, soil contamination as well as species differences in the tolerance to environmental changes might uncover the mechanisms how offsite effects of mines affect understory diversity and composition.

2.6.2 Determinants the footprint of offsite effects and their total impacts

Understory diversity indexes increased with the distance from the edge of mine sites following as expected for a point-source of pollution where disturbance and potential pollutants (such as acidic depositions, heavy metals, and other toxins) extend from the mine sites to offsite landscapes. Higher heavy metal content, for example, has been reported in soil and dust near mines, and their bioavailability decreased when moving away from the disturbance sources (Elmayel et al., 2020; Tang et al., 2020).

Unfortunately, in our study, information on wind direction or dust emissions by mine sites, which would have allowed us to assess correlations between the spread of contaminants and the observed offsite effects on plant communities, was not available, but should be considered in future studies.

As hypothesized in H3, the changes in understory diversity caused by offsite effects, although statistically significant, impacted a relatively small area (strongest effects occurred within 0.2 km) and weak (e.g., only 1-3 species varied along the distance). Offsite effects are considered enigmatic impacts as they are often overlooked in ecological evaluations of mining projects (Raiter et al., 2014). However, for some sensitive groups or species such as bryophytes, which had their cover reduced by about 15 % in forests near mines compared to distant plots (Fig. 2.5I), the offsite effects could be large.

Furthermore, when individual offsite effects are considered together, the cumulative impacts could be significant. The total disturbance footprint was estimated to up about 1,660 ha (Fig. 2.6) for the six selected mine sites (calculated with the Google Earth Engine Platform based on 0.2 km areas surrounding each site). The total offsite footprint could reach up to about 0.4 million ha (using the average offsite footprint value 280 ha of the six mine sites) across Canada's boreal forest, based on previous reports indicating that there were approximately 1,405 mine sites (105 operating and 1,300 abandoned mines within Canada's boreal forest, Brandt et al., 2013; Wells et al., 2020). Since mine sites were only selected in Canada's boreal forest, future work in other regions will be needed to predict the total footprint of offsite effects at a larger scale. Offsite effects of mines are a disturbance (like roads and gas pipelines) that have small effects at the scale of individual sites but that occur in many sites across the globe. Another concern is that the cumulative effects with other disturbances (e.g., logging, climate changes and forest fire) could be greater.

2.6.3 Management implications

Based on the results obtained in this study, we propose that mine offsite landscapes should be included in future ecological evaluations, and ecosystem types and mining lifecycle should be considered in the process. Furthermore, the predicted shifts in forest composition from coniferous to deciduous dominance with global change in boreal areas could exacerbate the offsite effects of mines on understory communities. Given that the strongest offsite effects mainly occurred within 0.2 km from mine sites, a buffer zone around mines is recommended to offset these effects. Keeping and planting conifer trees surrounding mine sites might be a reasonable option considering coniferous forest is the dominant stand type in boreal forest and showed a relatively high resistance to the offsite effects. Future work should determine the width of buffer zone and forest management strategies within the zone.

2.7 Conclusion

Offsite effects of mine sites were confirmed in boreal understory communities and their severity was associated with ecosystem type and mine lifecycle stage. Mine offsite effects on understory diversity and composition were generally small and seemed negligible in the overall biodiversity of boreal forest, especially in coniferous forest. However, key species loss, non-native species invasion, and total impacts at a large scale may increase the impact of mine offsite effects on ecosystem services. With global climate change and the predicted changes in boreal forest ecosystems, including the encroachment of deciduous species into coniferous forests, the areas that are affected by offsite effects of mines could grow in the future given the increased sensitivity of mixed and deciduous forests compared to coniferous forests. Furthermore, the increasing number of mining projects will lead to more landscapes being exposed to offsite effects. Our findings encourage future studies to evaluate how these changes by offsite effects affect ecological services of the boreal forest and

their cumulative impacts with other anthropogenic and natural disturbances at a large scale.

2.8 Acknowledgements

This research was financially supported by the NSERC-UQAT Industrial Chair in Northern Biodiversity in a mining context. We thank Enrique Hernández-Rodríguez for his assistance in creating the study area map. We deeply thank Julie Arseneault and Mélanie Jean for the establishment of the plots. We also thank Julie Arseneault, Mélanie Jean, Elizabeth Riendeau, Rémi Boisvert, Andréane Garant, Cyrielle Bigonneau, Simon Bégin, Matilde Offroy, Pamela Parisien, Jeffrey Raymond, Juliette Larrivée, Marie-Anyise Dubuc, Emmanuelle Richard, Maxime Thomas, Terresa Kim, Xavier Saint-Amant, Jolianne Garand and Natalie Vuong for their valuable assistance in the fieldwork. We also thank Josée Noël, Lucienne Anctil, Stéphanie Lafrenière and other people at the mines of Canadian Malartic, Casa Berardi, LaRonde, Lapa, Akasaba and Joutel for their help with access to the mine sites.

CHAPTER III

EFFETS SYNERGIQUES DANS PAYSAGE HORS SITE DU SITES MINIER :
LES CHANGEMENTS PRÉVUS DANS LES ÉCOSYSTÈMES POURRAIENT
EXACERBER LES EFFETS DE L'EXPLOITATION MINIÈRE SUR LA
STRUCTURE DES COMMUNAUTÉS DE BRYOPHYTES.

SYNERGISTIC EFFECTS IN MINE OFFSITE LANDSCAPES: PREDICTED
ECOSYSTEM SHIFTS COULD EXACERBATE MINING EFFECTS ON
BRYOPHYTE COMMUNITY STRUCTURE

Ecological Indicators, DOI: 10.1016/j.ecolind.2022.109555

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3.1 Résumé

Le changement global modifie l'abondance relative des types d'écosystèmes dans les forêts boréales, tandis que la transition énergétique verte entraîne une augmentation des activités minières dans le monde entier. L'interaction et les effets conséquents de ces deux tendances sur la biodiversité n'ont pas été examinés en profondeur. Les espèces de bryophytes peuvent être utilisées comme indicateurs pour mesurer ces effets car elles jouent des rôles écologiques clés dans les forêts boréales. Nous avons identifié et évalué l'interaction entre le type d'écosystème (c.-à-d. conifère, feuillu, forêt mixte et canopée ouverte) et l'exploitation minière sur la diversité et la composition des bryophytes à l'échelle du microhabitat dans des paysages d'un kilomètre autour de six sites miniers à différentes étapes du cycle de vie minier dans la forêt boréale canadienne. Indépendamment du type de microhabitat, les effets combinés du type d'écosystème et du stade d'exploitation minière étaient interactifs sur les bryophytes. La richesse en bryophytes et la composition de la communauté ont été affectées négativement par les effets hors site des mines uniquement dans les forêts de feuillus et mixtes. Les effets interactifs sur la richesse en bryophytes se sont produits principalement dans les microhabitats du sol. Nous avons également constaté que les forêts de feuillus, les forêts mixtes (forêt de conifères comme référence) et les mines avaient un impact négatif sur l'abondance des mousses à plumes et des sphaignes. En outre, des espèces indicatrices ont été identifiées pour les zones affectées par les mines (*Pohlia nutans* et *Dicranum polysetum*) et pour les zones témoins (*Sphagnum angustifolium* et *Plagiomnium cuspidatum*). Nos résultats suggèrent que les changements d'écosystème prévus avec les changements globaux, des forêts de conifères aux forêts de feuillus, pourraient potentiellement augmenter les effets de l'exploitation minière sur la résistance de l'écosystème forestier à travers les changements de la structure de la communauté de bryophytes. L'ajout de microhabitats (c'est-à-dire l'ajout de débris ligneux grossiers) près des sites miniers est une stratégie potentielle pour maintenir la richesse des espèces. Collectivement,

ces résultats font progresser notre compréhension de la façon dont l'exploitation minière affecte la biodiversité et soulignent l'importance de prendre en compte les paysages hors des sites miniers dans les futures évaluations environnementales des projets de développement dans le contexte des changements mondiaux.

Mots-clés : Activités minières, changements climatiques, impacts indirects, épiphytes, microhabitats, taïga.

3.2 Abstract

Global change is shifting ecosystem type relative abundance in boreal forests, while the green energy transition results in increased mining activities around the globe. The interaction and consequent effects of these two trends on biodiversity have not been examined in depth. Bryophytes species can be used as indicators to measure these effects because they play key ecological roles in boreal forests. We identified and evaluated the interaction between ecosystem type (i.e., coniferous, deciduous, mixed forest and open canopy) and mining on microhabitat scale bryophyte diversity and composition in 1-km landscapes surrounding six mine sites at different stages of the mining lifecycle in the Canadian boreal forest. Irrespective of microhabitat type, the combined effects of ecosystem type and mining stage were interactive on bryophytes. Bryophyte richness and community composition were negatively affected by offsite effects of mines in only deciduous and mixed forests. The interacted effects on bryophyte richness mainly occurred on the ground r microhabitats. We also found that deciduous, mixed forests (coniferous forest as a reference) and mines had a negative impact on the abundance of feather mosses and sphagna. Furthermore, indicator species were identified for areas affected by mines (*Pohlia nutans* and *Dicranum polysetum*) and for control areas (*Sphagnum angustifolium* and *Plagiomnium cuspidatum*). Our results suggest the predicted ecosystem shifts with global changes, from coniferous to deciduous forests, could

potentially increase the effects of mining on forest ecosystem resistance through the changes in bryophyte community structure. Adding microhabitats (i.e., adding coarse woody debris) near mine sites is a potential strategy in maintaining species richness. Collectively, these findings advance our understanding of how mining affects biodiversity and highlight the importance of considering mine offsite landscapes in future environmental evaluations of development projects in the context of global changes.

Key words: Mining activities, climate changes, indirect impacts, epiphytes, microhabitats, Taiga

3.3 Introduction

Global changes are modifying the distribution of ecosystem types across the globe. Changes include shifts from boreal coniferous to deciduous forest and from northern peatlands to forests (Hirota et al., 2011, van der Velde et al., 2021). How the effects of these global changes will interact with accelerated ongoing human activities is an important area of study. A synergistic effect appears when the combined effects of two disturbances is greater than their sum (Coors and De Meester, 2008, Raiter et al., 2014). An example is the interactions between global changes and forest fires (Hessburg et al., 2021), logging (Anderson et al., 2017, Cusack et al., 2016), and agricultural activity (Anderson et al., 2017, Cusack et al., 2016, Danneyrolles et al., 2019), which all present synergistic results in effects on biodiversity and the provision of ecosystem services. However, the effects of mining and their potential interactions with global changes have received little attention despite the fact that mines are a significant source of disturbances in forest ecosystems globally (Martins and Lima, 2020, Maynard et al., 2014). Numerous mines and mineral deposits are located in the boreal forest (Fig. 3.1) and more mining projects will be established in this area in coming decades as the energetic transition increases the world's demand

for minerals such as rare earths for clean technologies and uranium for power generation (Sonter et al., 2020, Watari et al., 2021, Yin et al., 2021). This will lead to more intact boreal landscapes being exposed to mining disturbances while they are also facing the consequences of global changes.

Coniferous, deciduous and mixed forests as well as open canopy (primarily peatlands) are four main ecosystem types in boreal landscapes. Predicted shifts from coniferous needle-leaved to deciduous broad-leaved tree dominance and from peatlands to forests are two scenarios under global change (Boisvert-Marsh and de Blois, 2021, Mack et al., 2021). The shifts could affect the response of forest biodiversity and function to mining disturbances through changes in plant community structure, given the plant communities of different ecosystems generally differ in tolerance and resilience capacities under stressors (Barbier et al., 2008, Jean et al., 2017). For example, low light intensity below coniferous trees appear to support high cover of bryophyte (Király and Ódor, 2010), while bryophyte cover in deciduous and mixed forest is constrained by the fall of broadleaf litter (Bartels et al., 2018). Furthermore, coniferous canopies buffer environmental variations (e.g., temperature, moisture, irradiance and wind in the understory) in the understory to a greater degree than deciduous forests, as persistent foliage and dense branching patterns; (Barbé et al., 2020, De Jalón et al., 2019). Boreal forests, including coniferous forest and peatlands, store large amounts of carbon (Bradshaw and Warkentin, 2015), while predicted ecosystem shifts and increasing mining activities (Fig. 3.1) could interact and affect ecological services via plant diversity and composition, including the bryophyte layer.

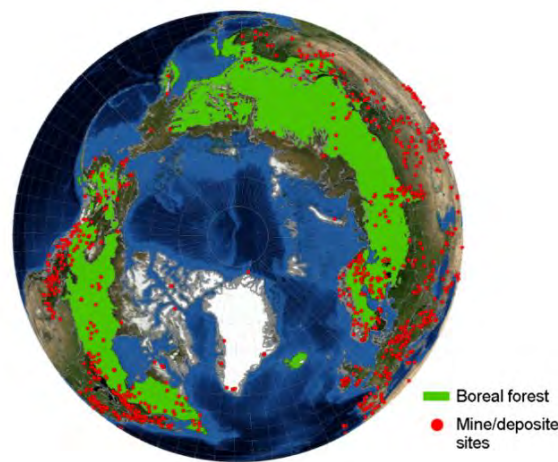


Fig. 3.1. The distribution of mines and deposits of major mineral commodities in global boreal forests (March 9th, 2022). Datasets for mine sites was retrieved from Mineral resources online spatial data, United States Geological Survey (Labay et al., 2017).

Bryophytes are widely distributed globally and particularly contribute ecological services in boreal ecosystems such as carbon and nitrogen cycling, water balance and forest succession (Arróniz-Crespo et al., 2014, Nilsson and Wardle, 2005, Rousk et al., 2013, Xia et al., 2020). Particularly, two most abundant groups, feather mosses and *Sphagnum*, shape ecological function and resilience. Feather mosses can account for as much as 30 % of net primary productivity and contribute over 50 % of total nitrogen sources in boreal coniferous forests (Wardle et al., 2012, Jean et al., 2021). *Sphagnum* is the ecosystem engineer of peatlands (Rydin and Jeglum, 2013), which store around 25 % of global soil carbon and represent important resources to mitigate climate changes (Turetsky et al., 2015). Despite the critical roles bryophytes play in the ecosphere, they lack true roots, vascular systems and thick cuticles and are therefore highly sensitive to the surrounding environment. These characteristics result in bryophytes being widely used as indicators of natural and human disturbances including climate changes and heavy metal deposition (Balabanova et al., 2017, Mahapatra et al., 2019, Printarakul and Meeinkuirt, 2022).

The type and abundance of microhabitats have been shown to be more important than mesohabitats for bryophyte diversity (Cole et al., 2008, Király and Ódor, 2010). Microhabitats for bryophytes are generally defined as the smallest subunit of forest habitat including pieces of deadwood and rocks (Barbé et al., 2020). Microhabitats can protect bryophytes from the effects of global changes and anthropogenic disturbances (Paquette et al., 2016, Scheffers et al., 2014). Usually, microhabitats provide a relatively stable substrate in terms of temperature and moisture to effectively buffer extreme climate events and environmental variations, for example bryophyte composition was more dependent on the characteristics of coarse woody debris than environmental features in boreal coniferous stands (Barbé et al., 2020).

The negative effects of mining on biodiversity and ecological services have been well documented in directly disturbed areas (Maus et al., 2020, Odell et al., 2018, Sonter et al., 2018), but offsite effects that occur in relatively intact landscapes surrounding mine sites (mine offsite landscapes) have received little attention (Raiter et al 2014). The main mechanisms of offsite effects associated with mines are dust, salt, excess nutrients, or other contaminants that move from mines to surrounding ecosystems via air, water or human activities (e.g., vehicle movement, Raiter et al., 2014). An increase in the number of mining projects will lead to the exposure of more landscapes to offsite effects. Furthermore, the disturbance of biodiversity and ecological services associated with mining is also dependent on the mining lifecycle as operating mine sites with more activities (e.g., digging, blasting, transportation) usually have a larger influence on surrounding biodiversity than non-operating sites (Adesipo et al., 2020; Bartels et al., 2019).

The aim of this study is to evaluate whether the effects of mining and predicted ecosystem shifts could be synergistic in mine offsite landscapes, resulting in changes in the bryophyte community at the microhabitat scale. Bryophytes were sampled from various microhabitats (i.e. ground, trees, logs, snags, stumps and rocks) in the boreal

forest of Québec (Canada) inside 1-km intact landscapes surrounding six mine sites (from operating and non-operating stages). The offsite landscapes were grouped into coniferous, deciduous, mixed forest and open canopy ecosystems. We hypothesized that mine sites at the operating stage have more offsite effects on bryophyte community structure (diversity and composition) than those at the non-operating stage (H1) and that ecosystem type affects the presence and magnitude of the offsite effects of mines (H2). The interactions between offsite effects of mines and ecosystem type on bryophytes are affected by microhabitats (H3). We expect a reduced abundance of important functional groups (i.e., feather mosses and sphagna) as a consequence of offsite effects of mines and the predicted shifts in ecosystem type (H4). How bryophyte structure is altered in boreal offsite mine landscapes and potential factors under global changes have not been addressed in previous studies; therefore, our study provides a first look at these industrial impacts.

3.4. Material and methods

3.4.1 Study area

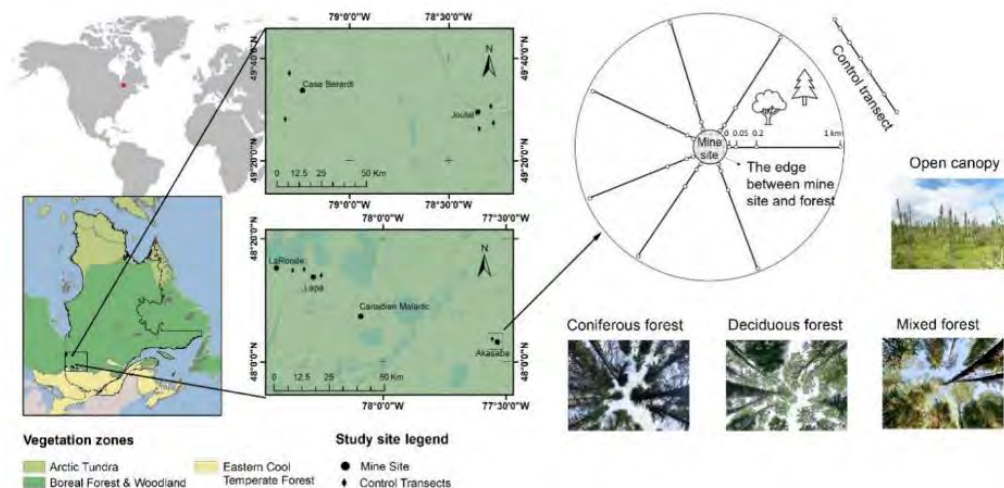


Figure 3.2 Study area. Left, map of Canada and Quebec showing the main vegetation zones. Center, map of study region indicating the position of sampling sites at six gold mine sites and 9 control transects (August 9th, 2021).

The study was conducted in six mine sites in the regions of Abitibi-Témiscamingue and Nord-du-Québec in the province of Québec, Canada (Figure 3.2): Akasaba (48°3'12"N; 77°32'8"W, Agnico Eagle Mines Ltd), Canadian Malartic (48°7'21"N; 78°5'23"W, Canadian Malartic Corp.), Casa Berardi (49°33'43"N; 79°14'8"W, Hecla Québec Inc.), Joutel (49°29'28"N; 78°21'8"W, Agnico Eagle Mines Ltd), Lapa (48°13'45"N; 78°17'1"W, Agnico Eagle Mines Ltd), LaRonde (48°15'14"N; 78°25'59"W, Agnico Eagle Mines Ltd). They differ in size, mining method, life stage and bioclimate domain. Lapa is a small underground mine, while Canadian Malartic is one of the largest open pit mines in North America. Three of them (Casa Berardi, LaRonde and Canadian Malartic) are operating and the other three (Akasaba, Joutel and Lapa) are non-operating. Furthermore, Casa Berardi and Joutel are located in the *Picea mariana*- moss bioclimatic domain, while the other four mine sites are all located in the *Abies balsamea* - *Betula papyrifera* bioclimatic domain. This region is characterized by a cold and humid continental climate with the mean temperature between -14 to -10°C in winter and between 16 to 18°C in summer (Environment Canada, 2021). Mean annual precipitation ranges from 790 to over 1,300 mm (Wang et al., 2013). More detailed information about the six mines is available in Table S2.1.

3.4.2 Fieldwork design and sampling

Fieldwork design

In each mine site, 6-8 transects (the number of transects was selected based on the size of each mine area, see Fig. S1.1) were established perpendicular to each mine periphery through different ecosystem types (See Table S2.2) in 2017-2019. Nine control transects (over three 3 km from the mine sites) were established in undisturbed natural sites, avoiding down wind from the mines, five in the *Picea mariana* – moss bioclimatic domain and four in the *Abies balsamea* – *Betula papyrifera* bioclimatic domain. As the environmental effects of mines (deposition of

dust or heavy metals) are usually felt farther than 1 km from mines (Gillings et al., 2022; Mattielli et al., 2009), the 1 km long transects were designed to be within the potentially influenced areas. In order to avoid other non-mining disturbances (like highways and towns), transects were not established in all cardinal directions at each mine. This is particularly true for two sites, Canadian Malartic and Lapa, as north of Canadian Malartic mine is the town of Malartic and south of the Lapa mine is a main highway, natural gas pipeline and hydro-corridor (See Fig. S1.1).

Seven vegetation plots (3 m in radius) were established along each transect at distances from the edge of the area directly disturbed by mining activities at 0, 0.02, 0.05, 0.1, 0.2, 0.5, 1 km. Three 1-m² quadrats per plot were randomly established to determine the cover of each macroscopic terricolous bryophyte species and different substrates (leaf litter, woody debris, organic matter, water, rock, lichen). The forest stand of each plot was categorized into one of four types based on tree species (diameter at breast height (DBH) ≥ 10 cm) composition in a prism plot with a 2M basal area factor centered on each vegetation plots: coniferous forest (> 70% of coniferous trees, N=173), deciduous forest (>70% deciduous trees, N=47), mixed (coniferous tree stems between 69% and 31%, n=37), and open ecosystems (DBH of all trees below 10 cm, primarily peatlands in our study area, N=107). In total, 371 plots were established, but 364 plots were analyzed because two control plots were destroyed by logging and some missing data.

Bryophyte sampling

The bryophyte community was sampled using a modified floristic habitat sampling technique (Newmaster et al., 2005) in the summer of 2019 and 2020. All bryophytes on each microhabitat (ground, trees, logs, snags, stumps and rocks, classification criteria in Table S2.2) within each vegetation plot were sampled and stored in individual paper bags with plot and microhabitat information. Samples were identified to the species level based on the “Flore des bryophytes du Québec-

Labrador” (Faubert, 2012) at the bryophyte lab of Université du Québec en Abitibi-Témiscamingue. Species Latin names in our database were checked using The Plant List database in the R package Taxon stand version 2.4 (Cayuela et al., 2012). In total, 186 bryophyte species (132 moss species and 54 liverwort species) belonging to 89 genera were identified (species list see Table S2.3) and the feather moss *Pleurozium schreberi* (Willd. ex Brid.) Mitt. and *Sphagnum angustifolium* (Warnst.) C.E.O. Jensen were the dominant species with 8.96 % and 7.09% ground cover, respectively.

Environmental variable sampling

Canopy openness and organic soil depth were measured three times in each plot, and shrub diameter at breast height (DBH, measured 1.3 m from ground level) was measured in the same plot as the tree basal area. Conifer proportion (%) was calculated by the percentage number of individuals of coniferous trees in the prism plot. The live crown ratio of each shrub (vegetation plot) and trees (prism plot) were measured and recorded (Table S2.4). Furthermore, environmental variables were compared among mine sites in Table S2.5.

3.4.3 Statistical analysis

3.4.3.1 Effect of ecosystem type and mining stage on bryophyte community structure

The sum of species occurring in all microhabitats per plot was used as the species richness of the bryophyte community at each plot. This study took the species richness for diversity indices as its high sensitivity to environmental changes (Wilsey and Stirling, 2007; Andersen et al., 2020). Bryophyte species were classified by taxonomy (moss and liverwort), so in each plot (N=364), we obtained the species richness and composition for three groups: total bryophytes, mosses and liverworts.

Species richness. Generalized linear mixed models (GLMMs, “glmmTMB” function from *glmmTMB* package, Brooks et al., 2017) were used to determine if mining stage,

ecosystem type and their interaction influenced total bryophyte, moss and liverwort richness. The richness, referring to the species richness per plot, is the total number of bryophytes/moss/liverwort species identified in each plot. Considering the nested structure of the sampling design, “plot” was nested in “site” as random effects, as microhabitat was the smallest sampling unit. All models were first performed with Poisson error distribution and a log-link function, but when overdispersion was detected, a negative binomial distribution was used in final models. One model (moss richness) showed convergence errors and was corrected by optimization of the model via the Broyden–Fletcher–Goldfarb–Shanno (BFGS) algorithm (Dai, 2002) in the function of the “glmmTMBControl” from the *glmmTMB* package. Significance of the predictors (mining stage, ecosystem type and their interaction) was tested by type II Wald chi-square (χ^2) tests with the “Anova” function in *car* package (Fox et al., 2012). A Tukey post hoc test (“emmeans” function in the *emmeans* package, Lenth et al., 2018) was performed to assess interaction terms (the differences between mining stage in each ecosystem type).

Community composition. Permutational multivariate analysis of variance with 999 permutations (PERMANOVA, “adonis” function from the *vegan* package, Oksanen et al., 2010) were run to assess the interacting effects of ecosystem type and mining stage on bryophyte community composition. In the PERMANOVA analysis, non-significant interaction terms were removed in final analysis. Results were visualized using principal coordinates analysis (PCoA, “PCOA” function in the *vegan* package) with Bray–Curtis dissimilarity matrices. The “envfit” function in *vegan* was then used to fit vectors of environmental parameters (microhabitat number, stand structure and soil properties) onto the ordinations with p-values (>0.05) derived from 999 permutations. When significant interactions between mining stage and ecosystem type were detected, the interaction effects on bryophyte composition shifts were qualified by calculating the Euclidean distances between centroids of PCoA cluster for each mining stage (distances between centroids of operating mines and controls,

non-operating mines and controls as well as operating and non-operating mines, respectively) within each ecosystem type following a changed method by Martineau et al. (2020). The calculated distances were used to determine whether the combined effects of predicted ecosystem shifts and mining stage were synergistic or not (coniferous forest as references, details in Fig. S2.1). In addition, to identify species that characterized the communities of each mining stage, point biserial-correlation coefficient (ϕ) was calculated for indicator species analysis (“multipatt” function in the *indicspecies* package, De Caceres et al., 2016). The value of “stat” (the point-biserial-serial-correlation-coefficient) was used to measure the strength of the co-occurrence pattern observed between a bryophyte species and each ecological state of mining stage. Moreover, generalized linear mixed models were also used to test the differences in 17 environmental variables between mining stages in each ecosystem type (Table S2.6).

3.4.3.2 Role of microhabitats in mediating the effects of ecosystem type and mining stage on bryophyte structure

Bryophyte richness for each microhabitat in each plot was calculated as the sum of all species detected on each microhabitat of a given type in the plot. Composition on each microhabitat type (trees, logs, snags, stumps and rocks) per plot was determined using relative species frequency, where $F_m = n_m / N_t * 100$, F_m is the relative frequency of species m on each microhabitat type; n_m is the number of occurrences of species m in each microhabitat per plot; N_t is the total number of all microhabitat types per plot (ground counted as one individual microhabitat. Considering it is impossible to count the number of ground microhabitat elements, the bryophyte composition on the ground was determined using species raw cover measured in the 1m² quadrats.

The analysis process for richness and composition per microhabitat was similar to the description in 3.4.3.1. However, there are some differences in models for richness associated with the characteristics of each dataset. Considering the lack of certain

microhabitats in some plots, zero-inflated (ZI) models (“zi=~.” used in “glmmTMB” function) were used in all models for bryophyte richness on each microhabitat. The model for bryophyte richness on rocks was analyzed with Poisson error distribution instead of negative binomial distribution because overdispersion was not detected. Furthermore, non-significant interaction terms (ecosystem type * mining stage) were removed from the final models (richness on trees, logs, stumps, snags and rocks).

3.4.3.3 Does a shift in forest composition and mining disturbance affect the ecological roles of bryophytes through abundance of important functional groups?

Structural equation modeling (SEM, “sem” function in *lavaan* package, Rosseel, 2012) was used to obtain a mechanistic understanding of how differences in ecosystem type, mining stage and microhabitats mediated changes the abundance of feather mosses and *Sphagnum*. A good model fit was evaluated using: 1) a Root Mean Square Error of Approximation (RMSEA) equal to or higher than 0.07 (Steiger, 2007) and 95% confidence interval (CI) = 0.054–0.086; 2) Comparative Fit Index (CFI) equal to or higher than 0.96; and 3) Tucker-Lewis Index (TLI) equal to or lower than 0.96 (Hu and Bentler, 1999). All analyses were performed with R 4.0.5 (2021-03-31). Results were visualized with the *ggplot2* package (Wickham, 2011).

3.5. Results

3.5.1 Synergistic effects of ecosystem type and mining stage on bryophyte richness and composition

Bryophyte richness in mine offsite landscapes was affected by the interaction between ecosystem type and mining stage (the results of GLMMs for total bryophyte, moss and liverwort richness, ANOVA type II sums, $P < 0.05$, Table S2.7). Generally, mines present a positive effect on bryophyte richness in coniferous forests and open canopies, but a negative influence in deciduous and mixed forests (the results of

Tukey HSD pairwise comparisons, Fig. 3 A-C). Specifically, moss richness was higher near both operating and non-operating sites than in controls in coniferous forest and open canopy, while it was significantly lower near operating sites than controls in mixed forest (Figure 3.3 B). Furthermore, more liverwort species were also found near operating sites than controls in open canopy, but less liverwort species were observed near both operating and non-operating sites in deciduous forests (Figure 3.3 A).

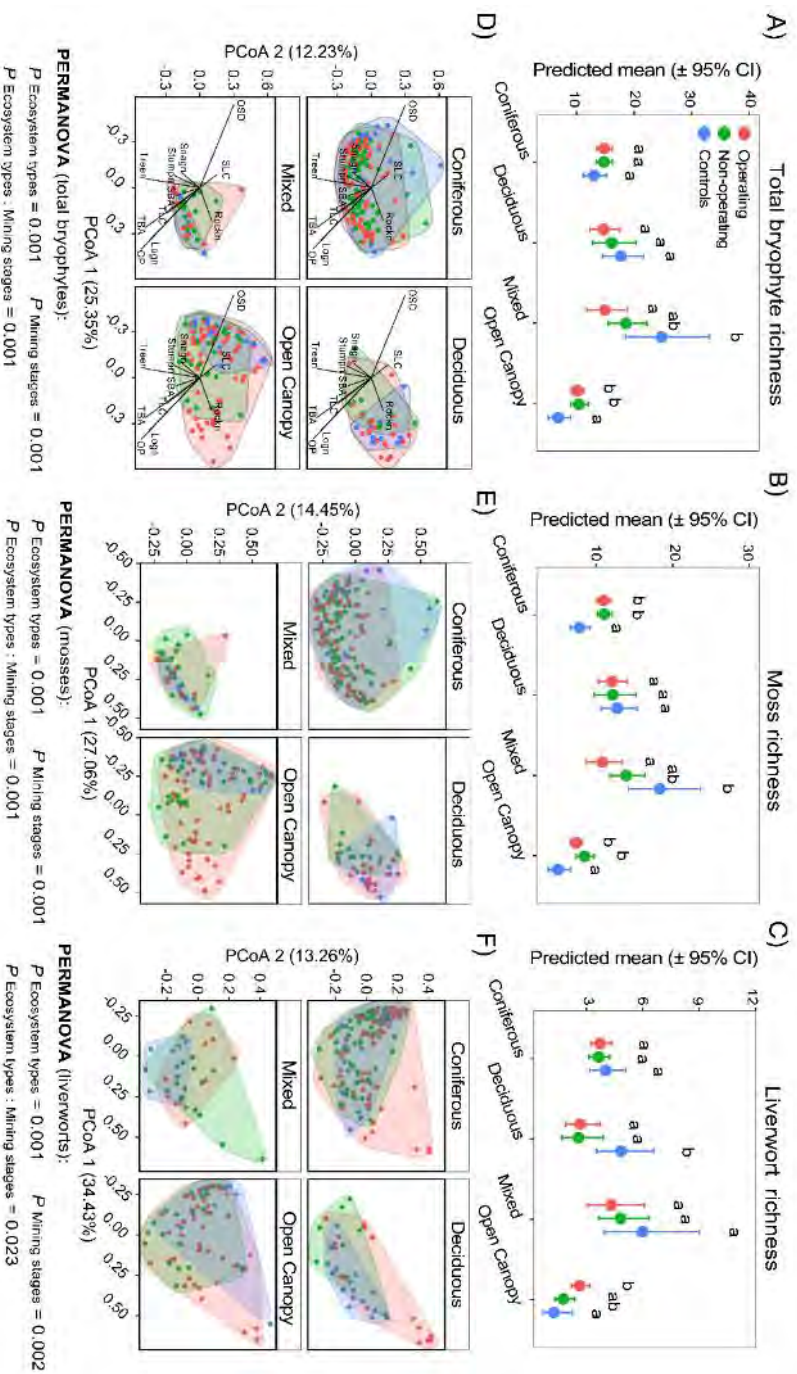


Figure 3.3 Comparison of bryophyte community structure between mining stages for each ecosystem type. A) Total bryophyte richness; B) Moss richness; C) Liverwort richness; D) Total bryophyte composition; E) Moss composition; F) Liverwort composition. Significant effects based on the results of generalized linear mixed model. Different letters denote significant differences ($\alpha \leq 0.05$) between mining stages (plots near operating mines, $N=167$; plots near non-operating sites, $N=136$; plots in mixed forest, $n=37$; plots in open canopy, $N=107$) from generalized linear mixed models with Tukey HSD pairwise comparisons (A, B, C). Principal co-ordinate analysis is used to visualise composition structure and present the results of PERMANOVA based on Bray–Curtis dissimilarity in D, E, F.

Table 3.1. List of indicator species for each mining stage with their point biserial correlation coefficient and p value (only significant species shown, $P < 0.05$).

Species	Abbreviation	Type	stat	P value
Operating (6)				
<i>Calliergonella lindbergii</i> (Mitten) Hedenas	Cal.lind	Moss	0.144	0.03*
<i>Hygroamblystegium varium</i> (Hedwig) Mönkemeyer	Hyg.vari	Moss	0.144	0.018*
<i>Hypnum cupressiforme</i> Hedw.	Hyp.cupr	Moss	0.17	0.01**
<i>Mylia anomala</i> (Hook.) J.J. Engel & Braggins	Myl.anom	Liverwort	0.185	0.002**
<i>Plagiothecium laetum</i> Schimp.	Pla.laet	Moss	0.154	0.01**
<i>Pohlia nutans</i> (Hedw.) Lindb.	Poh.nuta	Moss	0.207	0.002**
Non-operating (2)				
<i>Brachythecium erythrorrhizon</i> Schimp.	Bra.eryt	Moss	0.165	0.006**
<i>Dicranum polysetum</i> Sw.	Dic.poly	Moss	0.175	0.009*
Controls (12)				
<i>Barbilophozia attenuata</i> (Nees) Loeske	Bar.atte	Liverwort	0.17	0.006**
<i>Barbilophozia barbata</i> (Schreb.) Loeske	Bar.barb	Liverwort	0.14	0.017*
<i>Brachythecium reflexum</i> (Starke) Schimp.	Bra.refl	Moss	0.138	0.036*
<i>Breidleria pratensis</i> (Koch ex Spruce) Loeske	Bre.prat	Moss	0.125	0.03*
<i>Mnium spinulosum</i> Bruch & Schimp.	Min.spin	Moss	0.163	0.01**
<i>Mylia taylorii</i> (Hook.) Gray	Myl.tayl	Liverwort	0.147	0.01**
<i>Plagiomnium cuspidatum</i> (Hedw.) T.J. Kop.	Pla.cusp	Moss	0.231	0.002**
<i>Scapania mucronata</i> H. Buch	Sca.mucr	Liverwort	0.141	0.024*
<i>Sphagnum angustifolium</i> (Warnst.) C.E.O. Jensen	Sph.angu	Moss	0.235	0.001***
<i>Sphagnum capillifolium</i> (Ehrh.) Hedw.	Sph.capi	Moss	0.151	0.024*
<i>Sphagnum magellanicum</i> Brid.	Sph.mage	Moss	0.185	0.005**
<i>Sphagnum quinquefarium</i> (Lindb.) Warnst.	Sph.squa	Moss	0.142	0.014*

Notes: *significance at 0.05 level, ** significance at 0.01 level and *** significance at 0.001 level.

Similarly, bryophyte community composition near mine sites was also affected by the interaction between ecosystem type and mining stage (PERMANOVA results in Figure 3.3 D-F and Table S2.8). Overall, ecosystem type (PERMANOVA, $8.97\% < R^2 < 13.42\%$, Table S2.8) explained more differences in richness of total bryophytes, moss and liverwort species than mining stage (PERMANOVA, $1.42\% < R^2 < 1.58\%$, Table S2.8) and the interaction ecosystem type - mining stage interaction (PERMANOVA, $2.43\% < R^2 < 3.1\%$, Table S2.9). Specifically, there were greater differences in bryophyte composition among mining stages in deciduous and mixed forests, and open canopy (pairwise PERMANOVA results for total bryophytes, mosses and liverworts, $3\% < R^2 < 11\%$, Table S2.9) than in coniferous forest (pairwise PERMANOVA, $R^2 < 3$, Table S2.9). Furthermore, liverwort composition shifts present more synergistic effects (Table S2.9). The Euclidean distances between centroids of mining stages of liverworts in deciduous and mixed forests almost were greater than those in coniferous forest, while the synergistic effects were only detected in deciduous forests for mosses (Table S2.10). Also, the results were supported in PCoA of liverworts where some plots near operating mines are in the top right corner far from the plots near non-operating sites in mixed and deciduous forests (Figure 3.3F), while in moss composition, differences between operating and non-operating sites were visible only in open canopy (Figure 3.3E).

Forest structure, the number of individual microhabitat types, and organic soil depth were all correlated with total bryophyte species composition (Figure 3.3D). Canopy openness (OP, $R^2=0.45$, $p=0.001$, Table S2.11), organic soil depth (OSD, $R^2=0.40$, $p=0.001$, Table S2.11) and number of logs (logn, $R^2=0.33$, $p=0.001$, Table S2.11) appeared three significant environmental gradients structuring the community. OSD was more related with bryophyte composition in open canopy, while OP, tree basal area (TBA) and logn was associated with the composition in coniferous and mixed forests (Figure 3.3D).

Six indicator species were identified near mine operating mine sites (Table 3.1, five moss species and one liverwort species), two indicator species (both moss species) were identified for non-operating sites and twelve indicator species (eight moss species and four liverwort species) for controls. Almost all indicators for mining sites (operating and non-operating) were moss species and *Pohlia nutans* (Hedw.) Lindb. (stat=0.207, $p = 0.002$) and *Dicranum polysetum* Sw. (stat=0.175, $p = 0.009$) were the best indicators for operating and non-operating sites, respectively, while *Sphagnum angustifolium* (Warnst.) C.E.O. (stat=0.235, $p = 0.001$, Table 3.1) was the best indicator for control sites.

3.5.2 Role of microhabitats in mediating bryophyte response to ecosystem type and mining stage

No statistically significant interactions were found for bryophyte richness on all microhabitats except ground (results of GLMMs, ANOVA type II sums, Table S2.12). Ecosystem type influenced bryophyte richness in most microhabitats (except stumps) with generally, more species in deciduous and mixed forest than that in coniferous and open canopy (the results of Tukey HSD pairwise comparisons, Fig. 3.4 A-F). In contrast to richness, bryophyte composition was affected by the interaction between mining stage and ecosystem type on all microhabitats except rocks (PERMANOVA results in Fig. 3.3 G-L and Table S2.13). Generally, the Euclidean distances between centroids of mining stage were greater in deciduous, mixed forest and open canopy ecosystems than those in coniferous forest for all microhabitat although some opposite patterns were detected (e.g., Euclidean distances between centroids of operating and non-operating stages in mixed forest were lower than that in coniferous forest for total bryophytes on ground, trees and logs, Table S3.13). The interaction explained a greater proportion of differences in bryophytes species composition on snags (PERMANOVA, $R^2 = 7.8\%$, Table S2.13) than on other microhabitats (PERMANOVA, $R^2 < 4.0\%$, Table S2.13). Bryophyte composition in coniferous

forest was only different between mining stages on the ground, while in deciduous forest and open canopy, differences between mining stages were found on the ground, trees, logs and stumps (pairwise PERMANOVA, Table S2.14). Bryophyte composition in mixed forest was only affected by mining stages on stumps (pairwise PERMANOVA, Table S2.14).

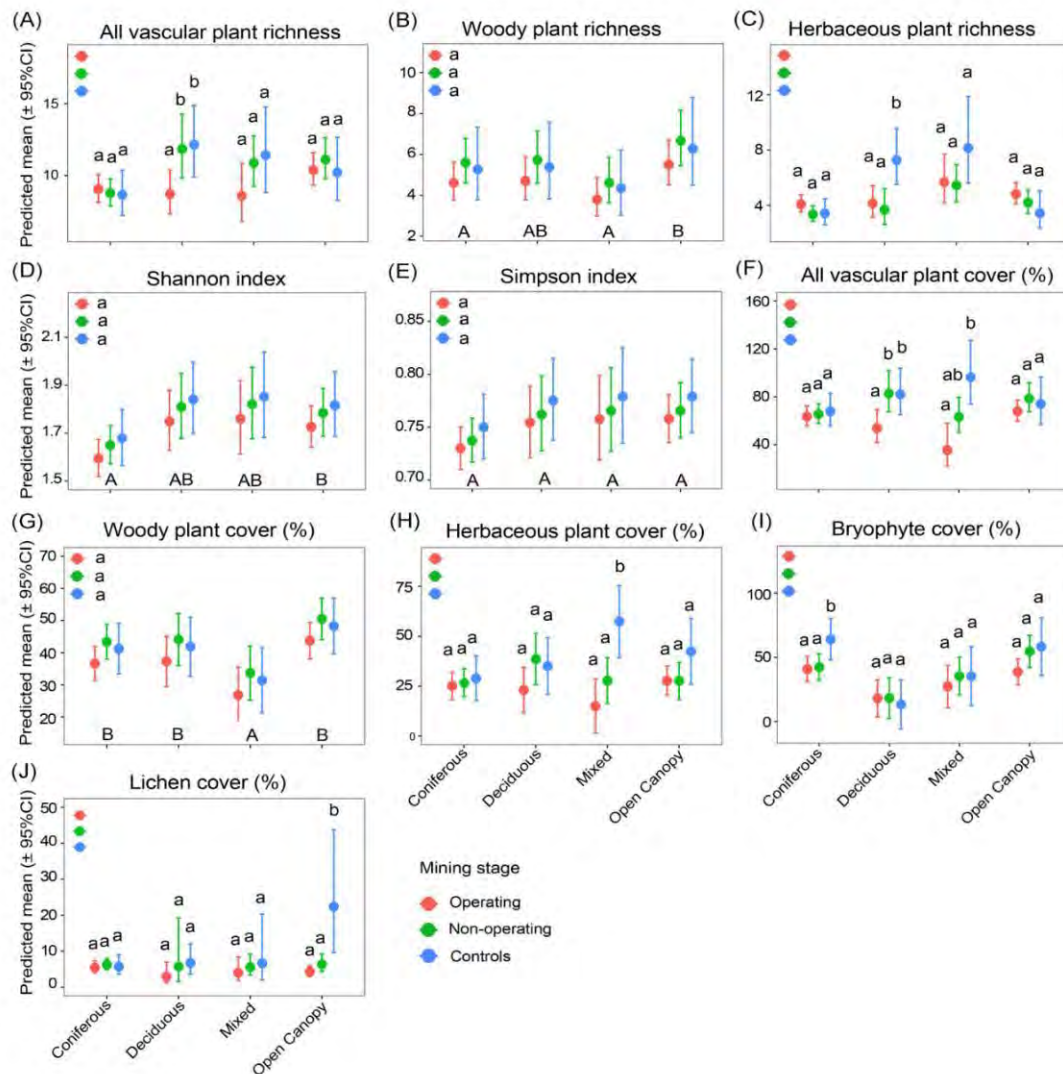


Figure 3.4 Comparison of the richness and composition of bryophytes on each microhabitat between mining stages for each ecosystem type. A) Total bryophyte richness on ground; B) Total bryophyte richness on trees; C) Total bryophyte richness

on logs; D) Total bryophyte richness on snags; E) Total bryophyte richness on stumps; F) Total bryophyte richness on rocks; G) Total bryophyte composition on ground; H) Total bryophyte composition on trees; I) Total bryophyte composition on logs; J) Total bryophyte composition on snags; K) Total bryophyte composition on stumps; L) Total bryophyte composition on rocks. Principal co-ordinates analysis and present the results of PERMANOVA using the Bray–Curtis distance in G, H, I, J, K, L.

3.5.3 Effects of ecosystem type and mining stage on the abundance of major bryophyte functional groups

The SEM explained 21.3% and 22% of the variation in the ground cover of feather mosses and sphagna, respectively (Figure 3.5). Deciduous (β (standardized coefficient) = -0.212) and mixed (β = -0.113) forests were negatively correlated with feathermoss ground cover, with coniferous forest as reference (Figure 3.5A). Operating mine sites (β = -0.183) also had a negative effect on surrounding feathermoss cover, while the numbers of trees (β = 0.234), snags (β = 0.185) and stumps (β = 0.126) were positively correlated with feathermoss cover but the number of logs (β = -0.148) was negatively correlated with feather moss cover (Figure 3.5A).

Similarly, compared to coniferous forest, *Sphagnum* cover was reduced in deciduous (β = -0.125) and mixed (β = -0.068, Figure 3.5B) forests. However, it was increased in open canopy (β = 0.288, Figure 3.5B). Both operating and non-operating sites had negative effects on *Sphagnum* cover in the surrounding landscape. Only log number (β = -0.190, Figure 3.5B) was associated with the ground cover of *Sphagnum* among microhabitat types. Open canopy and operating sites were the most parsimonious explanatory variables for *Sphagnum* cover near mine sites. Therefore, ecosystem type and microhabitats were more important than mining stages in determining feather moss cover near mine sites, while mining stages were the most important variable influencing *Sphagnum* cover in the study.

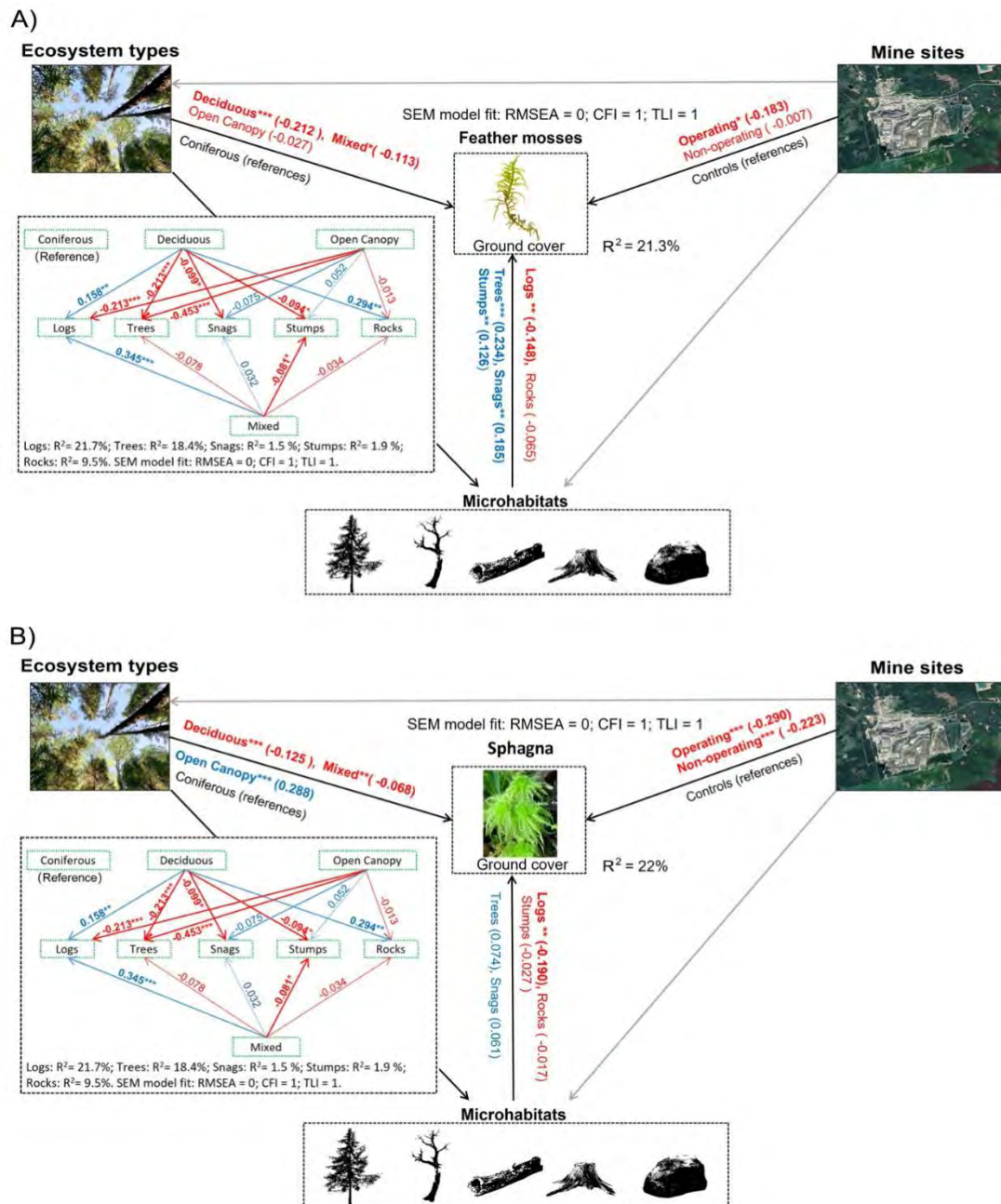


Figure 3.5 Structural equation models (SEM) of ecosystem type, mining stage and microhabitats as predictors of the ground cover of important functional bryophyte groups (i.e., feather mosses and sphagna). A) Feather moss ground cover; B) *Sphagnum* ground cover. Red text represents negative effects, blue text represents positive effects and significant parameters (≤ 0.05) are indicated in bold. The associations between ecosystem type and the number of individual microhabitats were evaluated in a separate structural equation model because of poor fit when all

elements were included in one model. Overall fit of piecewise SEM was evaluated using RMSEA (root mean square error of approximation), CFI (comparative fit index) and TLI (Tucker-Lewis index).

3.6 Discussion

3.6.1 Mine offsite effects and their synergistic interaction with ecosystem type on bryophytes

Offsite effects of mine sites on bryophytes were confirmed based on the differences in richness and composition of bryophyte communities in 1km radius landscapes near mine sites (operating or non-operating) compared to controls. The results extend offsite effects of mines from vascular plants (Boisvert et al., 2021; Chen et al., 2017; Dyer et al., 2001) to cryptogamic community structure. At the same time, operating mines had more effects on bryophyte structure (richness and composition) than non-operating ones (e.g., total bryophyte richness was affected by operating mines in mixed forest but not by non-operating sites in Fig. 3.3A). Compared with non-operating sites, more mining associated activities (e.g., blasting, digging, and transporting) occur in operating sites, which lead to more disturbances (e.g., atmospheric particulate matters and heavy metals) in offsite landscapes (Betancourt et al., 2005; Corriveau et al., 2011; Wang and Mulligan, 2006) and consequently bryophytes were more affected, confirming H1.

The presence and magnitude of the offsite effects is largely dependent on ecosystem type based on the significant interaction effects, which confirms H2. Offsite effects of mines reduced bryophyte richness in deciduous and mixed forest sites, however, the opposite pattern was observed in coniferous forest and open canopy sites. Combined with the microhabitat results (Fig. 3.4), we found that the increase in species richness mainly occurred on the ground in coniferous forest and in open canopy ecosystems. On the one hand, changes in species interactions may contribute to the positive impact of mine offsite effects. Reduced ground cover of “large” dominating moss species (i.e., feather mosses and sphagna Fig. 3.5) caused by the offsite effects in

coniferous and open canopy ecosystems might lead to less competition in disturbed habitats and provide opportunities for smaller bryophytes, such as *Pohlia nutans*, *Plagiothecium laetum* Schimp., *Hypnum cupressiforme* Hedw. and *Hygroamblystegium varium* (Hedw.) Mönk. to establish (Table 3.1). This finding is consistent with an earlier result where disturbed areas with less competitive species supported more bryophyte species (Zielińska et al., 2016). However, this mechanism did not apply to mixed and deciduous forests, possibly because larger bryophytes are not a limiting factor for other bryophyte species in these environments (Jean et al., 2020; Oechel and Van Cleve, 1986). The high ground cover of broadleaf litter (up to over 70 % near operating and non-operating sites, see Table S2.5) could mainly contribute to reduced bryophyte richness (Márialigeti et al., 2009; Sætre et al., 1997) near mined sites in deciduous and mixed forests. Low decomposition rates due to low activity of microorganisms in polluted soil near mine sites could lead to an increase in the ground cover and thickness of deciduous litter (Freedman and Hutchinson, 1980; Horodecki and Jagodziński, 2017; Strojan, 1978). On the other hand, the differences in resistance of each ecosystem type can be also attributed to the canopy structure and environmental characteristics. Conifers have higher canopy cover, leaf area index, and a more persistent foliage than deciduous trees, which may intercept more dust, heavy metals, and other pollutant emissions generated by mining (Augusto et al., 2002; Barbier et al., 2008; Nguyen et al., 2015). Furthermore, deciduous trees can translocate high amounts of heavy metals into the foliage which can then result in more heavy metal accumulation in topsoil than under coniferous trees (Van Nevel et al., 2011).

Bryophyte indicator species were identified for offsite landscapes at different mining stages. *Calliergonella lindbergii* (Mitt.) Hedenäs, *Hygroamblystegium varium*, *Hypnum cupressiforme*, *Mylia anomala* (Hook.) Gray, *Plagiothecium laetum* and *Pohlia nutans* were indicators of operating mines, and they are tolerant or pioneer species (Barrett and Watmough, 2015; Petschinger et al., 2021; Rydgren et al., 2004;

Salemaa et al., 2001). In particular, *P. nutans* had the highest indicator value in operating sites and has been reported as one of most tolerant moss species growing in the immediate vicinity of mines and associated disturbed areas (Gignac and Beckett, 1986; Helmisaari et al., 1995; Salemaa et al., 2001). More species (13 species) occurred more abundantly in undisturbed landscapes (controls), suggesting that they have low tolerance to mining disturbances and high indicator values for intact landscapes. Four *Sphagnum* species including *Sphagnum angustifolium*, which had the highest indicator value, suggests that the *Sphagnum* group could be particularly sensitive to offsite effects, which is consistent with previous results that *Sphagnum* was a key indicator of mining effects on vegetation (Mullan-Boudreau et al., 2017; Spratt and Wieder, 1988). Interestingly, only two indicator species were detected for non-operating sites, which suggests that bryophyte species composition near non-operating sites was generally similar with that either near operating mines or in controls and few specific species were associated with this condition where neither tolerant nor sensitive groups were particularly supported.

Therefore, ecosystem types and mining stages are two drivers for offsite effects of mines on bryophyte richness and composition. The combined effects of predicted ecosystem shift from coniferous to deciduous forest dominance and mining on bryophytes could be synergistic in the offsite boreal landscapes. Meanwhile, indicators for the offsite effects have potential value for future ecological assessment in mining projects

3.6.2 Microhabitats mitigated the synergistic effects of ecosystem type and mining on bryophyte structures

Whether microhabitats can mitigate effects from predicted global change on bryophytes was still unclear, but our results did confirm their potential buffer roles in the scenario of predicted ecosystem shifts and increases of mining projects (H3). Some types of microhabitats mitigated the offsite effects of mines and their

synergistic effects with predicted ecosystem shifts, which indicates that they could play the role of mining-refugia in offsite landscapes. The possible explanation is that microhabitats, and especially coarse woody debris (logs, snags and stumps), are specialised substrates which retain moisture, offering a more stable microclimate environment than soil (Haughian and Frego, 2017; Jönsson and Jonsson, 2007). Furthermore, offsite effects of mines on bryophyte richness were only found on the ground and rocks (Fig. 3.4A, F). These results indicate that bryophytes growing in microhabitats located near or on the forest ground are more vulnerable to offsite effects of mines than those colonizing vertical microhabitats (i.e., trees, stumps, snags). Microhabitats in the forest ground layer could accumulate higher amounts of dust deposition than standing microhabitats considering that dust deposited on tree and shrub leaf surfaces will finally deposit on the forest ground through the action of wind, rainwater and defoliation (Sase et al., 2012). Meanwhile, loss of canopy cover by the edge affects mainly communities on the ground, where the level of incident light, soil moisture and nutrient availability can be changed by canopy openness, while non-ground microhabitats having vertical dimensions might lead to less exposure probabilities to the changes in environmental variables (e.g., lightness, treefalls and soil moisture).

In contrast to richness, microhabitats generally did not mitigate the interaction and synergistic effects on bryophyte composition, indicating that community composition is generally much more vulnerable to environmental changes than species richness, a response that has been found after other disturbances (Oldén et al., 2014). Species richness is an emergent property of ecosystems and it is relatively constant after disturbances if local compensatory colonisations occurred (Legendre et al., 2005; Parody et al., 2001). Nevertheless, our results firstly confirmed the buffer roles of microhabitats for bryophytes in mine offsite landscapes, although the interaction and synergistic effects could still be detectable in community composition.

3.6.3 The offsite effects of mine sites on important functional bryophyte groups

Predicted shifts in ecosystem composition and mining could both limit the abundance of feather mosses and sphagna in mine offsite landscapes, which supports H4. Lower feather moss and *Sphagnum* abundances in deciduous and mixed forests than coniferous forest and open canopy ecosystems can be attributed to their strong environmental preferences. Feather mosses occur in closed-canopy, well-drained coniferous forests and sphagna occur in the more open-canopy, poorly-drained coniferous forests and peatlands (Bisbee et al., 2001; Peckham et al., 2009). Furthermore, compared with feather moss abundance, sphagna was more effected by offsite effects of mines. It indicates that *Sphagnum* abundance should be highly sensitive to mining activities, which might be attributed to the fact that *Sphagnum* growth is more susceptible to water and peat depth than feather mosses (Bergeron et al., 2009; Bisbee et al., 2001) since the landscapes near mine sites are characterized by low water table and depletion of organic soil content (Glina et al., 2019). Feather mosses occurred in a wider range of microhabitats than sphagna. The dominating habitat of sphagna is the ground while feather mosses can colonize various microhabitats (e.g. logs and rocks) which can serve as shelter from environmental changes (e.g. water content and temperature) by providing microclimatic buffering for species growing on them (Dražina et al., 2016; Haughian, 2018). The results reveal predicted ecosystem shifts and increasing mining activities might affect ecological services of boreal forests through reducing abundances of important functional groups.

Although the offsite effects of mines on bryophyte diversity, community structure and composition and drivers (ecosystem types, mining stages and microhabitats) were confirmed, further research is required to identify the source of the stress and that other studies (e.g., measuring heavy metal concentrations in the soil, installing dust

samplers to see the amounts of dust and level of contamination of the dust coming from the mines) will be needed in order to do that.

3.7 Conclusion

Our study provides the evidence that mine offsite landscapes could be a growing threat to boreal biodiversity of sensitive groups such as bryophytes as offsite effects extended mining effects beyond the areas originally degraded. Overall, our results revealed that ecosystem type and mining stage are the predictor of bryophyte community structure in mine offsite landscapes and their combined effects are generally interacted, although microhabitats have the potential to mitigate the interaction effects in maintaining bryophyte richness. Bryophyte communities in coniferous forest seemed more resistant to the offsite effects than other ecosystem types. Reduced abundance of important functional groups in offsite landscapes could be a challenge for ecological services in boreal forests, especially in coniferous forest and open canopy ecosystems. With global changes, the predicted shifts in ecosystem type might exacerbate the effects of mining on biodiversity and ecological services in mine offsite landscapes. Cumulative areas exposed to mine offsite effects could be large at global scales, and therefore mine offsite landscapes should be considered in future impact evaluations and landscape management. A certain level of coarse woody debris storage near mine sites could be an effective conservation strategy. Our findings further encourage future studies to evaluate how these changes in bryophyte structure affect ecological services (e.g., carbon and nitrogen cycle) of the boreal forest and the microhabitat refugia under the global change.

3.8 Acknowledgements

This research was financially supported by the NSERC-UQAT Industrial Chair in Northern Biodiversity in a mining context. We thank Enrique Hernández-Rodríguez

and Carlos Cerrejon Lozano for their assistance in creating the study area map. We deeply thank Julie Arseneault and Mélanie Jean for the establishment of the plots. We also thank Julie Arseneault, Mélanie Jean, Elizabeth Riendeau, Andréane Garant, Cyrielle Bigonneau, Simon Bégin, Matilde Offroy, Pamela Parisien, Jeffrey Raymond, Juliette Larrivée, Marie-Anyse Dubuc, Emmanuelle Richard, Maxime Thomas, Terresa Kim, Xavier Saint-Amant, Jolianne Garand and Natalie Vuong for their valuable assistance in the fieldwork. We also thank Josée Noël, Lucienne Anctil, Stéphanie Lafrenière and other people at the mines of Canadian Malartic, Casa Berardi, LaRonde, Lapa, Akasaba and Joutel for their help with access to the mine sites.

CHAPTER IV

OUT OF SITE, OUT OF MIND: CHANGES IN FEATHER MOSS
PHYLLOSPHERE MICROBIOTA IN MINE OFFSITE BOREAL LANDSCAPES

HORS DU SITE, HORS DE L'ESPRIT : CHANGEMENTS DANS LE
MICROBIOTE DE LA PHYLLOSPHÈRE DES MOUSSES À PLUMES DANS LES
PAYSAGES BORÉAUX HORS SITE DES MINES

Ce chapitre est en préparation pour *Environmental Microbiology*

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Highlights:

- Impact of mines on microbial communities of the feather moss phyllosphere were studied in mine offsite boreal landscapes.
- Mining lifecycle and ecosystem type both affected the diversity, community structure and composition of feather moss phyllosphere microbiota.
- Ecosystem type did not interact with the offsite effects of the mine lifecycle on the feather moss phyllosphere microbiota.
- The majority of effects occurred within 0.2 km surrounding mine sites, especially near operating mines.

4.1 Résumé

Les interactions plantes-microorganismes, incluant les associations entre les mousses à plumes des forêts boréales et le microbiote de leur phyllosphère, contribuent à la biodiversité et aux services écologiques dans les biomes forestiers. L'augmentation de l'exploitation minière pour répondre aux besoins en minéraux constitue une menace potentielle pour la biodiversité. Cependant, l'impact de l'exploitation minière sur la biodiversité dans les paysages hors sites des mines est encore peu connu, en particulier pour les microbiomes. Nous avons échantillonné *Pleurozium schreberi* (Brid.) Mitt. dans des sites témoins et le long de transects d'un kilomètre autour de six sites miniers à différents stades d'exploitation (en exploitation ou non) qui traversent quatre types d'écosystèmes (conifères, feuillus, forêts mixtes et canopée ouverte) dans la forêt boréale canadienne. Les effets hors site des mines sur le microbiote de la phyllosphère ont été détectés à l'aide de l'ADN microbien récupéré à partir de la phyllosphère de *P. schreberi*. Le stade d'exploitation et le type d'écosystème ont tous deux affecté la diversité alpha et bêta ainsi que l'abondance relative de certains taxons, généralement sans interaction entre les deux facteurs. Les taxons microbiens individuels ont été plus affectés par les effets hors site que la diversité au niveau de la

communauté. Les effets les plus forts se sont produits à moins de 0,2 km des mines en exploitation. Ces résultats indiquent que les sites miniers structurent les communautés microbiennes de la phyllosphère de la mousse à plumes dans les paysages boréaux et que ces effets sont modulés par le stade d'exploitation et le type d'écosystème. Les résultats impliquent également que les changements prédits de la dominance des conifères vers les feuillus dans la forêt boréale dans le contexte des changements globaux ne modifieraient pas l'impact des mines sur le microbiote de la phyllosphère dans les paysages hors site. D'autres études seront nécessaires pour évaluer comment ces effets hors site sur la diversité de la phyllosphère et la structure des communautés affectent le fonctionnement des écosystèmes boréaux et pour développer des stratégies d'atténuation appropriées.

Mots-clés : phyllosphère, écologie du paysage, indicateurs microbiens, bryophytes, taïga forestière

4.2 Abstract

Plant-microbe interactions, including associations between feather mosses from boreal forests and their phyllosphere microbiota, contribute to biodiversity and ecological services in forest biomes. Increases in mining for mineral requirements are potential threats to biodiversity. However, knowledge of impacts of mining on biodiversity in mine offsite landscapes is still poor, especially for microbiomes. We sampled *Pleurozium schreberi* (Brid.) Mitt. in control sites and within 1-km radial landscapes (away from the edge of each mine site) containing four ecosystem types of the Canadian boreal forest (i.e., coniferous, deciduous, mixed forests and open canopy peatlands) in six mines at different stages of the mine lifecycle. Offsite effects of mines on the phyllosphere microbiota were detected using microbial DNA recovered from the phyllosphere of *P. schreberi*. Mining stage and ecosystem type both affected alpha and beta diversity as well as the relative abundance of some taxa,

generally without interactions between the two factors. Individual microbial taxa were more affected by offsite effects than community-level diversity. The strongest effects occurred within 0.2 km from operating mines. These results indicate that mine sites are structuring microbial communities of the feather moss phyllosphere in boreal landscapes, and that these effects are driven by mining stage and ecosystem type. The results also imply that predicted shifts from coniferous to deciduous dominance in boreal forest under global changes would not modify the impact of mines on the phyllosphere microbiota in offsite landscapes. Further studies are needed to assess how these offsite effects on the phyllosphere diversity and community structure affect the functioning of boreal ecosystems and to develop appropriate mitigation strategies.

Keywords: phyllosphere, landscape ecology, microbial indicators, bryophytes, taiga forest

4.3 Introduction

The phyllosphere — the aerial surfaces of plants — represents a widespread and diverse habitat for various groups of microorganisms, such as bacteria and fungi (Bashir et al., 2021; Perreault and Laforest-Lapointe, 2021; Vorholt, 2012). Organisms in the phyllosphere not only promote host plant fitness and nutrient acquisition, but also play important roles in global biodiversity and biogeochemical cycles (Peñuelas and Terradas, 2014; Perreault and Laforest-Lapointe, 2021; Stone et al., 2018). In boreal forests, where nutrient availability is limited, the moss phyllosphere microbiota plays a major role in carbon and nitrogen cycling, as exemplified by the feather mosses-cyanobacteria (Rousk et al., 2013) and *Sphagnum*–methanotrophs (Putkinen et al., 2012) associations. *Pleurozium schreberi* (one of the most widespread boreal feather mosses) alone, through the activity of its phyllosphere nitrogen-fixing bacteria (e.g. Cyanobacteria), was shown to fix 1.5 to 2.0 kg nitrogen ha⁻¹ yr⁻¹ in boreal forests (DeLuca et al., 2002).

Increasing anthropogenic activities such as mining are main threats to biodiversity and ecosystem services in the Anthropocene (Chester et al., 2019; Corlett, 2015). Currently, the area occupied by mines extends to about 57,300 km² across the globe (Maus et al., 2020). Areas directly disturbed by mining (e.g., open pits, tailings and waste rock) have been extensively studied (Blackmore et al., 2018; Chung et al., 2019; Feng et al., 2019; Gagnon et al., 2020; Thavamani et al., 2017). Meanwhile, ecological degradation in mine offsite landscapes (i.e., relatively intact ecosystems surrounding mines) remains poorly characterized. Effects occurring in offsite landscapes are identified as offsite effects (Raiter et al., 2014) and are thought to be caused by the spread of contaminants (e.g., dust, salt, heavy metals and excess nutrients) through the air, water or human activities (e.g., vehicle movement). There is increasing evidence that these “out of site” landscapes impacts should not stay out of mind as both plant and animal species are affected by distance to mines (Boisvert et al., 2021; Duarte et al., 2015; Watkinson et al., 2021; Wu et al., 2021). However, to the best of our knowledge, changes in microbial communities, in particular those inhabiting plants in mine offsite landscapes, have not been studied. Understanding the effects of mines on microbial communities of the phyllosphere is necessary to fully understand the impacts of mining on ecosystem services.

The phyllosphere composition of feather moss (i.e., *P. schreberi*) has been described in previous studies. Proteobacteria, Acidobacteriota, Bacteroidota and Actinobacteriota were the most abundant bacterial phyla (Cutler et al., 2017; Holland-Moritz et al., 2021; Holland-Moritz et al., 2018; Jean et al., 2020a; Rodríguez-Rodríguez et al., 2022), while Ascomycota and Basidiomycota were dominant fungal phyla (Cutler et al., 2017). These studies found that forest type was the main factor affecting the community composition of feather mosses through the different environmental conditions shaped by canopy structure such as temperature, leaf litter input, pH, and availability of nitrogen, moisture, and light (Holland-Moritz et al., 2021; Jean et al., 2020a; Rodríguez-Rodríguez et al., 2022). Proteobacteria,

Acidobacteriota and WPS-2 (*Candidatus* phylum Eremiobacterota; Ji et al., 2017) were found at higher relative abundances in feather moss phyllosphere in coniferous than in deciduous forests, while deciduous forests supported higher relative abundances of Bacterioidota and Cyanobacteria (Jean et al., 2020a; Rodríguez-Rodríguez et al., 2022).

Deciduous, mixed, and coniferous forests, and open canopy areas (consisting mainly of peatlands) are four dominant ecosystem types in boreal landscapes. The resistance (i.e. the degree to which community composition remains unchanged in the face of a disturbance, Allison and Martiny, 2008) of the ecosystem types to environmental changes is conditioned by their structure and physicochemical characteristics. Generally, coniferous trees have higher canopy cover, leaf area index, less light transmission, and a more persistent foliage than deciduous trees, which leads to a relatively high resistance to environmental changes in coniferous forest and a relatively stable understory environment (Augusto et al., 2002; Eugster et al., 2000; Sánchez-Pinillos et al., 2019). Mixed forests usually have a more heterogeneous understory environment than monospecific forests (Chávez and Macdonald, 2012) and its resistance depends on the tree species composition of the mixture (Jactel et al., 2017). Although open-canopy peatlands receive more light, precipitation, solar radiation and atmospheric depositions than closed-canopy forests, they have a relatively high resistance to environmental changes (Łuców et al., 2020) because of humid microclimatic conditions generated by the accumulated peat (Augusto et al., 2002). From coniferous needle-leaved to deciduous broad-leaved tree dominance is the main ecosystem shift scenario in boreal forests in response to global changes (Boisvert-Marsh and de Blois, 2021; Mack et al., 2021; van der Velde et al., 2021). In a previous study, we have shown that this predicted shift could exacerbate mining effects on bryophyte community structure (Yin et al., 2022). Therefore, we predict that the effects of mines on the feather moss phyllosphere could also be mediated by the predicted shifts in ecosystem types.

Generally, the severity of mining disturbance varies with mine lifecycle from the establishment, operation to closure, and rehabilitation. The mine lifecycle might therefore also be a factor affecting the offsite effects. Emissions occur at each stage of the lifecycle but are especially abundant during operational activities where more pollutants (e.g., heavy metals, dust and acid mine drainage) are created and released into the surrounding environment (Punia, 2021). In contrast, the main risks in non-operating sites (i.e., establishment, closure and rehabilitation) are surface vegetation and soil clearing and residual pollution. Results from previous studies have indicated that offsite effects of mines were more obvious near operating mines than near non-operating sites in the community structure of understory vascular plants and bryophytes (Boisvert et al., 2021, Yin et al., 2022). Therefore, we expect to find the same pattern in the feather moss phyllosphere microbiota.

The objective of this study was to identify and characterize the impacts of mining on the feather moss phyllosphere microbiota in mine offsite boreal landscapes. To address the objective, we asked the following specific questions: Q1) Is there an offsite effect of mining on feather moss phyllosphere microbiota? Q2) What is the main driver influencing the phyllosphere microbiota in mine offsite landscapes: ecosystem type, mining stage or their interaction? Q3) At what distance can we detect the offsite effects of mines on the phyllosphere microbiota? To answer these questions, six mine sites at different stages of the mining lifecycle (grouped into operating mines and non-operating sites) were selected in the boreal forest of Quebec, Canada (Fig.4.1) and 1-km offsite landscapes around each mine site were used as study sites where *P. schreberi* shoots were collected. Plots in mine offsite landscapes were classified in four ecosystem types (i.e., coniferous, deciduous, mixed forests and open canopy areas). We expect that our results will help understand the impacts of mine sites on the spatial patterns of microbial communities in the moss phyllosphere and contribute to the identification of potential indicators of offsite effects of mines.

4.4 Materials and methods

4.4.1 Study sites and design

To conduct the study, six mine sites were selected in the Abitibi-Témiscamingue and Nord-du-Québec regions of the province of Québec, Canada (Fig. 4.1), including three operating mines (Casa Berardi, LaRonde and Canadian Malartic mine) and three non-operating sites (Akasaba, Joutel and Lapa mine sites). More detailed information about the six selected mines is available in Table S1.1. Using a stratified selection process, 6-8 transects (the number of transects was selected based on the size of each mine area, see Fig. S3.1) were established perpendicular to each mine periphery. Four plots were established along each transect with distance from the edge of intact forest at 0, 0.05, 0.2, and 1 km. Furthermore, six control transects with 7-8 plots per transect (44 plots in total) near the six mine sites were established in natural forests. The ecosystem of each plot was categorized into one of four types based on tree species (diameter at breast height (DBH) ≥ 10 cm) composition: coniferous forest ($> 70\%$ of coniferous tree stems), deciduous forest ($> 70\%$ deciduous tree stems), mixed (coniferous tree stems between 69% and 31%) and open canopy forests (DBH of all trees below 10 cm, peatlands in our studied areas). More detailed information about the design is shown in our previous studies (Boisvert et al., 2021; Yin et al., 2022).

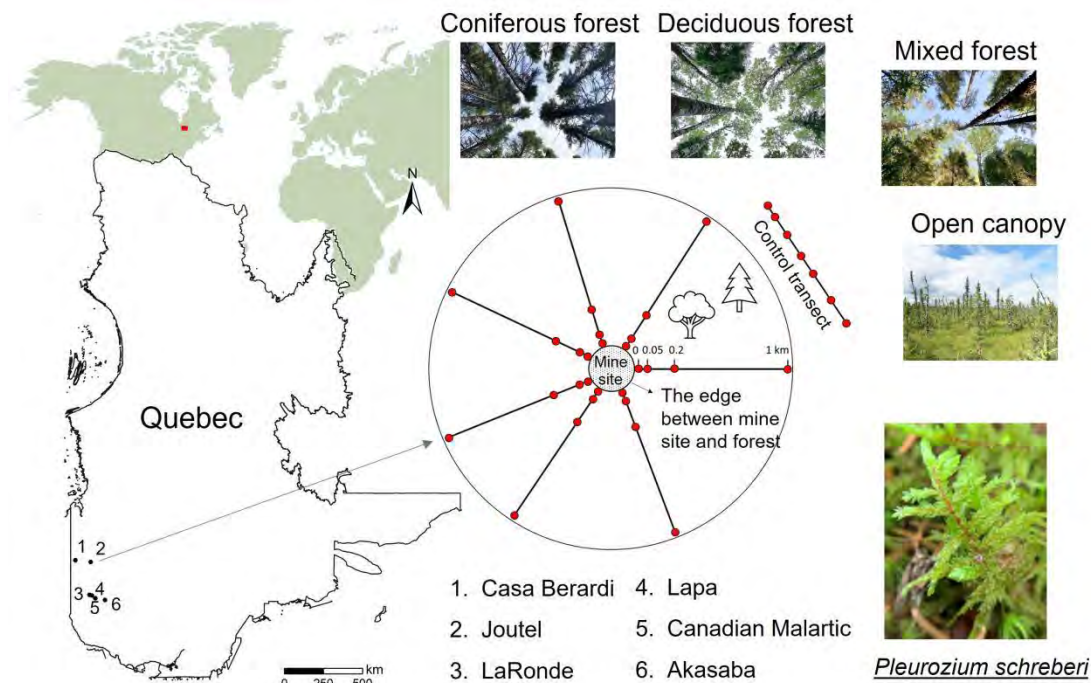


Figure 4.1 Study sites. Left, map of Quebec showing the location of the six gold mine sites. Right, sampling design. The four sampling plots at 0, 0.05, 0.2 and 1 km from the border of mine sites along transects at each study site are indicated by red circles. Six control transects were established near mine sites (see Figure S3.1).

4.4.2 Sampling

Between end of June and beginning of September of 2019, three colonies (a group of shoots of one species that live and interact closely with each other) of *P. schreberi* were randomly selected around the center point of each plot. The distance between colonies was approximately 10 m. At least 20 shoots per colony were picked, and debris and other plants were removed in the field. Only the top green parts (the last two years' growth, 0-3 cm from the tip of each shoot, Street et al., 2013) of each individual shoot were collected to avoid contamination from soil particles. All samples from one plot were pooled in sterile plastic roll bags and stored on ice in a cooler until reaching the laboratory, where samples were stored at -20°C until further

processing. In total, 211 samples were collected from the 220 plots (no *P. schreberi* was found in 9 plots). In addition, forest environmental variables were also measured in the field as previously described (Yin et al., 2022). Principal component analysis (PCA) analysis showed that the environmental variables differed significantly among ecosystem types (ANOVA, $p < 0.001$, Fig. S3.2), therefore, environmental variables were grouped into the four ecosystem types in further analyses.

4.4.3 DNA library preparation and sequencing

The DNA extraction process followed the protocol of Kembel et al. (2014), with some modifications. Briefly, 10-15 shoots were selected from the collected samples, placed into sterile 50 mL Falcon-tubes, and covered with phosphate buffered saline supplemented with Tween 20 (PBST) (30-35 ml). Shoots were removed from the tube after shaking on vortex for 30 seconds and on a stirrer plate for 5 minutes. Microbial cells were recovered from the PBST by centrifuging at $4000 \times g$ for 20 minutes at 4 °C. The supernatant was discarded and the pellet was transferred to a PowerBead tube of the DNeasy Power Soil DNA Isolation Kit (Qiagen, Valencia, CA) for DNA extraction, following the manufacturer's instructions. DNA extracts were quantified with the Qubit dsDNA HS assay kit using a Qubit Fluorometer 2.0 (Thermo Fisher Scientific, Waltham, MA, USA). Bacterial and fungal communities were characterized by amplifying and sequencing the V4-V5 regions of the 16S ribosomal RNA gene with the primer pair 515F-Y (5'-GTGYCAGCMGCCGCGGTAA-3') / 926R (5'-CCGYCAATTYMTTTRAGTTT-3') (Parada et al., 2016) and the ITS2 region of nuclear ribosomal DNA with the primer pair ITS9 (5'-GAACGCAGCRAAIIGYGA-3') (Menkis et al., 2012) / ITS4 (5'-TCCTCCGCTTATTGATATGC-3') (Bartels et al., 2018a), respectively. Library preparation for Illumina sequencing was performed according to the manufacturer's instructions for user-defined primers (Bokulich et al., 2013). The first amplification was performed in a 25 μ l reaction mix composed of 9 μ l of UltraPure™

DNase/RNase-Free distilled water (Gibco, Thermo Fisher Scientific), 200 μ M of each dNTP, 1.5mM of Mg^{2+} , 200 nM of each primer, 1 U of HotStarTaq Plus DNAPolymerase (Qiagen, Valencia, CA, USA), and 2.5 μ l of DNA extract. Thermocycling conditions were as follows: initial denaturation step at 95°C for 5 min, 34 (bacteria) or 40 (fungi) cycles at 94°C for 30 s, 50°C for 30 s, and 72°C for 1 min, and a final elongation step at 72°C for 10 min. PCR products were purified with magnetic beads (Agencourt AMPure XP, Beckman Coulter, Mississauga, ON, CA). Unique codes were added to each PCR product using the Nextera XT Index Kit following the manufacturer's instructions (Illumina). Indexed amplicons were purified with magnetic beads, quantified with the Qubit dsDNA BR Assay Kit (Thermo Fisher Scientific) and pooled at equimolar concentration. Sequencing was performed on an Illumina MiSeq platform with a MiSeq Reagent Kit v3 (600 cycles) at the Next Generation Sequencing Platform of the centre hospitaliers universitaires de Québec-Université Laval Research Centre. Sequence data from this study are available at the NCBI Sequence Read Archive (SRA) under the BioProject PRJNA800026.

4.4.4 Bioinformatic processing

The DADA2 Pipeline (1.16 version for 16S and ITS Pipeline Workflow 1.8 version for ITS, Callahan et al., 2016) on R platform was used to exclude primer sequences, filter and de-noise sequences, de-replicate unique amplicon sequence variants (ASVs, similar to 100%-identity operational taxonomic units), and remove chimeric sequences. Those ASVs were classified (minBoot=80) with the SILVA 138 (Quast et al., 2013) and UNITE 8.2 (Koljalg et al., 2005) taxonomic databases for 16S and ITS identification, respectively. Then, we identified and removed non-bacterial and non-fungal ASVs from bacterial and fungal ASV tables including ASVs classified as Archaea, Mitochondria, Chloroplast, Eukaryote and Rickettsioses. After the processing, 38,901 and 20,086 reads remained for bacteria and fungi, respectively.

The tables generated by the DADA2 Pipeline were imported into R (3.4) as a phyloseq object (*phyloseq* package version 1.12.3, McMurdie and Holmes, 2013) for further quality filtering steps. The removal of sequences identified as contaminants was used by the *decontam* R package (Davis et al., 2018), and then singletons, doubletons and ASV with less than 10 reads were removed to control for the potential influence of rare sequences. One sample from the LaRonde site showed a usual and much higher fungal richness than other samples based on visual inspection of rarefaction curves and was removed from further analyses. To analyze the alpha diversity of the phyllosphere microbiota, the datasets were rarefied to the minimum sample size (2,383 reads per sample for bacteria and 6,153 reads per sample for fungi, Weiss et al., 2017), whereas, for community similarity analysis, library size normalization was carried out using the geometric mean of pairwise ratios (Chen et al., 2018b).

4.5 Statistical analyses

To answer the three specific questions, statistical analyses were conducted on the bacterial and fungal dataset separately, and in two sections: 1) Examine offsite effects of mines on the phyllosphere microbiota as well as their potential drivers (ecosystem type and mine stage, Q1 and Q2); 2) Determine the distance influenced by the offsite effects (Q3). All analyses were performed with the statistical platform R 4.0.5 (2021-03-31) with R Studio software. Results were visualized with the *ggplot2* package (Wickham, 2011).

4.5.1 Effects of ecosystem type and mine stage on the phyllosphere microbiota

To test the presence of offsite effects of mines on the phyllosphere microbiota (Q1) and identify potential drivers of microbial diversity, community structure and composition (Q2), alpha, beta diversity and relative abundance of individual taxa

were analyzed. For alpha diversity analysis, generalized linear mixed models (GLMMs, “glmer” function from the *lme4* R package version 1.1-23, Bates et al., 2018) were applied on mining stage, ecosystem types and their interaction as predictors. Considering the nested structure of the sampling design, the nested terms site, transect and plot were included as random effects. Observed richness (number of ASVs), Shannon and InvSimpson indexes (Shannon and InvSimpson indexes were calculated with “estimate_richness” function from *phyloseq* package) were selected as the response variables for bacterial and fungal alpha diversity, separately. Based on the inherent statistical characteristics of response variables, Negative Binomial (overdispersed count data, observed bacterial and fungal richness, with “glmer.nb” function in *lme4* R package) and Gaussian (continuous number with normally distributed residuals, Shannon and InvSimpson Indexes with “glmer” function with family=gaussian and a log link) distribution were used in the models. Marginal R squared of GLMMs were calculated with the function “r.squaredGLMM” of the *MuMIn* R package R package 1.15.6 (Barton and Barton, 2015). Significant p-values for the effects of mining stage, ecosystem type and their interactions were calculated with ANOVA using Type II sum of squares (Type II sum of squares is more powerful than Type III for ANOVAs with unbalanced data, Langsrud, 2003, “Anova” function in the package “car” version 3.0-2 (Fox and Weisberg, 2019). When interaction terms were not significant, they were removed from the final models. Multiple pairwise comparisons for all response variables among mine stages and ecosystem types and when the interaction term was significant, pairwise comparisons were performed between mine stages in each ecosystem type. When the interactions were not significant, pairwise comparisons between each mine stage and between each ecosystem type, respectively were conducted with TukeyHSDtest (“emmeans” function in the *emmeans* package version 1.3. 5.1, Lenth et al., 2019).

Similarly, the effects of ecosystem type and mining stage on phyllosphere beta diversity were tested by permutational multivariate analysis of variance

(PERMANOVA, “adonis” function from the *vegan* package version 2, Oksanen et al., 2013) on Bray-Curtis distance matrices (“vegdist” function in the *vegan* package) with 999 permutations. PERMANOVA tests were performed for the bacteria and fungi ASV matrices as the response and ecosystem type, mining stage and their interactions as the explanatory variables and site factor as a ‘strata’ term to reduce randomizations across all sites. Bray-Curtis distance matrices were further visualized using non-metric Multi-dimensional Scaling (NMDS, “metaMDS” function in the *vegan* package) with three dimensions. We also used post-hoc pairwise PERMANOVAs (999 permutations, “pairwise.adonis” function in the package *RVAidemoire* version 0.9-45-2, Hervé, 2014) with a multiple comparison correction based on Benjamini-Hochberg method (Benjamini and Hochberg, 1995) to compare differences between mining stages.

Then, relative abundance of main bacterial and fungal phyla (average relative abundance across all samples > 0.5 %) was used as response variables to determine the effects of mining stage and ecosystem type (similar analysis process for alpha diversity). Microbiome relative abundances are compositional data that range between 0 and 1 and are also generally zero-inflated (Ho et al., 2019; Peng et al., 2016), therefore zero-inflated beta mixed models (ZIBMMs, and “glmmTMB” function with `ziformula=~.` and `family=beta_family` in *glmmTMB* package version 0.1.3, Brooks et al., 2017) were used instead of GLMMs here. The models that showed convergence errors were corrected by optimization of the model algorithm via Broyden–Fletcher–Goldfarb–Shanno (BFGS) algorithm (Dai, 2002) method in the function of the “glmmTMBControl” from *glmmTMB* package. Marginal R squared of ZIBMMs were calculated with the function “r2” of the *sjmisc* package version 2.4.0 (Lüdtke and Lüdtke, 2019). Furthermore, to identify the taxa (ASVs) driving differences in community composition between offsite landscapes of operating, non-operating mine sites and controls, indicator species analysis (“multipatt” function in the *indicspecies* package version 1.7. 6, De Cáceres et al.,

2016) with point biserial-correlation coefficient (ϕ) was applied.

4.5.2 Effects of the distance from mines on the phyllosphere microbiota

The effects of distance from the mines on alpha and beta diversity of bacterial and fungal communities (Q3) and on the relative abundance of individual taxa (i.e., main phyla, top 10 main genera through all samples, and common ASVs found in at least 90 % of samples for bacteria and 70 % of samples for fungi) were analysed. Control transects were removed from these analyses given that the control transects were designed for landscape-level analysis (Q1 and Q2). Firstly, to assess variations in diversity and community structure (alpha and beta diversity) along the transects, alpha diversity indices (Observed ASV richness, Shannon and InvSimpson) and Bray–Curtis dissimilarity matrices were used as response variables in GLMMs and Pairwise PERMANOVAs, respectively. The effect of distance and its interaction with mining stage (only operating and non-operating here) were used as predictors. The analysis process (error distribution, ANOVA test and post hoc test) was similar to the models we used in Q1 and Q2, but here transects were nested into sites, and ecosystem type were used as random factors. Distance influenced by offsite effects of mines were determined based on the results of TukeyHSD tests and graphs (showing the position of threshold point, “ggplot” function in *ggplot2* package). Considering the very large ranges of variation (orders of magnitude) between taxa and zeros in relative abundance datasets, row data were replaced by $\log(x + 1)$ transformation to present the spatial pattern with the distance in graphs. Finally, the offsite footprint of mines was evaluated and estimated using the influenced distance for alpha and beta diversity of bacterial and fungal communities as well as the relative abundance of individual taxa.

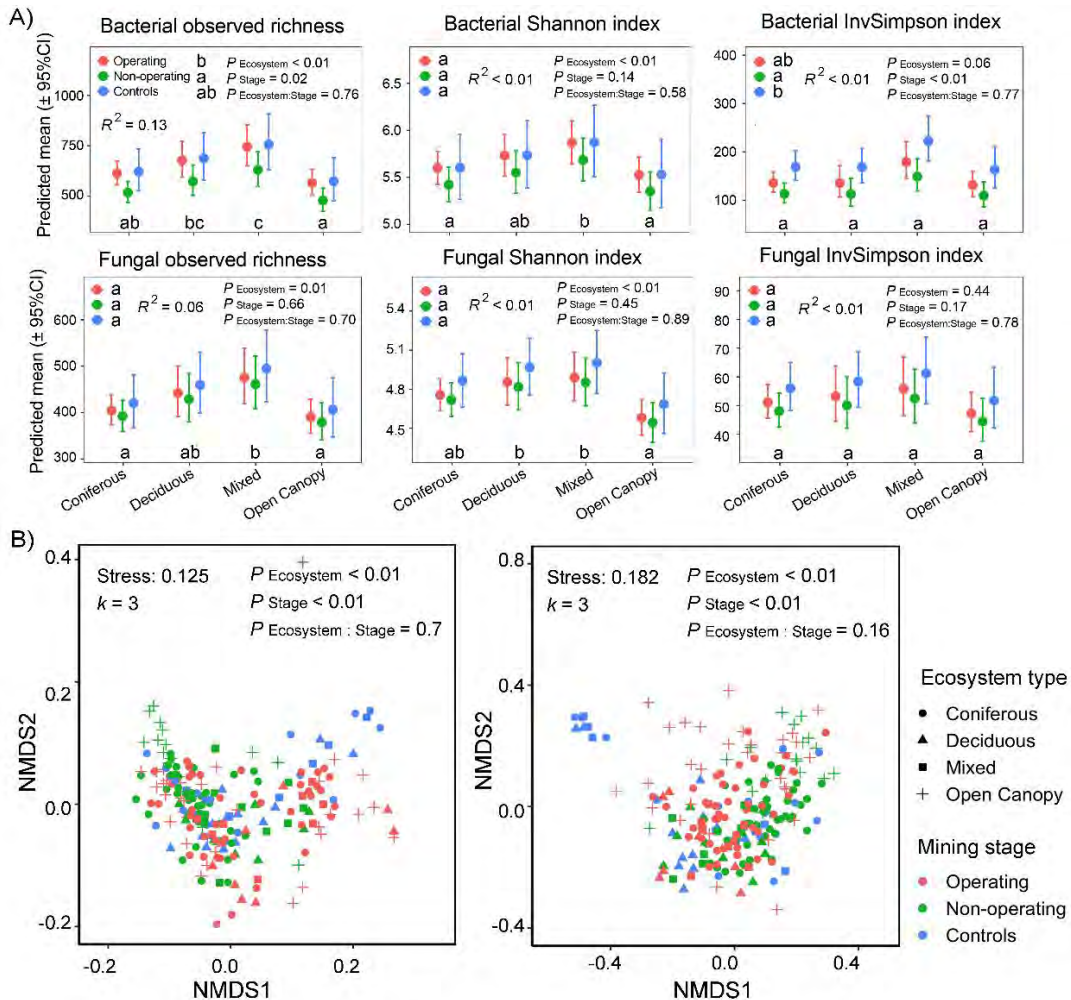


Figure 4.2 Changes in the phyllosphere alpha and beta diversity in mine offsite landscapes. A) Alpha diversity for both bacterial and fungal communities at different mining stages (plots surrounding operating sites, $N=92$; plots surrounding non-operating sites, $N=78$; plots in controls, $N=40$) in four ecosystem types (coniferous forest, $N=101$; deciduous forest, $N=33$; mixed forest, $N=26$; open canopy, $N=50$). Points show means for all samples; bars show a 95% confidence interval around the mean. Different lowercase letters indicate significantly different means across mining stage and ecosystem type, respectively. P value based on generalized linear mixed models (type II sum of squares) and Tukey HSD pairwise comparisons ($\alpha = 0.05$) were used to test the differences across ecosystem types and mining stages, respectively. B) Beta diversity of bacterial and fungal community structure. Non-metric Multi-dimensional Scaling (NMDS) are based on the Bray–Curtis dissimilarity. Tests in the top side of the panels present the results of PERMANOVA.

4.6 Results

4.6.1 Effects of mining stage and ecosystem type on the phyllosphere microbiota

Mining stage and ecosystem type both affected *P. schreberi* phyllosphere alpha and beta diversity in mine offsite landscapes without interaction between the two factors. Bacterial alpha diversity differed among mining stages (observed richness and InvSimpson index, ANOVA type II sums, $P_{\text{Stage}} < 0.05$, Fig. 4.2A). No significant differences in alpha diversity were found between operating sites and controls (post hoc test results in Fig. 4.2A), but both had higher alpha diversity values than non-operating sites (Fig. 4.2A). The effects of mining stage on bacterial alpha diversity did not depend on ecosystem type (ANOVA type II sums, $P_{\text{Ecosystem:Stage}} > 0.05$, Fig. 4.2A), but ecosystem type also affected the alpha diversity (bacterial observed richness and Shannon index, ANOVA type II sums, $P_{\text{Ecosystem}} < 0.05$, Fig. 4.2A). Generally, deciduous and mixed forests had a higher alpha diversity than coniferous forests and open canopy in the offsite landscapes (bacterial observed richness and Shannon index, Fig. 4.2A). Similar patterns were found in fungal alpha diversity with higher alpha diversity found in deciduous and mixed forests (fungal observed richness and Shannon index, Fig. 4.2A) and the lack of interaction between mining stage and ecosystem type. But in contrast to bacteria, none of the fungal diversity indices measured was significantly affected by the mining stage (ANOVA type II sums, $P_{\text{Stage}} > 0.05$, Fig. 4.2A). Furthermore, only 0-13% of shifts in bacterial and fungal alpha diversity were explained by fixed factors (mining stage and ecosystem types) in the models (see R^2 in Fig. 4.2A).

Similarly, the analysis of the beta diversity of phyllosphere bacterial and fungal communities showed a significant effect of both the mining stage (bacteria, $R^2 = 4.43\%$, fungi, $R^2 = 2.62\%$, PERMANOVAs in Fig. 4.2B and Table S3.1) and ecosystem type (bacteria, $R^2 = 5.83\%$, fungi, $R^2 = 4.63\%$, PERMANOVAs in Table S3.1) without interaction between the two factors (PERMANOVAs, $P_{\text{Ecosystem:Stage}} >$

0.05, Fig. 4.2B) Furthermore, all pair-wise comparisons for mining stage and ecosystem type showed significant differences in bacterial and fungal community structure (Pairwise PERMANOVA tests, $P < 0.01$, Table S3.2 and Table S3.3).

For most phyla detected in the moss phyllosphere, mining stage and ecosystem type both had a significant effect on the relative abundance. The effect of mining stage was variable depending on the phylum. The relative abundance of the bacterial phylum Acidobacteriota and the fungal phylum Ascomycota were lower in offsite landscapes of operating sites than in non-operating sites across all samples (see Tukey HSD pairwise comparisons in Fig. 4.3), while the relative abundances of the bacterial phyla Bacteroidota and Myxococcota were higher near operating sites than non-operating sites (Fig. 4.3). The relative abundance of the Armatimonadota was higher in both operating and non-operating sites than in the controls (Fig. 4.3). The relative abundance of dominant phyla also differed between ecosystem types. Higher relative abundance of Acidobacteriota was detected in coniferous forest and open canopy ecosystems than in deciduous and mixed forests, while deciduous and mixed forests had a higher relative abundance of Bacteroidota. A significant interaction effect between the mining stage and ecosystem type on the relative abundance of four bacterial phyla (i.e., Proteobacteria, Cyanobacteria, WPS-2 and Olpidiomyces, Fig. 4.3) was observed. Interestingly, the relative abundance of two groups of photosynthetic microorganisms, the Cyanobacteria phylum and the WPS-2 candidate phylum, were negatively affected by operating mines in either coniferous forests or open canopy, but not in deciduous and mixed forests (Fig. 4.3). Overall, 5-25% of the shifts in the relative abundance of bacterial and fungal phyla were explained by fixed factors (mining stage and ecosystem type) in the models (see R^2 in Fig. 4.3).

Indicator species analysis was performed to identify fungal and bacterial taxa driving the differences in microbial communities based on mining stage. Phyllosphere of *P. schreberi* near operating sites, non-operating sites and control sites were characterized respectively by 42 (8 phyla), 44 (6 phyla) and 22 (4 phyla, Fig. 4 and

Table S3.5) bacterial indicator species, and 54 (4 phyla), 23 (4 phyla) and 78 (5 phyla) fungal indicator species (Fig 4.5 and Table S3.6) (in this case, the term ‘species’ refers to ASVs). Bacterial ASV_84 (stat=0.318, Table S3.5) and ASV_101 (stat=0.318, Table S3.5) assigned to the genera *Acidiphilium* and *Candidatus Solibacter*, respectively, were the best indicator species for operating sites (Table S3.6), whereas bacterial ASV_8 (stat=0.279, family Acetobacteraceae, Table S3.5) and ASV_361 (stat=0.514, genus *Cupriavidus*, Table S3.5) were the best indicators of non-operating sites and controls, respectively (Table S3.5). Fungal ASV_10 (stat=0.326, genus *Phenoliferia*, Table S3.6) was the best indicator species for operating mines, while fungal ASV_136 (stat=0.314, order Orbiliales, Table S3.6) and ASV_429 (stat=0.397, genus *Thelephora*, Table S3.6) were the best indicator species of non-operating sites and controls, respectively.

4.6.2 Effects of distance from the mine on the phyllosphere and offsite footprint

Phyllosphere alpha and beta diversity were associated with the distance from mines without interaction with the mining stage (ANOVA type II sums for GLMMs in bacterial and fungal observed richness and Shannon indices, $P_{\text{Distance}} < 0.05$, $P_{\text{Stage}} < 0.05$ and $P_{\text{Distance:Stage}} > 0.05$; PERMANOVAs in bacteria and fungi, $P_{\text{Distance}} < 0.001$, $P_{\text{Stage}} < 0.001$ and $P_{\text{Distance:Stage}} > 0.05$, Table S3.7 and S3.8). Bacterial alpha diversity (observed richness and Shannon index) showed a decreasing trend with the distance from mine sites. Observed richness and Shannon index were respectively about 13% and 4% higher at 0 km than at 1 km, although only the 0 km value was significantly higher compared to the 0.2 and 1 km values (TukeyHSDtest results in Table S3.7 and Fig. 4.5). In contrast, fungal alpha diversity (observed richness and Shannon index) followed a non-linear pattern with the distance, with no significant difference between 0 and 1 km but with significant differences between 0 and 0.05 or 0.2 km (TukeyHSD tests in Table S3.7 and Fig. 4.5). For beta diversity, bacterial community structure at 0 and 0.05 km differed from that at 1 km (Fig. 4.5B), while fungal

community structure differed between 0 and 1 km, but no significant difference was found between 0.05 and 1 km (Fig. 4.5B). Therefore, in most cases, the variation of phyllosphere alpha and beta diversity was only detectable until 0.05 km into the forest interior.

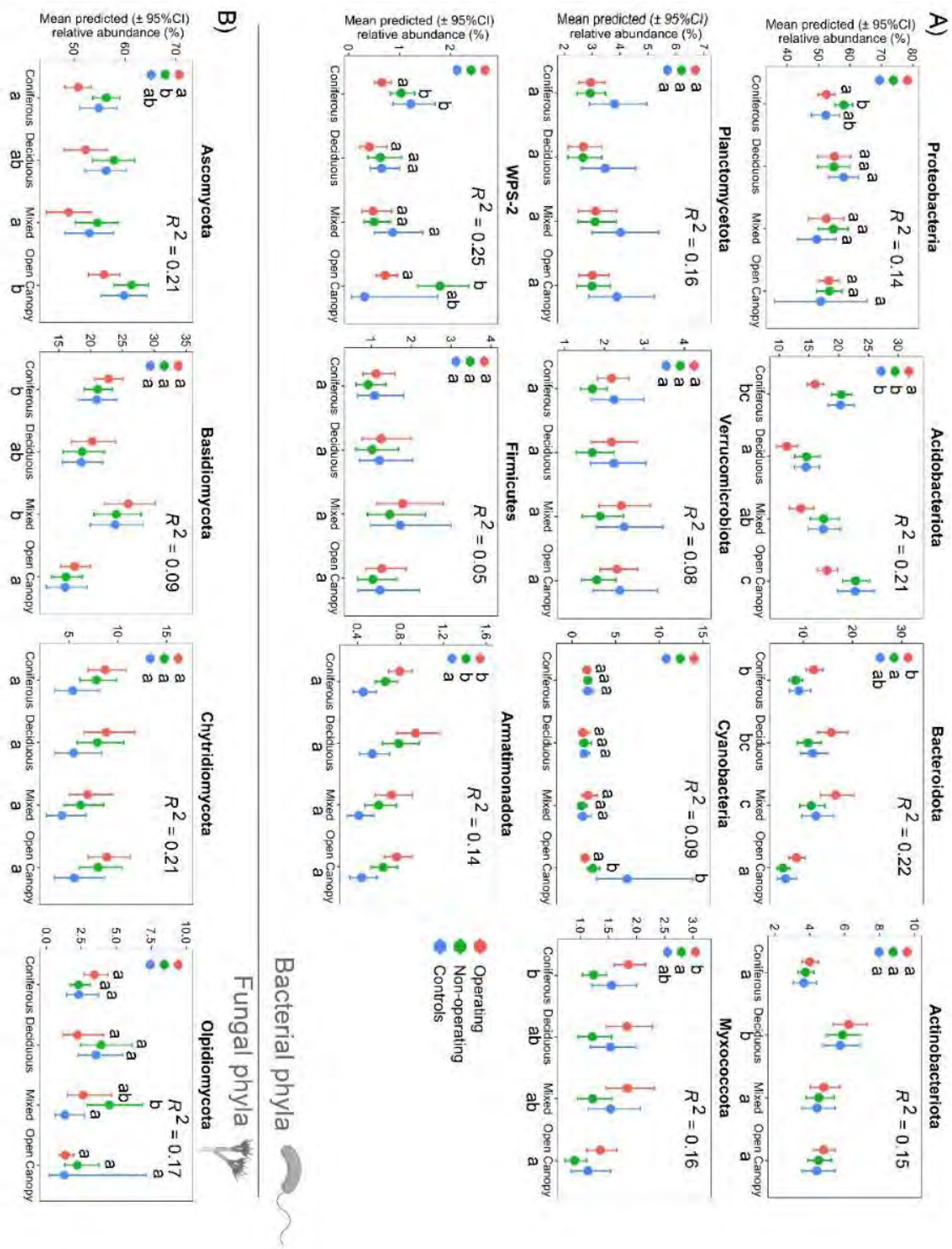


Figure 4.3 Differences in relative abundance of main phyllosphere phyla (relative abundance > 0.5 %) at each mining stage in four ecosystem types in mine offsite landscapes. Zero-inflated beta regression models were used to determine responses of relative abundance of main bacterial and fungal phyla to ecosystem types, mining stages and their interaction. Points show means for all samples; bars show a 95% confidence interval around the mean. Different lowercase letters indicate significantly different means across groups based on Tukey HSD pairwise comparisons ($\alpha = 0.05$). When interactions between ecosystem type and mining stage were significant, pairwise comparisons were used between mine stage in each ecosystem type, while when the interactions were not significant, the interaction terms were removed from the models and pairwise comparison were used between ecosystem type and mining stage, separately.

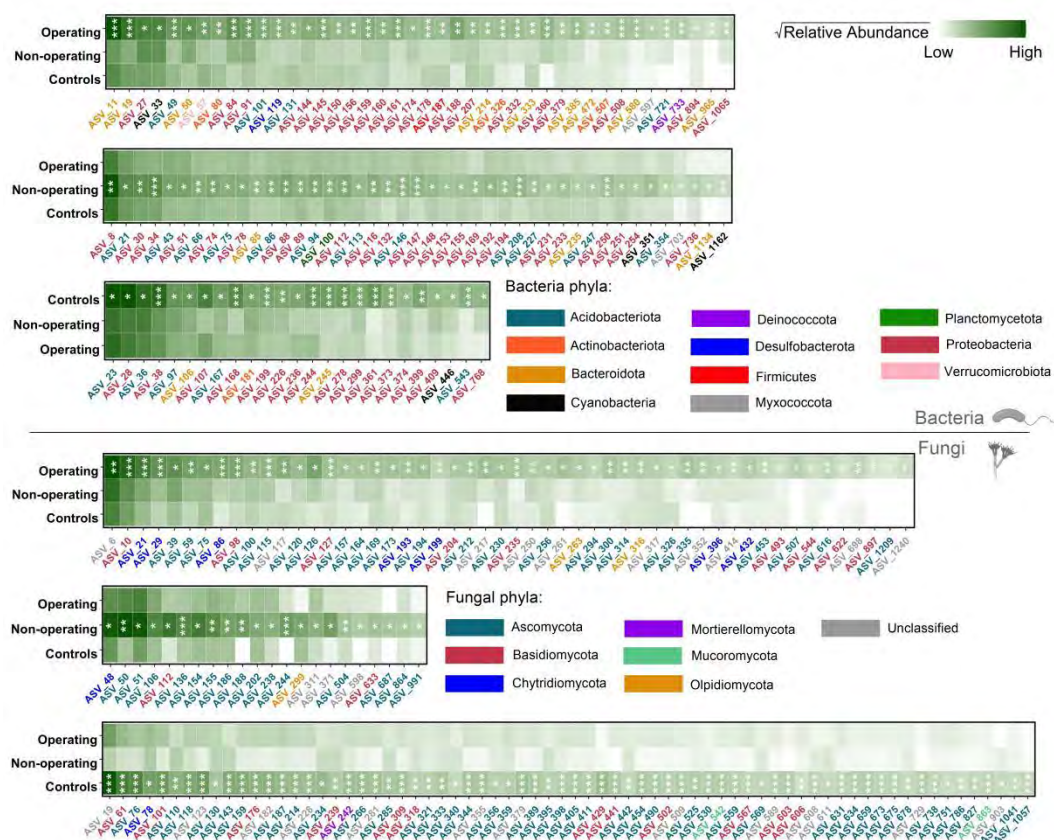


Figure 4.4 Heatmaps of the average relative abundance of indicator ASVs for mining stages. Only ASVs with total relative abundance > 0.05 % were included in the indicator analysis. * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$. Note relative abundance of each bacterial and fungal ASV is shown for each mine stage when not an indicator ASV in that stage.

In contrast to alpha and beta diversity results, analysis of the relative abundance of individual taxa detected significant interactions between the distance and the mining stage as well as significant effects of mining up to over 0.2 km. In total, for five phyla, four genera and four ASVs among bacterial taxa, as well as one fungal phylum and two fungal ASVs, the effect of distance on the relative abundance based on GLMMs depended on whether the mine site was operating (Fig. 4.5C, Table S3.9, S3.10 and S3.11). Relative abundance of three bacterial phyla (i.e., Bacteroidota, Actinobacteriota and Armatimonadota), one bacterial genus (i.e., *Sphingomonas*) and one fungal ASV (ASV 10, *Phenoliferia* see Table S3.7) were positively influenced by mines, with highest values (between 1.4 and 3.14-fold higher than those at 1 km) detected at 0 km and decreasing with the distance near operating mines. No linear associations with distance were found near non-operating sites for these taxa (Fig. 4.5C). For another four ASVs (bacterial ASV 15, *Sphingomonas*, and fungal ASV 21, Spizellomycetales order, ASV 22, *Cladosporium*, ASV 66, *Aureobasidium*, Fig. 4.5C and Table S3.11), the relative abundance was positively affected by mines near both operating and non-operating sites (1.63-7 fold changes). Furthermore, two bacterial phyla (i.e., Acidobacteriota and WPS-2), three bacterial genera (i.e., *Granulicella*, *Leptothrix* and *Terriglobus*) and six bacterial ASVs, as well as two fungal genera (i.e., *Epithamnolia* and *Thoreauomyces*) and four fungal ASVs were negatively affected by either operating mines or all mine sites (both operating and non-operating, Fig. 4.5C).

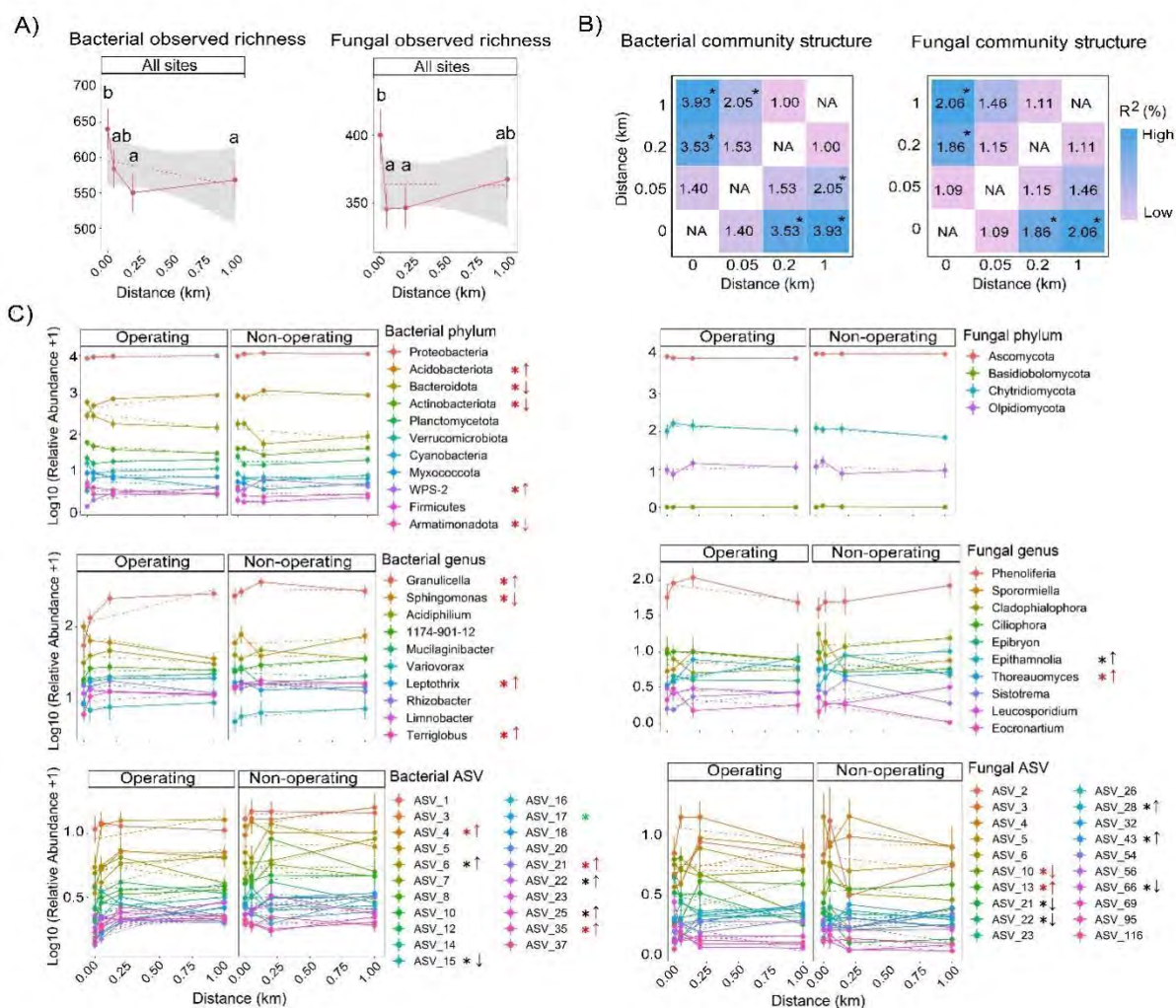


Figure 4.5 Changes in phyllosphere microbial communities with distance from mine sites. A) Bacterial and fungal observed richness. Different lowercase letters indicate significantly different means between different distances at 0 (N=39), 0.05 (N=44), 0.2 (N=43) and 1 km (N=44) from mine sites (results based on generalized linear mixed models and Tukey HSD pairwise comparisons, $\alpha = 0.05$). B) Bacterial and fungal community structure differences between distances based on the results of pairwise PERMANOVA on Bray-Curtis dissimilarity matrices, * $P < 0.05$. C) Relative abundance of bacterial and fungal taxa along the distance at phylum, genus and ASV levels. Points show means for relative abundance after log (x + 1) transformation; bars show standard error (SE). Black asterisks indicate significant association with distance at both mining stages based on the zero-inflated beta mixed regression models and post-hoc tests with Tukey HSD pairwise comparisons. Red

asterisks indicate the interaction between the distance and mining stage was significant and the effects occurred near operating mines, while green asterisks indicate the effects occurred near non-operating sites. Arrows used for linear patterns: up arrows, relative abundance is increasing with the distance; down arrows relative abundance is decreasing with the distance.

4.7 Discussion

4.7.1 The presence of offsite effects of mines on the feather moss phyllosphere microbiota

Our results confirmed the presence of offsite effects of mines on the feather moss phyllosphere microbiota (Q1) based on the differences in alpha and beta diversity, relative abundance of individual taxa, and indicator species of mined sites (operating or non-operating) and controls. These findings extend the offsite effects of mines on surrounding ecosystems to microbiomes, as only vegetation and animals were used to determine the effects in previous studies (Boisvert et al., 2021; Dyer et al., 2001; Watkinson et al., 2021; Wu et al., 2021, Yin et al., 2022). Surprisingly, similar levels of bacterial alpha diversity were detected in operating mines and controls, while the diversity in non-operating sites was lower (Fig. 4.2), which is contrary to our expectations. Integrating these results with those obtained for individual taxa, we suggest that this result is explained by taxon-specific responses to offsite effects. Some taxa (e.g., Armatimonadota, Fig. 4.3) were favored near mine sites, although some other (e.g., Acidobacteriota, Fig. 4.3) were suppressed, leading to similar levels of diversity as in the control sites. These results seem to support the novel ‘niche flip’ mechanism in microbial ecology, where alpha diversity is shaped by both disturbance frequency and intensity (Mancuso et al., 2021). In this mechanism, the microbial communities follow a U-shaped diversity dependence on the disturbance intensity, with the lowest level of diversity occurring at an intermediate level of disturbance (Mancuso et al., 2021). Here, controls, non-operating and operating sites represent zero, low and intermediate mining disturbance intensity, respectively, and followed this high-low-high (U-shaped) pattern in alpha diversity. Further work will be needed

to confirm the mechanisms underlying these observations, especially regarding the fluctuation and intensity of the offsite effects, which were not characterized in this study.

Differences in phyllosphere fungal and bacterial beta diversity between mining stages confirmed that the mining stage is a driver of microbial community structure in mine offsite landscapes. Previous studies have identified pH, total organic carbon, total phosphorus and heavy metal content as the main influencing factors in the variation of soil microbial community structure in mining polluted areas (Dimitriu and Grayston, 2010; Kane et al., 2020; Wu et al., 2022). These factors could also partly explain the variation observed in the moss phyllosphere in this study considering that feather moss grows in close contact with the soil. The concentration of metals (aluminium, copper) was also significantly correlated with variations in microbial functions in a study comparing the phyllosphere microbiota of various tree species (Lajoie et al., 2020). Considering that metal concentrations could be higher in the phyllosphere near operating than non-operating mines and that mosses are known to accumulate metals (Blagnytė and Paliulis, 2010; Stanković et al., 2018), we can expect that metal concentration in the moss phyllosphere could be responsible for the differences in community structure observed here between mining stages. For example, the drastic decline of Acidobacteriota near operating mines compared with non-operating and control sites indicated the sensitivity of Acidobacteria to heavy metals (Guo et al., 2019; Macdonald et al., 2011; Zheng et al., 2019). In contrast, the higher relative abundances of Bacteroidota near operating mines suggests that these landscapes could provide favorable conditions for some microorganisms. Bacteroidota usually harbour the majority of heavy metal resistance genes (Yan et al., 2020), and is able to quickly adapt and endure high heavy metal stresses in mining polluted areas (Zhao et al., 2020a; Zhao et al., 2020b; Zou et al., 2021). Characterizing soil properties, assessing dust emissions from mines, and measuring heavy metal concentrations in mosses would be important next steps to further

identify the environmental drivers responsible for changes in phyllosphere community structure observed in mine offsite landscapes.

Furthermore, operating mines supported some specific species, as highlighted by the indicator species analysis. Bacterial ASVs 101 and 84 as well as the fungal ASV 10 were the best indicators of operating mines (Table S3.5 and S3.6) and belonged to *Candidatus Solibacter*, *Acidiphilium* and *Phenoliferia*, respectively. The ability of members from the first two bacterial taxa to overcome stressful conditions (including high metal concentrations) have been reported in mining environments (Böhmer et al., 2020; Li et al., 2020). Furthermore, members of the fungal genus *Phenoliferia* can degrade phenolic compounds (Pakarinen et al., 2021; Perini et al., 2019) and its higher relative abundance near operating mines might be driven by a higher phenolic compound content of feather moss, as these compounds are usually produced by plants as a defense mechanism under heavy metal stress (Maleki et al., 2017). More studies are needed to explore which environmental variable or plant traits (such as heavy metals and phenolic compounds content) lead to the increased relative abundance of Fungal ASV 10 (*Phenoliferia*) near mines. Nevertheless, fungal ASV 10, with its relatively high abundance (about 1.5 % at 0 m near operating mines, Table S3.11) and its sensitivity to mining stage and distance, could be a reliable indicator to assess the presence and magnitude of offsite effects of mines. Surprisingly, one of the best indicator ASV for control sites (bacterial ASV361) was assigned to the *Cupriavidus* genus, which contains several species or strains known for their heavy metal resistance and adaptation to metal-contaminated environments (Mijnendonckx et al., 2013; Reith et al., 2009; Van Houdt et al., 2018). The species or strain identified here in the moss phyllosphere may lack the specific genes required for heavy metal resistance, which could be confirmed by further characterizing the moss phyllosphere DNA using shotgun metagenomics approaches.

Our results confirmed that offsite effects are usually cryptic because inherent limitations of impact evaluation (e.g., experimental design, detection methods, spatial

scales, statistical power) led to those effects being overlooked. If our experimental design had included operating mines only, and if the focus of our statistical analyses had been limited to alpha diversity, substantial offsite effects could have been undetected because taxa-specific responses were masked in the alpha diversity response (especially for fungi, Fig. 4.2). This study also highlights that the use of environmental DNA (eDNA) tools can reveal impacts of disturbances on groups of living organisms (here bacteria and fungi, but many others could be targeted) that are typically overlooked in impact evaluations because of the absence of efficient methods to detect them. The performance of these tools to assess biodiversity and measure the environmental impacts of disturbances is becoming widely accepted by the scientific community (Edge et al., 2020; Pawlowski et al., 2021) and hopefully, this will lead to a wider application of these tools in impact assessment in the near future.

4.7.2 Ecosystem type and mining stage affected phyllosphere microbiota in mine offsite landscapes

Our results provide further evidences that feather moss phyllosphere diversity and composition differed between ecosystem types (Jean et al., 2020b; Rodríguez-Rodríguez et al., 2022). However, no differences in alpha diversity (bacterial and fungal observed richness and bacterial Shannon index, Fig. 4.2B) were found between deciduous and coniferous forests in contrast with Rodríguez-Rodríguez et al. (2022) who found higher bacterial alpha diversity of the feather moss phyllosphere in coniferous stands (black spruce) than in deciduous stands (trembling aspen). To avoid a loss of statistical power by creating too many categories, coniferous forests and deciduous forests in this study were not monospecific stands, which may have led to these differences. Our observations of a higher relative abundance of the bacterial phylum Acidobacteria in coniferous forests and open canopy and a higher relative abundance of Bacteroidota in deciduous and mixed forests were, however, consistent

with findings from Rodríguez-Rodríguez et al. (2022). Despite the limitation that most forest stands in this study were located near mine sites and not in natural conditions, our results still provide insights into the relationships between ecosystem types and moss phyllosphere communities in boreal forests.

Surprisingly, the effects of mining stage on the alpha, beta diversity and relative abundance of most phyla were not ecosystem type-dependent. It suggests that ecosystem types did not influence the ability of the moss phyllosphere to resist to the offsite effects, which is in contrast with a previous study indicating that bryophyte diversity was more affected by mining stage in deciduous forest than in coniferous forest (Yin et al., 2022). Compared with deciduous forest, coniferous trees with high canopy cover, leaf area index and persistent foliage intercept more pollutants in atmospheric deposition (Barbier et al., 2008; Nguyen et al., 2015). Here, this mechanism did not play an obvious role in reducing the impact of mines on the moss phyllosphere microbiota as the effects were similar in coniferous and deciduous forests. To our knowledge, no study to date has directly compared the sensitivity of bryophyte communities to that of phyllosphere microbial communities to dust, but microorganisms colonizing the leaf surface are known to be strongly exposed to adverse conditions, including air pollutants (Bringel and Couée, 2015), and have been found to be more sensitive to environmental conditions than leaf endophytes (Sivakumar et al., 2020). A study also found that the overall reduction of dustfall by coniferous trees was between 38-42%, while deciduous trees reduced dustfall by 27-30% (Dochinger, 1980). So over 50 % of dustfall still could pass through the canopy structure, leading to loads of heavy metals and other toxins that may have impacted the phyllosphere microbial communities without reaching, in coniferous forests, the level at which bryophyte diversity would also be significantly impacted. On the other hand, it is also possible that an interaction between ecosystem types and mining would have emerged if the phyllosphere was sampled later in the season after leaf fall, when the differences in canopy structure were more striking

than in the summer. Further studies should be done to examine these two potential explanations. Overall, the results indicate that predicted shifts in ecosystem types would not generally modify the offsite effects of mines on the phyllosphere microbiota in boreal landscapes.

Notably, the offsite effects on the relative abundance of two photosynthetic bacterial phyla, WPS-2 and Cyanobacteria, were affected by mining stage-ecosystem type interactions. Indeed, no effect of the mining stage on the relative abundance of WPS-2 and Cyanobacteria was detected in both deciduous and mixed forests, which can be expected as these ecosystem types are known to be unfavorable for the two groups (Jean et al., 2020a; Qu et al., 2020; Rousk et al., 2013) because of relatively low air moisture, high soil pH and broadleaf litter inputs (Augusto et al., 2002; Qian et al., 2003). However, the relative abundance of these phyla was lower near operating mines in coniferous forest and open canopy peatland, which is in contrast with previous studies highlighting the resistance of Cyanobacteria and WPS-2 to metal contamination (Al-Amin et al., 2021; Cui et al., 2021; Grasby et al., 2013; Kavehei et al., 2021). This may be attributable to the differences between the soil and phyllosphere habitats (Leveau, 2019). However, there is almost no literature to explain these findings. The current knowledge on how environmental stress affects phyllosphere communities is very limited, especially for mosses, and more studies on this topic are needed. Overall, although mechanisms are still unclear, our results suggest that the ecological services supported by these taxonomic groups, such as the fixation of carbon and nitrogen, could be impacted in offsite landscapes of operating mines, especially in coniferous forests and open canopy peatlands. We also found fungi were more resistant to the offsite effects than bacteria. According to previous studies, fungi appear to be more tolerant to heavy metals than bacteria (Kelly et al., 2003; Zeng et al. 2020), and extreme conditions under mining disturbances imposing selection pressure may trigger additional evolutionary adaptation in the fungal communities (Adriaensen et al. 2005).

4.7.3 The footprint of offsite effects of mines on feather moss phyllosphere

Determining the offsite footprint of mines is challenging but necessary for policy maker and governments to assess ecological impacts of mining projects and, subsequently, develop mitigation strategies. Our findings indicated that phyllosphere alpha and beta diversity were affected by the distance from the mine site, but the effect was only detectable up to a distance between 0 - 0.05 km from mine sites, which is much lower than what was detected using the relative abundance of individual taxa (up to 1 km, such as fungal ASV10 near operating mines, Table S3.11). Therefore, if the offsite footprint is determined at the community-level, sizes of the offsite footprint and their impacts on ecological services could be underestimated.

Based on all results of alpha, beta diversity and relative abundance of individual taxa, we found that significant effects are detectable within 0.2 km of mine sites for most taxa, especially for operating mines. Therefore, if the average width of the offsite footprint surrounding mines is set to 0.2 km, the total area for footprint calculation would be about 4.5 km² (average 1.5 km²/mine, measured through Google Earth Pro 7.1) for the three operating mines in our study. With more than 6,000 active mines across the globe (Maus et al., 2020), the estimated global offsite footprint of active mines on surrounding phyllosphere microbiota could be up to 9,000 km² (about the land area of Cyprus). Considering non-operating sites still have effects on surrounding phyllosphere structure, the actual offsite effects of mines could be even larger. Although the mining method, size, shape and local environmental conditions could affect the range of offsite effects, our results still provide justification to include offsite footprints of mines when evaluating the total footprint of mining on landscapes.

4.8 Conclusion

Our findings highlight the presence of offsite effects of mines on microbiomes, as exemplified here in the feather moss phyllosphere. Although both mining lifecycle and forest composition were identified as factors affecting the offsite effects of mines, our study indicates that these effects are cryptic and can easily be ignored at the community level. Specific taxa such as Bacteroidota and fungal ASV10 (*Phenoliferia*) have higher indicator roles than community-level diversity to determine the presence and footprint of offsite effects of mines. Though the average distance influenced by the offsite effects is small (about 0.2 km) near individual mines, the sum might be large and impactful across the globe. Moreover, as more mining projects will be established in the coming decades with the energy transition, more landscapes will be exposed to the offsite effects of mines. The global offsite footprint of mines is therefore expected to expand and become a problem over larger areas. Further studies are needed to assess how these changes in microbial communities of the phyllosphere in mine offsite landscapes affect ecosystem functioning, including carbon and nitrogen cycling in boreal ecosystems, and to develop appropriate mitigation strategies.

4.9 Acknowledgements

This research was financially supported by the NSERC-UQAT Industrial Chair in Northern Biodiversity in a mining context. Thanks for Mebarek Lamara for helps in DNA extraction, bioinformatic analysis. We thank Enrique Hernández-Rodríguez and Carlos Cerrejon Lozano for their assistance in creating the study area map. We deeply thank Julie Arseneault and Mélanie Jean for the establishment of the plots. We also thank Julie Arseneault, Mélanie Jean, Elizabeth Riendeau, Andréane Garant, Cyrielle Bigonneau, Simon Bégin, Matilde Offroy, Paméla Parisien, Jeffrey Raymond, Juliette Larrivée, Marie-Anyse Dubuc, Emmanuelle Richard, Maxime Thomas, Teresa Kim,

Xavier Saint-Amant, Jolianne Garand and Natalie Vuong for their valuable assistance in the fieldwork, and Marie-Josée Morency for the metabarcoding library preparation. We thank Josée Noël, Lucienne Ancil, Stéphanie Lafrenière and other people at the mines of Canadian Malartic, Casa Berardi, LaRonde, Lapa, Akasaba and Joutel for their help with access to the mine sites. We also thank the two reviewers whose suggestions increased the quality of the manuscript.

CHAPITRE V

CONCLUSION GÉNÉRALE

5.1 Résumé des principaux chapitres

Cette thèse a été conçue pour identifier et caractériser les effets hors site des mines sur la biodiversité des paysages boréaux environnants et pour déterminer si la composition de la forêt et le cycle de vie des mines sont des moteurs de l'effet hors site. Dans le chapitre II, différents groupes de végétation de sous-bois de plantes ligneuses, herbacées, bryophytes et lichens ont d'abord été utilisés pour déterminer les effets hors site. Les résultats ont confirmé la présence d'effets hors site de la mine sur les communautés de sous-bois, et le stade d'exploitation et le type d'écosystème ont été des facteurs déterminants pour l'ampleur des effets hors site. Cependant, nous avons également constaté que les différents groupes de végétation ont réagi différemment aux effets hors site et, par rapport aux autres catégories, les bryophytes semblent être plus touchées en raison de la réduction de la couverture végétale près des sites miniers. Compte tenu des rôles des bryophytes dans les services écologiques de la forêt boréale et de leur grande sensibilité aux changements dans les microhabitats et la pollution, les bryophytes ont été sélectionnées pour déterminer les effets des microhabitats sur les effets hors site et les impacts potentiels sur les groupes fonctionnels (c'est-à-dire les mousses à plumes et les sphaignes) dans le chapitre III. Dans les conclusions du chapitre III, il a été observé que les bryophytes au sol étaient plus affectées par les effets hors site que les groupes sur les autres types de microhabitats. En outre, la couverture au sol de la mousse à plumes a présenté une réponse significative aux types d'écosystèmes et au stade d'exploitation. Sur la base des résultats du chapitre III, nous pensons que les mousses à plumes poussant sur le sol pourraient recevoir plus de contaminants, nous nous sommes donc demandés si le microbiote de la phyllosphère des mousses est également touché ? Pour répondre à cette question, dans le chapitre IV, nous avons sélectionné une espèce de mousse à plumes, *Pleurozium schreberi* pour identifier et déterminer les effets hors site des mines sur le microbiote de la phyllosphère. La thèse a permis d'élargir la

compréhension de différents groupes vivants sur les effets des mines sur la biodiversité et a fourni des implications pour la gestion forestière autour des sites miniers. Les trois chapitres présentent les principales informations relatives aux effets hors site des mines sur les plantes et les microbiomes associés de la phyllosphère dans le sous-étage de la forêt boréale : 1) l'effet hors site individuel est relativement faible et limité dans l'espace, mais son accumulation dans l'espace pourrait être substantielle ; 2) les effets sont plus nombreux à proximité des mines en exploitation que des sites non exploités ; 3) les effets hors site dépendent également du type de forêt et des communautés. Les plantes vasculaires (couverture totale et richesse) étaient plus touchées dans les forêts de feuillus que de conifères, tandis que les bryophytes étaient plus touchées dans les forêts de conifères. Les communautés bactériennes et fongiques de la phyllosphère ont été affectées de manière similaire par les mines, quel que soit le type de forêt ; 3) Les microhabitats peuvent atténuer les effets hors site. Les effets hors site des mines affectent principalement les microhabitats proches du sol ; 4) les effets hors site offrent des opportunités pour certaines espèces tolérantes (y compris la tolérance aux métaux lourds et à la luminosité) de se développer dans les paysages hors site des mines ; 5) les capacités indicatrices (sensibilité) des différents groupes de sous-bois aux effets hors site ont été trouvées : pour les plantes de sous-bois, bryophytes > plantes herbacées > plantes ligneuses ; pour le microbiote, bactéries > champignons. Par conséquent, nos résultats ont atteint les objectifs généraux et spécifiques que nous avons proposés et ont caractérisé les effets hors site des mines dans les paysages boréaux.

5.2 Synthèse des résultats

Les objectifs généraux et spécifiques du projet ont été atteints dans les trois chapitres centraux (chapitre II, III et IV), mais en examinant les résultats des chapitres ensemble pourrait nous aider à mieux comprendre les caractéristiques de ces effets dans les paysages hors site minier. Nous avons comparé les différentes réponses des

trois groupes aux effets hors site et leurs facteurs influencés, le type d'écosystème, le stade d'exploitation minière et les microhabitats. De plus, l'empreinte hors site des mines sur les communautés de sous-bois a finalement été déterminée sur la base de tous les résultats des trois groupes. Selon l'objectif global de notre recherche, la synthèse des résultats comprend deux parties : 1) les effets du stade d'exploitation et du type d'écosystème sur les communautés de sous-bois hors site et le rôle des microhabitats sur ces effets ; 2) la détermination de l'empreinte hors site des mines ; 3) la sensibilité de chaque groupe aux effets hors site des mines.

5.1.1 Les effets du stade d'exploitation et du type d'écosystème

L'interaction des effets combinés du stade d'exploitation et du type d'écosystème dépend des différentes communautés et microhabitats. Des effets significatifs des deux paramètres et de leur interaction ont généralement été trouvés dans les communautés de plantes vasculaires et de bryophytes du sous-bois en termes de composition de la communauté, mais les effets d'interaction sur la richesse des espèces ont été médiés par les microhabitats (Figure 5.1). En effet, seule la richesse des communautés de sous-étage (c'est-à-dire les plantes vasculaires et les bryophytes) sur le sol a démontré des effets d'interaction du stade d'exploitation et du type d'écosystème, tandis que les effets sur la richesse des bryophytes des microhabitats comme des rochers, des débris ligneux, des souches, des chicots et des arbres étaient tamponnés. Nous avons également constaté que seules les communautés des microhabitats inférieurs (c.-à-d. le sol, les rochers et la phyllosphère de la mousse de plumes) ont démontré les effets hors site des mines sur la richesse et la composition des communautés (Figure 5.1). Ces résultats suggèrent que les macrohabitats (type d'écosystème) pourraient être un moteur des effets hors site pour les communautés de sous-bois au sol, tandis que les microhabitats pourraient être le principal moteur pour ceux qui colonisent les microhabitats hors sol, ce qui est cohérent avec des études précédentes montrant que les communautés de sous-bois au sol et les microhabitats

hors sol avaient des réponses différentes aux changements environnementaux (Barbé et al., 2020 ; Caners et al., 2013). Ce phénomène pourrait être attribué à une combinaison de dépôts de poussière accumulée et de changements dans les structures de la canopée. Les communautés du sous-étage sur les microhabitats inférieurs étaient plus exposées aux effets hors site en raison de la poussière atmosphérique (y compris les métaux lourds, les métaux traces et autres polluants) déposée sur les microhabitats supérieurs et les surfaces des feuilles qui sont finalement déposées sur le sol de la forêt par le vent, l'eau de pluie et la défoliation (Sase et al., 2012). Tandis que les microhabitats hors sol ayant des dimensions verticales, pourraient être moins exposés aux changements des variables environnementales (par exemple, la disponibilité de la lumière, la chute des arbres et l'humidité du sol). Par conséquent, nous concluons que l'interaction des effets du stade d'exploitation et du type d'écosystème sur les effets hors site dépend largement des types de microhabitats.

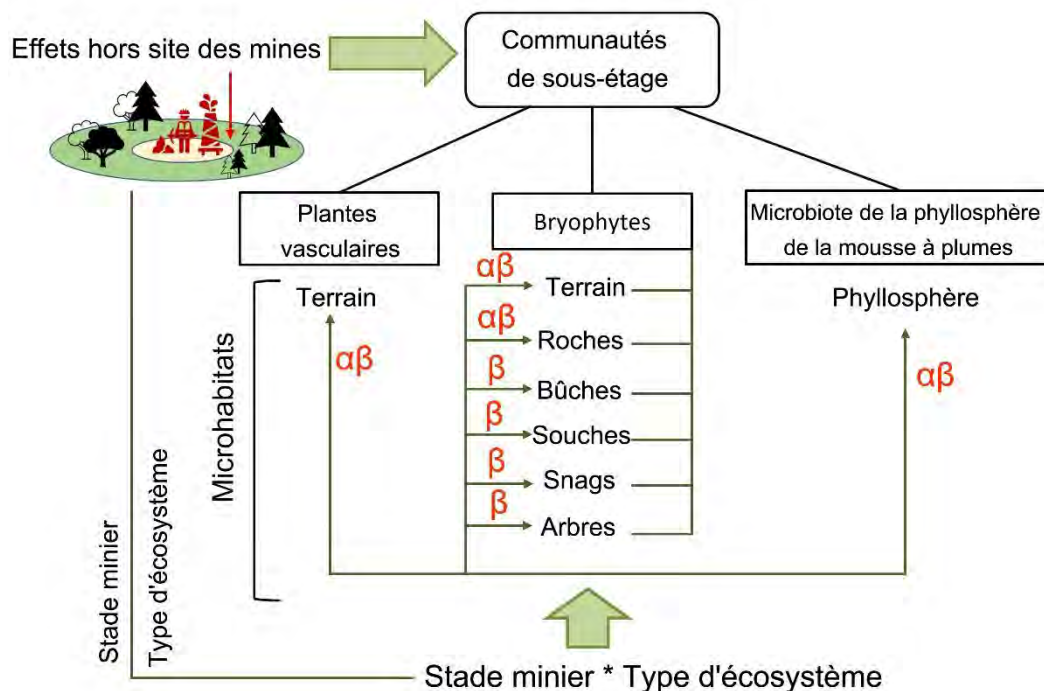


Figure 5.1 Effets de l'interaction stade d'exploitation - type d'écosystème sur les effets

hors site sur les communautés du sous-étage. α signifie que la richesse a été affectée et β signifie que la composition de la communauté a été affectée.

5.1.2 Détermination de l'empreinte hors site des mines

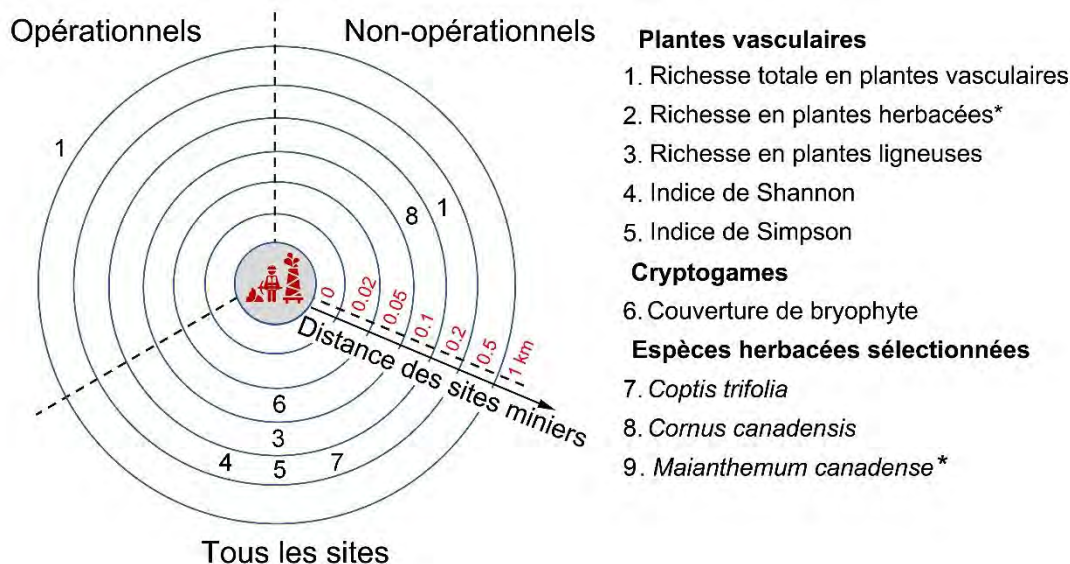


Figure 5.2 La distance d'influence des mines sur les communautés du sous-étage. Seuls les indices qui ont été associés à la distance sont présents dans les graphiques. * Indique qu'aucune distance influencée n'a été détectée

L'évaluation de l'empreinte hors site des mines est un défi dans les évaluations de l'impact écologique des mines, car les effets hors site sont subtils et hors de vue (Raiter et al., 2014). Dans cette étude, l'empreinte hors site a été déterminée en utilisant la distance influencée sur la diversité et la composition des plantes vasculaires, des bryophytes et du microbiote de la phyllosphère des mousses à plumes dans les couches de sous-bois de la forêt boréale. La distance d'influence des effets hors site pouvait atteindre plus d'un kilomètre, en particulier à proximité des mines en exploitation, mais les effets les plus forts se situaient à moins de 0,2 km des mines (Figures 5.2 et 5.3). Par conséquent, les résultats combinés confirment la taille de l'empreinte hors site des mines dans les forêts boréales, ce qui est cohérent avec les

résultats des chapitres 2 et 3. Les résultats indiquent que les mesures d'atténuation des effets hors site devraient se concentrer sur la plage de 0 à 0,2 km de la lisière de la forêt.

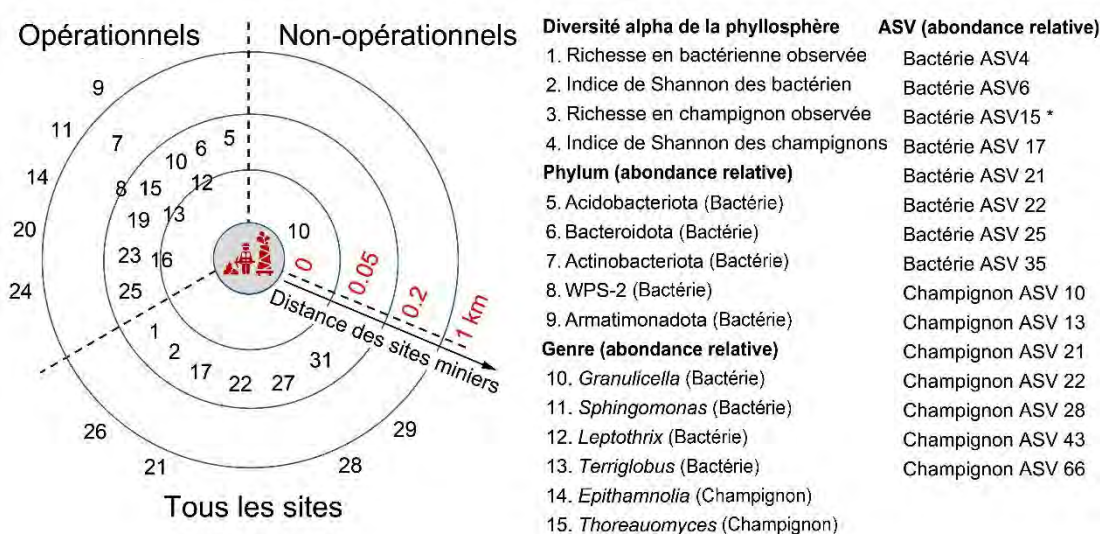


Figure 5.3 Influence de la distance des mines sur les communautés microbiennes de la phyllosphère de *Pleurozium schreberi*. Seuls les groupes/taxa associés à la distance sont présents. * Indique qu'aucune distance influencée n'a été détectée.

5.1.3 Structure des communautés bryophytes et microbiote de la phyllosphère des mousses à plumes

La structure de la communauté bryophyte (couverture et richesse) pourrait être associée au microbiote de la phyllosphère d'après les résultats des chapitres II, III et IV. Premièrement, le couvert total des mousses (chapitre II) et la couverture végétale de la mousse de plumes (chapitre III) ont été réduites dans la forêt de conifères et l'écosystème de canopée ouverte près des mines en exploitation, tandis que certains groupes bactériens fonctionnels de WPS-2 et de Cyanobactéries habitant dans la phyllosphère de la mousse de plumes ont également été réduits dans ces deux écosystèmes près des mines en exploitation. Deuxièmement, la richesse en bryophytes au sol a présenté une forme de "Ω" dans la forêt de conifères et

l'écosystème de canopée ouverte avec une valeur plus élevée près des sites non exploités que des sites d'exploitation et de contrôle (chapitre III), tandis que la richesse bactérienne de la phyllosphère a présenté une forme de "U" avec une valeur plus élevée près des sites non exploités. Ces deux liens suggèrent que la structure de la communauté voisine (par exemple, la couverture végétale et la richesse) pourrait être un moteur pour la structure de la communauté microbienne de la phyllosphère. Une étude a révélé que le taxon bactérien sur l'érable à sucre était positivement corrélé avec l'abondance relative de cet hôte dans le paysage (Lajoie et Steven, 2021). Une autre étude a révélé que l'abondance bactérienne de la phyllosphère était négativement associée à la richesse de la communauté voisine (Meyer et al., 2022). Dans cette étude, pour la diversité alpha de la phyllosphère (par exemple, la richesse), la couverture végétale de l'espèce hôte (*Pleurozium schreberi*) pourrait être un facteur positif, tandis que la richesse totale en bryophytes (richesse de la communauté voisine) pourrait être un facteur négatif. Les liens potentiels indiquent une voie indirecte des effets hors site des mines sur le microbiote de la phyllosphère par le biais de la structure des communautés voisines. D'autres analyses seront effectuées pour confirmer les liens entre la structure de la communauté de la phyllosphère de la mousse à plumes et son voisinage de bryophytes.

5.2 Contributions

Cette thèse propose une approche analytique et méthodologique pour identifier les effets hors site des mines et déterminer leur empreinte dans les paysages boréaux. La contribution de l'étude comprend :

- 1) Confirmation de la présence d'effets hors site des mines sur les communautés du sous-étage des forêts boréales. Les paysages hors site sont relativement intacts et sont facilement et souvent négligés dans les évaluations des impacts (Raiter et al., 2014), mais nos résultats ont quantifié les effets se produisant dans les paysages hors site des

mines. Bien que ces effets soient subtils, ils sont détectables à l'aide des plantes de sous-bois et des microbiomes associés, en particulier pour certains groupes spécifiques, tels que les espèces végétales introduites et les bactéries tolérantes aux métaux. Par conséquent, nos résultats présentent non seulement les effets hors site qui ne doivent pas être oubliés, mais fournissent également des indicateurs potentiellement utiles pour surveiller les effets hors site.

2) Une compréhension préliminaire des caractéristiques clés des effets hors site des mines dans les forêts boréales. Les effets hors site sont une sorte d'impact "énigmatique", hors site et de faible ampleur, mais qui peuvent être importants à l'échelle mondiale. Le cycle de vie de l'exploitation minière, la composition de la forêt et les microhabitats peuvent tous affecter leur force, ce qui permet de déterminer les éléments à inclure dans le futur cadre d'évaluation des effets hors site des mines.

3) Quantification de l'empreinte hors site des mines. L'identification des zones d'influence est essentielle pour évaluer et atténuer les effets hors site des mines. Nos résultats montrent que les effets les plus importants se produisent à moins de 0,2 km des sites miniers. Nous avons estimé l'empreinte hors site des mines à l'aide de ces données. Bien que les zones influencées puissent varier d'une mine à l'autre, notre estimation de l'empreinte constitue une référence pour les décideurs politiques et les parties prenantes dans les processus décisionnels.

4) Stratégies potentielles proposées pour réduire les effets hors site des mines. Étant donné que la forêt de conifères est le peuplement dominant de la forêt boréale et qu'elle présente, selon nos résultats, une résistance relativement élevée aux effets hors site, nous proposons d'utiliser des plantations de conifères indigènes comme zones tampons autour des sites miniers. De plus, un certain niveau de stockage de débris ligneux grossiers près des sites miniers est proposé pour compenser les effets hors site potentiels des mines sur les bryophytes, étant donné leur rôle dans le maintien de la richesse en bryophytes.

5.3 Limites et recherches futures

5.3.1 Limites de la recherche

- 1) Le plan d'échantillonnage nous a limité dans l'exploration des effets de la direction du vent, de la taille de la mine et des installations adjacentes (p. ex. bassin de résidus, amas de stériles, bâtiments et fosses) sur la distance d'influence.
- 2) Il est impossible d'explorer les effets du type d'écosystème sur la distance influencée dans les paysages réels car les paysages entourant les mines sélectionnées sont caractérisés par une mosaïque de parcelles de forêts de feuillus et de conifères.
- 3) La source de pollution des effets hors site des mines n'a pas été détectée dans l'étude. La concentration de poussière à la surface des plantes (en particulier pour les mousses à plumes), les métaux lourds et les nutriments du sol doivent être déterminés pour identifier la source de pollution.
- 4) Les associations entre les variables environnementales (par exemple, les métaux lourds) et la diversité du sous-étage près des sites miniers n'ont pas été évaluées.

5.3.2 Recherches futures

Sur la base des résultats de cette thèse, des recherches supplémentaires sont nécessaires pour mieux comprendre les effets hors site des mines sur les communautés du sous-étage dans les paysages boréaux :

- 1) Évaluer comment ces effets hors site des mines sur le sous-étage et la structure de la phyllosphère affectent les services écologiques des écosystèmes boréaux. En particulier, il est essentiel d'évaluer les effets hors site des mines sur la dynamique du carbone et de l'azote dans la forêt boréale, étant donné que la forêt boréale est manifestement limitée par l'azote et que la végétation et le sol de la forêt boréale

représentent l'un des plus grands réservoirs mondiaux de carbone (Näsholm et al., 1998 ; Strömgren et Mjöfors, 2012).

2) D'évaluer les effets combinés des effets hors site et d'autres perturbations dans les paysages hors site de la mine. L'interaction entre les effets hors site et les autres effets pourrait se faciliter mutuellement. Par exemple, nous avons déjà évalué les effets synergiques entre les effets hors site et les changements prévus dans la composition des forêts. Cependant, d'autres perturbations, notamment les incendies de forêt, la déforestation et les changements globaux de température et de précipitations, devraient être prises en compte dans les recherches futures sur les effets hors site des mines.

3) D'inclure davantage de sites miniers dans les recherches futures. Les sites miniers de différentes régions boréales, y compris l'Europe du Nord et l'Asie du Nord, sont essentiels pour l'acquisition de données complètes, qui peuvent fournir une estimation plus précise de l'empreinte hors site globale des mines dans la forêt boréale.

5.4 Sommaire

Les effets hors site des mines ont affecté les communautés boréales de plantes de sous-bois et le microbiote de la phyllosphère. Les effets ont été déterminés à la fois par le type d'écosystème et le cycle de vie des mines, et les effets les plus forts ont été discutables dans un rayon de 0,2 km des paysages hors site des mines. L'effet énigmatique (c'est-à-dire les effets hors site) a été confirmé et caractérisé pour la première fois dans les paysages boréaux. Ce projet a atteint l'objectif fixé, mais il ne représente que la "pointe de l'iceberg" des effets hors site des mines. Il reste encore beaucoup à découvrir sur les paysages hors site des mines et nos résultats devraient justifier d'autres études associées pour la biodiversité et les services écologiques.

“... This journey has a goal, but it does not stop at any goal, for it has come to realise that today's goal is only the starting point of tomorrow's journey.” - Sri Chinmoy

APPENDICE A

MATÉRIEL SUPPLÉMENTAIRE DU CHAPITRE II

Table S1.1 The information about the selected respective mines

	Akasaba	Canadian Malartic	Casa Berardi	LaRonde	Lapa	Joutel
Stage	Establishing Roads, deforestation and some human activities	Operation	Operation	Operation	Closing	Restored
Main disturbances		Activities associated with mining (e.g., blasting, digging and transportation)			Residual contaminants and some human activities	Residual contaminants (e.g., tailings)
Mine type	Open Pit	Open Pit	Underground & Open Pit	Underground	Underground	Open pit
Commodities	Gold, Copper	Gold, Silver	Gold, Silver	Gold, Silver, Copper, Zinc	Gold	Gold, Copper
Owned/operated by	Agnico Eagle Mines Ltd.	Canadian Malartic Corp.	Hecla Quebec Inc.	Agnico Eagle Mines Ltd	Agnico Eagle Mines Ltd	Agnico Eagle Mines Ltd
Num. Employees	-	699 (2017)	580 (2018)	833 (2017)	165 (2018)	-
Mine life	Preparation since 2014	Operation since 2005	Operation since 1988	Operation since 1988	Closed since 2018	Closed since 1988
Mining Method	-	▪ Truck & Shovel/ Loader	▪ Truck & Shovel/ Loader ▪ Longhole stopping ▪ Timbered stopping ▪ Longitudinal stopping	▪ Transverse open stopping ▪ Longitudinal retreat ▪ Cemented backfill ▪ Paste backfill	▪ Transverse open stopping ▪ Longitudinal retreat ▪ Cemented backfill	-

Notes: information from the platform of Mining Data Online (<https://miningdataonline.com/>).

Table S1.2 Description of the sampling design. The number of plots of each ecosystem type encountered at different distance along transect was included.

Distance (m)	Coniferous	Deciduous	Mixed	Open Canopy	Sum
0	20	9	2	13	44
20	19	7	2	16	44
50	20	4	8	12	44
100	27	2	4	11	44
200	24	1	5	14	44
500	15	5	6	18	44
1000	21	6	4	13	44
Controls	27	14	7	13	61
Sum	173	48	38	110	Total:

Table S1.3 The list of all vascular species identified in all samples (only plants identified at the species level shown)

ID	Species	Genus	Family	Group	Duration	Introduced	Habit
1	<i>Abies balsamea</i> (L.) Mill.	<i>Abies</i>	Pinaceae	Gymnosperm	Perennial	Native	Woody
2	<i>Acer rubrum</i> L.	<i>Acer</i>	Aceraceae	Dicot	Perennial	Native	Woody
3	<i>Acer spicatum</i> Lam.	<i>Acer</i>	Aceraceae	Dicot	Perennial	Native	Woody
4	<i>Achillea millefolium</i> L.	<i>Achillea</i>	Asteraceae	Dicot	Perennial	Introduced	Herbaceous
5	<i>Actaea pachypoda</i> Elliott.	<i>Actaea</i>	Ranunculaceae	Dicot	Perennial	Native	Herbaceous
6	<i>Actaea rubra</i> (Aiton) Willd.	<i>Actaea</i>	Ranunculaceae	Dicot	Perennial	Native	Herbaceous
7	<i>Agrostis gigantea</i> Roth	<i>Agrostis</i>	Poaceae	Monocot	Perennial	Introduced	Herbaceous
8	<i>Alnus alnobetula</i> (Ehrh.) K.Koch	<i>Alnus</i>	Betulaceae	Dicot	Perennial	Native	Woody
9	<i>Alnus incana</i> (L.) Moench	<i>Alnus</i>	Betulaceae	Dicot	Perennial	Native	Woody
10	<i>Alnus viridis</i> A.Gray	<i>Alnus</i>	Betulaceae	Dicot	Perennial	Native	Woody
11	<i>Amelanchier arborea</i> (F.Michx.) Fernald	<i>Amelanchier</i>	Rosaceae	Dicot	Perennial	Native	Woody
12	<i>Amelanchier bartramiana</i> (Tausch) M.Roem.	<i>Amelanchier</i>	Rosaceae	Dicot	Perennial	Native	Woody
13	<i>Amelanchier canadensis</i> Darl.	<i>Amelanchier</i>	Rosaceae	Dicot	Perennial	Native	Woody
14	<i>Amelanchier laevis</i> Wieg.	<i>Amelanchier</i>	Rosaceae	Dicot	Perennial	Native	Woody

Table S1.3 continued

	<i>Anaphalis margaritacea</i> (L.) Benth. & Hook.f.	<i>Anaphalis</i>	Asteraceae	Dicot	Perennial	Native	Herbaceous
15							
16	<i>Andromeda glaucophylla</i> Link	<i>Andromeda</i>	Ericaceae	Dicot	Perennial	Native	Woody
17	<i>Anemonastrum canadense</i> (Juz.) Holub	<i>Anemonastrum</i> <i>m</i>	Ranunculaceae	Dicot	Perennial	Native	Herbaceous
18	<i>Anemone quinquefolia</i> L.	<i>Anemone</i>	Ranunculaceae	Dicot	Perennial	Native	Herbaceous
19	<i>Apocynum androsaemifolium</i> L.	<i>Apocynum</i>	Apocynaceae	Dicot	Perennial	Native	Herbaceous
20	<i>Aralia hispida</i> Vent.	<i>Aralia</i>	Araliaceae	Dicot	Perennial	Native	Herbaceous
21	<i>Aralia nudicaulis</i> Blume	<i>Aralia</i>	Araliaceae	Dicot	Perennial	Native	Herbaceous
22	<i>Arctostaphylos uva-ursi</i> (L.) Spreng.	<i>Arctostaphylos</i>	Ericaceae	Dicot	Perennial	Native	Woody
23	<i>Athyrium filix-femina</i> (L.) Roth	<i>Athyrium</i>	Dryopteridaceae	Fern	Perennial	Native	Herbaceous
24	<i>Betula papyrifera</i> Marshall	<i>Betula</i>	Betulaceae	Dicot	Perennial	Native	Woody
25	<i>Betula pumila</i> L.	<i>Betula</i>	Betulaceae	Dicot	Perennial	Native	Woody
26	<i>Calamagrostis canadensis</i> (Michx.) P.Beauv.	<i>Calamagrostis</i>	Poaceae	Monocot	Perennial	Native	Herbaceous
27	<i>Caltha palustris</i> L.	<i>Caltha</i>	Ranunculaceae	Dicot	Perennial	Native	Herbaceous
28	<i>Capnoides sempervirens</i> (L.) Borkh.	<i>Capnoides</i>	Fumariaceae	Dicot	Perennial	Native	Herbaceous
29	<i>Carex disperma</i> Dewey	<i>Carex</i>	Cyperaceae	Monocot	Perennial	Native	Herbaceous
30	<i>Carex magellanica</i> Lam.	<i>Carex</i>	Cyperaceae	Monocot	Perennial	Native	Herbaceous
31	<i>Carex oligosperma</i> Michx.	<i>Carex</i>	Cyperaceae	Monocot	Perennial	Native	Herbaceous

Table S1.3 continued

32	<i>Carex platyphylla</i> J.Carey	<i>Carex</i>	Cyperaceae	Monocot	Perennial	Native	Herbaceous
33	<i>Carex retrorsa</i> Schwein.	<i>Carex</i>	Cyperaceae	Monocot	Perennial	Native	Herbaceous
34	<i>Carex stipata</i> Muhl. ex Willd.	<i>Carex</i>	Cyperaceae	Monocot	Perennial	Native	Herbaceous
35	<i>Carex tenera</i> Dewey	<i>Carex</i>	Cyperaceae	Monocot	Perennial	Native	Herbaceous
36	<i>Chamaedaphne calyculata</i> (L.) Moench	<i>Chamaedaphne</i>	Ericaceae	Dicot	Perennial	Native	Woody
37	<i>Cicuta bulbifera</i> L.	<i>Cicuta</i>	Apiaceae	Dicot	Perennial	Native	Herbaceous
38	<i>Cinna latifolia</i> (Trevir.) Griseb.	<i>Cinna</i>	Poaceae	Monocot	Perennial	Native	Herbaceous
39	<i>Circaea alpina</i> L.	<i>Circaea</i>	Onagraceae	Dicot	Perennial	Native	Herbaceous
40	<i>Clintonia borealis</i> (Aiton) Raf.	<i>Clintonia</i>	Liliaceae	Monocot	Perennial	Native	Herbaceous
41	<i>Comarum palustre</i> L.	<i>Comarum</i>	Rosaceae	Dicot	Perennial	Native	Herbaceous
42	<i>Coptis trifolia</i> (L.) Salisb.	<i>Coptis</i>	Ranunculaceae	Dicot	Perennial	Native	Herbaceous
43	<i>Cornus canadensis</i> L.	<i>Cornus</i>	Cornaceae	Dicot	Perennial	Native	Herbaceous
44	<i>Cornus sericea</i> L.	<i>Cornus</i>	Cornaceae	Dicot	Perennial	Native	Woody
45	<i>Corylus americana</i> Walter	<i>Corylus</i>	Betulaceae	Dicot	Perennial	Native	Woody
46	<i>Corylus cornuta</i> Marshall	<i>Corylus</i>	Betulaceae	Dicot	Perennial	Native	Woody
47	<i>Cylactis pubescens</i> (Raf.) W.A.Weber	<i>Cylactis</i>	Rosaceae	Dicot	Perennial	Native	Herbaceous

Table S1.3 continued

48	<i>Cyperus esculentus</i> L.	<i>Cyperus</i>	Cyperaceae	Monocot	Perennial	Native	Herbaceous
49	<i>Cypripedium acaule</i> Aiton	<i>Cypripedium</i>	Orchidaceae	Monocot	Perennial	Native	Herbaceous
50	<i>Danthonia compressa</i> Austin	<i>Danthonia</i>	Poaceae	Monocot	Perennial	Native	Herbaceous
51	<i>Delphinium carolinianum</i> Walter	<i>Delphinium</i>	Ranunculaceae	Dicot	Perennial	Native	Herbaceous
52	<i>Dendrolycopodium dendroideum</i> (Michx.) A. Haines	<i>Dendrolycopodium</i> <i>m</i>	Lycopodiaceae	Lycopod	Perennial	Native	Herbaceous
53	<i>Dendrolycopodium obscurum</i> (L.) A. Haines	<i>Dendrolycopodium</i> <i>m</i>	Lycopodiaceae	Lycopod	Perennial	Native	Herbaceous
54	<i>Diervilla Lonicera</i> Mill.	<i>Diervilla</i>	Caprifoliaceae	Dicot	Perennial	Native	Woody
55	<i>Drosera rotundifolia</i> L.	<i>Drosera</i>	Droseraceae	Dicot	Perennial	Native	Herbaceous
56	<i>Dryopteris carthusiana</i> (Vill.) H.P. Fuchs	<i>Dryopteris</i>	Dryopteridaceae	Fern	Perennial	Native	Herbaceous
57	<i>Elymus canadensis</i> L.	<i>Elymus</i>	Poaceae	Monocot	Perennial	Native	Herbaceous
58	<i>Elymus repens</i> (L.) Gould	<i>Elymus</i>	Poaceae	Monocot	Perennial	Introduced	Herbaceous
59	<i>Epigaea repens</i> L.	<i>Epigaea</i>	Ericaceae	Dicot	Perennial	Native	Woody
60	<i>Epilobium angustifolium</i> L.	<i>Epilobium</i>	Onagraceae	Dicot	Perennial	Native	Herbaceous
61	<i>Epilobium leptophyllum</i> Raf.	<i>Epilobium</i>	Onagraceae	Dicot	Perennial	Native	Herbaceous
62	<i>Epilobium palustre</i> L.	<i>Epilobium</i>	Onagraceae	Dicot	Perennial	Native	Herbaceous
63	<i>Epipactis helleborine</i> (L.) Crantz	<i>Epipactis</i>	Orchidaceae	Monocot	Perennial	Introduced	Herbaceous

Table S1.3 continued

64	<i>Equisetum arvense</i> L.	<i>Equisetum</i>	Equisetaceae	Horsetail	Perennial	Native	Herbaceous
65	<i>Equisetum fluviatile</i> L.	<i>Equisetum</i>	Equisetaceae	Horsetail	Perennial	Native	Herbaceous
66	<i>Equisetum pratense</i> Ehrh.	<i>Equisetum</i>	Equisetaceae	Horsetail	Perennial	Native	Herbaceous
67	<i>Equisetum sylvaticum</i> L.	<i>Equisetum</i>	Equisetaceae	Horsetail	Perennial	Native	Herbaceous
68	<i>Eriophorum vaginatum</i> L.	<i>Eriophorum</i>	Cyperaceae	Monocot	Perennial	Native	Herbaceous
69	<i>Eurybia macrophylla</i> (L.) Cass.	<i>Eurybia</i>	Asteraceae	Dicot	Perennial	Native	Herbaceous
70	<i>Fallopia cilioides</i> (Michx.) Holub	<i>Fallopia</i>	Polygonaceae	Dicot	Perennial	Native	Herbaceous
71	<i>Fragaria vesca</i> L.	<i>Fragaria</i>	Rosaceae	Dicot	Perennial	Native	Herbaceous
72	<i>Fragaria virginiana</i> Mill.	<i>Fragaria</i>	Rosaceae	Dicot	Perennial	Native	Herbaceous
73	<i>Galium aparine</i> L.	<i>Galium</i>	Rubiaceae	Dicot	Annual	Introduced	Herbaceous
74	<i>Galium asprellum</i> Michx.	<i>Galium</i>	Rubiaceae	Dicot	Perennial	Native	Herbaceous
75	<i>Galium sylvaticum</i> L.	<i>Galium</i>	Rubiaceae	Dicot	Perennial	Introduced	Herbaceous
76	<i>Gaultheria hispidula</i> (L.) Muhl. ex Bigelow	<i>Gaultheria</i>	Ericaceae	Dicot	Perennial	Native	Woody
77	<i>Gaultheria procumbens</i> L.	<i>Gaultheria</i>	Ericaceae	Dicot	Perennial	Native	Woody
78	<i>Gentiana linearis</i> Froel.	<i>Gentiana</i>	Gentianaceae	Dicot	Perennial	Native	Herbaceous
79	<i>Geocaulon lividum</i> (Richardson) Fernald	<i>Geocaulon</i>	Santalaceae	Dicot	Perennial	Native	Herbaceous

Table S1.3 continued

80	<i>Glyceria striata</i> (Lam.) Hitchc.	<i>Glyceria</i>	Poaceae	Monocot	Perennial	Native	Herbaceous
81	<i>Goodyera repens</i> (L.) R.Br.	<i>Goodyera</i>	Orchidaceae	Monocot	Perennial	Native	Herbaceous
82	<i>Gymnocarpium dryopteris</i> (L.) Newman	<i>Gymnocarpium</i>	Dryopteridaceae	Fern	Perennial	Native	Herbaceous
83	<i>Huperzia lucidula</i> (Michx.) Trevis.	<i>Huperzia</i>	Lycopodiaceae	Lycopod	Perennial	Native	Herbaceous
84	<i>Hypericum perforatum</i> L.	<i>Hypericum</i>	Clusiaceae	Dicot	Perennial	Introduced	Herbaceous
85	<i>Ilex mucronata</i> (L.) M. Powell, Savol. & S. Andrews	<i>Ilex</i>	Aquifoliaceae	Dicot	Perennial	Native	Woody
86	<i>Juncus effusus</i> L.	<i>Juncus</i>	Juncaceae	Monocot	Perennial	Native	Herbaceous
87	<i>Kalmia angustifolia</i> L.	<i>Kalmia</i>	Ericaceae	Dicot	Perennial	Native	Woody
88	<i>Kalmia microphylla</i> (Hook.) A. Heller	<i>Kalmiamicrophylla</i>	Ericaceae	Dicot	Perennial	Native	Woody
89	<i>Lactuca canadensis</i> L.	<i>Lactuca</i>	Asteraceae	Dicot	Perennial	Native	Woody
90	<i>Larix laricina</i> (Du Roi) K.Koch	<i>Larix</i>	Pinaceae	Gymnosperm	Perennial	Native	Woody
91	<i>Leucanthemum vulgare</i> (Vail.) Lam.	<i>Leucanthemum</i>	Asteraceae	Dicot	Perennial	Introduced	Herbaceous
92	<i>Linaria vulgaris</i> Mill.	<i>Linaria</i>	Scrophulariaceae	Dicot	Perennial	Introduced	Herbaceous
93	<i>Linnaea borealis</i> L.	<i>Linnaea</i>	Caprifoliaceae	Dicot	Perennial	Native	Herbaceous
94	<i>Lonicera caerulea</i> L.	<i>Lonicera</i>	Caprifoliaceae	Dicot	Perennial	Native	Woody
95	<i>Lonicera canadensis</i> Barttram ex Marshall	<i>Lonicera</i>	Caprifoliaceae	Dicot	Perennial	Native	Woody

Table S1.3 continued

96	<i>Lonicera hirsuta</i> Eaton	<i>Lonicera</i>	Caprifoliaceae	Dicot	Perennial	Native	Herbaceous
97	<i>Lotus corniculatus</i> L.	<i>Lotus</i>	Fabaceae	Dicot	Perennial	Introduced ^d	Herbaceous
98	<i>Lysimachia borealis</i> (Raf.) U.Manns & Anderb.	<i>Lysimachia</i>	bees	Herb	Perennial	Native	Herbaceous
99	<i>Lysimachia terrestris</i> (L.) Britton, Stems & Poggenb.	<i>Lysimachia</i>	Primulaceae	Dicot	Perennial	Native	Herbaceous
100	<i>Maianthemum canadense</i> Desf.	<i>Maianthemum</i>	Liliaceae	Monocot	Perennial	Native	Herbaceous
101	<i>Maianthemum trifolium</i> (L.) Sloboda	<i>Maianthemum</i>	Liliaceae	Monocot	Perennial	Native	Herbaceous
102	<i>Matteuccia struthiopteris</i> (L.) Tod.	<i>Matteucciastruit hiopteris</i>	Dryopteridaceae	Fern	Perennial	Native	Herbaceous
103	<i>Melilotus albus</i> Medik.	<i>Melilotus</i>	Fabaceae	Dicot	Perennial	Introduced ^d	Herbaceous
104	<i>Mitella nuda</i> L.	<i>Mitella</i>	Saxifragaceae	Dicot	Perennial	Native	Herbaceous
105	<i>Moneses uniflora</i> A.Gray	<i>Moneses</i>	Pyrolaceae	Dicot	Perennial	Native	Herbaceous
106	<i>Oclemena nemoralis</i> (Aiton) Greene	<i>Oclemena</i>	Asteraceae	Dicot	Perennial	Native	Herbaceous
107	<i>Oenothera biennis</i> L.	<i>Oenothera</i>	Onagraceae	Dicot	Biennial	Native	Herbaceous
108	<i>Onoclea sensibilis</i> L.	<i>Onoclea</i>	Dryopteridaceae	Fern	Perennial	Native	Herbaceous
109	<i>Oxalis acetosella</i> L.	<i>Oxalis</i>	Oxalidaceae	Dicot	Perennial	Native	Herbaceous
110	<i>Petasites frigidus</i> (L.) Fr.	<i>Petasites</i>	Asteraceae	Dicot	Perennial	Native	Herbaceous
111	<i>Phegopteris connectilis</i> (Michx.) Watt	<i>Phegopteris</i>	Thelypteridaceae ^e	Fern	Perennial	Native	Herbaceous

Table S1.3 continued

112	<i>Picea glauca</i> (Moench) Voss	<i>Picea</i>	Pinaceae	Gymnosperm	Perennial	Native	Woody
113	<i>Picea mariana</i> (Mill.) Britton, Sterns & Poggenb.	<i>Picea</i>	Pinaceae	Gymnosperm	Perennial	Native	Woody
114	<i>Pilosella aurantiaca</i> (L.) F.W.Schultz & Sch.Bip.	<i>Pilosella</i>	Asteraceae	Dicot	Perennial	Introduced	Herbaceous
115	<i>Pilosella caespitosa</i> (Dumort.) P.D.Sell & C.West	<i>Pilosella</i>	Asteraceae	Dicot	Perennial	Introduced	Herbaceous
116	<i>Pinus banksiana</i> Lamb.	<i>Pinus</i>	Pinaceae	Gymnosperm	Perennial	Native	Woody
117	<i>Poa compressa</i> L.	<i>Poa</i>	Poaceae	Monocot	Perennial	Introduced	Herbaceous
118	<i>Populus balsamifera</i> L.	<i>Populus</i>	Salicaceae	Dicot	Perennial	Native	Woody
119	<i>Populus deltoides</i> Marshall	<i>Populus</i>	Salicaceae	Dicot	Perennial	Native	Woody
120	<i>Populus tremuloides</i> Michx.	<i>Populus</i>	Salicaceae	Dicot	Perennial	Native	Woody
121	<i>Potentilla norvegica</i> L.	<i>Potentilla</i>	Rosaceae	Dicot	Perennial	Native	Herbaceous
122	<i>Prunus pensylvanica</i> L.f.	<i>Prunus</i>	Rosaceae	Dicot	Perennial	Native	Woody
123	<i>Prunus virginiana</i> L.	<i>Prunus</i>	Rosaceae	Dicot	Perennial	Native	Woody
124	<i>Pteridium aquilinum</i> (L.) Kuhn	<i>Pteridium</i>	Dennstaedtiaceae	Fern	Perennial	Native	Herbaceous
125	<i>Pyrola asarifolia</i> Michx.	<i>Pyrola</i>	Pyrolaceae	Dicot	Perennial	Native	Woody
126	<i>Rhamnus alnifolia</i> L.Hér.	<i>Rhamnus</i>	Rhamnaceae	Dicot	Perennial	Native	Woody
127	<i>Rhododendron canadense</i> (L.) Torr.	<i>Rhododendron</i>	Ericaceae	Dicot	Perennial	Native	Woody
128	<i>Rhododendron groenlandicum</i> (Oeder) Kron & Judd	<i>Rhododendron</i>	Ericaceae	Dicot	Perennial	Native	Woody

Table S1.3 continued

129	<i>Ribes hirtellum</i> Michx.	<i>Ribes</i>	Grossulariaceae	Dicot	Perennial	Native	Woody
130	<i>Ribes hudsonianum</i> Richardson	<i>Ribes</i>	Grossulariaceae	Dicot	Perennial	Native	Woody
131	<i>Ribes lacustre</i> (Pers.) Poir.	<i>Ribes</i>	Grossulariaceae	Dicot	Perennial	Native	Woody
132	<i>Ribes ruizii</i> Rehder	<i>Ribes</i>	Grossulariaceae	Dicot	Perennial	Native	Woody
133	<i>Ribes triste</i> Pall.	<i>Ribes</i>	Grossulariaceae	Dicot	Perennial	Native	Woody
134	<i>Rosa acicularis</i> Lindl.	<i>Rosa</i>	Rosaceae	Dicot	Perennial	Native	Woody
135	<i>Rosa blanda</i> Aiton	<i>Rosa</i>	Rosaceae	Dicot	Perennial	Native	Woody
136	<i>Rubus chamaemorus</i> L.	<i>Rubus</i>	Rosaceae	Dicot	Perennial	Native	Herbaceous
137	<i>Rubus idaeus</i> L.	<i>Rubus</i>	Rosaceae	Dicot	Perennial	Native	Woody
138	<i>Salix balsamifera</i> Barratt ex Andersson	<i>Salix</i>	Salicaceae	Dicot	Perennial	Native	Woody
139	<i>Salix bebbiana</i> Sarg.	<i>Salix</i>	Salicaceae	Dicot	Perennial	Native	Woody
140	<i>Sambucus racemosa</i> L.	<i>Sambucus</i>	Caprifoliaceae	Dicot	Perennial	Native	Woody
141	<i>Sarracenia purpurea</i> L.	<i>Sarracenia</i>	Sarraceniaceae	Dicot	Perennial	Native	Herbaceous
142	<i>Scheuchzeria palustris</i> L.	<i>Scheuchzeria</i>	Scheuchzeriaceae	Dicot	Perennial	Native	Herbaceous
143	<i>Scirpus microcarpus</i> J.Presl & C.Presl	<i>Scirpus</i>	Cyperaceae	Monocot	Perennial	Native	Herbaceous
144	<i>Scirpus pendulus</i> Muhl.	<i>Scirpus</i>	Cyperaceae	Monocot	Perennial	Native	Herbaceous
145	<i>Scutellaria galericulata</i> L.	<i>Scutellaria</i>	Lamiaceae	Dicot	Perennial	Native	Herbaceous

Table S1.3 continued

146	<i>Solidago brendae</i> Semple.	<i>Solidago</i>	Asteraceae	Dicot	Perennial	Native	Herbaceous
147	<i>Solidago canadensis</i> L.	<i>Solidago</i>	Asteraceae	Dicot	Perennial	Native	Herbaceous
148	<i>Solidago rugosa</i> Mill.	<i>Solidago</i>	Asteraceae	Dicot	Perennial	Native	Herbaceous
149	<i>Solidago uliginosa</i> Nutt.	<i>Solidago</i>	Asteraceae	Dicot	Perennial	Native	Herbaceous
150	<i>Sonchus oleraceus</i> (L.) L.	<i>Sonchus</i>	Asteraceae	Dicot	Annual	Introduced	Herbaceous
151	<i>Sorbus americana</i> Marshall	<i>Sorbus</i>	Rosaceae	Dicot	Perennial	Native	Woody
152	<i>Sparganium americanum</i> Nutt.	<i>Sparganium</i>	Typhaceae	Monocot	Perennial	Native	Herbaceous
153	<i>Spinulum annotinum</i> (L.) A. Haines	<i>Spinulum</i>	Lycopodiaceae	Lycopod	Perennial	Native	Herbaceous
154	<i>Spiranthes romanzoffiana</i> Cham.	<i>Spiranthes</i>	Orchidaceae	Dicot	Perennial	Native	Herbaceous
155	<i>Symphovotrichum puniceum</i> (L.) A.Löve & D.Löve	<i>Symphovotrichum</i>	Asteraceae	Dicot	Perennial	Native	Herbaceous
156	<i>Symphovotrichum ciliolatum</i> (Lindl.) A.Löve & D.Löve	<i>Symphovotrichum</i>	Asteraceae	Dicot	Perennial	Native	Herbaceous
157	<i>Taraxacum officinale</i> (L.) Weber ex F.H. Wigg.	<i>Taraxacum</i>	Asteraceae	Dicot	Perennial	Native	Herbaceous
158	<i>Taxus canadensis</i> Marshall	<i>Taxus</i>	Taxaceae	Gymnosperm	Perennial	Native	Woody
159	<i>Thalictrum pubescens</i> Pursh	<i>Thalictrum</i>	Ranunculaceae	Dicot	Perennial	Native	Herbaceous
160	<i>Thuja occidentalis</i> L.	<i>Thuja</i>	Cupressaceae	Gymnosperm	Perennial	Native	Woody
161	<i>Trifolium pratense</i> L.	<i>Trifolium</i>	Fabaceae	Dicot	Perennial	Introduced	Herbaceous

Table S1.3 continued

162	<i>Trillium grandiflorum</i> (Michx.) Salisb.	<i>Trillium</i>	Liliaceae	Monocot	Perennial	Native	Herbaceous
163	<i>Tussilago farfara</i> L.	<i>Tussilago</i>	Asteraceae	Dicot	Perennial	Introduced	Herbaceous
164	<i>Typha latifolia</i> L.	<i>Typha</i>	Typhaceae	Monocot	Perennial	Native	Herbaceous
165	<i>Vaccinium angustifolium</i> Aiton	<i>Vaccinium</i>	Ericaceae	Dicot	Perennial	Native	Woody
166	<i>Vaccinium myrtilloides</i> Michx.	<i>Vaccinium</i>	Ericaceae	Dicot	Perennial	Native	Woody
167	<i>Vaccinium oxycoccos</i> L.	<i>Vaccinium</i>	Ericaceae	Dicot	Perennial	Native	Woody
168	<i>Viburnum lentago</i> L.	<i>Viburnum</i>	Caprifoliaceae	Dicot	Perennial	Native	Woody
169	<i>Viburnum nudum</i> L.	<i>Viburnum</i>	Caprifoliaceae	Dicot	Perennial	Native	Woody
170	<i>Vicia cracca</i> L.	<i>Vicia</i>	Fabaceae	Dicot	Perennial	Introduced	Herbaceous
171	<i>Viola macloskeyi</i> F.E. Lloyd	<i>Viola</i>	Violaceae	Dicot	Perennial	Native	Herbaceous
172	<i>Viola rostrata</i> Pursh	<i>Viola</i>	Violaceae	Dicot	Perennial	Native	Herbaceous

Table S1.4 The list of environmental and stand structure variables (continuous variables with minimum, maximum, and average values; categorical variables with count number).

Variables/Treatment	Variable Type	Unite	Range
Distance from mine edge	Continuous	m	0, 20, 50, 100, 200, 500, 1000 m (N=44 for each distance)
Live crown ratio for shrubs ¹	Continuous	%	Min=0; Max=100; Average=21.54
Live crown ratio for trees ¹	Continuous	%	Min=0; Max=100; Average=24.14
Base areas for shrubs	Continuous	m ² /ha	Min=0; Max=7.084746; Average=1.27
Base areas for trees	Continuous	m ² /ha	Min=0; Max=58; Average=14.34
Canopy openness ²	Categorical	-	Min=1; Max=4; Average=1.13; 4 levels *
Leave litter ground cover	Continuous	%	Min=0; Max=100; Average=34.60
Woody debris ground cover	Continuous	%	Min=0; Max=92.67; Average=14.43
Water ground cover	Continuous	%	Min=0; Max=24.33; Average=2.60
Rock ground cover	Continuous	%	Min=0; Max=88; Average=7.17
Organic matter ground cover	Continuous	%	Min=0; Max=70.33; Average=5.44
Depth of organic soil	Continuous	cm	Min=0; Max=128.333; Average=30.92
Conifer proportion ³	Continuous	%	Min=0; Max=100; Average=44.30
Humidity	Continuous	-	Min=1; Max=10; Average=2.58
Bryophyte cover	Continuous	%	Min=0; Max=100; Average=34.83
Lichen cover	Continuous	%	Min=0; Max=78.33; Average=8.08
Ecosystem types	Categorical	-	Coniferous (N=173), Mixed (N=38), Deciduous (N=48), Open Canopy (N=110)
Mining stages	Categorical	-	Operating (N=168), Non-operating (N=140), Controls (N=61)

Notes: 1. Live crown ratio (%) was calculated as $(H-HCB)/H$, where H is the total height of the tree/shrub, and HCB is the height from crown top to base (Kuprevicius, Auty et al. 2013). 2. Canopy openness was measured with a spherical densiometer at 1 m height above ground and recorded as 1-4 levels (L1, 0 % (0 grid); L2, 1% - 40 % (1 - 9.6 grids); L3, 40 % - 70 % (9.7 - 16.8 grids); L4: > 70% (>16.8 grids). 3. Conifer proportion (%) was calculated by the number of individuals of coniferous trees in each plot.

Table S1.5 Comparison and differences in measured forest environmental and stand structure variables at different distance along transects (mean (SD)).

	0 m	20 m	50 m	100 m	200 m	500 m	1000 m	Controls	chi-squared	p-value
	N=44	N=44	N=44	N=44	N=44	N=44	N=44	N=61		
Soil humidity	5.22(3.124)	4.752(3.152)	4.917(2.801)	4.5(2.429)	4.345(2.292)	4.235(2.025)	4.322(2.403)	4.454(2.323)	2.518	0.926
Live crown ratio for shrubs	38.824 (17.744)	42.385 (19.591)	45.211 (21.197)	43.787 (20.909)	44.189 (24.894)	49.132 (23.896)	43.558 (20.764)	44.031 (22.458)	4.102	0.768
Base areas for shrubs	1.868 (1.45)	2.016 (1.509)	1.815 (1.026)	1.687 (1.236)	1.746 (1.289)	1.619 (1.423)	1.497 (1.177)	1.317 (1.002)	10.666	0.154
Live crown ratio for trees	29.648 (26.468)	31.876 (28.039)	27.893 (22.837)	30.45 (23.771)	27.869 (25.679)	23.498 (22.214)	29.247 (22.444)	26.961 (22.563)	3.006	0.885
Base areas for trees	11.636 (12.592)	11.773 (12.819)	15.473 (15.14)	16.209 (16.335)	15.302 (16.914)	12.338 (14.899)	14.697 (15.387)	12.689 (11.248)	3.799	0.803
Depth of organic soil	27.455 (25.116)	37.944 (33.116)	37.032 (28.279)	39.109 (32.347)	39.051 (31.301)	36.465 (31.706)	36.92 (32.281)	46.97 (31.251)	12.840	0.076
Leave litter ground cover	53.83 (34.017)	62.242 (33.076)	51.318 (30.767)	46.432 (31.485)	48.389 (33.662)	45.909 (36.674)	47.788 (35.97)	41.216 (37.617)	13.305	0.065
Rock ground cover	3.22 (13.553)	0.587 (3.669)	2.617 (8.866)	1.182 (5.812)	0.023 (0.151)	2.182 (9.742)	0.773 (4.775)	0.17 (1.062)	19.610	0.006
Water ground cover	0.833 (2.936)	0.417 (1.802)	0.561 (3.668)	0.811 (3.235)	0.235 (1.558)	0.106 (0.537)	0.235 (1.07)	0.992 (3.536)	7.485	0.380
Woody debris ground cover	17.799 (18.719)	16.836 (12.862)	18.654 (15.126)	17.773 (16.277)	14.183 (11.22)	15.623 (16.448)	14.872 (14.429)	15.179 (10.321)	4.935	0.668
Organic matter ground cover	1.595 (3.952)	0.288 (0.852)	1.547 (4.041)	1.102 (3.531)	0.492 (1.729)	2.386 (8.372)	2.318 (11.238)	0.563 (1.727)	5.291	0.625
Conifer proportion	43.612 (42.937)	48.438 (45.266)	54.454 (43.808)	62.561 (44.798)	57.841 (44.687)	41.635 (43.498)	49.795 (44.67)	51.78 (44.181)	6.969	0.432

Notes: ANOVA analysis with significant at $P < 0.05$.

Table S1.6 The list of understory diversity indexes (continuous variables with minimum, maximum and average values; Categorical variables with count number).

Responses	Variable	Type	Unite	Range
All vascular plant richness	Count	-		Min=2; Max=26; Average=3.89
Woody plant richness	Count	-		Min=0; Max=13; Average=2.43
Herb plant richness	Count	-		Min=0; Max=14; Average=2.93
Shannon	Continuous	-		Min=0.03; Max=2.62; Average=0.46
Simpson	Continuous	-		Min=0.03; Max=2.61; Average=1.67
All vascular plant cover	Continuous	%		Min=3.33; Max=171.33; Average=33.30
Woody plant cover	Continuous	%		Min=0; Max=121; Average=24.38
Herb plant cover	Continuous	%		Min=0; Max=127; Average=24.82
Bryophyte cover*	Continuous	%		Min=0; Max=100; Average=34.83
Lichen cover*	Continuous	%		Min=0; Max=78.33; Average=8.08

* The cover of bryophytes and lichens are both response and explanatory variables (for vascular plants)

Table S1.7 Results of mixed models for the plant richness and diversity indices testing for the effect of mine stage, ecosystem type and their interaction. Non-significant interaction terms were determined by Anova (Type=II) and deleted from the final models. Significant parameters (≤ 0.05) are indicated in bold.

	All vascular richness		Simpson		Shannon		Woody richness		Herbaceous richness	
	Estimate	<i>p</i>	Estimate	<i>p</i>	Estimate	<i>p</i>	Estimate	<i>p</i>	Estimate	<i>p</i>
(Intercept)	2.159	<0.001	-0.288	<0.001	0.518	<0.001	1.661	<0.001	1.230	<0.001
Mining stages (Reference: Controls)										
Operating	0.043	0.684	-0.027	0.215	-0.051	0.177	-0.133	0.493	0.181	0.254
Non-operating	0.013	0.904	-0.017	0.452	-0.018	0.648	0.060	0.756	-0.019	0.906
Ecosystem types (Reference: Coniferous)										
Deciduous	0.338	<0.001	0.033	0.165	0.092	0.014	0.023	0.725	0.755	<0.001
Mixed	0.277	0.037	0.038	0.168	0.098	0.025	-0.191	0.021	0.870	<0.001
Open Canopy	0.166	0.136	0.038	0.033	0.079	0.006	0.177	<0.001	0.009	0.966
Mining stages: Ecosystem types										
Deciduous: Operating	-0.375	0.007	-	-	-	-	-	-	-0.744	0.001
Deciduous: Non-operating	-0.038	0.791	-	-	-	-	-	-	-0.658	0.008
Mixed: Operating	-0.328	0.068	-	-	-	-	-	-	-0.543	0.043
Mixed: Non-operating	-0.062	0.692	-	-	-	-	-	-	-0.384	0.129
Open Canopy: Operating	-0.027	0.829	-	-	-	-	-	-	0.156	0.508
Open Canopy: Non-operating	0.072	0.584	-	-	-	-	-	-	0.215	0.383
Anova (Type=II)										
Factors	Chisq.	<i>P</i>	Chisq.	<i>P</i>	Chisq.	<i>P</i>	Chisq.	<i>P</i>	Chisq.	<i>P</i>
Mining stages	1.03	0.596	1.59	0.451	2.28	0.319	1.98	0.371	4.53	0.104
Ecosystem types	25.71	<0.001	5.79	0.122	12.72	0.005	22.70	<0.001	32.74	<0.001
Mining stages: Ecosystem types	12.67	0.049	5.05	0.538	5.09	0.533	5.93	0.431	18.80	0.005

Table S1.8 Results of mixed models on the plant cover testing for the effect of mining stages, ecosystem types and their interactions. Non-significant interaction terms were determined by Anova (Type=II) and deleted from the final models. Significant parameters (≤ 0.05) are indicated in bold.

	All vascular cover		Woody cover		Herbaceous cover		Bryophyte cover		Lichen cover	
	Estimate	<i>p</i>	Estimate	<i>p</i>	Estimate	<i>p</i>	Estimate	<i>p</i>	Estimate	<i>p</i>
(Intercept)	-0.385	<0.001	0.408	<0.001	0.272	<0.001	0.642	<0.001	-2.802	<0.001
Mining stages (Reference: Controls)										
Operating	-0.067	0.582	-0.036	0.591	-0.034	0.611	-0.233	0.016	-0.052	0.851
Non-operating	-0.038	0.750	0.014	0.835	-0.021	0.754	-0.215	0.027	0.087	0.750
Ecosystem types (Reference: Coniferous)										
Deciduous	0.191	0.161	0.021	0.562	0.056	0.410	-0.508	<0.001	0.163	0.678
Mixed	0.353	0.021	-0.114	0.005	0.278	0.021	-0.288	0.008	0.164	0.809
Open Canopy	0.088	0.570	0.052	0.059	-0.002	0.978	-0.059	0.608	1.556	0.006
Mining stages: Ecosystem types										
Deciduous:	-0.356	0.068			-0.100	0.243	0.280	0.013	-0.806	0.184
Deciduous: Non-operating	-0.047	0.791			0.102	0.259	0.264	0.022	-0.252	0.751
Mixed: Operating	-0.929	0.002			-0.394	0.004	0.152	0.257	-0.491	0.539
Mixed: Non-operating	-0.385	0.050			-0.287	0.030	0.215	0.091	-0.278	0.705
Open Canopy: Operating	-0.020	0.909			0.048	0.559	0.036	0.772	-1.827	0.002
Open Canopy: Non-operating	0.095	0.593			0.002	0.982	0.179	0.160	-1.534	0.011
Anova (Type=II)										
Factors	Chisq.	<i>P</i>	Chisq.	<i>P</i>	Chisq.	<i>P</i>	Chisq.	<i>P</i>	Chisq.	<i>P</i>
Mining stages	5.86	0.053	0.75	0.687	1.08	0.583	7.29	0.026	6.94	0.031
Ecosystem types	6.25	0.100	14.92	0.002	3.40	0.335	44.40	<0.001	2.70	0.439
Mining stages: Ecosystem types	16.13	0.013	1.812	0.936	17.44	0.008	18.21	0.006	12.90	0.045

Notes: in each case the interaction term was not significant and was dropped from the model.

Table S1.9 Influence of mining stage, ecosystem type, and their interaction on understory vascular community structure. Permutational multivariate analysis of variance (PERMANOVA) are used based on the Bray-Curtis dissimilarity index for all vascular, woody, and herbaceous plants.

Assemblage	Variable	Df	Mean square	<i>F</i> statistics	R ²	<i>P</i>
All vascular plants	Ecosystem types	3	3.142	8.588	0.065	0.001
	Mining Stages	2	1.219	3.331	0.017	0.001
	Ecosystem: Stages	6	0.545	1.490	0.022	0.002
	Residuals	357	0.366		0.896	
Woody plants	Ecosystem types	3	3.388	9.370	0.071	0.001
	Mining Stages	2	1.048	2.898	0.015	0.001
	Ecosystem: Stages	6	0.522	1.445	0.022	0.007
	Residuals	348	0.362		0.892	
Herbaceous plants	Ecosystem types	3	2.228	5.714	0.046	0.001
	Mining Stages	2	1.642	4.209	0.023	0.001
	Ecosystem: Stages	6	0.606	1.555	0.025	0.002
	Residuals	331	0.390		0.906	

Table S1.10. PERMANOVA post hoc pair-wise comparisons of mine stages on understory structure. Community structure of woody (A), herbaceous (B) plants was tested in each ecosystem type based on 999 permutations with Benjamini-Hochberg adjustment (BH).

A) Woody plants

Pairs	Df	Sums of squares	F.Model	R ²	<i>p</i>	<i>p.adj</i>
Coniferous forest						
Operating vs Nonoperating	1	0.74	2.01	0.01	0.018	0.026
Operating vs Controls	1	0.82	2.32	0.02	0.015	0.023
Nonoperating vs Controls	1	0.64	1.73	0.02	0.045	0.057
Mixed forest						
Operating vs Nonoperating	1	0.40	1.09	0.04	0.328	0.328
Operating vs Controls	1	0.98	2.57	0.13	0.004	0.007
Nonoperating vs Controls	1	0.70	1.81	0.07	0.031	0.044
Deciduous forest						
Operating vs Nonoperating	1	0.71	1.71	0.05	0.041	0.054
Operating vs Controls	1	0.68	1.65	0.05	0.053	0.065
Nonoperating vs Controls	1	0.74	1.90	0.07	0.042	0.054
Open canopy ecosystem						
Operating vs Nonoperating	1	0.49	1.42	0.02	0.145	0.154
Operating vs Controls	1	0.59	1.70	0.02	0.069	0.083
Nonoperating vs Controls	1	0.47	1.57	0.03	0.106	0.121

(B) Herbaceous plants

Pairs	Df	Sums of squares	F.Model	R ²	<i>p</i>	<i>p.adj</i>
Coniferous forest						
Operating vs Nonoperating	1	0.99	2.61	0.02	0.004	0.006
Operating vs Controls	1	0.39	0.98	0.01	0.443	0.443
Nonoperating vs Controls	1	0.41	1.11	0.01	0.306	0.311
Mixed forest						
Operating vs Nonoperating	1	0.63	1.82	0.07	0.044	0.055
Operating vs Controls	1	0.52	1.44	0.09	0.132	0.145
Nonoperating vs Controls	1	0.48	1.33	0.06	0.179	0.194
Deciduous forest						
Operating vs Nonoperating	1	1.01	2.56	0.08	0.008	0.012
Operating vs Controls	1	0.46	1.18	0.04	0.270	0.278
Nonoperating vs Controls	1	1.10	3.15	0.11	0.003	0.005
Open canopy ecosystem						
Operating vs Nonoperating	1	1.65	3.91	0.04	0.001	0.002
Operating vs Controls	1	1.29	3.00	0.04	0.001	0.002
Nonoperating vs Controls	1	1.18	3.09	0.06	0.001	0.002

Table S1.11 Results of mixed models on the understory diversity testing for the effect of distance from mine sites as well as its interaction with mining stages. Non-significant interaction terms were deleted from the final model. Significant parameters (≤ 0.05) are indicated in bold.

	(Intercept)		Distance		Distance: Mining stage (Operating)	
	Estimate	<i>P</i>	Estimate	<i>p</i>	Estimate	<i>p</i>
All vascular richness	2.222	<0.001	0.322	<0.001	-0.349	0.001
Simpson	-0.346	<0.001	0.082	0.025	-	-
Shannon	0.487	<0.001	0.105	0.012	-	-
Woody richness	1.708	<0.001	0.168	0.006	-	-
Herbaceous richness	1.367	<0.001	0.381	0.001	-0.506	0.001
Bryophyte cover	0.347	<0.001	0.137	0.001	-	-
Lichen cover	-2.878	<0.001	0.167	0.346	-	-
All vascular cover	-0.421	<0.001	0.101	0.168	-	-
Woody cover	0.421	<0.001	0.036	0.356	-	-
Herbaceous cover	0.273	<0.001	0.036	0.319	-	-

Notes: Non-operating sites as references for mining stages.

Table S1.12 Disturbed areas by mines including directed mined areas and estimated offsite disturbance areas (ha).

	Akasaba	Canadian Malartic	Casa Berardi	Lapa	LaRonde	Joutel	Total	Average (percentage)
Mining Stage	Non- operating	Operating	Operating	Non- operating	Operating	Non- operating		
Directed disturbance areas	103.90	2046.37	449.39	13.06	552.59	229.31	3394.61	565.77 (67.18%)
Offsite disturbance areas (0.2 km	86.10	473.70	536.79	52.59	345.28	163.87	1658.32	276.39 (32.82%)
Total disturbance areas	189.99	2520.07	986.18	65.64	897.87	393.18	5052.94	842.16



Figure S1.1 The maps of six selected mine sites with the position of transects. Scale was not present because the length of transect is 1 km which could be as references to the maps.

APPENDICE B

MATÉRIEL SUPPLÉMENTAIRE DU CHAPITRE III

Table S2.1 Description of the sampling design. Only the number of plots of each ecosystem type encountered at different distance along transect was included.

Distance (m)	Coniferous	Deciduous	Mixed	Open Canopy	Sum
0	20	9	2	13	44
20	19	7	2	14	42
50	20	4	7	12	43
100	27	2	4	11	44
200	24	1	5	13	43
500	15	4	6	18	43
1000	21	6	4	13	44
Controls	27	14	7	13	61
Sum	173	47	37	107	Total: 364

Table S2.2 Criteria used to classify the six microhabitats investigated for bryophytes

Microhabitat	Criteria
Ground	Forest floor not occupied by other microhabitats
Tree (trunk)	DBH $\geq$ 8 cm
Logs	DBH $\geq$ 8 cm and length $\geq$ 1 m
Stump	Diameter $\geq$ 8 m (measured at top) and height $<$ 1.5 m
Snags	DBH $\geq$ 8 cm and height $\geq$ 1.5 m
Rocks	Shortest diameter $\geq$ 20 cm (measured at top)

Table S2.3 Species list in all sampled plots in all studied sites

ID	Species	Abbreviation	Group	Genus
1	<i>Amblystegium serpens</i> (Hedw.) Schimp.	Amb.serp	Moss	<i>Amblystegium</i>
2	<i>Anastrophyllum hellerianum</i> (Nees ex Lindenb.) R.M. Schust.	Ana.hell	Moss	<i>Anastrophyllum</i>
3	<i>Anastrophyllum michauxii</i> (F. Weber) H. Buch ex A. Evans	Ana.mich	Moss	<i>Anastrophyllum</i>
4	<i>Aneura pinguis</i> (L.) Dumort.	Ane.ping	Liverwort	<i>Aneura</i>
5	<i>Atrichum angustatum</i> (Brid.) Bruch & Schimp.	Atr.angu	Moss	<i>Atrichum</i>
6	<i>Aulacomnium palustre</i> (Hedw.) Schwägr.	Aul.palu	Moss	<i>Aulacomnium</i>
7	<i>Barbilophozia attenuate</i> (Nees) Loeske	Bar.atte	Liverwort	<i>Barbilophozia</i>
8	<i>Barbilophozia barbata</i> (Schreb.) Loeske	Bar.barb	Liverwort	<i>Barbilophozia</i>
9	<i>Barbilophozia kunzeana</i> (Huebener) K. Müller	Bar.kunz	Liverwort	<i>Barbilophozia</i>
10	<i>Bazzania denudata</i> (Torr. ex Gottsche, Lindenb. & Nees) Trevis.	Baz.denu	Liverwort	<i>Bazzania</i>
11	<i>Bazzania trilobata</i> (L.) Gray	Baz.tril	Liverwort	<i>Bazzania</i>
12	<i>Blepharostoma trichophyllum</i> (L.) Dumort.	Ble.tric	Liverwort	<i>Blepharostoma</i>
13	<i>Brachythecium acutum</i> (Mitt.) Sull.	Bra.acut	Moss	<i>Brachythecium</i>
14	<i>Brachythecium albicans</i> (Hedw.) Schimp.	Bra.albi	Moss	<i>Brachythecium</i>
15	<i>Brachythecium campestre</i> (Müll. Hal.) Schimp.	Bra.camp	Moss	<i>Brachythecium</i>
16	<i>Brachythecium curtum</i> (Lindb.) Limpr.	Bra.oedi	Moss	<i>Brachythecium</i>
17	<i>Brachythecium erythrorrhizon</i> Schimp.	Bra.eryt	Moss	<i>Brachythecium</i>
18	<i>Brachythecium falcatum</i> (Grout) H.A. Crum	Bra.falc	Moss	<i>Brachythecium</i>
19	<i>Brachythecium populeum</i> (Hedw.) Schimp.	Bra.popu	Moss	<i>Brachythecium</i>
20	<i>Brachythecium reflexum</i> (Starke) Schimp.	Bra.refl	Moss	<i>Brachythecium</i>
21	<i>Brachythecium rivulare</i> Schimp.	Bra.rivu	Moss	<i>Brachythecium</i>
22	<i>Brachythecium rotaeanum</i> De Not.	Bra.rota	Moss	<i>Brachythecium</i>

Table S2.3 continued

23	<i>Brachythecium rutabulum</i> (Hedw.) Schimp.	Bra.ruta	Moss	<i>Brachythecium</i>
24	<i>Brachythecium starkei</i> (Brid.) Schimp.	Bra.star	Moss	<i>Brachythecium</i>
25	<i>Brachythecium velutinum</i> (Hedw.) Schimp.	Bra.velu	Moss	<i>Brachythecium</i>
26	<i>Breidleria pratensis</i> (Koch ex Spruce) Loeske	Hyp.prat	Moss	<i>Breidleria</i>
27	<i>Bryum creberrimum</i> Taylor	Bry.creb	Moss	<i>Bryum</i>
28	<i>Bryhnia novae-angliae</i> (Sull. & Lesq.) Grout	Bry.nova	Moss	<i>Bryhnia</i>
29	<i>Bryum pseudotriquetrum</i> (Hedw.) P. Gaertn., B. Mey. & Scherb.	Bry.pseu	Moss	<i>Bryum</i>
30	<i>Bryoeruthrophyllum recurvirostrum</i> (Hedwig) P. C. Chen	Bry.recu	Moss	<i>Bryoeruthrophyllum</i>
31	<i>Calliergon cordifolium</i> (Hedw.) Kindb.	Cal.cord	Moss	<i>Calliergon</i>
32	<i>Calypogeia fissa</i> (L.) Raddi	Cal.fiss	Liverwort	<i>Calypogeia</i>
33	<i>Callicladium haldanianum</i> (Grev.) H.A. Crum	Cal.hald	Moss	<i>Callicladium</i>
34	<i>Calypogeia integristipula</i> Stephani	Cal.inte	Liverwort	<i>Calypogeia</i>
35	<i>Calypogeia muelleriana</i> (Schiffner) K. Müller	Cal.muel	Liverwort	<i>Calypogeia</i>
36	<i>Calypogeia neesiana</i> (C. Massal. & Carestia) K. Müller	Cal.nees	Liverwort	<i>Calypogeia</i>
37	<i>Calliergon richardsonii</i> (Mitt.) Kindb.	Cal.rich	Liverwort	<i>Calliergon</i>
38	<i>Calypogeia sphagnicola</i> (Arnell & J. Perss.) Warnst. & Loeske	Cal.spha	Liverwort	<i>Calypogeia</i>
39	<i>Campyliadelphus chrysophyllus</i> (Brid.) R.S. Chopra	Cam.chry	Moss	<i>Campyliadelphus</i>
40	<i>Campylophyllum halleri</i> (Sw. ex Hedw.) M. Fleisch	Cam.hall	Moss	<i>Campylophyllum</i>
41	<i>Campylophyllum hispidulum</i> (Brid.) Hedenäs	Cam.hisp	Moss	<i>Campylophyllum</i>
42	<i>Campylium stellatum</i> (Hedw.) C.E.O. Jensen	Cam.stel	Moss	<i>Campylium</i>
43	<i>Cephalozia bicuspidata</i> (L.) Dumort.	Cep.bicu	Liverwort	<i>Cephalozia</i>
44	<i>Cephalozia catenulata</i> (Huebener) Lindb.	Cep.cate	Liverwort	<i>Cephalozia</i>
45	<i>Cephalozia connivens</i> (Dicks.) Lindb.	Cep.conn	Liverwort	<i>Cephalozia</i>

Table S2.3 continued

46	<i>Cephalozia elachista</i> (J.B. Jack ex Gottsche & Rabenh.) Schiffn.	Cep.elac	Liverwort	<i>Cephalozia</i>
47	<i>Cephalozia loitlesbergeri</i> Schiffner	Cep.loit	Liverwort	<i>Cephalozia</i>
48	<i>Cephalozia lunulifolia</i> (Dumort.) Dumort.	Cep.lunu	Liverwort	<i>Cephalozia</i>
49	<i>Cephalozia planiceps</i> (Aust.) Lindb.	Cep.plen	Liverwort	<i>Cephalozia</i>
50	<i>Cephaloziella rubella</i> (Nees) Warnst.	Cep.rube	Liverwort	<i>Cephaloziella</i>
51	<i>Ceratodon purpureus</i> (Hedw.) Brid.	Cer.purp	Moss	<i>Ceratodon</i>
52	<i>Chiloscyphus coadunatus</i> (Sw.) R.M. Schust. & J.J. Engel	Chi.coad	Liverwort	<i>Chiloscyphus</i>
53	<i>Chiloscyphus cuspidatus</i> (Nees) J.J. Engel & R.M. Schust.	Chi.cusp	Liverwort	<i>Chiloscyphus</i>
54	<i>Chiloscyphus profundus</i> (Nees) J.J. Engel & R.M. Schust.	Chi.prof	Liverwort	<i>Chiloscyphus</i>
55	<i>Cladopodiella fluitans</i> (Nees) Jörg.	Cla.flui	Liverwort	<i>Cladopodiella</i>
56	<i>Climacium dendroides</i> (Hedw.) F. Weber & D. Mohr	Cli.dend	Moss	<i>Climacium</i>
57	<i>Dicranum flagellare</i> Hedw.	Dic.flag	Moss	<i>Dicranum</i>
58	<i>Dicranum fulvum</i> Hook.	Dic.fulv	Moss	<i>Dicranum</i>
59	<i>Dicranum fuscescens</i> Turner	Dic.fusc	Moss	<i>Dicranum</i>
60	<i>Dicranella heteromalla</i> (Hedw.) Schimp.	Dic.hete	Moss	<i>Dicranella</i>
61	<i>Dicranum leioneuron</i> Kindb.	Dic.leio	Moss	<i>Dicranum</i>
62	<i>Dicranum majus</i> Turner	Dic.maju	Moss	<i>Dicranum</i>
63	<i>Dicranum montanum</i> Hedw.	Dic.mont	Moss	<i>Dicranum</i>
64	<i>Dicranum ontariense</i> W.L. Peterson	Dic.onta	Moss	<i>Dicranum</i>
65	<i>Dichodontium pellucidum</i> (Hedw.) Schimp.	Dic.pell	Moss	<i>Dichodontium</i>
66	<i>Dicranum polysetum</i> Sw.	Dic.poly	Moss	<i>Dicranum</i>
67	<i>Dicranum scoparium</i> Hedw.	Dic.scop	Moss	<i>Dicranum</i>
68	<i>Dicranum spurium</i> Hedw.	Dic.spur	Moss	<i>Dicranum</i>
69	<i>Dicranum undulatum</i> Schrad. ex Brid.	Dic.undu	Moss	<i>Dicranum</i>

Table S2.3 continued

70	<i>Distichium capillaceum</i> (Hedwig) Bruch & Schimper	Dis.capi	Moss	<i>Distichium</i>
71	<i>Drepanocladus aduncus</i> (Hedw.) Warnst.	Dre.adun	Moss	<i>Drepanocladus</i>
72	<i>Eurhynchium pulchellum</i> (Hedw.) Jenn.	Eur.pulc	Moss	<i>Eurhynchium</i>
73	<i>Fissidens osmundoides</i> Hedw.	Fis.osmu	Moss	<i>Fissidens</i>
74	<i>Frullania oakesiana</i> Austin	Fru.oake	Liverwort	<i>Frullania</i>
75	<i>Geocalyx graveolens</i> (Schrader) Nees	Geo.grav	Liverwort	<i>Geocalyx</i>
76	<i>Gymnocolea inflata</i> (Huds.) Dumort.	Gym.infl	Liverwort	<i>Gymnocolea</i>
77	<i>Herzogiella striatella</i> (Brid.) Z. Iwats.	Her.stri	Moss	<i>Herzogiella</i>
78	<i>Herzogiella turfacea</i> (Lindb.) Z. Iwats.	Her.turf	Moss	<i>Herzogiella</i>
79	<i>Homalia trichomanoides</i> (Hedw.) Schimp.	Hom.tric	Moss	<i>Homalia</i>
80	<i>Hygroamblystegium varium</i> (Hedwig) Mönkemeyer	Hyg.vari	Moss	<i>Hygroamblystegium</i>
81	<i>Hylocomium brevirostre</i> (Brid.) Schimp.	Hyl.brev	Moss	<i>Hylocomium</i>
82	<i>Hylocomium splendens</i> (Hedw.) Schimp.	Hyl.sple	Moss	<i>Hylocomium</i>
83	<i>Hypnum cupressiforme</i> Hedw.	Hyp.cupr	Moss	<i>Hypnum</i>
84	<i>Hypnum curvifolium</i> Hedw.	Hyp.curv	Moss	<i>Hypnum</i>
85	<i>Hypnum fauriei</i> Cardot	Hyp.faur	Moss	<i>Hypnum</i>
86	<i>Hypnum imponens</i> Hedw.	Hyp.impo	Moss	<i>Hypnum</i>
87	<i>Hypnum lindbergii</i> Mitt.	Cal.lind	Moss	<i>Hypnum</i>
88	<i>Hypnum pallescens</i> (Hedw.) P. Beauv.	Hyp.pall	Moss	<i>Hypnum</i>
89	<i>Jamesoniella autumnalis</i> (DC.) Stephani	Jam.autu	Liverwort	<i>Jamesoniella</i>
90	<i>Jungermania crenuliformis</i> Austin	Jun.cren	Liverwort	<i>Jungermania</i>
91	<i>Jungermannia leiantha</i> Grolle	Jun.leia	Liverwort	<i>Jungermannia</i>
92	<i>Kurzia pauciflora</i> (Dicks.) Grolle	Kur.pauc	Liverwort	<i>Kurzia</i>
93	<i>Leiomylia anomala</i> (Hook.) J.J. Engel & Braggins	Myl.anom	Liverwort	<i>Leiomylia</i>

Table S2.3 continued

94	<i>Lepidozia reptans</i> (L.) Dumort.	Lep.rept	Liverwort	<i>Lepidozia</i>
95	<i>Leptodictyum riparium</i> (Hedw.) Warnst.	Lep.ripa	Moss	<i>Leptodictyum</i>
96	<i>Lophozia longidens</i> (Lindb.) Macoun	Lop.long	Liverwort	<i>Lophozia</i>
97	<i>Lophozia sudetica</i> (Nees ex Huebener) Grolle	Lop.sude	Liverwort	<i>Lophozia</i>
98	<i>Lophozia ventricosa</i> (Dicks.) Dumort.	Lop.vent	Liverwort	<i>Lophozia</i>
99	<i>Marchantia polymorpha</i> L.	Mar.poly	Liverwort	<i>Marchantia</i>
100	<i>Mnium spinulosum</i> Bruch & Schimp.	Mni.spin	Moss	<i>Mnium</i>
101	<i>Mnium marginatum</i> (Dicks. ex With.) P. Beauv.	Mni.marg	Moss	<i>Mnium</i>
102	<i>Mylia taylorii</i> (Hook.) Gray	Myl.tayl	Liverwort	<i>Mylia</i>
103	<i>Nowellia Curvifolia</i> (Dicks.) Mitt.	Now.curv	Liverwort	<i>Nowellia</i>
104	<i>Orthotrichum anomalum</i> Hedw.	Ort.anom	Moss	<i>Orthotrichum</i>
105	<i>Orthotrichum elegans</i> Schwägr. ex Hook. & Grev.	Ort.eleg	Moss	<i>Orthotrichum</i>
106	<i>Orthotrichum obtusifolium</i> Brid.	Ort.obtu	Moss	<i>Orthotrichum</i>
107	<i>Orthotrichum ohioense</i> Sull. & Lesq.	Ort.ohio	Moss	<i>Orthotrichum</i>
108	<i>Orthotrichum speciosum</i> Nees	Ort.spec	Moss	<i>Orthotrichum</i>
109	<i>Orthotrichum stellatum</i> Brid.	Ort.stel	Moss	<i>Orthotrichum</i>
110	<i>Palustriella falcata</i> (Brid.) Hedenäs	Pal.falc	Moss	<i>Palustriella</i>
111	<i>Pallavicinia lyellii</i> (Hook.) Gray	Pal.lyel	Liverwort	<i>Pallavicinia</i>
112	<i>Pellia epiphylla</i> (Dicks.) Dumort.	Pel.epip	Liverwort	<i>Pellia</i>
113	<i>Plagiothecium cavifolium</i> (Brid.) Z. Iwats.	Pla.cavi	Moss	<i>Plagiothecium</i>
114	<i>Platydictya confervoides</i> (Brid.) H.A. Crum	Pla.conf	Moss	<i>Platydictya</i>
115	<i>Plagiomnium cuspidatum</i> (Hedw.) T.J. Kop.	Pla.cusp	Moss	<i>Plagiomnium</i>
116	<i>Plagiothecium denticulatum</i> (Hedw.) Schimp.	Pla.dent	Moss	<i>Plagiothecium</i>

Table S2.3 continued

117	<i>Plagiomnium drummondii</i> (Bruch & Schimp.) T.J. Kop.	Pla.drum	Moss	<i>Plagiomnium</i>
118	<i>Plagiomnium ellipticum</i> (Brid.) T.J. Kop.	Pla.elli	Moss	<i>Plagiomnium</i>
119	<i>Plagiothecium laetum</i> Schimp.	Pla.laet	Moss	<i>Plagiothecium</i>
120	<i>Plagiothecium latebricola</i> Schimp.	Pla.late	Moss	<i>Plagiothecium</i>
121	<i>Plagiomnium medium</i> (Bruch & Schimp.) T.J. Kop.	Pla.medi	Moss	<i>Plagiomnium</i>
122	<i>Platygyrium repens</i> (Brid.) Schimp.	Pla.repe	Moss	<i>Platygyrium</i>
123	<i>Platydictya subtilis</i> (Hedw.) H.A. Crum	Pla.subt	Moss	<i>Platydictya</i>
124	<i>Pleurozium schreberi</i> (Willd. ex Brid.) Mitt.	Ple.schr	Moss	<i>Pleurozium</i>
125	<i>Pohlia nutans</i> (Hedw.) Lindb.	Poh.nuta	Moss	<i>Pohlia</i>
126	<i>Pohlia wahlenbergii</i> (F. Weber & D. Mohr) A.L. Andrews	Poh.wahl	Moss	<i>Pohlia</i>
127	<i>Polytrichastrum alpinum</i> (Hedw.) G.L. Sm.	Pol.alpi	Moss	<i>Polytrichastrum</i> <i>m</i>
128	<i>Polytrichum commune</i> Hedw.	Pol.comm	Moss	<i>Polytrichum</i>
129	<i>Polytrichum juniperinum</i> Hedw.	Pol.juni	Moss	<i>Polytrichum</i>
130	<i>Polytrichastrum longisetum</i> (Sw. ex Brid.) G.L. Sm.	Pol.long	Moss	<i>Polytrichastrum</i> <i>m</i>
131	<i>Polytrichum ohioense</i> (Renauld & Cardot) G.L. Sm.	Pol.ohio	Moss	<i>Polytrichum</i>
132	<i>Polytrichum piliferum</i> Hedw.	Pol.pili	Moss	<i>Polytrichum</i>
133	<i>Polytrichum strictum</i> Menzies ex Brid.	Pol.stri	Moss	<i>Polytrichum</i>
134	<i>Pseudotaxiphyllum elegans</i> (Brid.) Z. Iwats.	Pse.eleg	Moss	<i>Pseudotaxiphyllum</i> <i>um</i>
135	<i>Pseudoleskeella nervosa</i> (Brid.) Nyholm	Pse.nerv	Moss	<i>Pseudoleskeella</i>
136	<i>Pseudocampylium radicale</i> (P. Beauv.) Vanderp. & Hedenäs	Pse.radi	Moss	<i>Pseudocampylium</i> <i>um</i>
137	<i>Ptilidium ciliare</i> (L.) Hampe	Pti.cili	Liverwort	<i>Ptilidium</i>
138	<i>Ptilium crista-castrensis</i> (Hedw.) De Not.	Pti.cris	Moss	<i>Ptilium</i>

Table S2.3 continued

139	<i>Ptilidium pulcherrimum</i> (Weber) Hampe	Pti.pulc	Liverwort	<i>Ptilidium</i>
140	<i>Pylaisia intricata</i> (Hedw.) Schimp.	Pyl.intr	Moss	<i>Pylaisia</i>
141	<i>Pylaisia polyantha</i> (Hedw.) Schimp.	Pyl.poly	Moss	<i>Pylaisia</i>
142	<i>Pylaisia selwynii</i> Kindb.	Pyl.selw	Moss	<i>Pylaisia</i>
143	<i>Racomitrium microcarpum</i> (Hedw.) Bednarek-Ochyra & Ochyra	Buc.micr	Moss	<i>Racomitrium</i>
144	<i>Radula complanata</i> (L.) Dumort.	Rad.comp	Liverwort	<i>Radula</i>
145	<i>Rhizomnium gracile</i> T.J. Kop.	Rhi.grac	Moss	<i>Rhizomnium</i>
146	<i>Rhizomnium magnifolium</i> (Horik.) T.J. Kop.	Rhi.magn	Moss	<i>Rhizomnium</i>
147	<i>Rhizomnium pseudopunctatum</i> (Bruch & Schimp.) T.J. Kop.	Rhi.pseu	Moss	<i>Rhizomnium</i>
148	<i>Rhizomnium punctatum</i> (Hedw.) T.J. Kop.	Rhi.punc	Moss	<i>Rhizomnium</i>
149	<i>Rhynchostegium serrulatum</i> (Hedw.) A. Jaeger	Rhy.serr	Moss	<i>Rhynchostegiu m</i>
150	<i>Rhytidiadelphus subpinnatus</i> (Lindb.) T.J. Kop.	Rhy.subp	Moss	<i>Rhytidiadelphu s</i>
151	<i>Rhytidiadelphus triquetrus</i> (Hedw.) Warnst.	Rhy.triq	Moss	<i>Rhytidiadelphu s</i>
152	<i>Riccardia latifrons</i> (Lindb.) Lindb.	Ric.lati	Liverwort	<i>Riccardia</i>
153	<i>Riccardia multifida</i> (L.) Gray	Ric.mult	Liverwort	<i>Riccardia</i>
154	<i>Riccardia vitrea</i> Furuki	Ric.vitr	Liverwort	<i>Riccardia</i>
155	<i>Sanionia uncinata</i> (Hedw.) Loeske	San.unci	Moss	<i>Sanionia</i>
156	<i>Scapania apiculata</i> Spruce	Sca.apic	Liverwort	<i>Scapania</i>
157	<i>Scapania irrigua</i> (Nees) Nees	Sca.irri	Liverwort	<i>Scapania</i>
158	<i>Scapania mucronata</i> H. Buch	Sca.mucr	Liverwort	<i>Scapania</i>
159	<i>Schistidium agassizii</i> Sull. & Lesq.	Sch.agas	Moss	<i>Schistidium</i>
160	<i>Schistochilopsis incisa</i> (Schrad.) Konstantinova	Sch.inci	Liverwort	<i>Schistochilopsis</i>

Table S2.3 continued

161	<i>Schistochilopsis laxa</i> (Lindb.) Konstantinova	Sch.laxa	Liverwort	<i>Schistochilopsis</i>
162	<i>Sphagnum angustifolium</i> (Warnst.) C.E.O. Jensen	Sph.angu	Moss	<i>Sphagnum</i>
163	<i>Sphagnum capillifolium</i> (Ehrh.) Hedw.	Sph.capi	Moss	<i>Sphagnum</i>
164	<i>Sphagnum centrale</i> C.E.O. Jensen	Sph.cent	Moss	<i>Sphagnum</i>
165	<i>Sphagnum cuspidatum</i> Ehrh. ex Hoffm.	Sph.cusp	Moss	<i>Sphagnum</i>
166	<i>Sphagnum fallax</i> H. Klinggr.	Sph.fall	Moss	<i>Sphagnum</i>
167	<i>Sphagnum fuscum</i> (Schimp.) H. Klinggr.	Sph.fusc	Moss	<i>Sphagnum</i>
168	<i>Sphagnum girgensohnii</i> Russow	Sph.girg	Moss	<i>Sphagnum</i>
169	<i>Sphagnum magellanicum</i> Brid.	Sph.mage	Moss	<i>Sphagnum</i>
170	<i>Sphagnum quinquefarium</i> (Lindb.) Warnst.	Sph.quin	Moss	<i>Sphagnum</i>
171	<i>Sphagnum rubellum</i> Wilson	Sph.rube	Moss	<i>Sphagnum</i>
172	<i>Sphagnum russowii</i> Warnst.	Sph.russ	Moss	<i>Sphagnum</i>
173	<i>Sphagnum squarrosum</i> Crome	Sph.squa	Moss	<i>Sphagnum</i>
174	<i>Sphagnum subfulvum</i> Sjors	Sph.subf	Moss	<i>Sphagnum</i>
175	<i>Sphagnum wulfianum</i> Girg.	Sph.wulf	Moss	<i>Sphagnum</i>
176	<i>Splachnum ampullaceum</i> Hedw.	Spl.ampu	Moss	<i>Splachnum</i>
177	<i>Tetraphis pellucida</i> Hedw.	Tet.pell	Moss	<i>Tetraphis</i>
178	<i>Thuidium delicatulum</i> (Hedw.) Schimp.	Thu.deli	Moss	<i>Thuidium</i>
179	<i>Thuidium recognitum</i> (Hedw.) Lindb.	Thu.reco	Moss	<i>Thuidium</i>
180	<i>Tomenthypnum falcifolium</i> (Renauld ex Nichols) Tuom.	Tom.falc	Moss	<i>Tomenthypnum</i>
181	<i>Tomenthypnum nitens</i> (Hedw.) Loeske	Tom.nite	Moss	<i>Tomenthypnum</i>
182	<i>Tortella humilis</i> (Hedw.) Jenn.	Tor.humi	Moss	<i>Tortella</i>

Table S2.3 continued

183	<i>Trematodon ambiguus</i> (Hedw.) Hornsch.	Tre.ambi	Moss	<i>Trematodon</i>
184	<i>Ulota coarctata</i> (P. Beauv.) Hammar	Ulo.coar	Moss	<i>Ulota</i>
185	<i>Ulota crispa</i> (Hedw.) Brid.	Ulo.cris	Moss	<i>Ulota</i>
186	<i>Warnstorfia fluitans</i> (Hedw.) Loeske	War.flui	Moss	<i>Warnstorfia</i>

Table S2.4 The list of environmental and stand structure variables as well as treatment (continuous variables with minimum, maximum, and average values $\pm$ standard deviation; categorical variables with count number).

NO.	Variables/Treatment	Variable Type	Unite	Range
<i>Stand structure</i>				
1	Live crown ratio for shrubs ¹	Continuous	%	Min=0; Max=100; Average=21.6 $\pm$ 43.8
2	Live crown ratio for trees ¹	Continuous	%	Min=0; Max=100; Average=24.2 $\pm$ 28.3
3	Base areas for shrubs	Continuous	m ² /ha	Min=0; Max=7.084746; Average=1.27 $\pm$ 1.68
4	Base areas for trees	Continuous	m ² /ha	Min=0; Max=58; Average=14.4 $\pm$ 13.7
5	Canopy openness ²	Continuous	-	Min=1; Max=4; Average=1.12 $\pm$ 2.85
6	Conifer proportion ³	Continuous	%	Min=0; Max=100; Average=44.3 $\pm$ 52.4
7	Depth of organic soil	Continuous	cm	Min=0; Max=128.33; Average=31 $\pm$ 38.4
<i>The cover of ground substrates</i>				
8	Leave litter ground cover	Continuous	%	Min=0; Max=100; Average=34.6 $\pm$ 49.2
9	Woody debris ground cover	Continuous	%	Min=0; Max=92.67; Average=13.4 $\pm$ 15.8
10	Water ground cover	Continuous	%	Min=0; Max=24.33; Average=2.60 $\pm$ 0.553
11	Rock ground cover	Continuous	%	Min=0; Max=88; Average=6.36 $\pm$ 1.07
12	Organic matter ground cover	Continuous	%	Min=0; Max=70.33; Average=5.48 $\pm$ 1.27
<i>The number of microhabitats</i>				
13	Number of trees per plot	Continuous	-	Min=0; Max=20; Average=3.29 $\pm$ 4.11
14	Number of logs per plot	Continuous	-	Min=0; Max=15; Average=8.13 $\pm$ 3.39
15	Number of stumps per plot	Continuous	-	Min=0; Max=78.33; Average=2.41 $\pm$ 2.02
16	Number of snags per plot	Continuous	-	Min=0; Max=5; Average=0.82 $\pm$ 0.44
17	Number of rocks per plot	Continuous	-	Min=0; Max=8; Average=0.854 $\pm$ 0.24

Notes: 1. Live crown ratio (%) was calculated as (H-HCB)/H, where H is the total height of the tree/shrub, and HCB is the height from crown top to base. 2. Canopy openness was measured with a spherical densiometer at 1 m height above ground and recorded as 1-4 levels (L1, 0 % (0 grid); L2, 1%-40 % (1-9.6 grids); L3, 40 %-70 % (9.7 - 16.8 grids); L4: > 70% (>16.8 grids). 3. Conifer proportion (%) was calculated by the number of individuals of coniferous trees in each plot.

Table S2.5: the variables grouped by mine sites (mean with standard deviation) with the results of Kruskal-Wallis Test.

Variables	Akasaba N=42	Casa Berardi N=56	Joutel N=49	Lapa N=49	LaRonde N=56	Canadian Malartic N=56	Controls N=61	chi- squared	p- value
<i>Stand structure</i>									
Live crown ratio for shrubs	52.125 (26.571)	49.911 (23.057)	49.876 (18.601)	38.482 (14.138)	45.632 (22.015)	30.428 (14.772)	44.031 (22.458)	45.631	<0.001
Live crown ratio for trees	23.682 (27.286)	20.447 (29.206)	31.502 (24.358)	37.322 (16.691)	29.399 (28.061)	29.673 (15.494)	26.961 (22.563)	21.281	0.001
Base areas for shrubs	1.545 (1.21)	1.448 (1.282)	2.303 (1.409)	1.639 (1.156)	1.611 (1.114)	1.896 (1.464)	1.317 (1.002)	18.174	0.006
Base areas for trees	5.984 (7.146)	3.13 (4.608)	19.959 (18.45)	22.98 (14.359)	12.264 (15.125)	18.75 (13.165)	12.689 (11.248)	84.59	<0.001
Canopy openness	2.29 (1.2)	2.16 (1.01)	2.96 (1.1)	3.36 (0.77)	3.09 (1.09)	3.34 (0.86)	2.77 (1.2)	52.648	<0.001
Conifer proportion	53.571 (49.869)	35.268 (47.14)	54.917 (42.572)	69.494 (33.382)	38.728 (43.466)	58.516 (41.006)	51.78 (44.181)	19.525	0.002
Depth of organic soil	52.472 (18.524)	73.406 (23.289)	36.194 (30.283)	12.088 (10.18)	14.944 (9.351)	30.024 (27.111)	47.336 (31.383)	165.01	<0.001
<i>The cover of ground substrates</i>									
Leave litter ground cover	25.278 (20.365)	21.598 (24.432)	51.204 (33.28)	68.687 (25.027)	77.723 (20.028)	56.458 (33.775)	41.216 (37.617)	111.92	<0.001
Woody debris ground cover	11.81 (10.362)	13.727 (14.112)	15.756 (11.233)	19.005 (18.964)	22.083 (20.09)	15.854 (10.286)	15.179 (10.321)	15.369	0.012
Water ground cover	0.079 (0.514)	0.476 (1.812)	0.918 (4.305)	0.211 (1.476)	0.238 (1.401)	0.75 (2.718)	0.992 (3.536)	8.8924	0.18
Rock ground cover	0 (0)	0.089 (0.668)	0.122 (0.604)	4.493 (12.313)	3.899 (13.613)	0.289 (1.232)	0.17 (1.062)	39.411	<0.001

Table S2.5 continued

Organic matter ground cover	1.063 (4.209)	1.193 (2.219)	3.063 (8.506)	0.765 (2.689)	0.274 (1.39)	2.03 (10.011)	0.563 (1.727)	19.345	0.004
<i>The number of microhabitats</i>									
Number of trees per plot	3.76(2.38)	3.95 (2.81)	3.84 (3.09)	3.09 (2.17)	3.38 (3.02)	6.82 (4.53)	3.64 (2.76)	29.981	<0.001
Number of logs per plot	1.29 (1.47)	0.71 (1.24)	1.9 (2.56)	2.31 (2.6)	2.45 (2.35)	2.77 (2.25)	2.54 (3.08)	37.718	<0.001
Number of stumps per plot	1.86 (1.42)	1.51 (1.77)	1.24 (1.36)	0.62 (0.83)	0.87 (1.11)	1.27 (1.45)	0.92 (1.14)	24.84	<0.001
Number of snags per plot	0.38 (0.62)	0.62 (1.01)	0.37 (0.7)	0.31 (0.7)	0.35 (0.87)	0.54 (0.83)	0.49 (0.89)	6.5313	0.366
Number of rocks per plot	0.05 (0.22)	0.07 (0.54)	0.02 (0.14)	0.31 (0.6)	0.67 (1.62)	0.34 (0.94)	0.16 (0.61)	26.949	<0.001

Table S2.6 Mean values of environmental variables near mine sites for each mining stages in four identified ecosystem types.

Variables	Coniferous			Deciduous			Mixed forest			Open Canopy		
	Operating	Non-operating	Controls	Operating	Non-operating	Controls	Operating	Non-operating	Controls	Operating	Non-operating	Controls
<i>Stand structure</i>												
Live crown ratio for shrubs	37.8 ± 2.5 a	43.5 ± 2.5 ab	40.2 ± 4.0 ab	36.3 ± 4.8 a	43.0 ± 5.8 ab	52.8 ± 5.6 ab	32.6 ± 6.1 a	41.2 ± 4.9 ab	41.7 ± 7.9 ab	50.4 ± 2.9 ab	57.2 ± 3.7 b	43.7 ± 5.8 ab
Live crown ratio for trees	41.4 ± 1.8 bc	44.2 ± 1.8 bc	37.4 ± 3.0 bc	48.9 ± 3.5 c	40.3 ± 4.3 bc	30.6 ± 4.1 b	31.8 ± 4.4 bc	32.4 ± 3.6 bc	29.4 ± 5.8 bc	0.0 ± a	0.0 ± 2.7 a	0.0 ± 4.3 a
Base areas for shrubs	1.9 ± 0.1 b	2.1 ± 0.1 b	1.4 ± 0.2 ab	2.3 ± 0.3 b	1.8 ± 0.3 ab	1.6 ± 0.3 ab	1.8 ± 0.4 ab	1.8 ± 0.3 ab	1.1 ± 0.5 ab	1.1 ± 0.2 a	1.5 ± 0.2 ab	1.0 ± 0.3 ab
Base areas for trees	16.7 ± 1.3 b	18.9 ± 1.2 b	15.0 ± 2.0 b	18.3 ± 2.4 b	21.1 ± 2.9 b	15.9 ± 2.8 b	27.0 ± 3.0 bc	36.3 ± 2.5 c	20.9 ± 4.0 b	0.0 ± a	0.0 ± 1.4 a	0.0 ± 1.8 a
Canopy openness	3.3 ± 0.1 bc	3.1 ± 0.1 b	2.9 ± 0.2 b	3.9 ± 0.2 c	3.4 ± 0.2 bc	3.5 ± 0.2 bc	3.5 ± 0.2 bc	3.7 ± 0.2 bc	3.6 ± 0.3 bc	1.9 ± 0.1 a	1.7 ± 0.1 a	1.2 ± 0.2 a
Conifer proportion	94.6 ± 1.0 d	95.5 ± 1.0 d	96.2 ± 1.6 d	10.4 ± 1.9 b	17.7 ± 2.4 b	10.9 ± 2.3 b	54.5 ± 2.5 c	54.2 ± 2.0 c	53.6 ± 3.2 c	0.0 ± a	0.0 ± 1.1 a	0.0 ± 1.5 a
Depth of organic soil	36.9 ± 3.1 bcd	33.4 ± 3.2 abcd	46.6 ± 5.2 cde	12.9 ± 6.2 a	10.4 ± 7.5 ab	26.8 ± 7.2 abcd	19.2 ± 8.1 abc	15.6 ± 6.3 ab	19.9 ± 10.2 abcd	54.8 ± 3.5 c	52.0 ± 4.7 de	84.0 ± 7.5 f
<i>The cover of ground substrates</i>												
Leave litter ground cover	51.7 ± 3.6 cd	50.6 ± 3.6 cd	52.4 ± 5.9 cd	86.5 ± 6.9 e	71.9 ± 8.5 de	50.8 ± 8.2 bcd	75.3 ± 8.9 de	72.2 ± 7.3 de	38.0 ± 11.6 abcd	36.7 ± 3.9 abc	25.5 ± 5.4 ab	9.3 ± 8.5 a
Woody debris ground cover	14.9 ± 1.56 a	14.4 ± 1.6 a	15.8 ± 2.6 a	18.4 ± 3.0 a	16.9 ± 3.7 a	15.0 ± 3.6 a	21.8 ± 3.9 a	19.7 ± 3.2 a	13.6 ± 5.1 a	17.6 ± 1.7 a	12.7 ± 2.3 a	14.9 ± 3.7 a

Table S2.6 continued

Water ground cover	0.7 ± 0.3	0.2 ± 0.3 a	0.0 ± 0.5 a	0.6 ± 0.6 a	0.0 ± 0.7 a	0.0 ± 0.1 a	0.7 ± 0.7 a	0.0 ± 0.6 a	0.0 ± 0.1 a	0.0 ± 0.3 a	0.3 ± 0.3 a	1.3 ± 0.4 a	4.7 ± 0.7 b
Rock ground cover	0.2 ± 0.7	0.4 ± 0.7 a	0.3 ± 1.2 a	3.7 ± 1.4 a	2.2 ± 1.8 a	0.17 ± 1.7 a	0.0 ± 1.5 a	4.3 ± 2.4 a	0.0 ± 2.5 a	0.8 ± 1.1 a	0.1 ± 1.8 a	0.0 ± 1.1 a	0.0 ± 1.8 a
Organic matter ground cover	1.0 ± 0.6	0.7 ± 0.6 a	0.8 ± 1.0 ab	0.3 ± 1.2 ab	0.8 ± 1.5 ab	1.0 ± 1.4 ab	0.2 ± 2.2 ab	1.6 ± 1.3 a	0.0 ± 2.0 ab	1.8 ± 2.0 ab	0.7 ± 0.9 b	5.2 ± 0.9 b	0.0 ± 1.5 ab
<i>The number of microhabitats</i>													
Number of trees per plot	6.8 ± 0.3	4.4 ± 0.3 cd	4.6 ± 0.5 bcd	3.7 ± 0.6 abc	2.8 ± 0.8 abc	3.4 ± 0.8 abc	7.2 ± 0.8 de	3.4 ± 0.7 abc	3.4 ± 1.1 abcd	2.1 ± 0.4 a	2.3 ± 0.5 ab	2.0 ± 0.8 abc	2.0 ± 0.8 abc
Number of logs per plot	2.4 ± 0.2	1.6 ± 0.2 abc	1.9 ± 0.4 abcd	2.9 ± 0.5 cde	2.2 ± 0.6 abcd	4.0 ± 0.6 def	3.7 ± 0.6 cdef	4.9 ± 0.5 ef	6.1 ± 0.8 f	0.9 ± 0.3 a	0.6 ± 0.4 a	0.4 ± 0.6 ab	0.4 ± 0.6 ab
Number of stumps per plot	1.4 ± 0.2	1.2 ± 0.2 a	1.0 ± 0.3 a	0.9 ± 0.4 a	0.9 ± 0.4 a	0.8 ± 0.4 a	0.8 ± 0.4 a	0.9 ± 0.3 a	1.0 ± 0.5 a	1.3 ± 0.2 a	1.6 ± 0.2 a	0.7 ± 0.4 a	0.7 ± 0.4 a
Number of snags per plot	0.7 ± 0.1	0.4 ± 0.1 a	0.4 ± 0.2 a	0.3 ± 0.2 a	0.2 ± 0.2 a	0.3 ± 0.2 a	0.3 ± 0.2 a	0.4 ± 0.2 a	1.3 ± 0.3 a	0.4 ± 0.1 a	0.2 ± 0.1 a	0.5 ± 0.2 a	0.5 ± 0.2 a
Number of rocks per plot	0.2 ± 0.1	0.2 ± 0.1 a	0.0 ± 0.2 a	1.6 ± 0.2 b	0.3 ± 0.2 a	0.4 ± 0.2 a	0.3 ± 0.2 a	0.1 ± 0.2 a	0.0 ± 0.3 a	0.2 ± 0.1 a	0.0 ± 0.1 a	0.3 ± 0.2 a	0.3 ± 0.2 a

Notes: Significant differences ($p = 0.05$) between mining stages crossing ecosystem types are indicated by different letters (generalized linear mixed model followed by Tukey HSD test).

Table S2.7 Results of generalized linear mixed models (GLMM) on the bryophyte richness testing for the effect of mine stages, ecosystem types and their interactions. Non-significant interaction terms were deleted from the final model. Significant parameters (≤ 0.05) are indicated in bold.

	All bryophyte richness (N=364)		Moss richness (N=364)		Liverwort richness (N=364)	
	Estimate	<i>p</i>	Estimate	<i>p</i>	Estimate	<i>p</i>
(Intercept)	2.54	<0.01	2.00	<0.01	1.40	<0.01
Mining Stage (Reference: Controls)						
Operating	0.13	0.15	0.35	<0.01	-0.08	0.59
Non-operating	0.13	0.17	0.36	<0.01	-0.10	0.50
Ecosystem type (Reference: Coniferous)						
Deciduous	0.31	0.02	0.50	<0.01	0.18	0.35
Mixed	0.65	<0.01	0.87	<0.01	0.40	0.10
Open Canopy	-0.68	<0.01	-0.46	0.01	-1.15	<0.01
Mining Stage: Ecosystem type						
Deciduous: Operating	-0.32	0.06	-0.41	0.01	-0.52	0.05
Deciduous: Non-operating	-0.22	0.22	-0.40	0.03	-0.54	0.07
Mixed: Operating	-0.65	<0.01	-0.89	<0.01	-0.25	0.42
Mixed: Non-operating	-0.42	0.04	-0.64	<0.01	-0.12	0.68
Open Canopy: Operating	0.28	0.13	0.05	0.80	0.81	0.02
Open Canopy: Non-operating	0.32	0.10	0.19	0.34	0.43	0.22
Anova (Type=II)						
Factors	Chisq.	P	Chisq.	P	Chisq.	P
Mining stage	0.91	0.63	7.36	0.03	3.98	0.14
Ecosystem type	88.12	<0.01	95.34	<0.01	44.89	<0.01
Mining stage: Ecosystem type	17.70	0.01	27.89	<0.01	16.40	0.01

Table S2.8 Influence of mining stage, ecosystem type and their interactions on bryophyte community composition using permutational multivariate analysis of variance (PERMANOVA) based on the Bray-Curtis dissimilarity index for total bryophyte, moss and liverwort species.

Assemblage	Variable	Df	Mean square	F statistics	R ² (%)	P
Total bryophyte species	Mining Stages	2	0.64	3.32	1.41	0.001
	Ecosystem types	3	3.81	19.84	12.60	0.001
	Sites	4	2.13	11.09	9.39	0.001
	Mining Stages: Ecosystem types	6	0.44	2.31	2.93	0.001
	Residuals	348	0.19		73.67	
Moss species	Mining Stages	2	0.73	3.57	1.49	0.001
	Ecosystem types	3	4.36	21.45	13.45	0.001
	Sites	4	2.25	11.06	9.25	0.001
	Mining Stages: Ecosystem types	6	0.50	2.46	3.09	0.001
	Residuals	348	0.20		72.73	
Liverwort species	Mining Stages	2	0.51	3.23	1.58	0.002
	Ecosystem types	3	1.92	12.27	8.97	0.001
	Sites	4	1.45	9.30	9.06	0.001
	Mining Stages: Ecosystem types	6	0.26	1.66	2.43	0.023
	Residuals	320	0.16		77.97	

Table S2.9 PERMANOVA (Bray-Curtis) post hoc pair-wise comparisons of mine stages in each ecosystem type on composition of total bryophytes, mosses and liverworts respectively based on 999 permutation and Benjamini-Hochberg adjustment (BH).

Pairs	Total bryophytes		Mosses		Liverworts	
	R ² (%)	p.adjusted	R ² (%)	p.adjusted	R ² (%)	p.adjusted
Coniferous						
Operating vs Nonoperating	0.91	0.20	0.96	0.17	1.57	0.05
Operating vs Controls	2.08	0.04	2.55	0.02	1.14	0.32
Nonoperating vs Controls	2.26	0.02	2.80	0.01	1.17	0.33
Deciduous						
Operating vs Nonoperating	6.25	0.03	6.53	0.03	9.49	0.04
Operating vs Controls	6.40	0.01	6.72	0.01	5.48	0.13
Nonoperating vs Controls	9.60	0.01	10.86	0.01	6.42	0.14
Mixed						
Operating vs Nonoperating	6.07	0.03	7.08	0.01	4.40	0.28
Operating vs Controls	8.40	0.06	8.92	0.06	7.62	0.21
Nonoperating vs Controls	5.48	0.20	4.89	0.28	7.01	0.13
Open Canopy						
Operating vs Nonoperating	3.39	<0.01	2.95	0.01	4.23	0.01
Operating vs Controls	4.25	<0.01	4.78	<0.01	1.96	0.30
Nonoperating vs Controls	6.07	0.01	6.50	0.01	3.31	0.23

Table S2.10 The results of the Euclidean distances (D) between mining stage centroids [operating(op), non-operating (nop) and controls (con)] for an ecosystem type. Composition dataset of liverwort species used in the diagram to describe the methodology.

	Bryophyte groups			Total bryophytes on each microhabitat				
	Total	Moss	liverworts	Ground	Trees	Logs	Snags	Stumps
D_{op-con}								
Coniferous	0.099	0.115	0.082	0.095	0.146	0.125	0.113	0.100
Deciduous	0.029↓	0.035↓	0.160↑	0.234↑	0.256↑	0.235↑	0.604↑	0.216↑
Mixed	0.067↓	0.053↓	0.160↑	0.270↑	0.220↑	0.221↑	0.071↓	0.067↓
Open Canopy	0.208↑	0.225↑	0.136↑	0.370↑	0.376↑	0.464↑	0.379↑	0.391↑
$D_{non-con}$								
Coniferous	0.123	0.150	0.072	0.047	0.113	0.073	0.113	0.114
Deciduous	0.172↑	0.182↑	0.196↑	0.425↑	0.520↑	0.355↑	0.498↑	0.414↑
Mixed	0.073↓	0.089↓	0.179↑	0.168↑	0.118↑	0.249↑	0.283↑	0.097↓
Open Canopy	0.183↑	0.203↑	0.139↑	0.636↑	0.477↑	0.388↑	0.572↑	0.487↑
D_{op-non}								
Coniferous	0.040	0.067	0.089	0.134	0.127	0.057	0.003	0.049
Deciduous	0.145↑	0.147↑	0.277↑	0.193↑	0.272↑	0.153↑	0.147↑	0.234↑
Mixed	0.007↓	0.036↓	0.070↓	0.104↓	0.109↓	0.033↓	0.273↑	0.118↑
Open Canopy	0.132↑	0.107↑	0.168↑	0.371↑	0.101↓	0.125↑	0.251↑	0.104↑

Notes: only groups affected significantly by the interactions between mining stage and ecosystem type were present in the table see PERMANOVA results in Table S2.7 and S2.12. “↑” means synergistic effects and “↓” means antagonistic effects.

Table S2.11. Nonmetric multi-dimensional scaling (NMDS) showing the associations between the composition of all bryophytes and selected variables. The significant relationships are in bold, and R^2 is the fit index of the model.

Variables	Abbreviation	PCOA1	PCOA2	R^2	P
Canopy openness	OP	0.61	-0.79	0.45	0.001
Live crown ratio for shrubs	SLC	-0.48	0.88	0.03	0.005
Base areas for shrubs	SBA	0.24	-0.97	0.05	0.003
Live crown ratio for trees	TLC	0.50	-0.87	0.12	0.001
Base areas for trees	TBA	0.47	-0.88	0.32	0.001
Depth of organic soil	OSD	-0.87	0.48	0.40	0.001
Number of trees per plot	Treen	-0.12	-0.99	0.25	0.001
Number of logs per plot	Logn	0.60	-0.80	0.33	0.001
Number of stumps per plot	Stumpn	-0.34	-0.94	0.06	0.001
Number of snags per plot	Snagn	-0.52	-0.85	0.04	0.002
Number of rocks per plot	Rockn	0.90	0.45	0.06	0.001

Table S2.12 Results of generalized linear mixed models (GLMM) on total bryophyte richness on each microhabitat testing for the effect of mine stages, ecosystem types and their interactions. Non-significant interaction terms were deleted from the final models. Significant parameters (≤ 0.05) are indicated in bold.

	Ground		Trees		Logs		Stumps		Snags		Rocks	
	Estimat	<i>P</i>	Estimat	<i>P</i>	Estimat	<i>P</i>	Estimat	<i>P</i>	Estimat	<i>P</i>	Estimat	<i>P</i>
(Intercept)	1.80	<0.0	1.99	<0.0	2.11	<0.	1.72	<0.	1.48	<0.0	1.00	<0.0
Mining Stage												
Operating	0.20	0.01	0.12	0.20	-0.55	0.24	<0.01	1.00	-0.22	0.37	0.62	0.02
Non-operating	0.27	0.03	0.04	0.70	-0.19	0.69	0.04	0.73	-0.31	0.21	0.36	0.23
Ecosystem type												
Deciduous	0.18	0.27	-0.02	0.85	0.31	0.02	0.08	0.55	0.03	0.90	0.54	<0.0
Mixed	0.26	0.23	0.20	0.04	0.54	<0.	0.09	0.54	0.60	<0.0	0.57	0.03
Open Canopy	-0.40	0.04	-0.77	<0.0	-0.98	<0.	-0.20	0.05	-0.48	0.02	-0.19	0.58
Mining Stage: Ecosystem type												
Deciduous:	-0.44	0.04										
Deciduous: Non-	-0.72	<0.0										
Mixed: Operating	-0.25	0.35										
Mixed: Non-	-0.24	0.34										
Open Canopy:	0.18	0.41										
Open Canopy:	0.25	0.26										
Anova (Type=II)												
Factors	Chisq.	<i>P</i>	Chisq.	<i>P</i>	Chisq.	<i>P</i>	Chisq.	<i>P</i>	Chisq.	<i>P</i>	Chisq.	<i>P</i>
Mining stage	4.01	0.13	2.33	0.31	4.73	0.19	0.25	0.88	1.71	0.43	0.19	0.91
Ecosystem type	16.08	<0.0	88.67	<0.0	52.73	<0.	5.92	0.12	18.90	<0.0	16.07	<0.0
Mining stage:	14.29	0.03	6.28	0.39	4.55	0.60	6.31	0.39	11.13	0.08	9.64	0.14

Table S2.13 Influence of mining stages, ecosystem types and their interactions on bryophyte community dissimilarity on different microhabitat using permutational multivariate analysis of variance (PERMANOVA) based on the Bray-Curtis dissimilarity index for trees, logs, stumps, snags and rocks.

Microhabitats	Variable	Df	Mean square	F statistics	R ² (%)	P
Ground (N=345)	Mining Stages	2	1.26	3.66	1.91	0.001
	Ecosystem types	3	3.55	10.33	8.06	0.001
	Mining Stages: Ecosystem types	6	0.71	2.07	3.22	0.001
	Residuals	334	0.34	0.87		
Trees (N=312)	Mining Stages	2	0.57	2.45	1.4	0.003
	Ecosystem types	3	3.09	13.32	11.1	0.001
	Mining Stages: Ecosystem types	6	0.50	2.16	3.6	0.001
	Residuals	301	0.23	0.84		
Logs (N=226)	Mining Stages	2	0.59	2.37	1.9	0.004
	Ecosystem types	3	1.74	6.96	8.4	0.001
	Mining Stages: Ecosystem types	6	0.34	1.37	3.3	0.035
	Residuals	215	0.25	0.86		
Stumps (N=212)	Mining Stages	2	0.52	1.79	1.6	0.01
	Ecosystem types	3	1.58	5.45	7.1	0.001
	Mining Stages: Ecosystem types	6	0.42	1.46	3.8	0.012
	Residuals	201	0.29	0.88		
Snags (N=103)	Mining Stages	2	0.50	1.90	3.3	0.017
	Ecosystem types	3	1.00	3.77	9.7	0.001
	Mining Stages: Ecosystem types	6	0.40	1.51	7.8	0.017
	Residuals	92	0.26	0.79		
Rocks (N=39)	Mining Stages	2	0.58	1.78	8.3	0.015
	Ecosystem types	3	0.59	1.81	12.6	0.003
	Residuals	34	0.33	0.79		

Table S2.14. PERMANOVA (Bray-Curtis) post hoc pair-wise comparisons of mine stages in each ecosystem type on composition of all bryophyte species on each microhabitat respectively based on 999 permutation and Benjamini-Hochberg adjustment (BH).

Pairwise PERMANOVA	Ground			Trees			Logs			Stumps			Snags			
	R ² (%)	p-value	p-adjusted	R ² (%)	p-value	p-adjusted	R ² (%)	p-value	p-adjusted	R ² (%)	p-value	p-adjusted	R ² (%)	p-value	p-adjusted	
Coniferous	Operating vs Nonoperating Controls	0.62	0.59	0.59	0.90	0.22	0.24	0.68	0.71	0.70	0.85	0.87	4.10	0.02	0.08	
	Operating vs Controls	2.90	<0.01	<0.01	1.38	0.22	0.24	1.22	0.48	0.55	1.04	0.84	0.87	5.54	0.02	0.07
	Nonoperating vs Controls	2.46	0.01	0.01	2.06	0.06	0.07	1.33	0.54	0.59	0.93	0.89	0.90	6.09	0.09	0.17
Deciduous																
Operating vs Nonoperating Controls	7.91	<0.01	0.01	3.87	0.37	0.38	5.27	0.24	0.30	5.25	0.69	0.74	17.73	0.26	0.33	
Operating vs Controls	5.33	0.06	0.07	5.69	0.06	0.07	8.89	<0.01	0.01	12.11	0.01	0.03	20.45	0.70	0.72	
Nonoperating vs Controls	9.97	<0.01	<0.01	13.12	0.01	0.01	11.62	0.01	0.01	11.94	0.08	0.11	16.01	0.27	0.34	
Mixed																
Operating vs Nonoperating Controls	4.88	0.13	0.14	7.05	0.05	0.06	6.37	0.06	0.08	8.56	0.15	0.18	23.49	0.10	0.17	
Operating vs Controls	5.16	0.55	0.56	9.27	0.11	0.12	4.63	0.66	0.70	19.32	0.04	0.06	18.53	0.60	0.63	
Nonoperating vs Controls	5.44	0.19	0.21	7.76	0.05	0.07	4.38	0.55	0.59	13.04	0.03	0.05	18.07	0.12	0.18	
Open Canopy																
Operating vs Nonoperating Controls	2.88	0.01	0.01	5.37	<0.01	<0.01	9.24	0.01	0.01	3.82	0.02	0.04	8.57	0.18	0.25	
Operating vs Controls	8.50	<0.01	<0.01	5.00	0.02	0.02	5.56	0.48	0.55	3.89	0.08	0.11	5.99	0.71	0.72	
Nonoperating vs Controls	11.37	<0.01	<0.01	3.91	0.23	0.25	5.84	0.42	0.50	6.12	0.13	0.16	10.31	0.50	0.53	

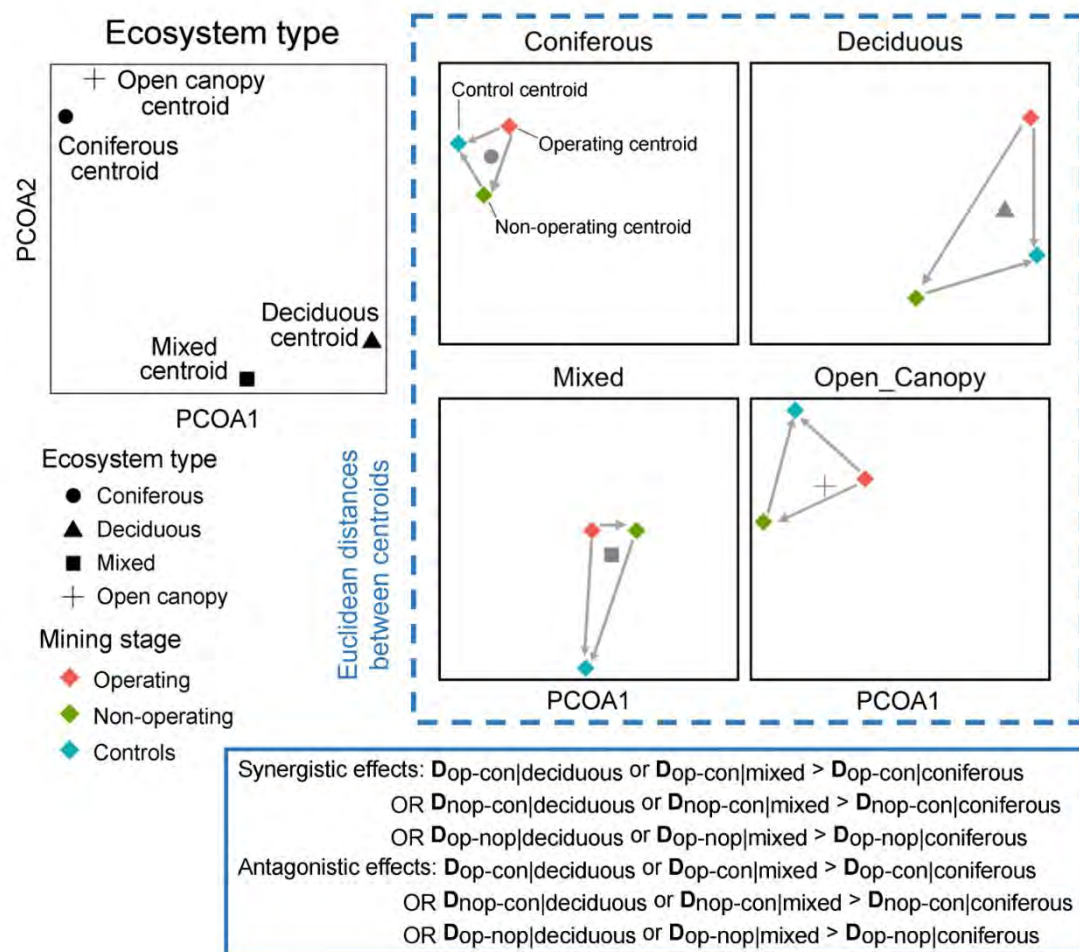


Figure S2.1 Testing for Interaction effects for community composition shifts in the study. op, operating centroid; nop, non-operating centroid; con, control centroid. $D_{op-con|deciduous}$ means Euclidean distances between operating centroid and control centroids in deciduous forest.

APPENDICE C

MATÉRIEL SUPPLÉMENTAIRE DU CHAPITRE IV

Table S3.1. Results of PERMANOVA showing the relative importance of forest type, mining stage and their interaction in structuring the phyllosphere communities based on Bray-Curtis dissimilarity.

	Df	Sums Of Sqs	Mean Sqs	F.Model	R ² (%)	P
Bacterial community						
Mining stage (Stage)	2	2.32	1.16	4.57	4.43	< 0.001
Forest type (Forest)	3	3.04	1.01	3.98	5.83	< 0.001
Forest: Stage	6	1.56	0.26	1.02	2.72	0.70
Residuals	198	50.31	0.25	0.88		
Total	209	57.22	1.00			
Fungal community						
Mining stage (Stage)	2	2.39	1.20	3.38	2.62	< 0.001
Forest type (Forest)	3	4.10	1.37	3.86	4.63	< 0.001
Forest: Stage	6	2.41	0.40	1.13	3.06	0.16
Residuals	197	69.77	0.35	0.89		
Total	208 ¹	78.67	1.00			

Notes: 1, one outlier was removed from analysis (LR2A plot)

Table S3.2 Results of pairwise PERMANOVA for the community composition between mining stages with Benjamini-Hochberg adjustment.

Pairwise PERMANOVA	Bacterial community			Fungal community		
	<i>F</i>	R ² (%)	<i>P</i> .adj	<i>F</i>	R ² (%)	<i>P</i> .adj
Operating vs. Non-operating	5.18	2.99	0.001	2.34	1.38	0.001
Operating vs. Controls	4.50	3.35	0.001	3.27	2.47	0.001
Non-operating vs. Controls	4.38	3.64	0.001	3.24	2.72	0.001

Table S3.3 Results of pairwise PERMANOVA for the community composition between forest types with Benjamini-Hochberg adjustment.

Pairwise PERMANOVA	Bacterial community			Fungal community		
	<i>F</i>	R ² (%)	<i>P</i> .adj	<i>F</i>	R ² (%)	<i>P</i> .adj
Coniferous vs. Open Canopy	4.48	2.92	0.002	4.30	2.80	0.002
Coniferous vs. Mixed	2.51	1.97	0.005	1.84	1.45	0.007
Coniferous vs. Deciduous	5.10	3.72	0.002	3.60	2.67	0.002
Deciduous vs. Open Canopy	6.10	7.00	0.002	3.72	4.44	0.002
Deciduous vs. Mixed	1.78	3.03	0.019	1.95	3.37	0.004
Mixed vs. Open Canopy	4.74	6.03	0.002	4.14	5.30	0.002

Table S3.5 List of bacterial indicators for mining stages with their taxonomic information (only significant species are shown, $P < 0.05$ and $\text{stat} \geq 0.2$). “Stat” is the association statistic (the square root of the indicator value) and “P value” is the p-value of the permutational test.

ASV	stat	P value	Kingdom	Phylum	Class	Order	Family	Genus
<i>Operating</i>								
ASV_11	0.29	0.002	Bacteria	Bacteroidota	Bacteroidia	Chitinophagales	Chitinophagaceae	NA
ASV_19	0.276	0.002	Bacteria	Bacteroidota	Bacteroidia	Cytophagales	Microscillaceae	<i>OLB12</i>
ASV_84	0.318	0.001	Bacteria	Proteobacteria	Alphaproteobacteria	Acetobacterales	Acetobacteraceae	<i>Acidiphilium</i>
ASV_101	0.318	0.001	Bacteria	Acidobacteriota	Acidobacteriae	Solibacterales	Solibacteraceae	<i>Candidatus Solibacter</i>
ASV_159	0.317	0.001	Bacteria	Proteobacteria	Alphaproteobacteria	Rhizobiales	Beijerinckiaceae	<i>Methylobacterium -Methylorubrum</i>
ASV_119	0.297	0.002	Bacteria	Desulfobacterota	Desulfuromonadia	Geobacterales	Geobacteraceae	<i>Geobacter</i>
ASV_91	0.29	0.003	Bacteria	Proteobacteria	Alphaproteobacteria	Rhizobiales	Beijerinckiaceae	<i>1174-901-12</i>
ASV_145	0.287	0.002	Bacteria	Proteobacteria	Alphaproteobacteria	Sphingomonadales	Sphingomonadaceae	<i>Sphingomonas</i>
ASV_326	0.284	0.001	Bacteria	Actinobacteriota	Actinobacteria	Frankiales	Nakamurellaceae	<i>Nakamurella</i>
ASV_49	0.276	0.004	Bacteria	Acidobacteriota	Acidobacteriae	Acidobacteriales	Acidobacteriaceae	<i>Occallatibacter</i>
ASV_178	0.268	0.001	Bacteria	Proteobacteria	Alphaproteobacteria	Caulobacterales	Hyphomonadaceae	<i>SWB02</i>
ASV_156	0.267	0.003	Bacteria	Proteobacteria	Alphaproteobacteria	Rhizobiales	Xanthobacteraceae	<i>Pseudorhodoplan es</i>
ASV_360	0.266	0.004	Bacteria	Proteobacteria	Alphaproteobacteria	Acetobacterales	Acetobacteraceae	<i>Acidiphilium</i>
ASV_80	0.26	0.003	Bacteria	Actinobacteriota	Thermoleophilina	Solirubrobacterales	67-14	NA
ASV_721	0.258	0.001	Bacteria	Acidobacteriota	Acidobacteriae	Acidobacteriales	Acidobacteriaceae	<i>Bryocella</i>
ASV_160	0.251	0.004	Bacteria	Proteobacteria	Alphaproteobacteria	Sphingomonadales	Sphingomonadaceae	<i>Sphingomonas</i>
ASV_385	0.249	0.001	Bacteria	Bacteroidota	Bacteroidia	Chitinophagales	Chitinophagaceae	<i>Aurantisolimonas</i>

Table S3.5. continued

ASV_150	0.246	0.005	Bacteria	Proteobacteria	Gammaproteobacteria	Burkholderiales	Nitrosomonadaceae	<i>IS-44</i>
ASV_507	0.246	0.001	Bacteria	Actinobacteriota	Actinobacteria	Micromonosporales	Micromonosporaceae	<i>Actinoplanes</i>
ASV_161	0.245	0.001	Bacteria	Proteobacteria	Gammaproteobacteria	Burkholderiales	Oxalobacteraceae	<i>Massilia</i>
ASV_187	0.244	0.006	Bacteria	Firmicutes	Desulfitobacteria	Desulfitobacteriales	Desulfitobacteriaceae	<i>Desulfosporosinus</i>
ASV_214	0.244	0.005	Bacteria	Bacteroidota	SLA-28	NA	NA	NA
ASV_580	0.244	0.001	Bacteria	Bacteroidota	Bacteroidia	Cytophagales	Spirosomaceae	<i>Huanghella</i>
ASV_472	0.239	0.001	Bacteria	Bacteroidota	Bacteroidia	Cytophagales	Spirosomaceae	<i>Spirosoma</i>
ASV_57	0.238	0.006	Bacteria	Verrucomicrobiota	Verrucomicrobiae	Opitutales	Opitutaceae	<i>Lacunisphaera</i>
ASV_207	0.238	0.005	Bacteria	Proteobacteria	Gammaproteobacteria	Burkholderiales	Comamonadaceae	NA
ASV_332	0.23	0.001	Bacteria	Proteobacteria	Alphaproteobacteria	Rhizobiales	Beijerinckiacae	<i>1174-901-12</i>
ASV_131	0.226	0.011	Bacteria	Acidobacteriota	Acidobacteriae	Bryobacterales	Bryobacteraceae	<i>Bryobacter</i>
ASV_333	0.226	0.006	Bacteria	Bacteroidota	Bacteroidia	Chitinophagales	Chitinophagaceae	<i>Ferruginibacter</i>
ASV_379	0.221	0.003	Bacteria	Proteobacteria	Alphaproteobacteria	Sphingomonadales	Sphingomonadaceae	<i>Sphingomonas</i>
ASV_733	0.218	0.002	Bacteria	Deinococota	Deinococci	Deinococcales	Deinococcaceae	<i>Deinococcus</i>
ASV_508	0.217	0.004	Bacteria	Proteobacteria	Alphaproteobacteria	Rhizobiales	Beijerinckiacae	<i>1174-901-12</i>
ASV_1065	0.215	0.007	Bacteria	Proteobacteria	Alphaproteobacteria	Acetobacterales	Acetobacteraceae	<i>Acidiphilium</i>
ASV_188	0.209	0.002	Bacteria	Proteobacteria	Alphaproteobacteria	Acetobacterales	Acetobacteraceae	<i>Acidiphilium</i>
ASV_27	0.204	0.017	Bacteria	Proteobacteria	Alphaproteobacteria	Sphingomonadales	Sphingomonadaceae	<i>Sphingomonas</i>
<i>Non-operating</i>								
ASV_8	0.279	0.001	Bacteria	Proteobacteria	Alphaproteobacteria	Acetobacterales	Acetobacteraceae	NA
ASV_34	0.275	0.001	Bacteria	Proteobacteria	Alphaproteobacteria	Caulobacterales	Caulobacteraceae	PMMR1
ASV_147	0.266	0.001	Bacteria	Proteobacteria	Alphaproteobacteria	Acetobacterales	Acetobacteraceae	NA

Table S3.5 continued

ASV_208	0.266	0.001	Bacteria	Acidobacteriota	Acidobacteriae	Acidobacteriales	Acidobacteriaceae	<i>Granulicella</i>
ASV_132	0.262	0.001	Bacteria	Proteobacteria	Alphaproteobacteria	Sphingomonadales	Sphingomonadaceae	<i>Novosphingobium</i>
ASV_146	0.26	0.003	Bacteria	Acidobacteriota	Acidobacteriae	Acidobacteriales	Acidobacteriaceae	<i>Byocella</i>
ASV_116	0.25	0.003	Bacteria	Proteobacteria	Alphaproteobacteria	Acetobacterales	Acetobacteraceae	NA
ASV_194	0.245	0.004	Bacteria	Proteobacteria	Alphaproteobacteria	Acetobacterales	Acetobacteraceae	NA
ASV_30	0.243	0.004	Bacteria	Proteobacteria	Alphaproteobacteria	Acetobacterales	Acetobacteraceae	NA
ASV_227	0.241	0.001	Bacteria	Acidobacteriota	Acidobacteriae	Acidobacteriales	Acidobacteriaceae	<i>Granulicella</i>
ASV_88	0.233	0.006	Bacteria	Proteobacteria	Alphaproteobacteria	Acetobacterales	Acetobacteraceae	NA
ASV_169	0.231	0.004	Bacteria	Proteobacteria	Alphaproteobacteria	Sphingomonadales	Sphingomonadaceae	<i>Novosphingobium</i>
ASV_85	0.23	0.004	Bacteria	Bacteroidota	Bacteroidia	Chitinophagales	Chitinophagaceae	<i>Aurantisolimonas</i>
ASV_66	0.228	0.01	Bacteria	Acidobacteriota	Acidobacteriae	Acidobacteriales	Acidobacteriaceae	<i>Granulicella</i>
ASV_250	0.222	0.001	Bacteria	Proteobacteria	Alphaproteobacteria	Caulobacterales	Caulobacteraceae	NA
ASV_86	0.221	0.011	Bacteria	Acidobacteriota	Acidobacteriae	Acidobacteriales	Acidobacteriaceae	<i>Granulicella</i>
ASV_112	0.221	0.011	Bacteria	Proteobacteria	Alphaproteobacteria	Acetobacterales	Acetobacteraceae	<i>Acidisona</i>
ASV_89	0.217	0.007	Bacteria	Proteobacteria	Alphaproteobacteria	Caulobacterales	Caulobacteraceae	NA
ASV_94	0.214	0.007	Bacteria	Acidobacteriota	Acidobacteriae	Acidobacteriales	Acidobacteriaceae	<i>Byocella</i>
ASV_51	0.213	0.005	Bacteria	Proteobacteria	Alphaproteobacteria	Acetobacterales	Acetobacteraceae	<i>Acidiphilium</i>
ASV_100	0.213	0.01	Bacteria	Planctomycetota	Planctomycetes	Isophaerales	Isophaeraceae	<i>Tundritsphaera</i>
ASV_74	0.209	0.009	Bacteria	Proteobacteria	Alphaproteobacteria	Acetobacterales	Acetobacteraceae	<i>Acidisona</i>
ASV_354	0.204	0.013	Bacteria	Acidobacteriota	Acidobacteriae	Acidobacteriales	Acidobacteriaceae	<i>Granulicella</i>
ASV_148	0.203	0.022	Bacteria	Proteobacteria	Alphaproteobacteria	Acetobacterales	Acetobacteraceae	<i>Acidisphaera</i>
ASV_192	0.203	0.015	Bacteria	Proteobacteria	Alphaproteobacteria	Acetobacterales	Acetobacteraceae	NA

Table S3.5 continued

Controls								
ASV_361	0.514	0.001	Bacteria	Proteobacteria	Gammaproteobacteria	Burkholderiales	Burkholderiaceae	Cupriavidus
ASV_543	0.386	0.001	Bacteria	Acidobacteriota	Acidobacteriae	Subgroup 2	NA	NA
ASV_373	0.364	0.001	Bacteria	Proteobacteria	Alphaproteobacteria	Rhizobiales	Beijerinckiaceae	Roseiarcus
ASV_244	0.358	0.001	Bacteria	Proteobacteria	Gammaproteobacteria	Salinisphaerales	Solimonadaceae	Nevskia
ASV_278	0.358	0.001	Bacteria	Proteobacteria	Gammaproteobacteria	Xanthomonadales	Xanthomonadaceae	Stenotrophomonas
ASV_245	0.347	0.001	Bacteria	Bacteroidota	Bacteroidia	Chitinophagales	Chitinophagaceae	Puia
ASV_299	0.307	0.001	Bacteria	Proteobacteria	Gammaproteobacteria	Xanthomonadales	Rhodanobacteraceae	Rhodanobacter
ASV_38	0.299	0.001	Bacteria	Proteobacteria	Alphaproteobacteria	Rhizobiales	Xanthobacteraceae	Bradyrhizobium
ASV_199	0.299	0.001	Bacteria	Proteobacteria	Gammaproteobacteria	WD260	NA	NA
ASV_226	0.258	0.001	Bacteria	Proteobacteria	Gammaproteobacteria	Burkholderiales	Comamonadaceae	Variovorax
ASV_168	0.241	0.001	Bacteria	Proteobacteria	Gammaproteobacteria	Pseudomonadales	Pseudomonadaceae	Pseudomonas
ASV_167	0.212	0.006	Bacteria	Acidobacteriota	Acidobacteriae	Acidobacteriales	Acidobacteriaceae	Granulicella
ASV_106	0.208	0.01	Bacteria	Bacteroidota	Bacteroidia	Sphingobacteriales	Sphingobacteriaceae	Mucilaginibacter

Table S3.6 List of fungal indicators for mining stages with their taxonomic information (only significant species are shown $P < 0.05$ and $\text{stat} \geq 0.2$). “Stat” is the association statistic (the square root of the indicator value) and “P value” is the p-value of the permutational test.

ASV	stat	P	Kingdom	Phylum	Class	Order	Family	Genus
<i>Operating</i>								
ASV_10	0.326	0.001	Fungi	Basidiomycota	Microbotryomycetes	Kriegeriales	Kriegeriaceae	<i>Phenoliferia</i>
ASV_21	0.32	0.001	Fungi	Chytridiomycota	Spizellomycetes	Spizellomycetales	NA	NA
ASV_86	0.32	0.001	Fungi	Chytridiomycota	Spizellomycetes	Spizellomycetales	Powellomycetaceae	<i>Thoreauomyces</i>
ASV_115	0.314	0.001	Fungi	Ascomycota	Dothideomycetes	Mytiliniidiales	Mytiliniaceae	<i>Lophium</i>
ASV_212	0.259	0.001	Fungi	Ascomycota	Dothideomycetes	Mytiliniidiales	Mytiliniaceae	<i>Lophium</i>
ASV_98	0.256	0.001	Fungi	Basidiomycota	Tremellomycetes	Tremellales	NA	NA
ASV_235	0.254	0.001	Fungi	Basidiomycota	Microbotryomycetes	Sporidiobolales	Sporidiobolaceae	<i>Rhodospordiobolus</i>
ASV_453	0.253	0.002	Fungi	Ascomycota	Dothideomycetes	Capnodiales	Teratosphaeriaceae	<i>Devriesia</i>
ASV_29	0.247	0.003	Fungi	Chytridiomycota	Rhizophyidiomycetes	Rhizophydiales	Rhizophyidiaceae	<i>Rhizophydium</i>
ASV_193	0.245	0.004	Fungi	Chytridiomycota	Rhizophyidiomycetes	Rhizophydiales	Rhizophyidiaceae	<i>Rhizophydium</i>
ASV_314	0.245	0.001	Fungi	Ascomycota	Dothideomycetes	Pleosporales	Phaeosphaeriaceae	<i>Setomelanomna</i>
ASV_250	0.241	0.004	Fungi	NA	NA	NA	NA	NA
ASV_616	0.24	0.004	Fungi	Ascomycota	Dothideomycetes	Dothideales	Dothideaceae	<i>Rhizosphaera</i>
ASV_414	0.237	0.003	Fungi	NA	NA	NA	NA	NA
ASV_127	0.233	0.004	Fungi	Basidiomycota	Tremellomycetes	Cystofilobasidiales	Mrakiaceae	<i>Mrakia</i>
ASV_59	0.228	0.002	Fungi	Ascomycota	NA	NA	NA	NA

Table S3.6 continued

ASV_100	0.226	0.01	Fungi	Ascomycota	Dothideomycetes	Capnodiales	Neodevriesiaceae	NA
ASV_217	0.223	0.005	Fungi	NA	NA	NA	NA	NA
ASV_39	0.221	0.011	Fungi	Ascomycota	Leotiomycetes	Leotiales	Mniaciacae	<i>Epithamnolia</i>
ASV_199	0.219	0.004	Fungi	Chytridiomycota	Spizellomycetes	Spizellomycetales	NA	NA
ASV_698	0.219	0.006	Fungi	NA	NA	NA	NA	NA
ASV_169	0.218	0.009	Fungi	Ascomycota	Dothideomycetes	Capnodiales	Teratosphaeriaceae	<i>Capnobotryella</i>
ASV_897	0.218	0.012	Fungi	Basidiomycota	Tremellomycetes	Tremellales	NA	NA
ASV_75	0.217	0.006	Fungi	Ascomycota	Pezizomycotina_ cls_Incertae_sedis	Pezizomycotina_ ord_Incertae_sedis	Pezizomycotina_ fam_Incertae_sedis	<i>Ciliophora</i>
ASV_6	0.217	0.013	Fungi	NA	NA	NA	NA	NA
ASV_117	0.214	0.003	Fungi	NA	NA	NA	NA	NA
ASV_164	0.209	0.005	Fungi	Ascomycota	Dothideomycetes	Venturiales	Venturiaceae	<i>Venturia</i>
ASV_622	0.208	0.003	Fungi	Basidiomycota	Tremellomycetes	Filobasidiales	Piskurozymaceae	NA
ASV_1240	0.207	0.007	Fungi	NA	NA	NA	NA	NA
ASV_352	0.204	0.007	Fungi	NA	NA	NA	NA	NA
ASV_544	0.204	0.011	Fungi	Basidiomycota	Agaricomycetes	Cantharellales	Ceratobasidiaceae	<i>Ceratobasidium</i>
ASV_316	0.2	0.012	Fungi	Olpidiomycota	GS18	NA	NA	NA
<i>Non-operating</i>								
ASV_136	0.314	0.001	Fungi	Ascomycota	Orbiliomycetes	Orbiliales	NA	NA
ASV_186	0.245	0.004	Fungi	Ascomycota	Eurotiomycetes	Chaetothyriales	NA	NA
ASV_50	0.222	0.005	Fungi	Ascomycota	Dothideomycetes	Dothideales	NA	NA
ASV_244	0.222	0.001	Fungi	Ascomycota	Eurotiomycetes	Chaetothyriales	Herpotrichiellaceae	<i>Cladophialophora</i>
ASV_155	0.22	0.005	Fungi	Ascomycota	Eurotiomycetes	Chaetothyriales	Herpotrichiellaceae	NA

Table S3.6 continued

ASV_504	0.212	0.009	Fungi	Ascomycota	Eurotiomycetes	Chaetothyriales	Herpotrichiellaceae	<i>Cladophialophora</i>
ASV_188	0.208	0.01	Fungi	Ascomycota	Leotiomycetes	Helotiales	NA	NA
<i>Controls</i>								
ASV_429	0.397	0.001	Fungi	Basidiomycota	Agaricomycetes	Thelephorales	Thelephoraceae	<i>Thelephora</i>
ASV_634	0.395	0.001	Fungi	Ascomycota	Pezizomycotina_ cls_Incertae_sedis	Pezizomycotina_ ord_Incertae_sedis	Pezizomycotina_ fam_Incertae_sedis	<i>Ciliophora</i>
ASV_729	0.395	0.001	Fungi	NA	NA	NA	NA	NA
ASV_659	0.391	0.001	Fungi	Ascomycota	Pezizomycotina_ cls_Incertae_sedis	Pezizomycotina_ ord_Incertae_sedis	Pezizomycotina_ fam_Incertae_sedis	<i>Ciliophora</i>
ASV_502	0.389	0.001	Fungi	Basidiomycota	Tremellomycetes	Filobasidiales	Piskurozymaceae	<i>Solicoceozyma</i>
ASV_673	0.388	0.001	Fungi	Ascomycota	Leotiomycetes	Helotiales	NA	NA
ASV_675	0.386	0.001	Fungi	Ascomycota	Eurotiomycetes	Eurotiales	Trichocomaceae	<i>Talaromyces</i>
ASV_441	0.384	0.001	Fungi	Basidiomycota	Agaricomycetes	Agaricales	Inocybaceae	<i>Inocybe</i>
ASV_490	0.384	0.001	Fungi	Ascomycota	Dothideomycetes	Pleosporales	Phaeosphaeriaceae	<i>Paraphoma</i>
ASV_542	0.384	0.001	Fungi	Mucromycota	Umbelopsidomycetes	Umbelopsidales	Umbelopsidaceae	<i>Umbelopsis</i>
ASV_509	0.383	0.001	Fungi	NA	NA	NA	NA	NA
ASV_603	0.383	0.001	Fungi	Basidiomycota	Agaricomycetes	Russulales	Russulaceae	<i>Lactifusus</i>
ASV_559	0.382	0.001	Fungi	Ascomycota	Leotiomycetes	Helotiales	Hyaloscyphaceae	NA
ASV_405	0.376	0.001	Fungi	Ascomycota	Leotiomycetes	Helotiales	Helotiaceae	<i>Ascocoryne</i>
ASV_76	0.373	0.001	Fungi	Ascomycota	Pezizomycetes	Pezizales	Pyrenomataceae	<i>Sphaerosporella</i>
ASV_389	0.373	0.001	Fungi	Ascomycota	Leotiomycetes	Helotiales	NA	NA
ASV_266	0.372	0.001	Fungi	Ascomycota	Leotiomycetes	Helotiales	Helotiales_fam_ Incertae_sedis	<i>Leptodontidium</i>
ASV_123	0.371	0.001	Fungi	NA	NA	NA	NA	NA

Table S3.6 continued

ASV_355	0.371	0.001	Fungi	NA	NA	NA	NA	NA	NA
ASV_379	0.371	0.001	Fungi	NA	NA	NA	NA	NA	NA
ASV_19	0.37	0.001	Fungi	NA	NA	NA	NA	NA	NA
ASV_454	0.369	0.001	Fungi	Ascomycota	Leotiomycetes	Helotiales	Hyaloscyphaceae	<i>Hyaloscypha</i>	
ASV_159	0.368	0.001	Fungi	Ascomycota	Leotiomycetes	Helotiales	Helotiaceae	<i>Meliniomyces</i>	
ASV_61	0.365	0.001	Fungi	Basidiomycota	Microbotryomycetes	Leucosporidiales	Leucosporidiaceae	<i>Leucosporidium</i>	
ASV_738	0.364	0.001	Fungi	Ascomycota	Leotiomycetes	Thelebolales	Pseudoeurotiaceae	NA	
ASV_101	0.363	0.001	Fungi	Basidiomycota	Agaricomycetes	Agaricales	Strophariaceae	<i>Hypopholoma</i>	
ASV_176	0.363	0.001	Fungi	Basidiomycota	Agaricomycetes	Agaricales	Tricholomataceae	<i>Mycena</i>	
ASV_411	0.362	0.001	Fungi	Ascomycota	Leotiomycetes	Thelebolales	NA	NA	
ASV_182	0.359	0.001	Fungi	NA	NA	NA	NA	NA	
ASV_530	0.359	0.001	Fungi	Ascomycota	Eurotiomycetes	Chaetothyriales	Hepotrichiellaceae	NA	
ASV_589	0.358	0.001	Fungi	NA	NA	NA	NA	NA	
ASV_228	0.354	0.001	Fungi	NA	NA	NA	NA	NA	
ASV_281	0.353	0.001	Fungi	NA	NA	NA	NA	NA	
ASV_118	0.352	0.001	Fungi	Ascomycota	Sordariomycetes	Xylariales	Xylariaceae	<i>Hypoxylon</i>	
ASV_214	0.351	0.001	Fungi	Ascomycota	Sordariomycetes	Xylariales	Xylariaceae	<i>Hypoxylon</i>	
ASV_309	0.349	0.001	Fungi	Basidiomycota	Microbotryomycetes	Kriegeriales	Kriegeriaceae	<i>Phenoliferia</i>	
ASV_242	0.347	0.001	Fungi	Mortierellomycota	Mortierellomycetes	Mortierellales	Mortierellaceae	<i>Mortierella</i>	
ASV_631	0.345	0.001	Fungi	Ascomycota	Saccharomycetes	Saccharomycetales	Saccharomycetales_fam_Incertae_sedis	<i>Nadsonia</i>	
ASV_567	0.333	0.001	Fungi	Basidiomycota	Agaricomycetes	Agaricales	Hymenogastreae	<i>Hebeloma</i>	
ASV_187	0.296	0.001	Fungi	Ascomycota	Leotiomycetes	Helotiales	Hyaloscyphaceae	<i>Hyaloscypha</i>	

Table S3.6 continued

ASV_398	0.287	0.001	Fungi	Ascomycota	Leotiomycetes	Helotiales	Helotiales_fam_Incertae_sedis	<i>Leptodontidium</i>
ASV_525	0.273	0.001	Fungi	Ascomycota	Eurotiomycetes	Chaetothyriales	NA	NA
ASV_143	0.259	0.001	Fungi	Ascomycota	Leotiomycetes	Helotiales	Helotiaceae	<i>Collophora</i>
ASV_344	0.256	0.001	Fungi	Ascomycota	Leotiomycetes	Helotiales	Helotiales_fam_Incertae_sedis	<i>Cadophora</i>
ASV_395	0.252	0.003	Fungi	Ascomycota	Eurotiomycetes	Chaetothyriales	Herpotrichiellaceae	<i>Rhinochlaetella</i>
ASV_333	0.246	0.001	Fungi	Ascomycota	NA	NA	NA	NA
ASV_442	0.244	0.005	Fungi	Ascomycota	Dothideomycetes	Capnodiales	NA	NA
ASV_110	0.241	0.005	Fungi	Ascomycota	Eurotiomycetes	Chaetothyriales	NA	NA
ASV_318	0.239	0.003	Fungi	Basidiomycota	Agaricomycetes	Thelephorales	Thelephoraceae	<i>Thelephora</i>
ASV_678	0.232	0.002	Fungi	Ascomycota	Dothideomycetes	Venturiales	Venturiaceae	<i>Venturia</i>
ASV_321	0.231	0.002	Fungi	Ascomycota	NA	NA	NA	NA
ASV_608	0.229	0.001	Fungi	NA	NA	NA	NA	NA
ASV_606	0.22	0.007	Fungi	Basidiomycota	Tremellomycetes	Cystofilobasidiales	Mrkaniaceae	<i>Mrakia</i>
ASV_239	0.218	0.006	Fungi	Basidiomycota	Agaricomycetes	Polyporales	Ganodermataceae	<i>Ganoderma</i>
ASV_356	0.207	0.013	Fungi	Ascomycota	Leotiomycetes	Helotiales	Leotiaceae	<i>Pezoloma</i>
ASV_285	0.202	0.01	Fungi	Ascomycota	Eurotiomycetes	Chaetothyriales	NA	NA
ASV_78	0.2	0.012	Fungi	Chytridiomycota	Spizellomycetes	Spizellomycetales	NA	NA

Table S3.7 Phyllosphere alpha diversity at each distance level from mine sites. Mean $\pm$ SD are shown. Significant p-values for the effects of the distance, mining stage and their interactions were calculated with ANOVA using Type II sum of squares based on the results of GLMMs. Different lowercase letters represent significant multiple compositions with Tukey's post hoc tests at the 0.05 level. Significant relationships are highlighted in bold.

Alpha diversity	ANOVA (type II sums)		Distance from mine sites (km)			
	Distance	Distance* Stage	0 (N=39)	0.05 (N=44)	0.2 (N=43)	1 (N=44)
Bacteria						
Observed	p<0.01	p=0.63	640.85 $\pm$ 180.97 b	584.09 $\pm$ 180.41 ab	550.28 $\pm$ 183.82 a	568.25 $\pm$ 190.34 a
Shannon	p<0.01	p=0.86	5.72 $\pm$ 0.43 b	5.55 $\pm$ 0.47 ab	5.45 $\pm$ 0.50 a	5.51 $\pm$ 0.43 a
InvSimpson	p=0.03	p=0.81	153.42 $\pm$ 78.92 a	127.28 $\pm$ 68.16 a	115.77 $\pm$ 59.95 a	119.45 $\pm$ 63.43 a
Fungi						
Observed	p<0.01	p=0.47	462.50 $\pm$ 142.32 b	382.16 $\pm$ 104.14 a	396.40 $\pm$ 129.17 a	404.80 $\pm$ 113.62 ab
Shannon	p=0.01	p=0.64	4.88 $\pm$ 0.38 b	4.62 $\pm$ 0.47 a	4.69 $\pm$ 0.41 ab	4.69 $\pm$ 0.48 ab
InvSimpson	p=0.33	p=0.42	55.31 $\pm$ 21.65 a	45.84 $\pm$ 25.13 a	47.96 $\pm$ 22.99 a	48.34 $\pm$ 25.71 a

Table S3.8 PERMANOVA results showing the relative importance of distance from mines, mining stage and their interactions in structuring the phyllosphere communities based on Bray-Curtis dissimilarity.

	Df	SumsOfSqs	MeanSqs	F.Model	R ² (%)	Pr(>F)
Bacterial community						
Mining stage (Stage)	1	1.34	1.34	5.09	3%	< 0.001
Distance	1	0.66	0.66	2.50	1%	< 0.001
Distance: Stage	1	0.33	0.33	1.26	1%	0.07
Residuals	166	43.58	0.26	0.95		
Total	169	45.91	1.00			
Fungal community						
Mining stage (Stage)	1	0.97	0.97	2.67	2%	< 0.001
Distance	1	0.64	0.64	1.74	1%	< 0.001
Distance: Stage	1	0.42	0.42	1.14	1%	0.06
Residuals	165	60.17	0.36	0.97		
Total	168	62.20	1.00			

Table S3.9 Relative abundance of main phyllosphere phyla (i.e. relative abundance > 0.5 through all samples) at each distance from mine sites. Mean $\pm$ SD are shown. Zero-inflated beta mixed regression models were used to compare mean relative abundance of phyllosphere phylum. Lowercase letters represent significant differences based on Tukey's test at the 0.05 level. Significant relationships (significant differences in Tukey's test at 0.05 level) are highlighted in bold.

Phylum	Mining Stage	Distance (km)			
		0	0.05	0.2	1
Bacteria					
Proteobacteria		52.28 ± 9.7 a	54.49 ± 8.56 a	55.43 ± 7.89 a	55.2 ± 8.01 a
Acidobacteriota	Operating	11.96 ± 5.69 a	15.28 ± 5.63 ab	18.07 ± 6.02 b	19.52 ± 4.8 b
	Non-Operating	19.82 ± 7.07 a	18.48 ± 6.53 a	22.36 ± 6.78 a	20.13 ± 7.3 a
Bacteroidota	Operating	16.74 ± 5.98 b	12.68 ± 6.77 a	10.34 ± 6.06 a	9.57 ± 6.06 a
	Non-Operating	10.09 ± 5.52 a	10.13 ± 5.9 a	5.68 ± 3.83 a	7.43 ± 5.16 a
Actinobacteriota	Operating	5.33 ± 2.22 b	5.22 ± 3.58 ab	4.29 ± 2.02 ab	3.82 ± 1.75 a
	Non-Operating	4.38 ± 1.88 a	4.39 ± 1.62 a	3.53 ± 1.45 a	4.45 ± 2.04 a
Planctomycetota		3.45 ± 1.86 a	2.75 ± 1.65 a	2.77 ± 1.4 a	3.14 ± 1.69 a
Verrucomicrobiota		2.46 ± 1.71 a	1.86 ± 1.51 a	1.84 ± 1.5 a	2.14 ± 1.61 a
Cyanobacteria		0.98 ± 1.06 a	1.8 ± 1.95 a	2.33 ± 4.04 a	1.33 ± 1.46 a
Myxococcota		1.59 ± 0.95 a	1.74 ± 1.82 a	1.22 ± 0.75 a	1.4 ± 0.71 a
WPS-2	Operating	0.17± 0.15 a	0.40 ± 0.31 ab	0.77 ± 0.80 b	1.01 ± 0.88 b
	Non-Operating	0.89 ± 0.79 a	0.971 ± 0.92 a	1.54 ± 1.20 a	1.13 ± 0.86 a
Firmicutes		1.04 ± 1.27 a	0.72 ± 1.19 a	0.71 ± 1.16 a	0.86 ± 1.35 a
Armatimonadota	Operating	0.96 ± 0.6 b	0.94 ± 0.52 b	0.81 ± 0.41 ab	0.61 ± 0.36 a
	Non-Operating	0.92 ± 0.38 b	0.56 ± 0.22 a	0.53 ± 0.35 a	0.61 ± 0.28 ab
Fungi					
Ascomycota		56.42 ± 10.23 a	55.21 ± 12.86 a	54.72 ± 10.61 a	54.32 ± 12.08 a
Basidiomycota		18.53 ± 9.75 a	19.84 ± 10.94 a	21.31 ± 9.72 a	22.41 ± 10.26 a
Chytridiomycota		8.98 ± 7.1 a	9.02 ± 5.84 a	9.95 ± 8.6 a	7.59 ± 6.4 a
Olpidiomycota		2.76 ± 2.94 a	2.61 ± 2.68 a	2.63 ± 2.87 a	3.06 ± 4.21 a

Table S3.10 Relative abundance of top 10 phyllosphere genera at each distance from mine sites. Mean $\pm$ SD are shown. Zero-inflated beta mixed regression models were used to compare mean relative abundance of phyllosphere genera. Lowercase letters represent significant differences based on Tukey's test at the 0.05 level. Significant relationships (i.e. significant differences in Tukey test at 0.05 level) are highlighted in bold.

Genus	Mining Stage	Distance (km)			
		0	0.05	0.2	1
Bacterial genus					
<i>Granulicella</i>	Operating	6.33 ± 5.3 a	8.57 ± 4.71 b	11.07 ± 4.56 bc	11.77 ± 4.32 c
	Non-Operating	11.91 ± 5.12 a	11.94 ± 4.06 a	13.75 ± 4.08 a	12.15 ± 4.28 a
<i>Sphingomonas</i>	Operating	7.03 ± 3.61 b	5.85 ± 3.39 ab	5.55 ± 3.32 ab	4.19 ± 2.63 a
	Non-Operating	5.61 ± 2.87 a	6.68 ± 3.74 a	4.43 ± 2.5 a	6.3 ± 3.53 a
<i>Acidiphilium</i>		4.36 ± 2.68 a	4.17 ± 2.03 a	4.85 ± 2.92 a	3.8 ± 1.74 a
<i>g117490112</i>		3.06 ± 1.71 a	3.3 ± 1.54 a	3.44 ± 1.53 a	3.5 ± 1.39 a
<i>Mucilaginibacter</i>		2.88 ± 1.48 a	3.01 ± 1.42 a	2.64 ± 1.2 a	2.99 ± 1.6 a
<i>Variovorax</i>		1.72 ± 2.2 a	1.84 ± 2.49 a	2.38 ± 3.71 a	2.7 ± 4.06 a
<i>Leptothrix</i>	Operating	1.72 ± 1.22 a	2.74 ± 1.75 b	2.68 ± 1.22 b	3.1 ± 2.47 b
	Non-Operating	2.63 ± 1.43 a	2.56 ± 1.32 a	2.26 ± 1.4 a	2.41 ± 1.48 a
<i>Rhizobacter</i>		2.51 ± 1.69 a	2.37 ± 1.44 a	2.63 ± 1.46 a	2.09 ± 1.26 a
<i>Limnobacter</i>		2.27 ± 1.52 a	2.37 ± 1.36 a	2.25 ± 1.14 a	2.19 ± 1.22 a
<i>Terriglobus</i>	Operating	1.3 ± 1.03 a	2.11 ± 2.12 b	2.06 ± 1.04 b	1.96 ± 1.1 b
	Non-Operating	2.23 ± 1.18 a	2.39 ± 0.96 a	2.62 ± 1.39 a	2.47 ± 1.2 a
Fungal genus					
<i>Phenoliferia</i>		6.08 ± 5.89 a	6.67 ± 4.9 a	7.65 ± 5.89 a	6.73 ± 5.14 a
<i>Sporormiella</i>		2.41 ± 3.79 a	4.19 ± 8.67 a	3.46 ± 6.64 a	3.93 ± 8.62 a
<i>Cladophialophora</i>		2.96 ± 3.03 a	1.98 ± 1.97 a	2.16 ± 3.34 a	2.19 ± 2.47 a
<i>Ciliophora</i>		2.22 ± 2.26 a	1.72 ± 1.45 a	1.63 ± 1.3 a	1.54 ± 1.43 a
<i>Epibryon</i>		1.28 ± 1.89 a	1.41 ± 1.52 a	1.47 ± 1.43 a	1 ± 0.89 a
<i>Epithamnolia</i>		0.93 ± 1.15 a	1.26 ± 1.99 ab	1.58 ± 1.55 ab	2.14 ± 1.99 b
<i>Thoreauomyces</i>	Operating	0.93 ± 1.2 a	0.98 ± 0.96 a	1.96 ± 2.14 b	1.61 ± 1.69 ab
	Non-Operating	1.25 ± 0.69 a	1.35 ± 1 a	1.03 ± 0.86 a	1.4 ± 1.62 a
<i>Sistotrema</i>		0.79 ± 1.95 a	0.62 ± 1.66 a	1.78 ± 5.23 a	1.92 ± 7.65 a
<i>Leucosporidium</i>		0.89 ± 1.74 a	0.6 ± 1 a	0.63 ± 0.79 a	0.96 ± 1.84 a
<i>Eocronartium</i>		1.17 ± 4.19 a	1.46 ± 5.17 a	0.52 ± 1.56 a	0.4 ± 1.56 a

Table S3.11 Relative abundance of phyllosphere common ASVs at each distance from mine sites. Mean $\pm$ SD are shown. Zero-inflated beta mixed regression models were used to compare mean relative abundance of phyllosphere ASVs. Lowercase letters represent significant differences based on Tukey's test at the 0.05 level. Significant relationships (significant differences in Tukey test at 0.05 level) are highlighted in bold. Common ASVs were identified based on the presence in at least in 90 % of the samples for bacteria and 70 % of the samples for fungi.

ASV	Taxonomic information			Mining Stage	Distance (km)			
	Order	Family	Genus		0	0.05	0.2	1
Bacterial ASV								
ASV1	Burkholderiales	Burkholderiaceae	<i>Limnobacter</i>		0.02 ± 0.01 a	0.02 ± 0.01 a	0.02 ± 0.01 a	0.02 ± 0.01 a
ASV3	Sphingomonadales	Sphingomonadaceae	<i>Sphingomonas</i>		1.45 ± 1.17 a	1.84 ± 1.64 a	1.88 ± 1.23 a	2.07 ± 1.53 a
ASV4	Burkholderiales	Comamonadaceae	<i>Leptothrix</i>	Operating	1.29 ± 1.02 a	2.23 ± 1.54 b	2.2 ± 1.12 b	2.38 ± 1.64 b
				Non-Operating	2.13 ± 1.21 a	2.15 ± 1.03 a	1.98 ± 1.2 a	1.96 ± 1.19 a
ASV5	Acetobacterales	Acetobacteraceae	<i>Acidiphilium</i>		1.35 ± 1.47 a	1.36 ± 0.98 a	1.94 ± 1.72 a	1.52 ± 1.04 a
ASV6	Burkholderiales	Burkholderiaceae	<i>Caballeronia</i>		0.91 ± 1.31 a	1.13 ± 0.89 ab	1.4 ± 1.18 b	1.65 ± 1.17 b
ASV7	Burkholderiales	Comamonadaceae	<i>Rhizobacter</i>		1.21 ± 0.81 a	1.07 ± 0.68 a	1.32 ± 0.74 a	0.97 ± 0.66 a
ASV8	Acetobacterales	Acetobacteraceae	<i>N/A</i>		0.61 ± 0.8 a	0.88 ± 0.77 a	1.37 ± 1.44 a	1.04 ± 0.91 a
ASV10	Acidobacteriales	Acidobacteriaceae	<i>Terriglobus</i>		0.67 ± 0.57 a	0.97 ± 1.59 a	0.93 ± 0.63 a	0.97 ± 0.71 a
ASV12	Burkholderiales	Comamonadaceae	<i>Rhizobacter</i>		0.85 ± 0.9 a	0.77 ± 0.58 a	0.97 ± 0.7 a	0.72 ± 0.56 a
ASV14	Acidobacteriales	Acidobacteriaceae	<i>Granulicella</i>		0.53 ± 0.38 a	0.76 ± 0.64 a	0.56 ± 0.33 a	0.7 ± 0.35 a
ASV15	Sphingomonadales	Sphingomonadaceae	<i>Sphingomonas</i>		0.59 ± 0.44 ab	0.78 ± 0.55 b	0.6 ± 0.5 ab	0.48 ± 0.37 a
ASV16	WD260	<i>N/A</i>	<i>N/A</i>		0.39 ± 0.37 a	0.39 ± 0.37 a	0.53 ± 0.46 a	0.66 ± 0.6 a
ASV17	Sphingomonadales	Sphingomonadaceae	<i>Sphingomonas</i>	Operating	0.51 ± 0.63 a	0.44 ± 0.33 a	0.78 ± 0.81 a	0.46 ± 0.35 a
				Non-Operating	0.57 ± 0.33 ab	0.53 ± 0.44 ab	0.38 ± 0.34 a	0.71 ± 0.45 b
ASV18	WD260	<i>N/A</i>	<i>N/A</i>		0.39 ± 0.39 a	0.37 ± 0.26 a	0.54 ± 0.42 a	0.7 ± 0.77 a

Table S3.11 continued

ASV20	Acidobacteriales	Acidobacteriaceae	<i>Granulicella</i>		0.39 ± 0.37 a	0.48 ± 0.42 a	0.41 ± 0.27 a	0.44 ± 0.32 a
ASV21	Acidobacteriales	Acidobacteriaceae	<i>Granulicella</i>	Operating	0.14 ± 0.16 a	0.21 ± 0.24 a	0.4 ± 0.44 ab	0.7 ± 0.69 b
				Non-Operating	0.49 ± 0.42 a	0.55 ± 0.43 a	0.6 ± 0.4 a	0.54 ± 0.42 a
ASV22	Rhizobiales	Beijerinckiaceae	<i>1174-901-12</i>		0.41 ± 0.36 a	0.49 ± 0.39 ab	0.6 ± 0.47 ab	0.66 ± 0.44 b
ASV23	Acidobacteriales	Acidobacteriaceae	<i>Granulicella</i>		0.46 ± 0.42 a	0.39 ± 0.27 a	0.38 ± 0.37 a	0.4 ± 0.35 a
ASV25	Acidobacteriales	Acidobacteriaceae	<i>Terriglobus</i>		0.41 ± 0.47 a	0.44 ± 0.44 a	0.66 ± 0.71 b	0.5 ± 0.51 ab
ASV35	Acidobacteriales	Acidobacteriaceae	<i>Granulicella</i>	Operating	0.16 ± 0.14 a	0.3 ± 0.26 ab	0.57 ± 0.4 c	0.44 ± 0.29 bc
				Non-Operating	0.36 ± 0.26 a	0.36 ± 0.22 a	0.29 ± 0.28 a	0.46 ± 0.39 a
ASV37	Acetobacterales	Acetobacteraceae	<i>NA</i>		0.31 ± 0.28 a	0.35 ± 0.42 a	0.44 ± 0.34 a	0.39 ± 0.38 a
Fungal ASV								
ASV2	Pleosporales	Sporormiaceae	<i>Sporormiella</i>		2.15 ± 3.66 a	3.76 ± 8.04 a	2.84 ± 6.2 a	3.53 ± 8.13 a
ASV3	Kriegeriales	Kriegeriaceae	<i>Phenoliferia</i>		1.64 ± 1.64 a	2.27 ± 1.67 a	2.71 ± 3.03 a	2.06 ± 2.36 a
ASV4	Chaetothyriales	<i>NA</i>	<i>NA</i>		2.38 ± 3.14 a	1.73 ± 2.5 a	2.51 ± 2.38 a	1.95 ± 1.99 a
ASV5	Kriegeriales	Kriegeriaceae	<i>Phenoliferia</i>		1.52 ± 2.08 a	1.39 ± 1.39 a	1.52 ± 1.28 a	1.44 ± 1.49 a
ASV6	<i>NA</i>	<i>NA</i>	<i>NA</i>		1.24 ± 1.43 a	1.56 ± 1.94 a	1.05 ± 1.09 a	1.13 ± 1.5 a
ASV10	Kriegeriales	Kriegeriaceae	<i>Phenoliferia</i>	Operating	1.57 ± 1.69 b	1.43 ± 1.12 b	1.15 ± 0.99 b	0.5 ± 0.53 a
				Non-Operating	0.73 ± 1.42 a	0.66 ± 0.91 a	0.33 ± 0.35 a	0.53 ± 0.52 a
ASV13	Spizellomycetales	Powellomycetaceae	<i>Thoreauomyces</i>	Operating	0.37 ± 0.47 ab	0.42 ± 0.52 a	1.05 ± 1.02 bc	1.05 ± 1.07 c
				Non-Operating	0.67 ± 0.53 a	1.01 ± 0.8 a	0.85 ± 0.87 a	1.04 ± 1.32 a
ASV21	Spizellomycetales	<i>NA</i>	<i>NA</i>		0.71 ± 0.76 a	0.65 ± 0.9 a	0.63 ± 1.13 ab	0.32 ± 0.74 b
ASV22	Capnodiales	Cladosporiaceae	<i>Cladosporium</i>		0.93 ± 1.42 b	0.61 ± 0.7 ab	0.37 ± 0.34 ab	0.4 ± 0.4 a
ASV23	Chaetothyriales	Epityraceae	<i>Epityron</i>		0.4 ± 0.38 a	0.63 ± 0.73 a	0.53 ± 0.51 a	0.51 ± 0.48 a

Table S3.11 continued

ASV26	Chaetothyriales	NA	NA	0.39 ± 0.68 a	0.4 ± 1.06 a	0.59 ± 1.04 a	0.58 ± 1.09 a
ASV28	Capnoidiales	NA	NA	0.42 ± 0.53 a	0.43 ± 0.49 ab	0.48 ± 0.43 ab	0.54 ± 0.52 b
ASV32	Dothideales	Dothioraceae	Hormonema	0.39 ± 0.54 a	0.34 ± 0.34 a	0.51 ± 0.63 a	0.51 ± 0.73 a
ASV43	Helotiales	Hyaloscyphaceae	Polydesmia	0.24 ± 0.7 a	0.29 ± 0.36 ab	0.47 ± 1.12 ab	0.72 ± 1.06 b
ASV54	Chaetothyriales	Herpotrichiellaceae	Capronia	0.29 ± 0.32 a	0.33 ± 0.46 a	0.26 ± 0.24 a	0.44 ± 0.44 a
ASV56	NA	NA	NA	0.22 ± 0.28 a	0.35 ± 0.45 a	0.43 ± 0.45 a	0.41 ± 0.59 a
ASV66	Dothideales	Aureobasidiaceae	Aureobasidium	0.77 ± 1.47 b	0.27 ± 0.73 ab	0.09 ± 0.13 a	0.11 ± 0.16 a
ASV69	NA	NA	NA	0.25 ± 0.43 a	0.24 ± 0.38 a	0.26 ± 0.36 a	0.26 ± 0.39 a
ASV95	Pleosporales	Pleosporaceae	Alternaria	0.27 ± 0.5 b	0.26 ± 0.64 ab	0.09 ± 0.1 ab	0.11 ± 0.17 a
ASV116	Orbiliales	Orbiliaceae	NA	0.13 ± 0.22 a	0.19 ± 0.24 a	0.18 ± 0.2 a	0.13 ± 0.18 a

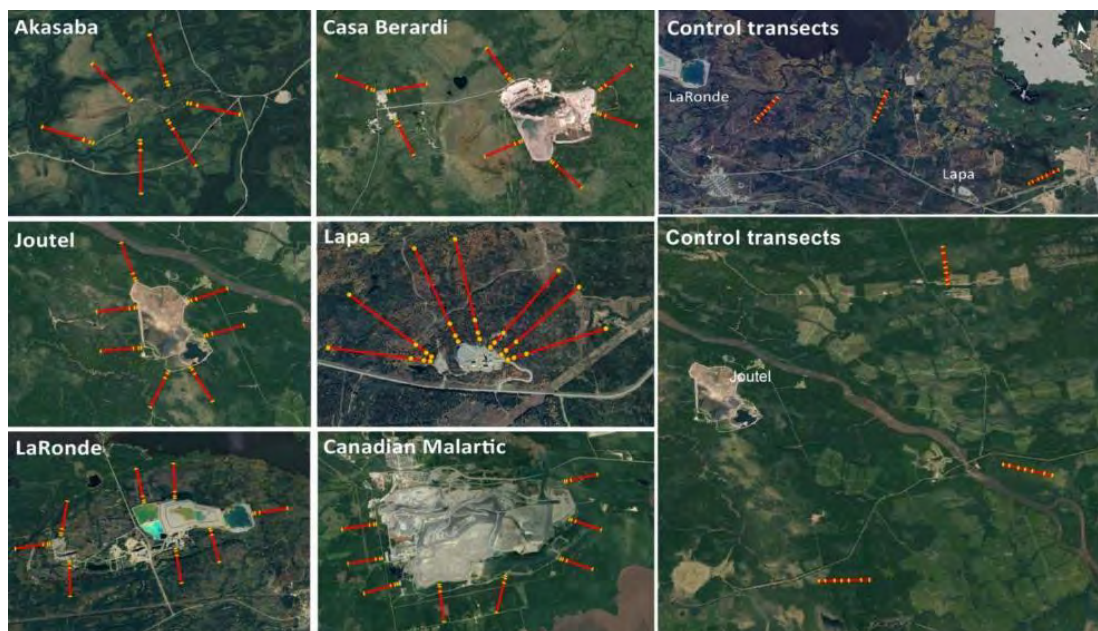


Figure S3.1 Maps of the study sites and six selected mine sites and controls showing the position of transects. Six selected mine sites and controls: Akasaba site, 6 transects, 24 plots; Casa Berardi site, 8 transects, 32 plots; Joutel site, 7 transects, 28 plots; Lapa site, 7 transects, 28 plots; LaRonde site, 8 transects, 32 plots; Canadian Malartic site, 8 transects, 32 plots; Control transects, 6 transects, 44 plots. Scales are not shown because the 1 km transects can be used as a reference in each map.

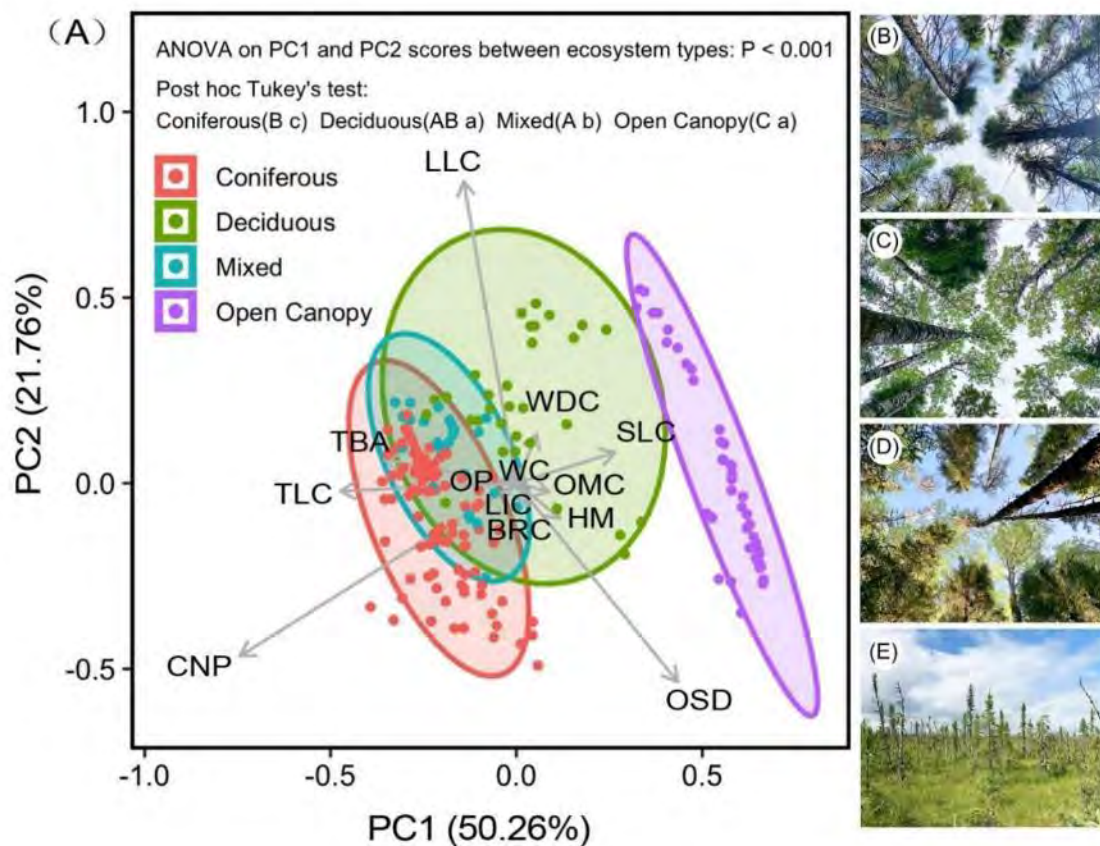


Figure S3.2 Biplot of principal component analysis (PCA) of forest structures and environmental variables across plots and forest types. Each ecosystem type (coniferous (B), deciduous (C), mixed forest (D) and open canopy ecosystem(E)) is represented by a different color and ellipse, and all plots are displayed (small circles). SLC: live crown ratio for sapling trees; TLC: live crown ratio for canopy trees; SBA: base areas for sapling trees; TBA: base areas for canopy trees; OP: Openness; LLC: leave litter ground coverage; WDC: woody debris ground coverage; WC: water ground coverage; RC: rock ground coverage; OMC: organic matter ground coverage; OSD: depth of organic soil; CNP: conifer proportion; HM: humility; BRC: bryophyte ground coverage; LIC: lichen ground coverage.

ANNEXE A

LES RISQUES ENVIRONNEMENTAUX POTENTIELS ASSOCIÉS AU
DÉVELOPPEMENT DE LA PRODUCTION D'ÉLÉMENTS DE TERRES RARES AU
CANADA

THE POTENTIAL ENVIRONMENTAL RISKS ASSOCIATED WITH THE
DEVELOPMENT OF RARE EARTH ELEMENT PRODUCTION IN CANADA

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Environmental Reviews, DOI: 10.1139/er-2020-0115

A1.1 Résumé

Le développement de la production de métaux du groupe des terres rares (MTR) au Canada pourrait générer des avantages économiques importants, mais il présente aussi des risques sérieux pour l'environnement. Les métaux du groupe des terres rares ont été largement utilisés dans la vie moderne et les industries, et ils sont même indispensables dans certaines technologies de pointe essentielles (par exemple, les aimants permanents). La demande croissante et les tensions commerciales actuelles entre les États-Unis et la Chine offrent des occasions de développement économique commercial au Canada, qui dispose de riches ressources en MTR, pour développer son propre secteur. Cependant, les problèmes d'environnement et de santé causés par la production de MTR constituent des défis auxquels le Canada doit faire face, puisque des impacts environnementaux importants ont été rapportés ailleurs (par exemple en Chine). Il existe peu de documentation sur les risques environnementaux potentiels associés au développement de la production de MTR au Canada. Il est important de savoir quels problèmes environnementaux, en particulier ceux générés par les MTR eux-mêmes, pourraient survenir au Canada dans le futur. Par conséquent, trois aspects majeurs sont évalués et résumés à partir de perspectives multidisciplinaires dans cet article : (1) un modèle conceptuel général du transport des MTR en tant que groupe dans l'environnement est établi ; (2) les niveaux de toxicité, les mécanismes biochimiques et les effets physiologiques des MTR sur différents organismes sont examinés, et des études de cas provenant de zones d'exploitation minière des MTR existantes sont brièvement mises en évidence ; et (3) en tenant compte des conditions environnementales spécifiques et des facteurs de risque, les risques environnementaux auxquels le Canada pourrait être confronté dans les développements futurs des MTR sont identifiés et discutés. Cette synthèse se termine par une macro-identification des risques environnementaux potentiels associés au développement de la production de MTR au Canada, en tenant compte de

la santé humaine et écologique. Les auteurs notent que l'ingestion, l'inhalation et l'exposition cutanée chez les travailleurs et les résidents avoisinants (y compris les communautés autochtones potentielles), ainsi que les conditions climatiques subarctiques et arctiques pourraient accroître les risques pour la santé humaine et écologique dans le développement futur de la production de MTR au Canada. Enfin, des orientations de recherche futures sont proposées qui pourraient être appliquées à la fois au contexte canadien et à d'autres contextes géographiques.

Mots-clés : lanthanides, industrie minière, contaminants émergents, risques écologiques, régions boréales.

A1.2 Abstract

The development of rare earth element (REE) production in Canada could generate significant economic benefits, but also poses serious potential risks to the environment. Rare earth elements have been widely used in modern life and industries, and even are indispensable in some crucial advanced technologies (e.g., permanent magnets). Increasing demand and the context of current US-China trade tensions provide a commercial economic development opportunity for Canada, which has rich resources of REEs, to develop its own sector. However, environmental and health issues caused by REE production are challenges Canada has to face, given that significant environmental impacts have been reported elsewhere (e.g., China). Little literature is available on the potential environmental risks associated with the development of REE production in Canada. It is important to know what environmental issues, particularly those generated by REEs themselves, may happen in Canada in the future. Therefore, three major aspects are evaluated and summarized from multidisciplinary perspectives in this paper: 1) a general conceptual model of the transport of REEs as a group in the environment is established; 2) toxicity levels, biochemical mechanisms, and physiological effects of REEs on different organisms

are reviewed, and case-studies from existing REE mining areas are briefly highlighted; and 3) Considering specific environmental condition and risk factors, environmental risks Canada may face in future REE developments are identified and discussed. This review concludes with a macro-identification of potential environmental risks associated with the development of REE production in Canada considering both human and ecological health. We note that ingestion, inhalation and dermal exposure for workers and surrounding residents (including potentially indigenous communities), and sub-arctic/arctic climate conditions could increase the risks to human and ecological health in future REE production development in Canada. Finally, future research directions are proposed that could be applied to both Canadian and other geographical contexts.

Key words: lanthanides, mining industry, emerging contaminants, ecological risks, boreal areas

A1.3 Introduction

Increasing demand for rare earth elements (REE) is stimulating an increasing number of production projects outside of China, which is currently the largest global REE producer. Several countries are planning to develop their own REE industry including Canada, which holds rich REE reserves representing about 7% of raw global resources (Zhou et al. 2017). There are large potential economic and strategic advantages for Canada in the future development of the REE industry, especially in the context of current US-China trade tensions and the economic crisis caused by COVID-19. However, REE industries can produce serious health and environmental issues, such as REE bioaccumulation (Li et al. 2014), radiation exposure (Ault et al. 2015), species invasion (Boojar and Tavakkoli 2011) and biodiversity loss (Degtjarenko et al. 2016) (see Figure A1.1). Many of these environmental issues are also present in other types of mining activities, but radioactive pollution and REE

toxicity are specific environmental risks associated with REE production that are attracting more attention (Gwenzi et al. 2018; Ma et al. 2019).

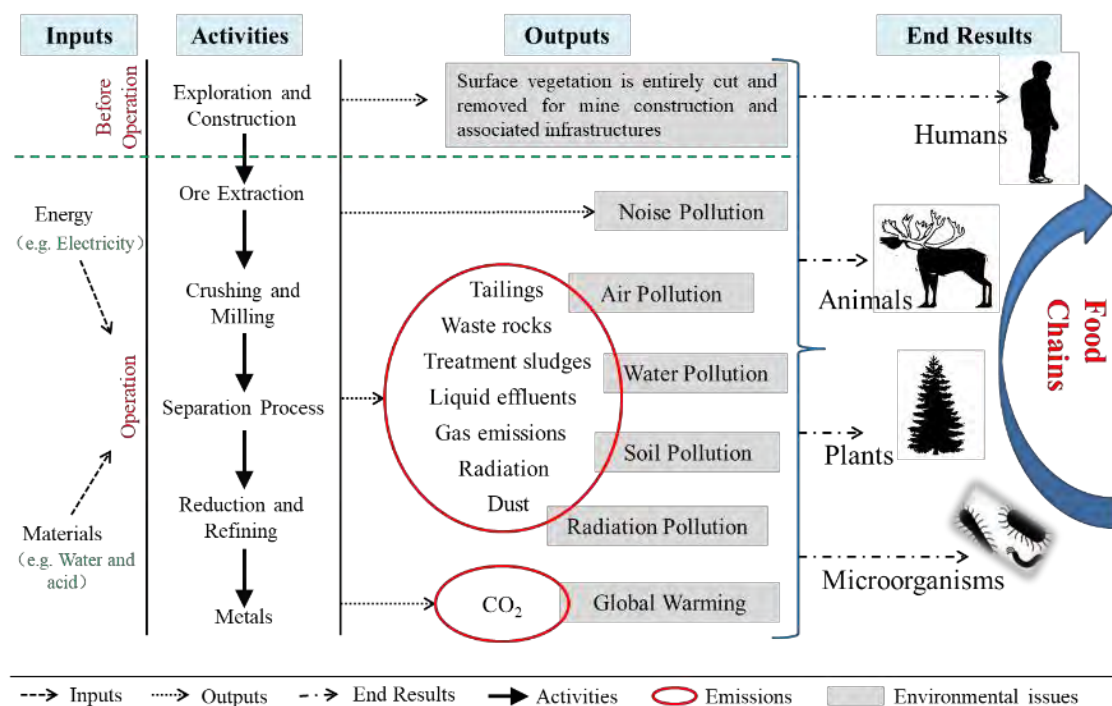


Figure A1.1 The conceptual model of the environmental issues associated with REEs production (Peiró and Méndez, 2013; Edahbi et al., 2019; Pell et al., 2019).

REE mining and processing affect the geochemical behaviour and fate of REEs in the environment, which can lead to them becoming toxicants (Figure A1.1). Firstly, products and tailings containing residual REEs generated production directly increase the content of REEs in surface environments. Secondly, mining activities could change natural soil and water physicochemical conditions (e.g., pH and redox states). Acidity can solubilize otherwise stable REE fractions, increasing their bioavailability and migration in the environment. As a result, high concentrations of REEs have been detected in soils (Liu et al. 2019), vegetables (Li et al. 2013) and other crops (Zhuang et al. 2017) and even human bodies (Li et al. 2014) near mining and processing operations. The number of REE-contaminated areas is expected to steadily increase

over the next decades with increasing REE production. Increasing REE accumulation in organisms has the potential to exceed organisms' toxicity thresholds, therefore causing health damage (e.g., growth inhibition or illness). Due to increasing numbers of exposure sources, such as tailings, wastewater and dust from mining, e-waste, REE-containing products and medical applications (Pagano et al. 2015b; Gwenzi et al. 2018; Pagano et al. 2019), the toxicity of REEs is receiving increased attention (Sauvé and Desrochers 2014; Gwenzi et al. 2018).

The development of REE production in Canada will increase REE concentrations in the environment in the coming decades. This review focuses on the associated potential risks to ecological and human health in Canada. Increased REE development adversely affecting ecosystem services has been reported in those countries with REE mining and associated processing and downstream industries (Balaram 2019, Ali 2014, Yang et al. 2013), thus the development of REE sectors will pose new environmental risks in Canada. However, few primary studies have been carried out in Canada related to the environmental concerns associated with REE production. Given that over 50 REE projects are in development in Canada (Orris et al. 2018), it is urgent to review and summarize knowledge on REE transfer and toxicity, with an integration of the characteristics of Canadian landscapes. Here, a broad range of studies on the environmental risks of REE production are reviewed, Canada-specific contexts are evaluated, potential health and ecological risks in Canada are presented, and future research directions are proposed.

A1.4 Introduction to REEs and their uses

REEs include 15 lanthanide elements in the periodic table as well as yttrium (Y) and scandium (Sc) as defined by the International Union of Pure and Applied Chemistry (IUPAC). REEs were considered as non-essential elements for organisms (Pang et al. 2002) until the discovery of a lanthanide-dependant enzyme for methane metabolism

of methanotrophic bacteria found in extreme acidic environments (Pol et al. 2014, Picone & den Camp, 2019). According to their atomic numbers, these 17 chemical elements can be divided into heavy rare earths (HREEs) of terbium (Tb), dysprosium (Dy), holmium (Ho), erbium (Er), thulium (Tm), ytterbium (Yb), lutetium (Lu), and yttrium (Y), as well as light rare earth elements (LREEs) of lanthanum (La), cerium (Ce), praseodymium (Pr), neodymium (Nd), promethium (Pm), samarium (Sm), europium (Eu), gadolinium (Gd) and scandium (Sc) (Haque et al. 2014). The REEs share chemical and physical characteristics and therefore, they are considered as a geochemically coherent group. The configuration of the valence electrons of the outermost shell is the same for lanthanides, and most of them are electropositive (+3), which results in their unique phosphorescent, magnetic, and catalytic properties (Balaram 2019). They are widely used in magnets, catalysts, alloys, phosphors, and glass production and in polishing (reviewed by Balaram, 2019; Haque et al. 2014; Weng et al. 2013). Furthermore, REEs are used as antibiotic feed additives in animal feed and fertilizers in agricultural production (Pang et al. 2002; Abdelnour et al. 2019).

A1.5 Specific environmental issues associated with REE production

REE mineral deposit types can strongly influence environmental impacts, as processing methods vary based on the physico-chemical properties of deposit (Weng et al. 2013). Significant use of chemicals (e.g., acids and alkalis), REE content in tailings and radioactive element (Th and U) concentration in ores are arguably the most common concerns (Weng et al. 2013). REE in mineral deposits are usually hosted by a wide and diverse range (more than 200) of REE-bearing minerals (the main REE-bearing minerals are reviewed by Weng et al. 2013 and Batapola et al. 2020). The REE deposits exist primarily in four geologic environments (Weng et al. 2013; Dostal, J. 2017): carbonatites, alkaline igneous systems, ion-absorption clay deposits (mainly in China), and monazite-bearing placer deposits. Carbonatite and

alkaline rocks constitute the majority of the world REE resources, but LREEs are relatively more abundant in carbonatites than HREEs, and alkaline igneous rocks have relatively low total REE grades and contain small amounts of U and Th. In Canada, the major REE-containing mineral types are carbonatite and alkaline igneous (the mineral types and geological province are listed in Table A1). Therefore, both the generation of mine tailings and the potential release of radionuclides are environmental risks for REE development in Canada.

The principal flow sheets of REE production includes five steps: 1) Mining of REE-bearing mineral ore including silicates, fluor carbonates, oxides/hydroxides, and phosphates; 2) Beneficiation of rare earth ore mixtures; 3) Separation of REE ores into individual REE minerals; 4) Reduction of REE minerals to individual REE metals; 5) Refining of individual REE metals to purify REEs (reviewed by Chi et al, 2004, Khan et al. 2017 and Wu et al. 2018). REE production and processing (mainly milling and smelting operations) are characterized by high energy and resource consumption, and high levels of pollution, as well as complicated sequential processes compared with other metals (Balomenos et al. 2017; Haque et al. 2014). For example, the extraction of 1 tonne of REE requires 4.41 tonnes of sulphuric acid, 12.32 tonnes sodium chloride, 1.64 tonnes of sodium hydroxide, 1.17 tonnes of hydrochloric acid, 1.90 tonnes of water, and milling of almost 50 tonnes of mineral ore at Bayan Obo in Mongolia, and generally, the energy used to obtain 1 tonne of an individual REE varies from 38 GJ to 48 GJ, except for scandium (148 GJ/tonne) and yttrium (75 GJ/tonne) (Peiró, & Méndez, 2013). Many potential environmental and health issues are generated by this processing with significant radionuclides, heavy metals and acids in water discharge, tailings and air emissions, as well as a high energy cost.

The radioactive pollution and REE toxicity caused by mining and processing are unique environmental risks that are different from most other mining contexts. The

radioactive pollution generated during REE production is mainly caused by the co-occurrence of REE deposits with radioactive elements such as U (uranium), Th (thorium), and their decay products (e.g., Radium) (Golev et al. 2014). The natural radionuclide content of the main REE minerals was summarized by García et al. (2020). Usually, there is a higher content of radioactive elements in the environment near a REE mining site. For example, the mean content of airborne Th in total suspended particulate ranged from 820 $\mu\text{Bq m}^{-3}$ in mining areas to 39,720 $\mu\text{Bq m}^{-3}$ in smelting areas in the Bayan Obo mining region, far higher than global background level of 0.5 $\mu\text{Bq m}^{-3}$ (Wang et al. 2016). As a result, biota near REE mines has higher health risks from exposure to high levels of radioactive elements. While radioactive pollution has been an important issue receiving attention in public health for some time, the direct toxicity of REEs is often overlooked because of the insignificant amounts found in the environment naturally, and the benefits of REEs in agricultural production. The growth in REE extraction and manufacturing has led to a strong increase of REE concentration in localized surface environments (Tepe et al. 2014; Hatje et al. 2016). For example, the concentrations of REEs in the Sidaosha River (passing through Baotou City where the Bayan Obo REE deposit is located) exceed 30,000 $\mu\text{g/L}$ of suspended particles and surface sediments, respectively, which was 200 times higher than the average value of REEs of other rivers in Northern China (Liang et al. 2014). As emerging contaminants, their toxicity is attracting more attention (Gwenzi et al. 2018; Ma et al. 2019), especially in countries like Canada planning major REE exploitation projects in the near future.

A1.6 The transport of REEs in the environment

The mobility of REEs and their movement in the environment are similar to other potentially toxic metals and metalloids (e.g., Arsenic and Mercury) driven by three main mechanisms: 1) sorption-desorption, 2) precipitation-dissolution and 3) reduction-oxidation processes (Violante et al. 2010; Cao et al. 2001). Soil and

solution components (e.g., ligands, anions, clay minerals, colloids and suspended matter), pH, and redox condition are considered as critical environmental parameters controlling these three mechanisms as they influence the transformations among REE fractions (see Table A1S1). Generally, REEs in the environment can be partitioned into six fractions based on their geochemical behaviours (see Table A1S2, Tessler et al. 1979; Cao et al. 2000): water soluble (R(ws)), ion-exchangeable (R(ie)), bound to carbonates (R(cb)), bound to iron and manganese oxides (R(fm)), bound to organic matter (R(om)), and residual (R(rd)). REE bioavailability and potential migratory capacity depends upon their fraction, ranked from strong to weak as follows: R(ws) and R(ie) > R(cb) > R(fm) > R(om) > R(rd) (Wen et al. 2013). R(ws) and R(ie) are quite mobile and bioavailable to biota (Lu et al. 2013). In natural environments, R(rd) is a major form, which is stable and cannot be absorbed directly by organisms (Pang et al. 2002), but REEs exist in more active states in mining areas, such as R(ie) in soils and dissolved R(ws) in water bodies (Liang et al. 2014). Mining activities disrupt the spatial distribution and natural state of minerals, including notably turning deposits into smaller unconsolidated particles with a relatively higher specific surface area. More REE minerals are transported from subsurface to surfaces and these surficial mining areas become enriched in total REEs. Those REE minerals significantly increase the source supply of REEs in the earth's surface by exposing them to much higher rates of weathering and erosion (e.g., rain, wind, microbial decomposition) than seen in more consolidated subsurface locations. A conceptual model based on recently published literature (mainly adapted from Gwenzi et al. 2018) is presented in Figure A1.2. This model shows that hydrological and wind-driven (e.g., weathering, dust and atmospheric fallout) processes, as well as human activities (e.g., REE mining and international trade) play a key role in environmental mobilization and transport of REEs. At last, additional REEs by mining mostly settle into soil and water, leading to increased exposure for terrestrial and aquatic food webs.

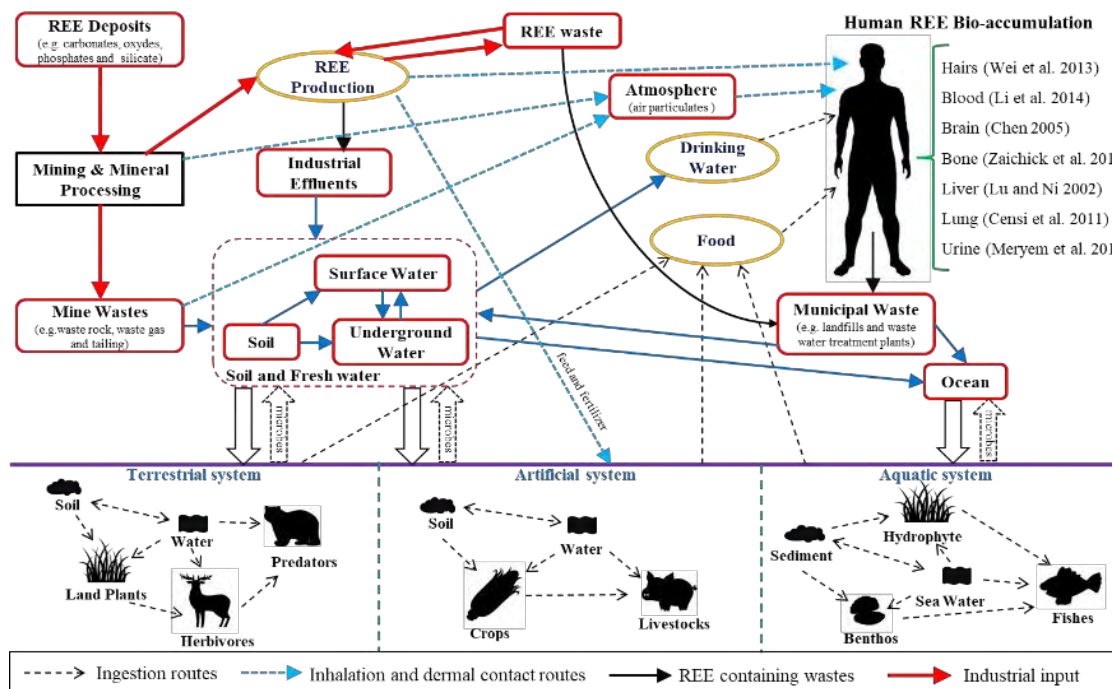


Figure A1.2 The conceptual model of the sources of REE inputs and their fate in the environment. Terrestrial system (MacMillan et al. 2017), Artificial System (Jiang et al. 2012), Marine system (Li et al., 2016).

REEs in soil can enter plant cells through soil-plant pathways. For most plants, REE concentrations generally follow the order of root > leaf > stem > flower > fruit (He et al. 2019a; Hu et al. 2004; Xu et al. 2002), but close to REE mining or other polluted areas, the REE content is higher in leaf surfaces because of dust fall (Zhuang et al. 2017). Water is the main exposure intermediary for aquatic organisms. Increased REEs by mining can enter adjacent aquatic systems through industrial and domestic wastewater as well as surface run-off, and then deposit in sediments or remain distributed in the water column depending on particle size and makeup and mixing regimes (Chua, 1998; Liang et al. 2014). For example, increased usage in Gd-based contrast agents (Gd-CA) for magnetic resonance imaging contributed to a more than one order of magnitude increase in anthropogenic Gd concentrations in San Francisco

Bay (from 8.27 to 112 pmol kg⁻¹) over the two decades from 1993 to 2013 (Hatje et al. 2016).

Ingestion, inhalation and dermal contact are the three main pathways for the entry of REEs in humans (Figure A1.2). Ingestion is mainly associated with REE-contaminated food and drinking water and medicine containing REEs. Gd-CA are a prominent cause of REE ingestion, and some gadolinium accumulation in brain and liver tissues have been reported (Kanda et al. 2015; Frenzel et al. 2017). On the other hand, most of Gd-CA are excreted and released into waste water systems and then are discharged into surface and groundwater, potentially subsequently reaching back to drinking water supplies (Aime and Peter, 2009; Bichler et al. 2016; Schmidt et al. 2019). Inhalation of airborne dust containing REEs is another pathway by which they can enter the human body. High concentrations of REEs (56.9 to 165.4 µg L⁻¹) were found in lung fluids of people working in REE associated industries (Censi et al. 2011). Further, REEs can enter the human body through dermal absorption (Sun et al. 2017), for example, through bathing using water containing high REEs and exposure to REE dust. Current research shows that REEs can accumulate in various human tissues: hair (Wei et al. 2013), blood (Li et al. 2013), brain (Chen et al. 2005), bone (Zaichick et al. 2011), liver (Chen & Zhu, 2009) and lungs (Censi et al. 2011). Risks are not explicitly well understood though, and it is likely that there could be subtle positive and negative health effects due to these emerging contaminants, as well as the largely negative oxidative stress effects that have been reported (Pagano et al. 2015).

A1.7 A review of REE toxicity on organisms

A1.7.1 Method

A literature review of published, peer-reviewed studies on the REEs affecting the environment available in Scopus (<https://www.elsevier.com/solutions/scopus>) was

done in two steps, first in November 2019 and in January 2021 to update the review. Searches were conducted using the combined following keywords: “terbium” OR “dysprosium” OR “holmium” OR “erbium” OR “thulium” OR “ytterbium” OR “lutetium” OR “yttrium” OR “lanthanum” OR “cerium” OR “praseodymium” OR “neodymium” OR “promethium” OR “samarium” OR “europium” OR “gadolinium” OR “scandium” OR “lanthanide” OR “rare earth element*” AND “health risk*” OR “health impact*” OR “toxicity*” OR “animals*” OR “plant*” OR “human*” OR “microbiome*” OR “environment*” OR “bacteria*” OR “fungus*” Also, Google Scholar (<http://scholar.google.com>) was searched to ensure as many publications as possible were collected, and 27,580 references were acquired from these searches. In a second screening stage eligibility and exclusion criterion were determined. First with regards to literature type, only published peer-reviewed journal articles of primary research and reviews were selected. Second, in order to avoid any confusion and difficulties of translation, only on articles published in English were kept. Thirdly, with regards to timeframe, articles published up to 2020 were selected to see the potential growth/evolution of research and related publications. As the review process focused on REE toxicity to organisms, the subject areas were restricted in “biochemistry, genetics and molecular biology”, “environmental science”, “agricultural and biological sciences” and “immunology and microbiology”. Titles, keywords and abstracts were reviewed and screened following the above criteria. Finally, 1,228 articles remained and were used for the analysis of publication volume and subsequently classified into five topics of humans, plants, animals, microbiome and “others” (review articles and those about abiotic effects/environments).

A1.7.2 State-of-the-art in REE-associated environmental risks/effects

The number of published studies on REE environmental risks and impacts shows an increasing trend with time (Figure A1.3), with a faster pace since 2011, likely because of the higher number of REE projects recently undertaken outside of China.

The attention received by different target species or biotic groups in these studies is clearly uneven. The number of annual published articles generally followed the order of plants > animals > humans > microorganisms. Initially most articles were about plants and animals, then the effect of REEs on humans and microorganisms became the center of focus in an increasing number of publications. However, research on humans and microbes is still less abundant than for animals and plants up to 2020.

Review articles about environmental issues associated with REE production were focused on ecological risks (mainly animal and plant health), human health, and toxicity mechanisms of REEs. However, little information has been synthesized on environmental and individual factors that influence REE toxicity (Figure A1S1).

A1.7.3 The toxic effects of REEs on different target biota

Multifaceted problems caused by REEs have been reported by a body of literature (shown in Figure A1.3) and publications about REE-associated health effects mainly include acute toxicity and bio-accumulation studies. Globally, these studies have shown that when the content of REEs in organisms exceeds the background concentration or toxic threshold, there is a potential risk to the health of ecosystems and purportedly to humans.

REEs hormetic dose-response is the core of the toxicity research on various life forms (reviewed by Agathokleous et al. 2018; Agathokleous et al. 2019; Pagano et al. 2015a) and is characterized by biphasic concentration-related effects: low doses of REEs are beneficial for organisms, but inhibitory or toxic effects are present at high levels (Agathokleous et al. 2018; Pagano et al. 2015). For example, La^{3+} at doses of 0 - 25 μM increased shoot biomass of maize plants associated with increased photosynthetic rates and chlorophyll index values, but at a higher level (above 25 μM), photosynthesis was inhibited (Carolina et al. 2018). The mechanism of hormetic

effects of REEs is still not clear, but the chronic accumulation of these elements in the environment could exceed the toxicity threshold of REEs.

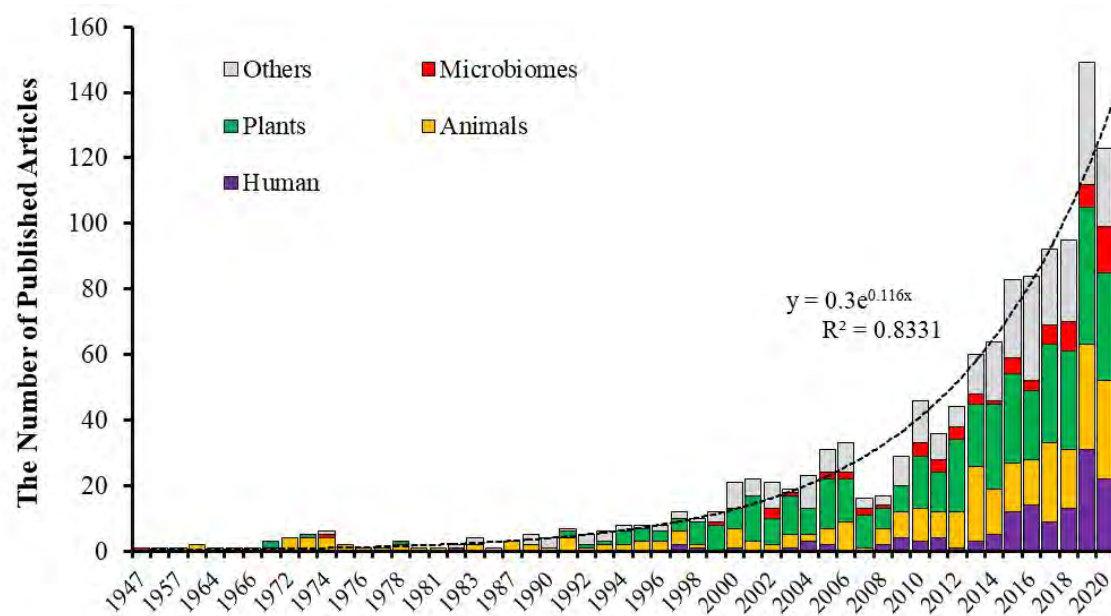


Figure A1.3 Annual rate of publication of journal articles on REE-associated environmental risks/effects for different groups of biotas.

LC50 (lethal concentration 50%) and EC50 (median effective concentration) are statistically derived doses used to evaluate toxicity threshold in the biotic ligand model (BLM) (Smith et al. 2015). Lower LD50 (Lethal Dose 50%)/LC₅₀ value indicates higher acute toxicity. The toxicity threshold (LC50 and EC50) of REEs on aquatic organisms has been reviewed by Malhotra et al. (2020) and toxicity-modifying factors such as pH, dissolved organic carbon (DOC), temperature and potential competitors (e.g., Ca²⁺) have a main impact on the bioavailability and toxicity of REEs to aquatic organisms. However, there is a significant gap in knowledge about LC50 and EC50 of REEs in some special exposure conditions (e.g., Canadian boreal and arctic environment) and in non-aquatic organisms. On the other hand, the lowest observable effect concentrations (LOECs) are used in chronic or

long-term ecotoxicological studies, which is more helpful to better understand the actual ecotoxicological risk induced by REEs on the environment (Cardon et al. 2019). The traditional toxicity model using dissolved concentrations as the dose descriptor without considering its bioavailability, therefore failed to accurately predict the toxicity of REEs (Zhao et al. 2015; Gong et al. 2019b). REEs in the 3+ oxidation state are considered to be the dominant toxic chemical species and its concentration is a good indicator of toxicity (Aharchaou et al. 2020), therefore the LC50 and EC50 based on free ion activities should be widely used instead of dissolved concentrations. For example, Gong et al. (2019b) determined the EC50 of Ce and Y on the root elongation of wheat (*Triticum aestivum*) and found the values of EC50 of dissolved concentration were far higher than those of free-ion concentration: $EC_{50}[Y_{diss}] = 6.08 \mu M$ and $EC_{50}[Ce_{diss}] = 7.11 \mu M$; $EC_{50}[Y_{ion}] = 2.05 \mu M$ and $EC_{50}[Ce_{ion}] = 2.28 \mu M$. Furthermore, the analytical techniques for chemical characterization of REEs are also a key factor in determining data quality (Filella and Rodushkin, 2018) and these techniques have been reviewed by Gorbatenko and Revina (2015). Inductively coupled plasma mass spectrometry (ICP-MS) and instrumental neutron activation analysis (INAA) are commonly used for REE determination in environmental samples because of their multi-element capability, ease of operation, high sensitivity and accuracy.

The significant toxicity of high concentrations of REEs to plants, animals and microorganisms has been reviewed by Abdelnour et al. (2019) and Adeel et al. (2019), and effects have been identified on cell division (Kotelnikova et al. 2019), gene expression (Dubé et al. 2019, Qi et al. 2019), blood variables (Adua et al. 2013), photosynthesis (Xu et al. 2019, Zicari et al. 2018), metabolic capacity (Henriques et al. 2019, Pinto et al. 2019), reproductive capacity (Zhang et al. 2014), oxidative stress (Oral et al. 2017; Henriques et al. 2019) and various growth indices (Xu et al. 2019, recent studies listed in Table A1S3).

Current literature has found some human health problems caused by REEs (reviewed by Gwenzi et al. 2018, Adeel et al. 2019). Excess REEs have adverse impacts on organ function, as illustrated in Figure A1.4. In addition, interactive effects between REEs and other heavy metals have been identified. For example, Y was shown to increase the toxicity of chromium on oxidative damage to DNA in the blood of workers in a chromate plant (Bai et al. 2019). REEs and other heavy metals are usually concomitantly present in the environment, therefore, their interactive effects should be considered in environmental risk assessments.

The existing data to date on REE toxicity are mainly based on controlled experiments or laboratory tests, and although useful, typically only show acute effects in isolation (i.e., without multiple stressors). REE toxicity cases from operating REEs mining projects are summarized based on Scopus and the China National Knowledge Infrastructure (CNKI) database (see Table A1A2), which may facilitate better identification of potential environmental and health risks for Canada. We found that nearly all cases are from China and concentrated in Inner Mongolia where the largest LREE mines are located and Jiangxi with the largest HREE mines on the globe. Most of these studies focused on soil, water, human health and food, but data about wildlife is limited.

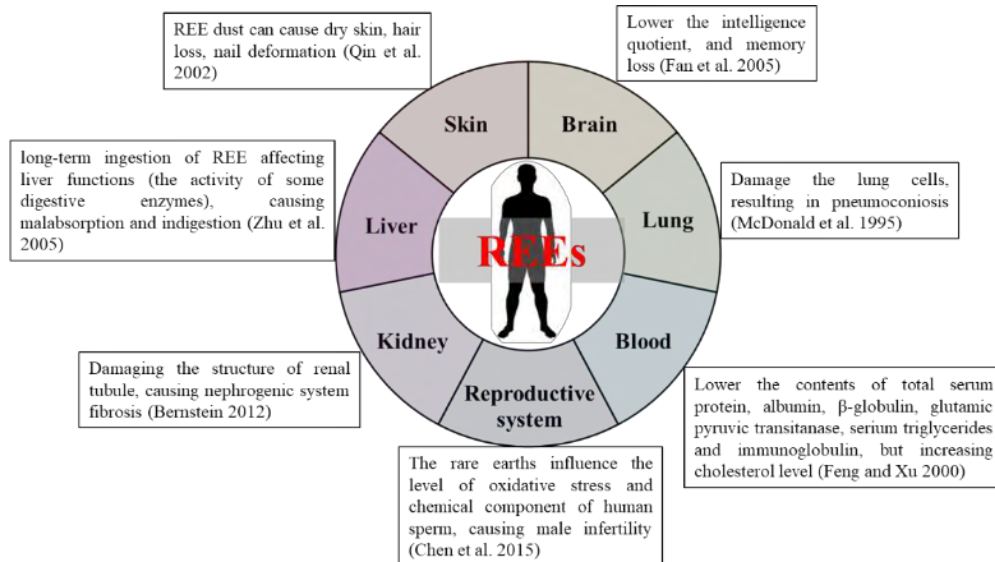


Figure A1.4 Effects of REEs on human organs/systems.

A1.7.4 The mechanisms and influential factors of REEs toxicity

Based on the collected published literature, the mechanisms of REEs toxicity and main environmental and individual factors influencing toxicity are summarized in Table A1.1 and A1.2, respectively. Except general metal-induced oxidative stress, the specific molecular mechanisms of REEs toxicity include: (1) acting as a substitute or antagonist for Ca^{2+} , (2) replacement of Mg^{2+} , (3) reaction with essential inorganic salt substances in the body, (4) altering endogenous hormonal balance, (5) effecting on expression level of genes. These mechanisms are described and summarized in Table A1.1. Compared with other heavy metals, the molecular mechanisms of REE toxicity to various life forms are still not well-defined and further work is required in this field. As discussed above, trivalent cationic REEs are the dominant toxic chemical species, and therefore drivers affecting the transfer of REE fractions to this more soluble/free form (e.g., pH) are also factors for REE toxicity by increasing bioavailability. The factors influencing REE toxicity can be divided into two

categories based on collected references: (1) external environmental factors including interaction between REEs, exposure time, dose, potential competitors (e.g., Ca^{2+}) and pH, as well as (2) individual characteristics of the receptors including species, growth stage, and sex (see Table A1.2).

A1.8 Potential environmental risks associated with REE development in Canada

A1.8.1 The state of REE development in Canada

Currently, there are no REE mines in operation in Canada (Natural Resources Canada, 2018) although in the past some REEs were generated as a by-product during uranium and niobium production. The exploration activity in Canada is, however, quite high, with over 50 projects in development (Orris et al. 2018), accounting for up to 7 percent of REE projects worldwide (Figure A1.5). The REE deposits in Canada are dominated by REE-bearing carbonates, silicate phases and phosphates (Kalvig and Machacek 2018). Some advanced REE exploration projects are ongoing in Canada, and some of them have high concentrations of HREEs that are globally less abundant than LREEs, and particularly needed for many renewable energy technologies (Karen 2015). For example, the deposits of Strange Lake and the Nechalacho project have a high HREE proportion (HREEs/REEs) at 38%. It is anticipated that Canada would be a global leader in the production of REEs (especially for HREEs) by 2030 (Kalvig and Machacek 2018).

However, the pollution prevention and investment cost of REE production may be greater in Canada compared to China because of characteristics of the ore. This is because REE mineral phases for HREE production are different, with solid state minerals in Canada and ion-adsorption clay minerals in China. Ion-adsorption type REE minerals, which supplies most of HREEs in the world are found only in China. REEs in a trivalent cationic state are adsorbed onto clay minerals, and they are extracted by leaching with an aqueous electrolyte solution (sodium chloride or

ammonium sulfate) via an ion-exchange process (Yang et al. 2013). The REE minerals in Canada are in solid state mineral phases, such as carbonates, and usually are associated with the radioactive elements uranium and thorium. The technological processes of extracting solid state minerals of REEs will be more complicated than ion-adsorption clay minerals. Therefore, limiting environmental risks and cost constitutes important challenges for future development of Canadian REE industry.

Table A1.1 Summary of the main molecular mechanisms of REE toxicity.

Mechanisms	Description	Reference
Substitute or antagonist for Ca^{2+}	REEs and Ca share similar chemical and physiological functions because of similar ionic radius. REEs can interfere with the normal physiological functions of Ca by replacing Ca in enzymatic reactions. For example, as a Ca^{2+} Channel antagonist, REEs can interact with membrane associated enzymes, and prevent the influx of other mineral elements, thereby disrupting elemental homeostasis.	Jakubek et al. (2009) Wang et al. (2008) Zeng et al., (2003)
Replacement of Mg^{2+}	Lanthanides could replace the magnesium atom of chlorophyll a and form a new REE-chlorophyll a, therefore influencing the photosynthesis of plants.	Chen et al. (2000) Wang et al. (2014b)
Reaction with essential inorganic salts in the body	Exogenous REEs can react with substances in the body, which can lead to physiological toxicity and even diseases. For example, inhalation of airborne particles including REEs can lead to pulmonary fibrosis because the reaction between the phosphate in lung fluid and REEs produces phosphate precipitation in lungs. Besides, REEs can lead to deposition by combination with negative ions (e.g. with phosphatic ion) in serum.	Censi et al. (2011) Wu et al. (2006)
Alter endogenous hormonal balance	REEs can alter endogenous hormone levels, which subsequently promote or inhibit the growth and physiological function. Generally, REEs at low concentration increase the level of Gibberellin A3 (GA3), trans-Zeatin-riboside (ZR) and indole-3-acetic acid (IAA) but decrease their level at high REE concentration because of reduced auxin carrier abundances.	Wang et al. (2014c) Liu et al. (2017) Cui et al. (2019) Luo et al. (2008)
Effecting expression level of genes	REEs can regulate the physiological function of organisms by affecting functional gene expression. For example, 1) La^{3+} can enhance endocytosis in root cells by regulating the expression level of genes encoding DNA methylases/demethylases; 2) Rare earth elements can activate the expression of silent or poorly expressed genes; 3) La^{3+} can increase the expression of auxin biosynthesis-related genes.	He et al. (2019b) Tanaka et al. (2010) Liu et al. (2016) Dubé et al. (2019)

Table A1.2 Key factors influencing REE toxicity to organisms.

Factors	Test models	Finding	References	
Individual characteristics	Growth stage	Rice	Changes in the photosynthesis indexes of rice under the the effects of acid rain and La ³⁺ were the booting stage > seedling stage > grain filling stage	Zhang et al. (2018)
		Mussel (<i>Mytilus galloprovincialis</i>)	La and Y are more toxic (mortality) to developing embryos and larvae than to juveniles	Mestre et al. (2019)
	Sex	Human	Total serum protein tended to decrease for male compared with female in a REE mining area in China	Zhang et al. (2000)
		Rat	REEs at high dose lead to renal and liver damage in rats. Female rats present a higher REE tolerance than males after 90-day exposure based on various metabolic profiles in the serum and urine.	Wu et al. (2006)
	Species	Rat and human	Kidney cell lines of rats were more sensitive than human in the strength of cell viability reduction, exposed to four lanthanides (La, Ce, Eu, and Yb).	Heller, et al. (2019)
Environmental factors	REEs interaction	Wheat (<i>Triticum aestivum</i>)	Additive toxicity was found for Y-La and La-Ce mixtures, whereas antagonistic interaction was observed for Y-Ce mixtures.	Gong et al. (2019)
	Potential competitors	Wheat (<i>Triticum aestivum</i>)	Ca ²⁺ and Mg ²⁺ increased the values of EC50 (Y ³⁺) by 19 and 2.24 folds respectively, but no significant increase with Na ⁺ , K ⁺ and H ⁺ .	Gong et al. (2019)
	Exposure time	Rare minnow (<i>Gobiocypris rarus</i>)	The mortality increased with exposure time from 24 to 96h for the same concentration of LaCl ₃ ·7H ₂ O.	Hua et al. (2017)
			The growth (e.g., levels of chlorophyll and carotenoid) of plants increased at lower Ce concentrations and decreased at higher Ce concentrations	Zicari et al. (2018)
	Dose	<i>Lemna minor</i>	Low-dose La ³⁺ (0.2, 2.0, and 4.0 μM) exposure increased the total content of microcystins produced by <i>Microcystis aeruginosa</i> (algae), while the production of microcystins was inhibited at high-dose La ³⁺ (8.0, 20, 40, and 60 μM) exposure.	Shen et al. (2018)
	pH	Rice	At 81.6 μM LaCl ₃ , the photosynthesis was increased at pH 4.5, but toxic effects were observed at pH 3.5.	Wang et al. (2014b)

A1.8.2 The REE background levels in Canada

The natural REE background levels before the mass production of REEs are key reference points for environmental and health risk assessment of both current and future REE projects in Canada. In this section, many studies that have made contributions to understanding Canadian baseline data of REEs in natural resources and organisms are discussed (see Table A1.3). Data from two important reviews (Adeel et al. 2019; Bayon et al. 2015) were integrated with publications about Canadian REE background concentrations in soil/sediments found during the literature search (Casse et al. 2019; Cheung et al. 2009; Chiarenzelli et al. 2001; Johannesson and Hendry, 2000; MacMillan et al. 2018; Rasmussen et al. 2017; Romero-Freire et al.). The Canadian background concentration of each REE in water and soils/sediments was not above the levels found in most of the regions in the world (Figure A1A1 and A1A2). For example, in Canada, the average concentrations of total REE (total REEs =88.8 mg/kg) in house dust sampled from 1025 urban homes (Rasmussen et al. 2017) were below the average level of REEs (137.1 mg/kg) found in global soil (Kabata-Pendias, 2011). Therefore, the health risk of current REEs in the environment is low.

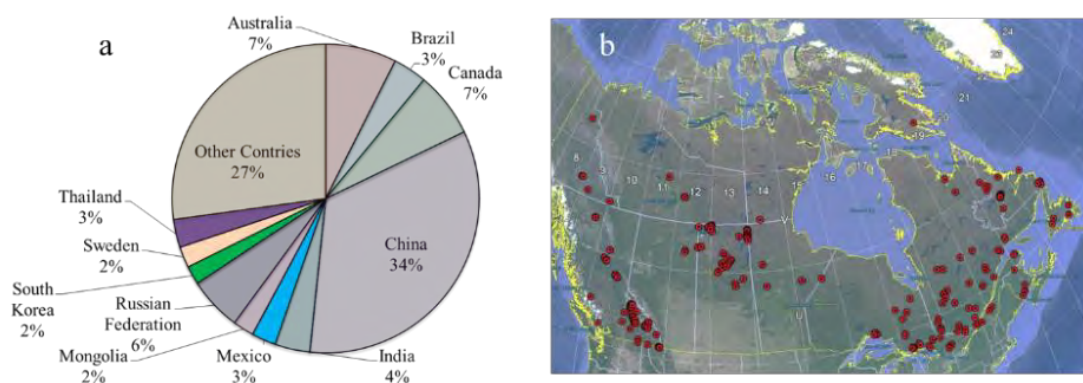


Figure A1.5 The global REE projects (a) and the REE deposits in Canada (b) (data from USGS 2019 edited with Google Earth).

The data of REE levels in biota in Canada was reviewed, and the selected recent studies are shown in Table A1.3. REE concentrations were significantly different among biotic groups, and higher REE levels were reported in mosses and lichens than other groups ($p < 0.05$, Figure A1A3). Also, it appears that aquatic organisms can absorb and accumulate more REEs than terrestrial organisms (Table A1.3 and Figure A1A3). Currently, a generally acknowledged allowable daily intake of total and individual REEs in humans was not available, but Zhu et al. (1997) have proposed a daily allowable intake of $70 \mu\text{g/kg}\cdot\text{bw}$ (4.9 mg/day , considering a 70 kg adult) for total REEs, which was determined from a human health study in REE mining areas and animal experiments. This value has been used in many subsequent studies (Zhuang et al. 2017a; Zhuang et al. 2017b. Yang et al. 2016), so it will be also used in this study to assess associated health risk. Some aquatic animals, plants and most lichens and mosses have a high health risk for human food intake. In Table A1.3, for example, the REE concentration of blue mussel is 4.86 mg/kg and if 1 kg blue mussel is eaten, the REE intake will be more than the allowable daily intake. Furthermore, some traditional food of Indigenous people should be noted, for example a study shows that the REE content of Labrador tea (*Ledumgroen landicum*) varies from 42.3 to 171.6 mg/kg in a REE rich area in northern Saskatchewan (Dunn and Hoffman, 1986), which is a level representing a high risk for human health if consumed excessively. More studies about the ability of REE accumulation in similar important species (e.g., indigenous medicines and food) should be included, and then their health risks can be analyzed and compared.

A1.8.3 The potential environmental risks of REE development in Canada

REE development is a new sector in Canada, and radioactive contamination and REE toxicity are unique potential risks compared with other types of mines. Those potential risks are cryptic and of high risk for public health because there are few proven mitigation strategies appropriate for Canada to reduce or minimize their

adverse impacts. For the identification of associated environmental risks in Canada in this review, risk is deemed to be “the probability of occurrence of a harmful event” (Scheer et al. 2014) and depends on both the level of exposure to the hazard and the vulnerability of the receptors (York et al. 2012). In this synthesis, “harmful events” are environmental issues caused by REE mining and processing such as elevated concentrations of REEs or other hazardous substances in the environment, and the receptors are biota living in Canada, therefore, potential environmental risks associated with REE development can be identified in different communities and ecosystems by analyzing the exposure routes, their influence factors, and vulnerability of various life forms in Canada. Then several typical REE projects are discussed as different scenarios.

As there are no Canadian federal water quality standards or guidelines for Rare Earth Elements, although a draft federal water quality guideline for Ce, La, Nd and Y will be presented soon (Dixit, 2020), the data from other countries is used to compare with available Canadian information to evaluate the risks of exposure to REEs via water. Illegal production in China represents more than 50 % of global REE production (Langkau and Erdmann 2020), which is much more likely to damage the local environment and people’s health (see Table A1.4) as there are no strict safety rules, regulated extraction procedures and accepted environmental strategies at these sites. However, illegal production is unlikely in Canada. Therefore, we follow the basic assumption that associated laws, regulations, rules and strategies will be strictly enforced at each REE project in Canada.

Table A1.3 REE background content (mg/kg) of living organisms in Canada.

location	Species	Group	Y	La	Ce	Pr	Nd	Sm	Eu	Gd	Tb	Dy	Ho	Er	Tm	Yb	Lu	References
1 Kuujuaarpiik-Whapmagooost ut, Quebec	Benthos	Aquatic animals /plants		1.3821	1.664	0.22	0.71	0.0908	0.0239	0.0995	0.0094	0.0357	0.0066	0.0209	0.0022	0.0147	0.0021	MaceMillan <i>et al.</i> , 2017
2 Kuujuaarpiik-Whapmagooost ut, Quebec	Blue Mussel	Aquatic animals		1.394	1.9618	0.24	0.85	0.1262	0.0199	0.1302	0.0122	0.0533	0.0097	0.0299	0.0034	0.0221	0.0035	MaceMillan <i>et al.</i> , 2017
3 Kuujuaarpiik-Whapmagooost ut, Quebec	Brook Trout_1	Aquatic animals		0.0001	0.0001	0	0	0.0002	0	0	0	0	0	0	0	0	0	MaceMillan <i>et al.</i> , 2017
4 Kuujuaarpiik-Whapmagooost ut, Quebec	Brook Trout_2	Aquatic animals		0.0008	0.0014	0	0	0.0002	0	0.0002	0	0.0002	0	0	0	0	0	MaceMillan <i>et al.</i> , 2017
5 Near Montreal, Quebec	Brook Trout_3	Aquatic animals	0.045	0.2074	0.1872	0.03	0.11	0.0146	0.0029	0.0207	0.0021	0.0101	0.0021	0.0057	0.0006	0.0028	0.0004	MaceMillan <i>et al.</i> , 2018
6 Near Montreal, Quebec	Mussel	Aquatic animals	0.052	0.0666	0.0697	0.01	0.05	0.0102	0.0024	0.013	0.0016	0.0075	0.0015	0.004	0.0004	0.0024	0.0004	MaceMillan <i>et al.</i> , 2018
7 Kuujuaarpiik-Whapmagooost ut, Quebec	Ringed Seal	Aquatic animals		0.0003	0.0004	0	0	0.0002	0	0.0002	0	0	0	0	0	0	0	MaceMillan <i>et al.</i> , 2017
8 Kuujuaarpiik-Whapmagooost ut, Quebec	Sea Urchin	Aquatic animals		0.6974	0.6052	0.1	0.34	0.0497	0.0081	0.0554	0.0056	0.0267	0.0053	0.0164	0.002	0.0128	0.0019	MaceMillan <i>et al.</i> , 2017
9 Kuujuaarpiik-Whapmagooost ut, Quebec	Whitefish	Aquatic animals		0.0001	0.0001	0	0	0.0002	0	0.0002	0	0	0	0	0	0	0	MaceMillan <i>et al.</i> , 2017
10 Nomans River, Quebec	Typha latifolia	Aquatic plants		0.2294	0.477	0.05	0.2	0.0299	0.016	0.0256	0.0033	0.0141	0.0025	0.006	0.0008	0.0047	0.0006	Romero-Freire <i>et al.</i> , 2019
11 Kuujuaarpiik-Whapmagooost ut, Quebec	Zooplankton_1	Aquatic plants		3.9405	4.87	0.75	2.59	0.3285	0.0643	0.3306	0.0334	0.146	0.0271	0.0867	0.0101	0.065	0.0095	MaceMillan <i>et al.</i> , 2017
12 Near Montreal, Quebec	Zooplankton_2	Aquatic plants	0.365	0.3775	0.7903	0.09	0.4	0.0854	0.0268	0.0981	0.0122	0.0667	0.0124	0.0366	0.0046	0.0304	0.0044	MaceMillan <i>et al.</i> , 2018
13 Deslauriers Island, Quebec	Chironomus riparius	Bird		0.0276	0.15	0.16	0.02	0.0728	0.0095	0	0.0087	0	0.0041	0.0009	0.0024	0.0016	0	Brown <i>et al.</i> , 2019

Table A1.3 continued

		Chirotopocp halus ridibundus_ male	Bird	0.0053	0.19	0.21	0.02	0.0843	0.0094	0	0.007	0	0.001	0	0	0	0	Brown <i>et al.</i> 2019
14	Deslauriers Island, Quebec		Bird															
15	Kuujuarapik- Whapmagooost ui, Quebec	Parnigian	Bird	0.0003	0.0003	0	0	0.0002	0	0.0002	0	0	0	0	0	0	0	MackMill an <i>et al.</i> 2017
16	Other Lake area, Nunavut	Alectoria pubescens	Lichen	0.73	1.81	5.89	0.42	1.7	0.314	0.068	0.231	0.031	0.176	0.082	0.075	0.015	0	Charenz elli <i>et al.</i> 2001
17	Other Lake area, Nunavut	Calciophlea saxicola	Lichen	0.38	0.88	1.71	0.21	0.77	0.155	0.027	0.114	0.017	0.099	0.041	0.038	0.004	0	Charenz elli <i>et al.</i> 2001
18	Other Lake area, Nunavut	Cetaria cucullata	Lichen	0.24	0.56	1.15	0.13	0.5	0.09	0.018	0.076	0.01	0.052	0.027	0.024	0.004	0	Charenz elli <i>et al.</i> 2001
19	Other Lake area, Nunavut	Cetaria nivalis	Lichen	0.05	0.1	0.2	0.02	0.09	0.02		0.011		0.011	0.004	0.005	0.002	0	Charenz elli <i>et al.</i> 2001
20	Other Lake area, Nunavut	Cladonia rangiferina	Lichen	2	3.55	11.84	0.8	3.45	0.644	0.131	0.532	0.076	0.431	0.199	0.195	0.023	0	Charenz elli <i>et al.</i> 2001
21	Other Lake area, Nunavut	Cladonia stelaris	Lichen	0.16	0.33	0.66	0.07	0.29	0.054	0.012	0.044	0.005	0.032	0.018	0.019	0.004	0	Charenz elli <i>et al.</i> 2001
22	Kuujuarapik- Whapmagooost ui, Quebec	Lichen spp.	Lichen	1.2467	2.4703	0.26	0.95	0.1406	0.029	0.1427	0.0142	0.0593	0.0109	0.0327	0.0247	0.0033	0	Charenz elli <i>et al.</i> 2001
23	Other Lake area, Nunavut	Masonhalea richardsonii	Lichen	0.6	1.46	3.24	0.35	1.3	0.212	0.052	0.17	0.022	0.117	0.068	0.056	0.009	0	Charenz elli <i>et al.</i> 2001
24	Other Lake area, Nunavut	Nephroma arcticum	Lichen	0.74	2.11	4.27	0.49	1.82	0.334	0.074	0.23	0.033	0.173	0.088	0.092	0.011	0	Charenz elli <i>et al.</i> 2001
25	Other Lake area, Nunavut	Parmelia fuliginosa	Lichen	2.21	4.92	16.73	1.06	4.44	0.89	0.24	0.58	0.115	0.506	0.275	0.25	0.037	0	Charenz elli <i>et al.</i> 2001
26	Other Lake area, Nunavut	Parmelia mouguetii	Lichen	1.55	4.38	10.17	0.79	3.23	0.582	0.14	0.464	0.067	0.366	0.194	0.19	0.026	0	Charenz elli <i>et al.</i> 2001
27	Other Lake area, Nunavut	Stereocaulo n paschale	Lichen	0.16	0.53	1.17	0.13	0.49	0.069	0.015	0.052	0.007	0.034	0.017	0.017	0.004	0	Charenz elli <i>et al.</i> 2001
28	Other Lake area, Nunavut	Umbilicaria polyphylla	Lichen	0.95	2.15	6.82	0.51	2.12	0.382	0.096	0.319	0.043	0.22	0.103	0.088	0.015	0	Charenz elli <i>et al.</i> 2001
29	Other Lake area, Nunavut	Brachytheci um velutinum	Mosse	1.56	3.54	8.1	0.67	2.72	0.527	0.141	0.44	0.067	0.355	0.201	0.205	0.043	0	Charenz elli <i>et al.</i> 2001
30	Other Lake area, Nunavut	Dicranum seoparum	Mosse	0.72	2.13	7.38	0.46	1.85	0.337	0.073	0.225	0.031	0.168	0.09	0.083	0.016	0	Charenz elli <i>et al.</i> 2001
31	Other Lake area, Nunavut	Hydrocomia splendens	Mosse	0.13	0.23	0.43	0.05	0.18	0.035	0.006	0.034	0.006	0.023	0.014	0.016	0.003	0	Charenz elli <i>et al.</i> 2001

Table A1.3 continued

32	Oter Lake area, Nunavut	<i>Polypodium juniperina</i>	Mosses	0.7	2.02	5.71	0.4	1.6	0.253	0.088	0.189	0.04	0.141	0.076	0.075	0.008	Chiarenzelli <i>et al.</i> 2001
33	Oter Lake area, Nunavut	<i>Grimmia apocarpa</i>	Mosses	3.38	4.44	15.55	1.01	4.24	0.872	0.23	0.817	0.129	0.778	0.443	0.413	0.061	Chiarenzelli <i>et al.</i> 2001
34	Kuujuaq-Wapmagooi, Quebec	Caribou	Terrestrial animal		0.0006	0.0004	0	0	0.0002	0	0.0002	0	0	0	0	0	MacMillan <i>et al.</i> 2017
35	Kuujuaq-Wapmagooi, Quebec	Eider	Terrestrial animal		0.001	0.0018	0	0	0.0005	0.0002	0.0005	0	0.0002	0	0.0002	0	MacMillan <i>et al.</i> 2017
36	Kuujuaq-Wapmagooi, Quebec	Hare	Terrestrial animal		0.0006	0.0003	0	0	0.0002	0	0.0002	0	0	0	0	0	MacMillan <i>et al.</i> 2017
37	Oter Lake area, Nunavut	<i>Betula glandulosa</i>	Terrestrial plants	0.03	0.05	0.07	0.01	0.03	0.007	0	0.006	0	0.005	0	0.005	0	Chiarenzelli <i>et al.</i> 2001
38	Oter Lake area, Nunavut	<i>Empetrum nigrum</i>	Terrestrial plants	0.05	0.08	0.16	0.02	0.07	0.014	0.004	0.009	0	0.01	0.007	0.007	0.001	Chiarenzelli <i>et al.</i> 2001
39	Nomans River, Quebec	<i>Equisetum arvense</i>	Terrestrial plants		1.1177	1.6159	0.19	0.68	0.0925	0.0464	0.0816	0.0105	0.0435	0.0077	0.0195	0.0025	Romero-Freire <i>et al.</i> 2019
40	Oter Lake area, Nunavut	<i>Ledum palustre</i>	Terrestrial plants	0.04	0.06	0.12	0.01	0.05	0.011	0	0.009	0	0.007	0.006	0.006	0	Chiarenzelli <i>et al.</i> 2001
41	Kuujuaq-Wapmagooi, Quebec	Plants spp.	Terrestrial plants		0.0317	0.0594	0.01	0.02	0.0035	0.0091	0.0033	0.0003	0.0015	0.0003	0.001	0	MacMillan <i>et al.</i> 2017
42	Oter Lake area, Nunavut	<i>Salix arctica</i>	Terrestrial plants	0.04	0.12	0.19	0.02	0.08	0.021	0	0.009	0	0.012	0.005	0	0	Chiarenzelli <i>et al.</i> 2001
43	Oter Lake area, Nunavut	<i>Setaria ssp.</i>	Terrestrial plants	0.19	0.62	0.91	0.11	0.47	0.074	0.013	0.069	0.008	0.039	0.022	0.015	0.001	Chiarenzelli <i>et al.</i> 2001

Note: ADW, acceptable daily sample weight; HHR, human health risks (Low: ADW > 1.2 kg; Mid: 0.6 kg ≤ ADW ≤ 1.2 kg; High: ADW < 0.6 kg). For animals, the REE content was from muscle samples.

Table A1.4 List of main health important aspects in REE industry with concerned groups.

Exposure	Risks	Groups Involved	Planning & Control
Ingestion	Food sources	Indigenous people near REE mines or processing plants	Avoid collecting food/medicine sources from contaminated areas;
	Traditional medicines	Indigenous people near REE mines or processing plants	Guidelines for recommended daily limits.
	Drinking water	Surrounding residents and indigenous people near REE mines or processing plants	Water quality monitoring
Inhalation and dermal contact	Dust	Surrounding residents and workers in mines and processing plants	Strict preventive measures (e.g., anti-dust respirators and); Air quality monitoring;
	Radiation	Surrounding residents and workers in mines and processing plants	Determining the distance influenced by REE mines based on content of REEs and radioactive elements and moving residents from influenced areas.

A1.8.3.1 Potential risks to human health

1) Workers

Occupational exposure is generally the most significant risk to human health associated with REE production. Those working in the REE mining and processing industry can be easily and chronically exposed to multiple hazardous substances (e.g., REE-containing dust, heavy metals, radioactive materials and process chemicals) by ingestion, inhalation and dermal contact (Figure A1.1 and A1.2), and this exposure then has the potential to damage human organs or tissues after long-term exposure and potential bioaccumulation. Radiation risk is usually considered as the greatest threat for workers' health in REE mining areas. Cases from the Bayan Obo mine show that the risk of death from leukemia for workers increased significantly over ten years (Li et al. 1986). Although Canada, as one of the biggest producer and supplier of uranium in the world (Winde et al. 2017), has experience in the management of

radioactive exposure during production and waste processing, new and increased exposure sources still have the potential to threaten workers' health if not managed properly. On the other hand, both inhalation and dermal contact are the main exposure route of REEs for people working in mining areas, especially for the inhalation of atmospheric particulates containing REEs, and long-term exposure of REE dust may cause rare earth pneumoconiosis and negative neurological effects (Wang et al. 2014a). Furthermore, snowfall in mining areas should be studied and monitored in Canada, because compared with raindrops, snowflakes with larger surface area and slower fall velocity may accumulate more REE dust (Cereceda-Balic et al. 2012), which increases the risk of REEs' dermal contact for workers and nearby residents. Environmental quality monitoring and strict occupational radiation protection should be able to effectively reduce the risk for workers and avoid acute damages.

2) Surrounding residents

REE mining and processing also pose significant potential risks for surrounding residents, primarily through the inhalation of REE dust and ingestion of food and water with higher REEs content, and some cases have been reported in Chinese REE mine areas (see Table A1A2). For example, Baotou City with the Bayan Obo REE mine, has suffered from considerable pollution issues from REE production and processing (Julio and Fu 2014). As most REE mine projects are far away from population centers and there strict mining waste management laws in Canada, health risks for the general public may be limited. However, there are still some potential risks: Food sources near REE mines or processing plants. Hazardous substances (e.g., REEs, heavy metals and radioactive elements) can disperse and leach into surrounding soil and water (for cases see Table A1A2). Vegetables growing in contaminated soil have the potential to increase the rate of REE bioaccumulation inside the human body, especially for root vegetables (such as potato and sugar beet)

because they absorb more REEs from polluted soil than other vegetables, based on data from China (Zhuang et al. 2017). Aquatic products are also a potential health risk as data collected in Canada (Table A1.3) indicate that benthic organisms (e.g., blue mussels) have higher background content of REEs and a large REE bioaccumulation capacity (Zhang et al. 2009; Squadrone et al. 2019). These and other food sources with a higher REE bioaccumulation capacity should be given special consideration during REE production development in Canada.

3) Indigenous People

Indigenous peoples near REE mining areas in Canada may be more affected by environmental issues associated with REE development than other groups, because they may rely more heavily on local biota for their daily lives (e.g., food, medicine and traditional activities on the land) (Sharma 2010; Wesche and Chan 2010). Potential health risks involve lands used for hunting, fishing, trapping, and food gathering being impacted by REE mines. For example, the Kipawa project (a proposed REE mine site) is located on the traditional lands of two Anicinabek First Nations in western Quebec. There are some potential risks for Indigenous peoples: 1) Traditional food from REE rich areas. An increase in REE projects will lead to more REE rich areas in Canada, which is a serious threat to their traditional territories and potentially health. The content of REEs and radioactive elements in traditional foods is expected to increase inside the influenced areas of REE mines 2) Traditional medicines. More than 500 medicinal plant taxa used by Indigenous peoples in Canada and the REE content of them near mines can be increased especially for those hyperaccumulator for REEs. For example, lichens (11 taxa from 4 families) and bryophytes (6 taxa from 2 families) with higher REE background levels (Table A1.3) and capacity of bioaccumulation (Agnan et al. 2014; Urošević et al. 2020) are used as traditional medicines (Upreti et al. 2012), which may increase the health risks for Indigenous people in the context of REE production. Under the basic assumption that

associated laws, regulations, rules and strategies will be strictly enforced, the health risk from drinking water and inhalation may be low for Indigenous peoples. Avoiding collecting food/medicine sources from contaminated areas and guidelines for recommended daily limits of REE content could lower the risks significantly. Very importantly too, regulators should pre-emptively consult with Indigenous stakeholders to avoid siting potentially damaging extraction and processing operations/facilities where they may do harm.

REE projects as they may impact on indigenous communities, have recently received increasing attention. For example, a community-based environmental monitoring program (IMALIRIJIT) was initiated prior to the start of a rare earth elements (REEs) mining project in Nunavik, Quebec (Gérin-Lajoie, et al., 2018). This will provide baseline data and reference points to better understand the diverse impacts of REE projects on socio-ecological systems, as well as to protect the health, well-being, and quality of life indigenous people in other as future REE projects arise near other northern communities

A1.8.3.2 Potential ecological risks in Canada

The geochemical behaviour of REEs in Canada is influenced by the geographical and geological location of the deposits (geological provinces of main REE projects in Table A1A1) The studies from Canada identified pH, dissolved organic carbon (DOC), free REE³⁺ concentration, active ligand concentration and temperature as the main drivers for REE transfer and bioaccumulation in Canadian aquatic ecosystems (MacMillan et al. 2018; Serrano et al. 2000). For example, a laboratory study on material from the Eldor REE project showed that low temperatures and freeze-thaw events may increase the concentration of elements leached from the tailings produced by the beneficiation of REE minerals in sub-arctic areas (Costis et al. 2020). The

main potential ecological risks of Canadian REE projects for ecosystems are discussed below.

1) Wetland and aquatic ecosystems

Wetlands and lakes cover about 13% and 9 %, respectively, of the land area of Canada (Wulder et al. 2018) and support many ecosystem services. In wetlands and open-water aquatic ecosystems, reducing conditions and chemical reduction are a main factor controlling REE behaviour (Davranche et al. 2015; Mihajlovic et al. 2017). Organic matter is a major sink for REEs due to its high binding capacity for cationic metallic contaminants through ligation or electrostatic interactions (Davranche et al. 2015). In sulfide containing ore and waste environments, acid mine drainage solubilizes REEs, however they can be subsequently adsorbed by soil organic matter in wetlands. Furthermore, the suspended particles including REEs from mining sites can settle in surrounding wetlands and forests. Those REEs will be released to the water under reducing conditions such as seasonal floods caused by spring snowmelt (Davranche et al. 2015; Shiller, 2010). Soil and solution components (e.g., colloids and suspended matter), pH, and redox condition are considered as critical environmental parameters controlling the geochemical behaviors of REEs in the environment because they can increase or decrease cation exchange capacity (CEC). However, more research is needed on geochemical behaviours of REEs because little is known about REEs in different environments (e.g., vast boreal and sub-arctic peatlands in Canada).

Low pH values of certain environments can increase the toxicity of REE for plants and animals by increasing the solubility and bioavailability of REEs. For example, at the Bayan Obo REE mine, a study found water pollution from mine tailings reduced local toad (*Strauchbufo raddei*) population density by affecting the mating rate and causing oxidative damage to tissues and DNA damage in tadpole blood cells (Huang et al. 2019). In Canadian boreal areas, the low pH of bogs and acidic lakes has the

potential to make them sinks for REEs following the release of REEs from future mining sites.

2) Boreal forest ecosystems

The Canadian boreal forests comprise about 30% of global boreal forests and play critical roles in the global carbon cycle and in biodiversity conservation (Peng et al. 2011; Venier et al. 2014). Many studies on fossil fuels and mineral mines (e.g., gold, coal and diamond) affecting boreal forest ecosystems have been published (Naeth and Wilkinson, 1997; Amalesh et al. 2018), however, little is known about REE mines. Dust and heavy traffic are probably the main REE risks for boreal forest ecosystems in Canada. In particular the significant number and quantity of radioactive elements (Wang et al. 2016), REEs (Wang and Liang 2014) and other toxic heavy metal elements (e.g., As, Cd and Pb) (Kexin et al. 2015) in atmospheric deposition from REE mining, and their impact may extend over 3-7 km from sources (Chang and Chang 2012; Degtjarenko et al. 2016). These dusts may enter the stomatal openings and impact physiological functions of plants notably because REEs can interfere with the normal physiological functions of Ca^{2+} and Mg^{2+} . Therefore, there is a potential risk to ecological services of forests near REE mines as photosynthesis and nutrient absorption may be disrupted (mechanisms see Table A1.1 and A1.2). The accumulation of these elements in plants also has potential health risks for other biota and humans via trophic transfers in local and regional food webs.

In addition, the large amount of chemicals and materials required for REE mining and processing increases traffic intensity (e.g., 7-10 tons of ammonium salts and over 10 tons of acid are used for the production of 1 ton rare-earth oxides based on the data from China, Huang et al. 2015), which will disturb roadside biota and animal movement by noise, dust, road accidents and fluid leakage, especially for some species at risks particularly sensitive to new linear disturbances (e.g., Caribou) in the boreal forest. For example, mountain woodland caribou avoided high-use roads and

mines with a buffer of 2 km and low-use roads by 1 km based on the data in British Columbia, Canada (Polfus et al. 2011). Different species have different habitats, physiological features and ecological services, and potential risks caused by REE mining for species at risk in Canada require more specialists involved in assessments prior, during, and after development.

3) Sub-arctic/arctic environment

Many REE projects are at an advanced stage of development in northern regions of Canada (e.g., Strange Lake, Montviel, Eldor and Foxtrot projects), especially in sub-arctic and arctic regions. Environmental concerns related to the development of REE projects in the sub-arctic/arctic climate (extreme temperatures, timing and duration of snow and rain, and freeze-thaw period) should have been raised. Recent studies present that climate condition of cold regions can promote the mobilization of REEs by affecting main driving factors (pH, DOC and redox condition) of REE mobility and bioavailability. Firstly, high-latitude sub-arctic and arctic surface/inland waters often contain sizeable reservoirs of DOC because of the activity of microbial communities at low temperatures (Pokrovsky et al. 2018; Ritson et al. 2014). Furthermore, poor fens and bogs common in watersheds in these regions are highly acidic (typically with pore water pH less than 4.5) (Zoltai and Vitt 1995). As a result, the mobility and bioavailability REEs could be promoted with high DOC content and low pH. Secondly, the redox environments in the lower active layer in permafrost affected soils leads to DOC and REEs that cannot be consumed by the biota or depleted by abiotic reactions (Ji et al. 2021). REEs persist in suprapermafrost water and can be transported to the adjacent rivers or lakes by high surface flows, such as spring thaws (Ji et al. 2021). Thirdly, repeated freeze-thaw cycles could release more REEs and other heavy metals from tailings through intensive erosion process (Pokrovsky et al. 2018). For example, the leaching potential of elements present in flotation tailings was assessed at the Eldor project (Costis et al. 2020), and the results

showed that freezing event increased leachate concentrations. As climate changes in high latitudes, the depth of active layer and freeze-thaw cycles are expected to be increased in the sub-arctic and arctic regions of Canada. Therefore, sub-arctic/arctic climate conditions may importantly influence the environmental risks caused by REE mining in Canada.

A1.9 Conclusions and next steps for research in Canada

The concentration of REEs in the environment has increased significantly in recent years, and this trend is expected to continue over the next decades as more REE projects are developed globally. REEs as emerging contaminants and radioactive elements (U and Th) are main ecological and health risks. Canada ranks among the top countries with rich REE deposits and is expected to be a global leader in the production of REEs in the coming decades. Although no REE mines are currently operating in Canada, many projects are at an advanced stage of development. To understand and assess potential environmental risks that Canada could face in the future, this paper examined environmental behaviours and fates of REEs and their toxicity on ecological and human health. Based on this overview, Canadian REE background studies and data from case studies in long standing REE mining areas, this paper integrated environmental conditions and Canadian specific risk factors. The risk assessment presented here suggests that ingestion exposure via traditional foods and medicines for Indigenous people as well as inhalation and dermal exposure to dust and radiation for workers and surrounding residents could be the main health risks. Considering the many advanced REE projects in sub-arctic/arctic regions in Canada, the environmental and climate condition (acidic waters, low temperature and repeated freeze-thaw cycles) promoting REE transportation is expected to influence ecological risks, which should be a matter of concern in REE mines. However, the environmental issues caused by REE development vary depending on deposit ore minerals, processing technology, waste management strategies, policies and

regulations, and vulnerabilities of the local environment. Ion-adsorption rare earth deposits, for example, provide 80% of HREEs in the world, but this type of REEs ores with high HREE content are currently mainly found in China (Yang et al. 2013; Ram et al. 2019), while the HREEs of Canada are mainly present in carbonates or peralkaline ores. Although some data are available for environmental issues caused by HREEs development in China and other countries, their experience may also be only of limited use for the identification of associated potential environmental risks in Canada. Furthermore, unique climate, ecological systems and species at risk could generate new risks in Canada. Therefore, the following points should be addressed in Canada to determine accurate risk assessment in actual REEs project development:

- (1) More REE baseline concentrations in Canada are required for soil, water, vegetation animals, and humans especially for species at risks, which will permit more detailed and accurate risk assessments
- (2) Determining the toxicity of REEs on target species for food sources in Canada and then providing guidelines for recommended daily limits of REE content. For example, REE uptake, bioaccumulation and toxicity threshold of those species that are traditional food and medicines for Indigenous people.
- (3) Mixed REEs, and not necessarily individual elements alone, should get more attention, because REEs usually transfer and accumulate as a group of contaminants. Additionally, multiple stressors, including effects of associated radionuclides and acidity (for the proposed sites with sulfidic substrates) with REEs should be concomitantly examined.
- (4) Due to the geochemical behaviour of REEs in different physical and chemical conditions, more studies should be performed to better understand the environmental behaviour and fate of REEs under Canadian environmental conditions and in our

most keystone ecosystems, such as wetlands, acid-sensitive Precambrian Shield lakes, boreal forests and tundra.

(5) Developing cleaner technologies for exploiting Canadian REE deposit minerals, reducing the materials and energy consumption and waste discharge during REE mining and processing, which is the most effective way to limit potential environmental risks.

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A1.10 Appendix

Table A1A1 Advanced Rare Earth Elements Exploration Projects in Canada.

Project Name	Province	Geological Province	Deposit Type	Size (Million tonnes)	Total REE%	HREE % (HREE/ REEs)
Montviel Project	Quebec	Superior	CAR	250	1.45%	3%
Hoidas Lake Project	Saskatchewan	Churchill	CAR	2.9	2.57%	4%
Two Tom Project,	Labrador	Grenville	CAR	40	1.18%	6%
Clay Howells Project,	Ontario	Superior	ALK	8.5	0.73%	10%
Eldor Project	Quebec	Churchill	CAR	250	1.90%	10%
Echo-Ridge Project	Ontario	Superior	PLA	60	0.16%	15%
Grande Vallee Project	Quebec	Appalachian Orogen	SLE	1,040	0.01%	20%
Foxtrot Project	Labrador	Grenville	ALK	10	1.10%	20%
Strange Lake Project	Quebec	Churchill	ALK & PEG	300	0.93%	38%
Nechalacho Project	North West Territories	Slave	ALK	320	1.70%	38%
Zeus-Kipawa Project	Quebec	Grenville	ALK	16	0.51%	37%

Notes: CAR – Carbonatite; ALK – Alkaline Complex; PEG – Pegmatites; SLE – Shale; ALU/PLA – Alluvial or Placers and modified Placers; G&R - Granites and Rhyolites. (Natural Resources Canada, 2014; Weng et al., 2013; Zhou et al., 2017)

Table A1A2 The summary of the REEs toxicity cases in REEs mining areas.

N O.	Country	Mine/Province	Component affected	Finding	Reference
1	China	Bayan mine / Mongolia	Atmospheric aerosol particles	These values of REEs associated with PM _{2.5} are much higher than those in non-REEs mining areas. Long-term exposure to PM _{2.5} has been consistently linked to cardiovascular and cardiopulmonary diseases.	Li et al. (2018)
2	China	Bayan mine / Mongolia	Obso Children' urine	The content of Y, La, Ce, Pr, Nd, and Sm in the urine of children in the RE mining area were significantly higher than that in the control group.	Liang et al. (2018)
3	China	Bayan mine / Mongolia	Obso Groundwater	The contamination of RE tailings impoundment leakage on the surrounding groundwater. The pollution dispersion plume in groundwater on the south side of the RE tailings was over a distance of about 5.7 km.	Si et al. (2016)
4	China	Bayan mine / Mongolia	Obso Human hair	The REE concentrations in the hair of people from mining are higher than those in control area.. The mean concentrations of REEs were higher in hair of males than that of females from the mining area.	Wei et al. (2013)
5	China	Weishan earth mine / Shandong	rare Vegetables	REE concentration in vegetables of mining area (94.08 µg/kg) was significant higher than that of control area (38.67 µg/kg). REE concentration declined in the order of leaf vegetable > taproot vegetable > allaceous vegetable > gourd vegetable.	Zhuang et al. (2017a)
6	China	-/Fujian	Soil, Water, Human blood and hair	The concentrations of REEs in soil, water and human blood and hair of mining area were higher than those of other regions.	Li et al. (2014)
7	China	Bayan mine / Mongolia	Obso Human Urine	Urinary heavy metals and uranium in the population increased concomitantly with increasing REE level. Sex, age, educational level, alcohol and smoking habit were major factors contributing to inter-individual variation. Males were more exposed to metals than females.	Hao et al. (2015)

Table A1A2 Continued

8	China	Bayan mine / Mongolia	Obo Inner	Soil	Tailing ponds are the main source of REE in the surrounding soil and the concentration of REEs, decreased with the distance from tailing ponds.	Zheng et al. (2016)
9	China	-/ Jiangxi		Human liver functions	Long-term intake of low-dosage REEs could affect liver function of people living in REE mining area, influencing the activity of a variety of enzymes necessary in the digestion, absorption, and metabolism.	Zhu et al. (2005)
10	China	Bayan mine / Mongolia	Obo Inner	Soil	1. The REE concentrations in soil from mining area were significantly higher than the reference. 2. The REE transportation led to the soil REE contamination along the railway, and its distribution was affected by the dominant wind direction.	Guo et al. (2013)
11	China	XinLong earth mine / Jiangxi	rare /	Water	The surface water and shallow groundwater of mining area were characterized by low pH, high total nitrogen, manganese and REEs. Samples investigated showed 88 % water of pond and stream were unsuitable for agricultural use and aquaculture water supply, and 50 % of well and spring water were unsuitable for drinking water	Hao et al. (2016)
12	China	Weishan earth mine / Shandong	rare /	Crops	The REE concentrations of cereals, wheat and maize from mining area were higher than control areas. The health risk assessment demonstrated that the estimated daily intakes of REEs through cereal consumption were considerably lower than the acceptable daily intake (70 µg/kg body weight). The damage to adults can be neglected	Zhuang et al. (2017b)

Table A1A2 Continued

13	China -/ Fujian	Soil, Vegetable, Human hair and blood	1. The farmlands in vicinity of mining site were contaminated with higher REEs level. 2. Accumulation levels of REEs differ significantly among vegetable species. 3. Elevated concentration of REEs in human hair and blood is associated with soil. 4. The distance from mining site determines the REEs exposure level of local residents.	Li et al. (2013a)
14	China -/ Jiangxi	Human brain	REE was difficult to accumulate in the brain stem, but it was susceptible to cerebral cortex, thus causing subclinical damage (sensory conduction was interfered at the cerebral cortex).	Zhu et al. (1997)
15	China Bayan mine / Mongolia Obo Inner	Dust	REEs are the most abundant elements, followed by Mo, Cd, Pb, and Ag in the road dust around the REE mine. Considerable proportions of polluted metals came directly from the mine, and pollution levels increased with decreasing distance and size of particles.	Tian et al. (2018)
16	China -/ Jiangxi	Human blood	Blood biochemical indices (total serum protein, albumin, β -globulin, glutamic pyruvic transaminase, serum triglycerides, and immunoglobulin) of people from REE-high regions were lower than normal region.	Zhang et al. (2000)
17	China -/ Jiangxi	Soil	High concentrations of REEs in tailings (392 mg/kg), stream waters (4460 $\mu\text{g/L}$), sediments (462 mg/kg) and farmland soils (928 mg/kg) were detected in comparison with control sites. REEs contamination in the farmland soils was mainly originated from the stream water contaminated by the leaching from the tailings.	Liu et al. (2019)
18	China -/ Jiangxi	Children's IQ and blood	1. The score of IQ was significantly lower in the exposure group than in the control. 2. The quantity of blood REEs of the children was lower in exposed group compared to control group.	Fan et al. (2004)

Table A1A2 Continued

19	China	-/ Fujian	Soil, Vegetables	1. The concentration of REEs and heavy metals of the farmland soil near REEs mining area were high, especially for Cd, Cu and As, exceeding 10, 1, 0.3 times higher than Chinese National Soil Quality Standard. 2. The concentrations of Gd and As concentration in vegetables were higher than references.	Li et al. (2013b)
20	China	-/ Shandong	Fruits, Chicken eggs	1. The REEs concentration of fruits from the mining area (142.5 µg/kg) were significantly higher than control area (16.1 µg/kg) and pome fruits and berry fruits were easier to accumulate REEs. 2. For chicken eggs, there were no significant differences between mining areas and control area.	Zhuang et al. (2016)
21	China	-/ Shandong	Crops	There were no significant differences between mining areas and non-mining area in the REEs concentration of crops. Rare earth elements ingested via wheat, corn, and soybeans do not endanger human health.	Liu et al. (2017)
22	China	-/ Mongolia	Toad (<i>Strauchbufo raddei</i>)	Water leakage pollution from tailings pond can reduce the tadpole population density by affecting the mating rate, and the main influencing mechanism of pollution on the <i>Strauchbufo raddei</i> postembryonic development is to cause oxidative damage of tissues and DNA damage of blood cells.	Huang et al. (2019)
23	China	-/ Fujian	Soil, Vegetable, Human blood, Water	1. The concentrations of REEs of cultivated soils, well water, vegetables from the mining area exceeded the references. 2. REE concentrations in local resident's blood and hair were nearly 156 and 10 times higher than those of the general population, respectively. 3. The average daily intake of REEs of local inhabitants was estimated to be 12.5 mg/kg.day, exceeding the maximum daily intake to maintain human health (6.0-6.7 mg/kg.day in China).	Li et al. (2013c)

Table A1A2 Continued

24	China	Bayan mine / Mongolia	Obo Inner	Human leukemia	The workers at REEs mine who die from leukemia significantly increased. The chromosomal aberration rate of lymphocytes of miners was higher than that of the control. The chromosomal aberration rate of the miners was higher than that of the maintenance workers, and the distortion rate increased with working time.	Li et al. (1986)
25	China	Bayan mine / Mongolia	Obo Inner	Human cancer	The dust-exposed miners after long-term (more than 30 years) exposure to high air concentrations of thorium containing rare -earth dusts and thoron short -lived progeny could induce excess lung cancer	Chen et al. (2002)

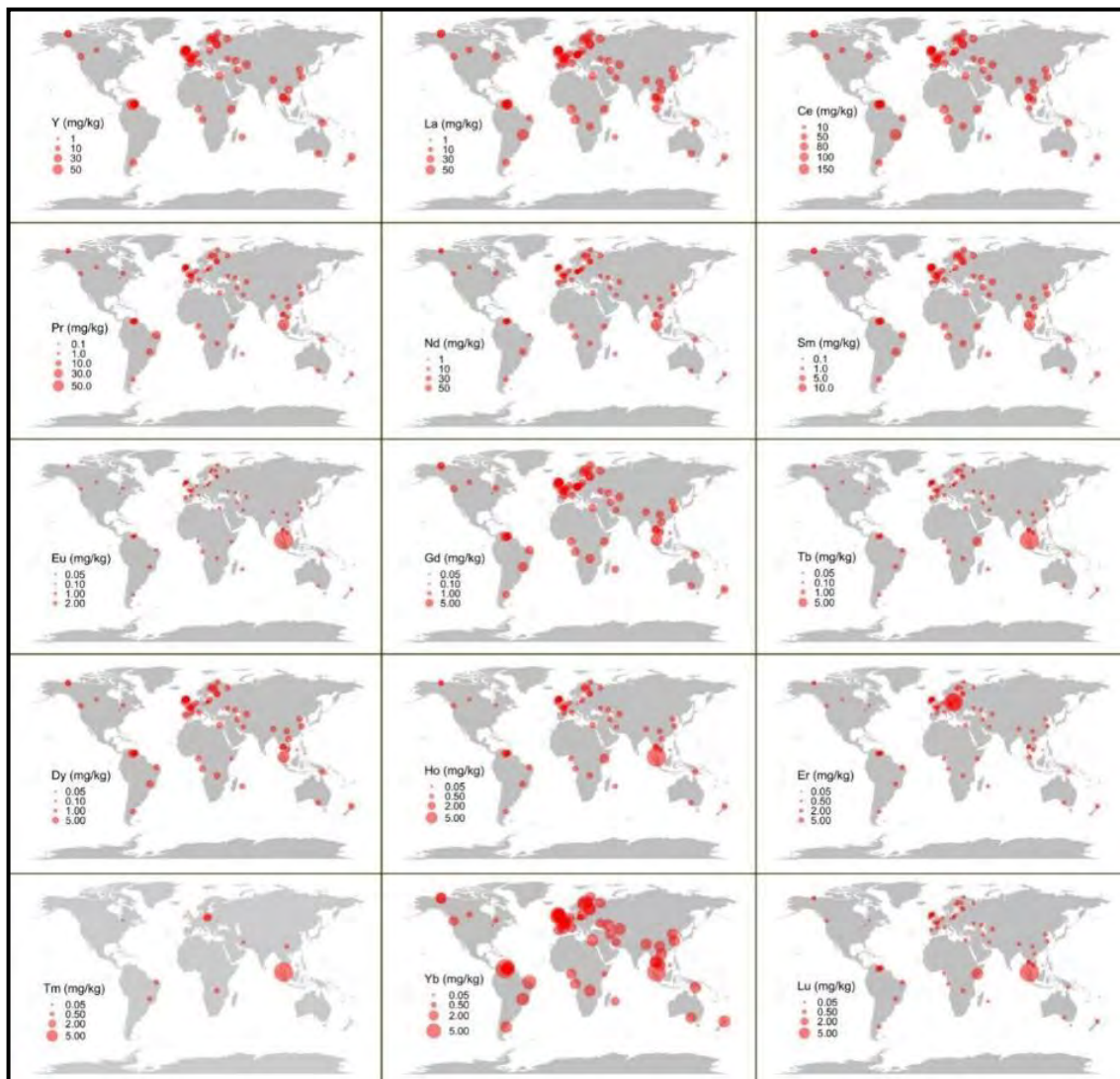


Figure A1A1 Spatial difference of REEs in soils/dusts and sediments in different countries. Data are compiled from two reviews (Adeel et al. 2019; Bayon et al. 2015) and collected Canadian data (Romero-Freire et al. 2019; Casse et al. 2019; MacMillan et al. 2018; Rasmussen et al. 2017; Chiarenzelli et al. 2001).

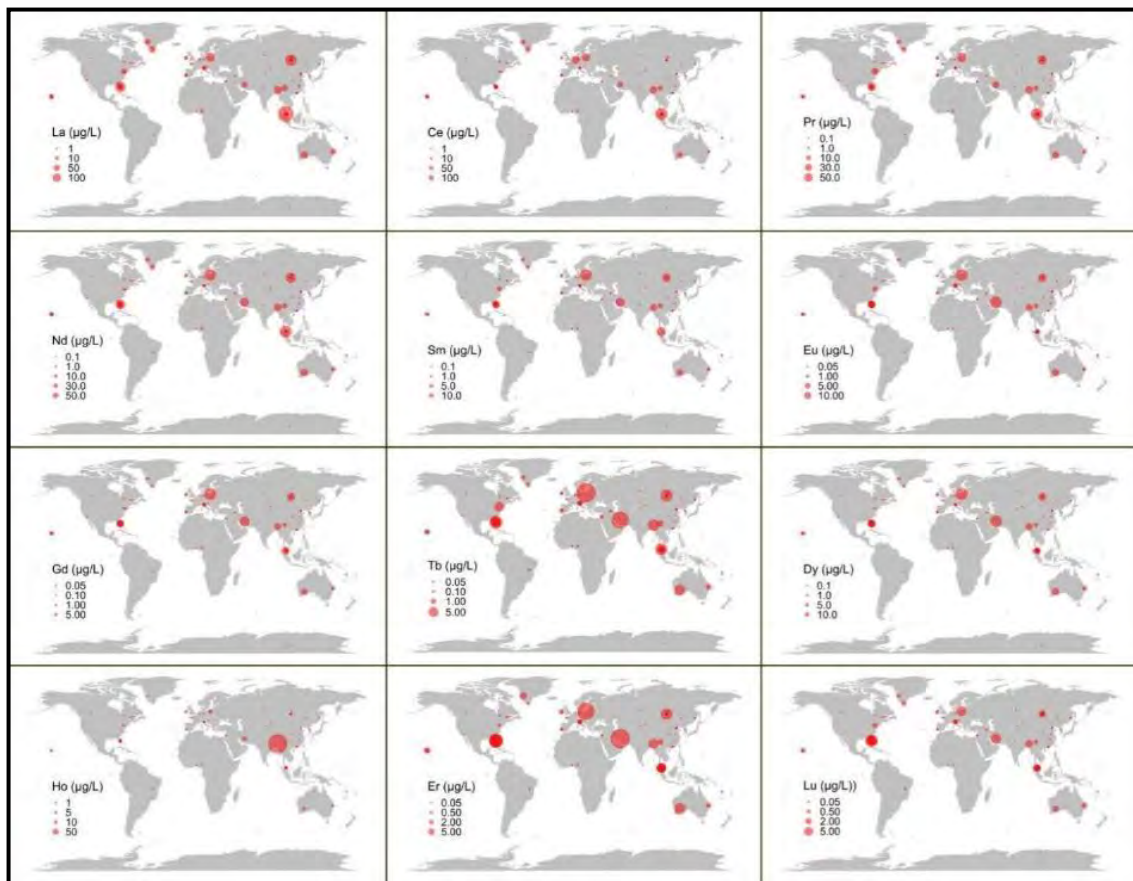


Figure A1A2 Spatial difference of REEs in various water matrices in different countries. Data are compiled from two reviews (Adeel et al. 2019; Bayon et al. 2015) and collected Canadian data (Romero-Freire et al. 2019; Casse et al. 2019; Cheung et al. 2009; Johannesson and Hendry 2000).

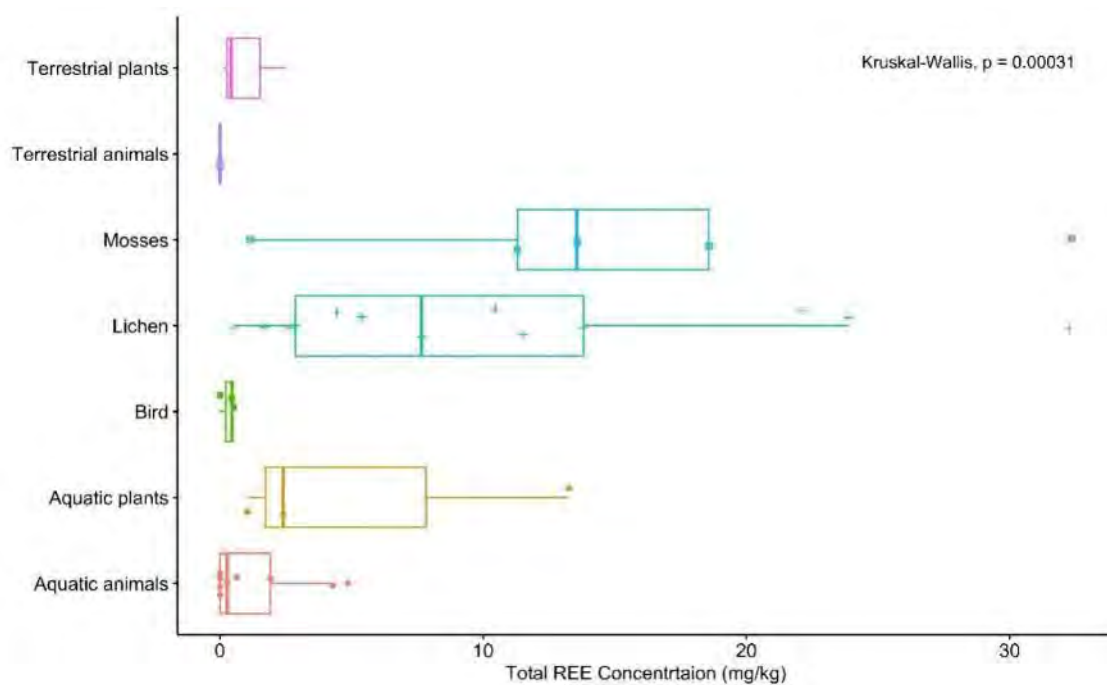


Figure A1A3 The REE concentration in different groups of living organisms in Canada based on Table A1.3.

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A1.11 Supplementary Materials

Table A1S1. The main driving factors (pH, DOC and Redox condition) for REEs mobility and bioavailability.

Factors	Finding	References
pH	Low pH and redox potential favored the REEs released from exchangeable fraction and Fe–Mn oxide fraction.	Cao et al. (2001)
	REEs in carbonate bound and Fe–Mn oxide bound fractions were negatively correlated with soil pH.	Zhang and Shan (2001)
	More of the REEs became adsorbed onto the organic matter with increased soil pH in acid sulfate soils.	Sukitprapanon et al. (2019)
Dissolved organic carbon	Humic substances enhanced REE adsorption at low pH range (i.e., pH < 4) and decreased REE adsorption when pH was above 4.	Tang and Johannesson (2010)
	There was a stronger connection between REEs and DOC than REEs and pH in a moderately acidic water (pH of samples: 3.61-6.42) of the Great Dismal Swamp watershed. Dissolved REEs were positively correlated to DOC concentrations and inversely correlated with pH.	Johannesson et al. (2004)
Redox condition	The major REE carriers were variable during alternating redox conditions. The REEs were bound to colloidal and dissolved organic matter under reducing conditions, while they were bound to big size particulates, colloidal, organic and Fe-enriched fractions in oxidizing conditions.	Guénet et al. (2018)
	Significant amounts of REEs were released during anoxic incubation (at pH=5)) of the wetland soil, while oxic incubation at pH=7 can stimulate soil desorption.	Davranche et al. (2011)

Table A1S2. The description of each REE fraction (Mseddi et al. 2010; Tessier et al. 1979)

Fractions	Description
Water soluble	Free REE^{3+} dissolved in water or soil solution.
Ion-exchangeable	The REEs adsorbed on clay minerals by electrostatic interaction. They can be released easily into leaching liquor upon ion exchange with NH_4^+ , Mg^{2+} , Fe^{2+} and Fe^{3+} .
Bound to Carbonates	Carbonate dissolution into soil porewaters increases soil pH, potentially enhancing the fixation of trace metals to carbonates via incorporation of cations in the carbonate unit cell. This fraction would be susceptible to changes of pH.
Bound to Iron and Manganese Oxides	Iron and manganese oxides exist as nodules, concretions, cement between particles, or simply as a coating on particles; these oxides are excellent scavengers for trace metals and are thermodynamically unstable under anoxic conditions.
Bound to Organic Matter	Trace metal may be bound to various forms of organic matter: living organisms, detritus, coating on mineral particles, etc. The complexation and peptization properties of natural organic matter (notably humic and fulvic acids) are well recognized, as is the phenomenon of bioaccumulation in certain living organisms. Under oxidizing conditions in natural waters, organic matter can be degraded leading to a release of soluble trace metals
Residual	Once the first four fractions have been removed, the residual solid should contain mainly primary and secondary minerals, which may hold trace metals within their crystal structure. These metals are not expected to be released in solution over a reasonable time span under the conditions normally encountered in nature.

Table AIS3 Recent studies about the REE toxicity to different groups of biotas.

Life forms	Species	Elements	REE Compound and Treatment	Toxicity	References
Vertebrate	Rat	Y	YCl ₃ (12mmol/L and 24mmol/L)	Neuronal cell death (+)	Ding et al. (2017)
	Rat	La	LaCl ₃ (0, 0.125, 0.25, 0.5, and 1%)	Spatial learning and memory (-), lactate transport (-)	Jin et al. (2017)
	Rat	La	La(NO ₃) ₃ (0, 1.5, 6.0, 24.0, and 144.0 mg/kg body weight)	Growth index: body weight (-), organ weight (-) and food consumption (-)	Fang et al. (2018)
Invertebrate	Rainbow trout (<i>Oncorhynchus mykiss</i>)	Ce, Er, Gd, La, Nd, Sm, Y	CeCl ₃ , ErCl ₃ , GdCl ₃ , LaCl ₃ , NdCl ₃ , SmCl ₃ and YCl ₃ (0.064, 0.32, 1.6, 8 and 40 mg/L)	Gene expression changes (- or +)	Dubé et al. (2019)
	Mussels (<i>Mytilus galloprovincialis</i>)	Gd	GdCl ₃ (0, 15, 30, 60, 120 µg/L)	Metabolic capacity (-), lipid peroxidation (+), antioxidant and biotransformation defences (+)	Henriques et al. (2019)
		La, Y	LaCl ₃ (0.01, 0.1, 1, 10, 100 mg/L) YCl ₃ (0.001, 0.01, 0.1, 1, 10 mg/L)	Embryogenesis success (-) Metabolic capacity (-), antioxidant and biotransformation defences (+), Neurotoxicity: activity of Acetylcholinesterase (-), Reproductive capacity: female spawning (-) Neurotoxicity: motor behaviors (-), dopaminergic and GABAergic neurons (-)	Mestre et al. (2019) Pinto et al. (2019)
Invertebrate	Nematode (<i>Caenorhabditis elegans</i>)	Nd, Sc	Pr, NdCl ₃ , PrCl ₃ and SeCl ₃ (0,10, 50, 75,100, 200 mg/L)	Metabolic capacity (-)	Xu et al. (2017a)
	Bivalve (<i>Corbicula fluminea</i>)	La	LaCl ₃ (0, 50, 100 and 200 µM)	Metabolic capacity (-)	Zhao and Liu (2018)
	Sea Urchin (<i>Paracentrotus lividus</i> and <i>Arbacia lixula</i>)	Dy, Er, Yb, Lu, Ce	REE salts (10-4 - 10-7 M)	Mitotic activity of embryos (-), nitric oxide (+), antioxidant defences (+), Fertilization success (-), offspring development (-)	Oral et al. (2017)
	Sea Urchin (<i>Sphaerechinus granularis</i>)	Dy, Er, Yb, Lu, Ce	DyCl ₃ , HoCl ₃ , ErCl ₃ , YbCl ₃ , LuCl ₃ and CeCl ₃ (10-4 - 10-7 M)	Fertilization success (-), offspring development (-)	Gravina et al. (2018)

Table A1S3 continued

		La	LaCl ₃ (0.40 and 1.20 mM)	Population respiration (-) and the respiratory key enzymes (-)	Zhang et al. (2019)
	Soybean (<i>Glycine max</i>)	La	LaCl ₃ (0.40 and 1.20 mM)	Root growth (-), photosynthesis parameters (-)	Zhang et al. (2018)
	Curlyleaf-pondweed (<i>Potamogeton crispus</i>)	Y	YCl ₃ (0, 10, 20, and 30 µM)	Photosynthetic pigments (-), growth rates (-), protein content (-), leaf area damage (+), elemental homeostasis (- or +)	Xu et al. (2019)
	<i>Brassica chinensis</i> and <i>Helianthus annuus</i>	La, Nd	La ₂ O ₃ and Nd ₂ O ₃ (20 mg/kg, 100 mg/kg, 200 mg/kg, and 300 mg/kg MS media)	plant biomass (-), total chlorophyll (-)	Rezaee et al. (2018)
Vascular Plant	Greater duckweed (<i>Spirodela polyrhiza</i>)	Ce	Ce(NO ₃) ₃ ·6H ₂ O (0, 10, 20, 40 and 60 µM)	Growth (-), photosynthetic pigment (-), protein contents (-), cell death (+)	Xu et al. (2017b)
	Duckweed (<i>Lemna minor</i>)	Ce	Ce nitrate (0, 2.5, 5, 10, 20 µM and 0.1, 0.5, 1 mM) Ce chloride and Ce sulphate (0.5 and 1mM)	Photosynthetic pigments (-), hydrogen peroxide (+), antioxidant activity (+)	Zicari et al. (2018)
	Onion (<i>Allium cepa</i>)	La, Ce	LaCl ₃ and CeCl ₃ (0, 10, 20, 50, 100, and 200 mg/kg air-dry soil)	Cell division (-), aberrant cells (+)	Kotelnikova et al. (2019)
	Sweetpotato (<i>Ipomoea batatas</i>)	Ce	CeCl ₃ (10, 20, 40 and 80 mg/L)	Photosynthesis (-), activities of antioxidant enzymes (+)	Jiang et al. (2017)
Non-vascular Plant	Algae (<i>Chlorella vulgaris</i> and <i>Phaeodactylum Tricornutum</i>)	La	La(NO ₃) ₃ ·6H ₂ O (0, 2, 4, 6, 8, and 10 mg/L)	Photosynthetic system (-), activities of antioxidant enzymes (+)	Sun et al. (2019)
Microbiome	Soil microbial communities	La ₂ O ₃ , Nd ₂ O ₃ , Gd ₂ O ₃	La ₂ O ₃ , Nd ₂ O ₃ , Gd ₂ O (10, 50, 100 mg/kg soil)	The proliferation of antibiotic resistance genes (+)	Qi et al. (2019)
	Fungus (<i>Shirata bambusicola</i>)	La	LaCl ₃ (0.0-1.4 g/L)	Hydrogen peroxide (+), hypocrellin A (+)	Lu et al. (2019)

Note: (-) means negative impacts; (+) means positive impact.

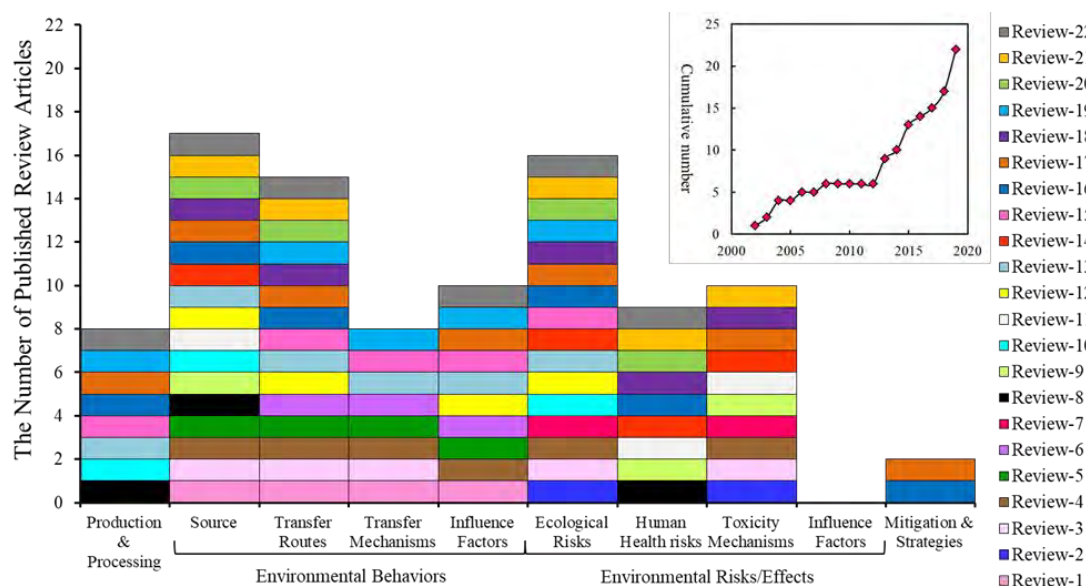


Figure A1S1 The topics reported in review literatures

Note: Review-1 (Pang et al. 2002), Review-2 (Tomonori 2003), Review-3 (Hu et al. 2004), Review-4 (Tyler 2004), Review-5 (Hu et al. 2006), Review-6 (Liang et al. 2008), Review-7 (Zhang et al. 2013), Review-8 (Weng et al. 2013), Review-9 (Rim et al. 2013), Review-10 (Haque et al. 2014), Review-11 (Pagano et al. 2015a), Review-12 (Pagano et al. 2015b), Review-13 (Migaszewski and Gałuszka 2015), Review-14 (Rim 2016), Review-15 (Khan et al. 2017), Review-16 (Gwenzi et al. 2018), Review-17 (Turra 2018), Review-18 (Adeel et al. 2019), Review-19 (Edahbi et al. 2019), Review-20 (Pagano et al. 2019), Review-21 (Abdelnour et al. 2019), Review-22 (Balaram 2019), Review-23 (Malhotra et al. 2020), Review-24 (Langkau et al. 2020).

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ANNEXE B

RICCARDIA VITREA (ANEURACEAE), UNE ESPÈCE D'HÉPATIQUE
NOUVELLE EN CHINE ET EN AMÉRIQUE DU NORD

RICCARDIA VITREA (ANEURACEAE), A LIVERWORT SPECIES NEW TO
CHINA AND NORTH AMERICA

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Herzogia, DOI: 10.13158/heia.34.2.2021.279

A2.1 Résumé

L'espèce d'hépatique *Riccardia vitrea* Furuki (Aneuraceae) est signalée pour la première fois en Chine et en Amérique du Nord (Canada). Ce rapport est principalement axé sur sa présence au Canada. *Riccardia vitrea* est caractérisé par de nombreux petits corps huileux dans les cellules épidermiques et internes, et par des pinnules ultimes prostrées, larges et peu développées, et se distingue ainsi facilement des autres espèces canadiennes du genre. Ceci constitue une extension significative de son aire de répartition connue, de l'Asie orientale à l'Amérique du Nord. Cette espèce pourrait être négligée en raison du fait que les spécimens secs sans corps gras sont souvent utilisés pour l'identification, et elle est probablement plus répandue que ce qui est actuellement connu dans les forêts boréales du Canada. Des illustrations et des descriptions basées sur les spécimens canadiens sont fournies avec une clé pour les espèces canadiennes du genre.

Mots-clés : Bryophytes, hepaticae, hépatiques thalloïdes simples, forêt boréale, Québec.

A2.2 Abstract

The liverwort species *Riccardia vitrea* Furuki (Aneuraceae) is reported for the first time for China and North America (Canada). This report is primarily focused on its occurrence in Canada. *Riccardia vitrea* is characterized by numerous small oil bodies in both epidermal and inner cells, and by prostrate, wide and poorly developed ultimate pinnules, and is thus easily distinguished from other Canadian species of the genus. This constitutes a significant extension of its known range from East Asia to North America. This species could be overlooked owing to the fact that dry specimens without oil bodies are often used in identification, and it is probably more widespread than currently known in the boreal forests of Canada. Illustrations and

descriptions based on the Canadian specimens are provided with a key for Canadian species of the genus.

Key words: Bryophytes, hepaticae, simple thalloid liverworts, boreal forest, Quebec.

A2.3 Introduction

Riccardia vitrea Furuki is a simple thalloid liverwort species in the family Aneuraceae, which was first recorded in Japan (Honshu, Shikoku, Kyushu, Furuki 1991). Its known range was subsequently expanded to China (Yin 2017, not published) and Russia (Bakalin 2018). To date, *R. vitrea* has not been reported from outside of East Asia. However, a new population was observed during recent fieldwork on bryophyte diversity in boreal forests at a post-mine site in north-west Quebec, Canada. Combining morphological and phylogenetic analyses, this new record was confirmed as an addition to the bryophyte flora of Canada and North America. Although *Riccardia* Gray is the largest genus of simple thalloid liverworts, with more than 250 names currently accepted (Gradstein & Cuvertino 2015, Söderström et al. 2016), only four species had been recorded in Canada, *R. chamedryfolia* (With.) Grolle, *R. latifrons* (Lindb.) Lindb., *R. palmata* (Hedw.) Carruth., and *R. multifida* (L.) Gray (Stotler & Crandall- Stotler 2017, Haughian et al. 2016, Faubert 2012, Damsholt 2007, Hong 2007, Brassard & Williams 1975), and they are all common species in the global temperate and boreal zones. *Riccardia vitrea* can be easily confused with these other species when it is dry, because of the great polymorphism of this genus (Reeb & Bardat 2014) and the loss of oil bodies which provide key characteristics for taxonomy in genus *Riccardia*. The shape, position, size and number of oil bodies are important in this genus for identification at the species level. Many bryologists have suggested the use of fresh material with oil bodies in this genus for more reliable identification (Reeb & Bardat 2014, Bakalin

2018, Schuster 1989). Short morphological and ecological descriptions of the species based on the specimens collected in Canada are provided in this study.

A2.4 Materials and Methods

The materials were collected during a field survey in Joutel, James Bay, Quebec, Canada, during the summers of 2019 and 2020. The study area is located in the *Picea mariana* feather moss bioclimatic domain and is influenced by cold and dry arctic air during the winter. The mean temperature is -20 °C in January and 16 °C in July. Mean annual precipitation is approximately 850mm (Bergeron et al. 2004). Collected samples were examined under a microscope (Olympus BX 43) and a stereomicroscope (Olympus SZX 16). The photos were taken with an Olympus DP 22 digital camera.



Figure A2.1 Habitat of *Riccardia vitrea* at Joutel, Quebec, Canada (Sample 201909 Joutel10 (UQAT)). A – Deciduous forest in which *R. vitrea* was collected. B–D – The microhabitat of *R. vitrea* at this site.

A2.5 Taxonomic description and discussion

Riccardia vitrea Furuki, J. Hattori Bot. Lab. 70: 327. 1991. (Figures A2.1 and A2.2)

Thallus medium to large, in dense radial patches, closely adjacent to the substratum, prostrate along main axis, irregularly pinnately or palmately branched, fresh-green to brown-green when fresh, whitish brown when dry, smooth and flat at apices. The main axis up to 9mm long, 0.3 – 0.8 mm wide, biconvex to plano-convex, in cross-section 5 – 7 cells (98 – 150µm) thick, obtuse at margins. Ultimate branches prostrate, poorly developed, wide, stuck to the substratum, up to 1.5mm long, 0.3 – 0.6mm wide, biconvex to plano-convex, only 5 – 7 cells (92–130 µm) thick in cross-section, acute to obtuse at margins. Epidermal cells of thallus thin walled, 28 – 92 × 22– 45 µm, about 1/3 – 1/2 size of inner cells (Figure A2.2F, G). Inner cells 53 – 107 × 20 – 61 µm. Oil bodies brownish to grayish brown, globose to ovoid, indistinctly granular. Oil bodies in epidermal cells (0 –)1–9/cell, 3.64 – 11.74 × 3.16 – 7.22 µm. Oil bodies in inner cells (0 –)2–14/cell, 4.44 – 9.97 × 5.58 – 14.98 µm. Mucilage hairs in 2 rows on the ventral surface of thallus only at apices, 16.66 – 46.31µm long, 7.56 – 18.58 µm thick. Rhizoids are often present on the ventral surface of the thallus. Dioicous. Female and male branches lateral on the main axis or base of primary. Few mature calyptrae and developed male branches were found in collected samples. Based on collected samples, paraphyses fringed, scale-like, 5– 8 cell high (100 – 200 µm), 2– 4 cells wide at the base. Elaters 85 – 230 µm. Spores 11.74 – 13.30 µm in diam. These descriptions are based on collected Canadian samples of 202007-CT6E and 201909-Joutel10. The description of Chinese samples was presented in Yin (2017), but figures are attached in the supporting document (Figures A2S1 and A2S2 in Supplementary documents online).

Habitat: On wet sandy soil and rock at open sites (in Japan, China and Russia). On decaying logs in deciduous and mixed forests (in Russia and Canada). In Canada, this species is found on moist decaying logs without bark (Figure 1), growing with other

bryophyte species including *Syzygiella autumnalis* (DC.) K.Feldberg, Vána, Hentschel & Heinrichs, *Geocalyx graveolens* (Schrad.) Nees, *Lophocolea bidentata* (L.) Dumort., and *Sanionia uncinata* (Hedw.) Loeske.

Range: Japan (Furuki 1991, Furuki & Higuchi 2006), China (Yin 2017), Russia (Bakalin 2018) and Canada (present report)

Phylogenetic analysis: This new record of the species is strongly supported by a molecular phylogenetic analysis based on DNA sequences data of one plastid regions *rbcL* and available sequences for other species downloaded from GenBank (Bainard et al. 2013, Masuzaki et al. 2010, Forrest et al. 2006). The phylogenetic study includes *R. vitrea* samples from Japan (GenBank: AB476596, Masuzaki et al. 2010), China (GenBank: OK302921 and OK302922) and Canada (GenBank: OK302923) as well as for the similar species *R. latifrons* and *R. palmata* that are recorded in Canada. The Canadian sample was placed in the same clade with *R. vitrea* from China and Japan and has a clearly different topology from other morphologically similar species known from Canada (phylogenetic tree and primers used shown in Figure A2S3 and Table A2S1 in Supplementary documents online).

Specimens examined from Canada: CANADA. Quebec, James Bay, Joutel, 49°30'35.64"N/78°17'16.8"W, 300m, on decaying logs, in mixed forest, 8 July 2020, X. B. Yin, X. St-Amant & T. Kim 202007-CT6E (UQAT); Canada. Quebec, James Bay, Joutel, 49°29'2.04"N/78°23'13.92"W, 300 m, on decaying logs, in deciduous forest, 27 September 2019, X. B. Yin & R. Boisvert 201909-Joutel10 (UQAT).

Discussion: The main characters of *R. vitrea* include 1) thallus closely adjacent to the substratum (Figure A2.2A), 2) numerous small oil bodies in epidermal and inner cells (Figure A2.2F, G), 3) prostrate, wide and poorly developed ultimate pinnules (Figure A2.2D), 4) large epidermal cells, similar size with inner cells seen in the cross-section of the thallus (Figure A2.2I, J), 5) obtuse to acute margins in cross-section of thallus

(Figure A2.2J). This species was already recorded in Japan (Furuki 1991, Furuki & Higuchi 2006), China (Anhui, Fujian, Hainan, Yunnan provinces, Yin 2017) and Russia (Far East, Bakalin 2018). Although it was first believed to be endemic to Japan when found by Furuki (1991), more findings were recorded outside Japan. At the same time, its known habitat has been extended from open moist soil and rock to include decaying logs in forests, and its biogeographical range includes subtropical and tropical ecosystems to sub-arctic northern ecosystems (Bakalin 2018). This species could be widely overlooked due to the fact that dry specimens without oil bodies are often used in identification, and it is probably more widespread than currently known. Misidentification with *R. latifrons* may be the main reason for its current limited recorded distribution in Canada. In the field, we found *R. latifrons* and *R. vitrea* at the same place, but the presence of a brown-green thallus and prostrate, wide ultimate pinnules make *R. vitrea* significantly different from *R. latifrons*, with narrow and ascending to erect ultimate pinnules. Another unique character of *R. latifrons* is the apparent lack of oil bodies, which allows easy separation from other species in Canada. However, identification can be difficult when material is dry due to the loss of morphological traits (e.g., oil bodies, thallus shape). Examining fresh material is best for accurate identification in genus *Riccardia*. This species is also similar to *R. palmata* in having small and ovoid oil bodies. However, oil bodies in North American samples are 1–2 per inner cell, and mostly absent from the epidermal cells (Schuster 1987). It is worth mentioning that we found that oil bodies can be kept for more than five months in a 4 °C fridge. Based on collected samples and references (Furuki 1991, Faubert 2012, Schuster 1987), the key to *Riccardia* species present in Canada is as follows.

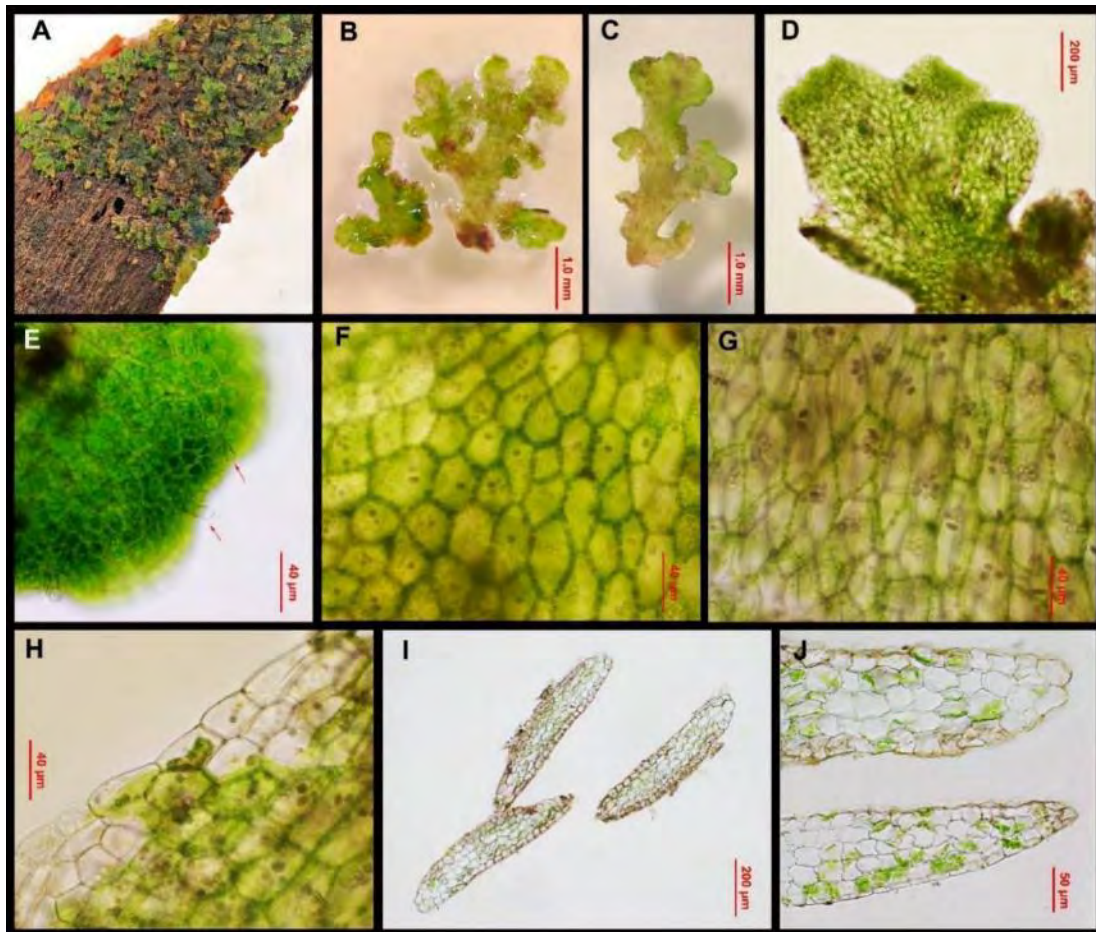


Figure A2.2 *Riccardia vitrea* Furuki. A – A colony of *R. vitrea* on decaying log. B–D – Thalli. E – Apex of ultimate pinnule with mucilage hairs. F– Epidermal cells with oil bodies. G – Inner cells with oil bodies. H – Marginal cells of ultimate pinnule. I–J – Cross sections of thallus (I – Cross sections of ultimate pinnules; J – Cross section of main axes). All from 201909-Joutel10 (UQAT).

1 Thallus regularly to irregularly pinnately to tripinnately branched; having moderately to well-developed pinnae; winged at the margin of the ultimate pinnule, unistratose alar portion more than 2 cells wide; ultimate pinnules cross-sections 3–5 cells thick; usually on wet soil or rocks, rarely on wet decaying logs 2

1* Thallus irregularly pinnately to tripinnately branched, having poorly to moderately developed pinnae; acute to winged at the margin of ultimate pinnule, unistratose alar

portion less than 2 cells wide; ultimate pinnules cross-sections 4 –7 cells thick; usually on wet decaying logs or peatland **3**

2 Autoicous; flat cells along margin; oil bodies (1–3) in each epidermal, inner and wing cell **R.**

chamedryfolia

2* Heteroicous; crenulate cells along margin; oil bodies (1–2) in each inner cell, absent or rare in epidermal and wing cells **R.**

multifida

3 Monoicous; oil bodies absent or rare in all cells **R. latifrons**

3* Dioicous; oil bodies present at least in inner cells **4**

4 Ultimate pinnules prostrate to ascending, moderately developed, less than 0.4mm wide; emarginate to bifid at apices; oil bodies usually 1–2 per inner cell, mostly absent from the epidermal cells, diameter over 15µm **R.**

palmata

4* Ultimate pinnules prostrate, poorly developed, more than 0.4mm wide; rounded at apices; oil bodies more than 2 per inner and epidermal cells, diameter less than 10µm **R. vitrea**

A2.6 Acknowledgements

This research was supported by Agnico Eagle Mines Ltd, NSERC-UQAT Industrial Research Chair on northern biodiversity in a mining context, and the National Natural Science Foundation of China (NSFC) (no. 31570206). We are grateful to Dr. Frank Müller and Dr. Marc Favreau for their helpful remarks on the manuscript, and Dr. Frank Müller for the preparation of the German abstract. We also thank the council of the First Nation of Abitibiwinini (Pikogan) on whose traditional land the colony was found.

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A2.7 Supplementary documents

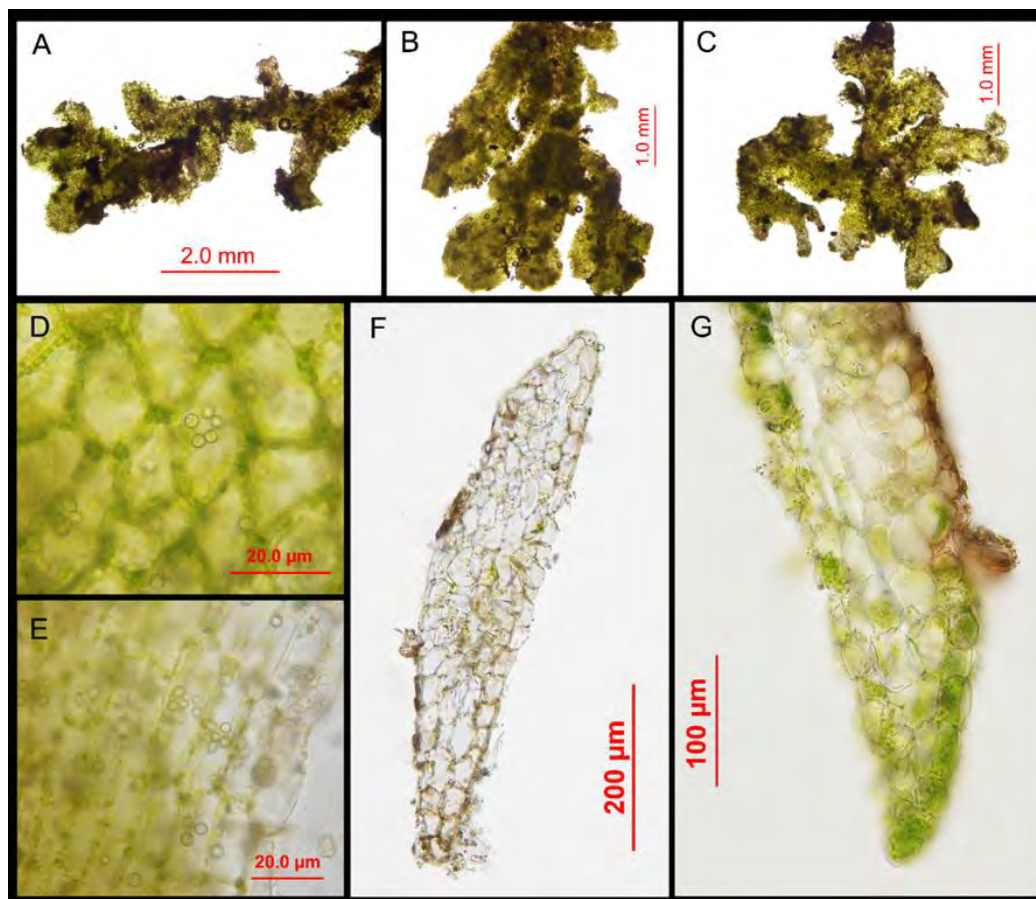


Figure A2S1 *Riccardia vitrea* Furuki. A–C–Thalli; D–E–Epidermal cells with oil bodies; F–G–Cross-sections of the thallus (F – Cross-sections of ultimate pinnules, G – Cross-section of main axes). D, E, F from Shi et al., 20150920-16 (HSNU); A, B, G from Shi et al., 20150922-7; C from Shi et al., 20150922-31. Voucher specimen for Chinese materials (Herbaria of East China Normal University, China. HSNU): 1 – China, Fujian Province, Dehua County, Jiuxian Mountain, Wang et al., 20090730-49; 2 – China, Anhui Province, Qingliangfeng National Nature Reserve, Shi et al., 20150920-16, 20150922-7, 20150922-31; 3 – China, Yunnan Province, Mengla County, Peng et al., 20150130-23, 2016042-10.

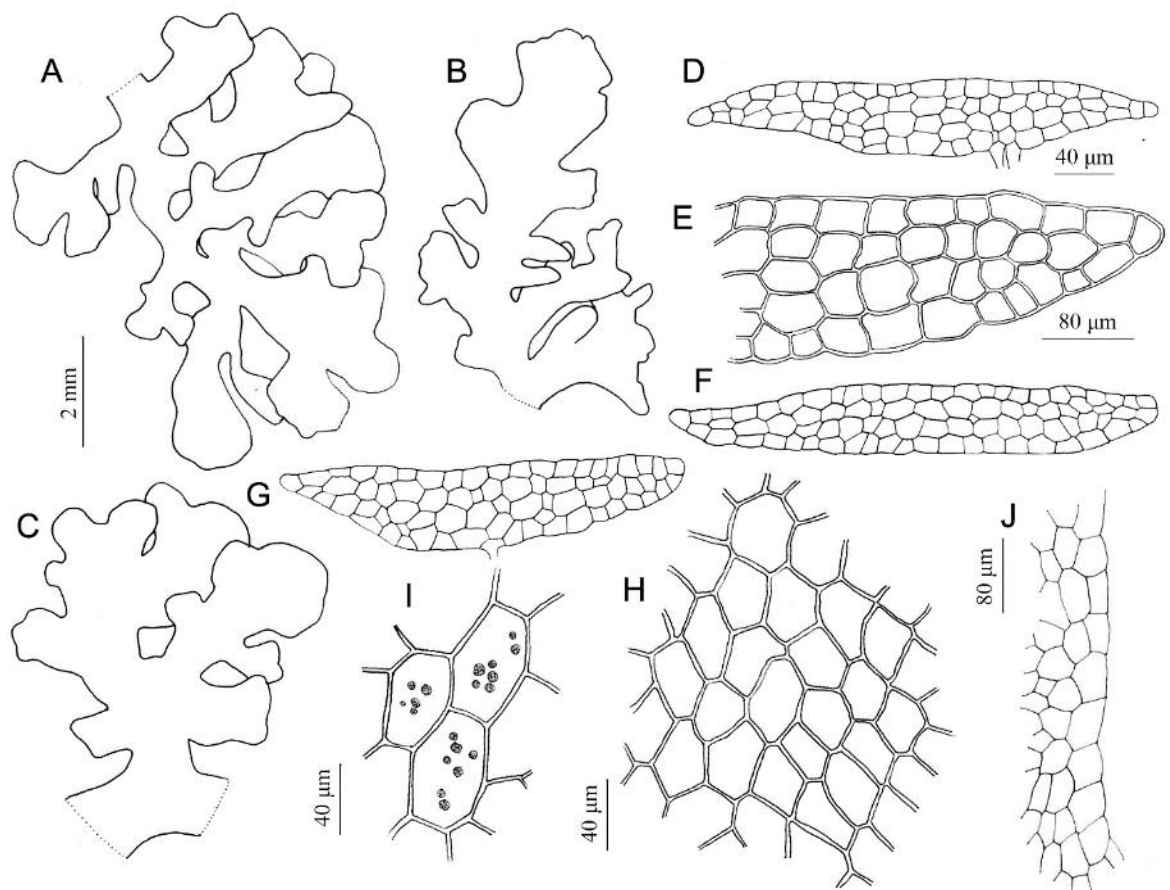


Figure A2S2. *Riccardia vitrea* Furuki. A–C – Thalli. D–G – Cross sections of thallus (D–E. ultimate pinnules; F–G. main axes). H – Epidermal cells of median part of ultimate pinnule. I – Epidermal cells of main axes, with oil bodies. J – Marginal cells of ultimate pinnule. All from Shi et al.20150920–16 (HSNU). The figure from Yin (2017).

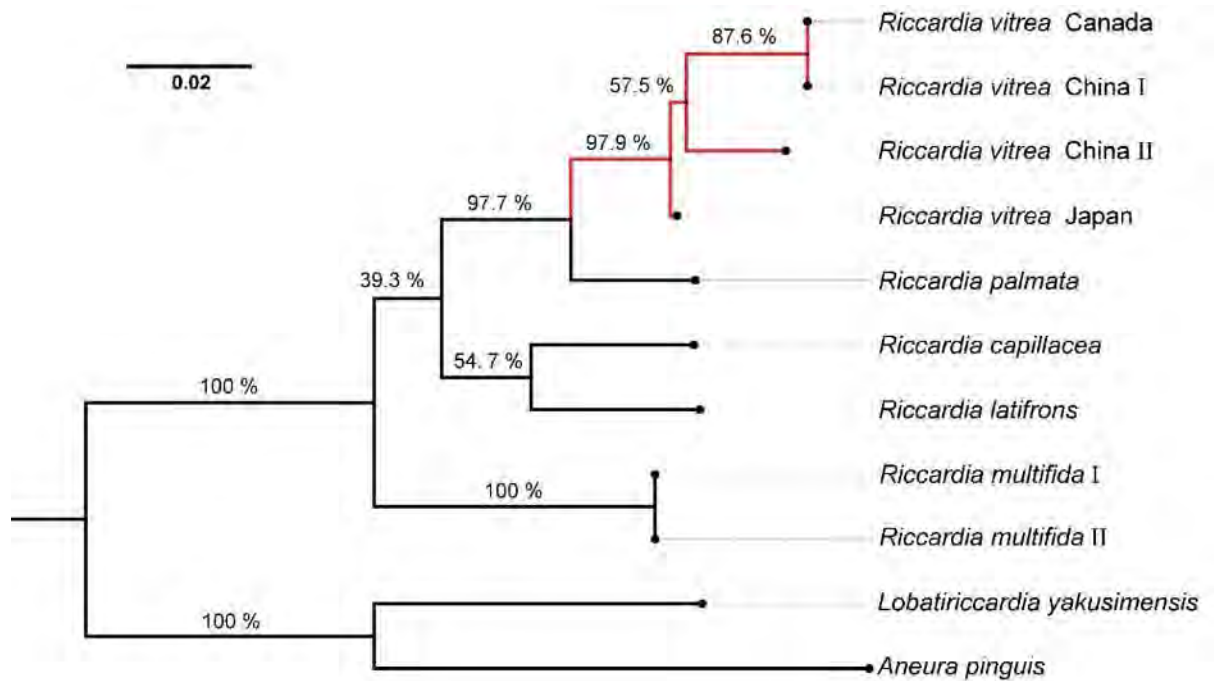


Figure A2S3 A simple maximum likelihood tree was constructed based on the *rbcL* gene of *R. vitrea* and similar species in Canada (*R. palmata* and *R. latifrons*). Primer sequences for *rbcL* are listed in Table A2S1. A phylogenetic tree with more species and sequences will be shown in a future article about the revision of Aneuraceae in China. Sequences ID from Genbank are listed in Table A2S2. The maximum likelihood (ML) method as implemented by PhyML version 3.0 (GUINDON et al. 2010). The best-fit model was GTR+G+I determined using ModelTest (POSADA & KEITH 1998). The proportion of invariant sites and the gamma distribution parameter alpha were estimated from the data. Likelihood bootstrap searches (MLBS) were performed with 1000 times.

Table A2S1 Primer sequences used for *rbcL*.

	Primer	Sequence (5'–3')	Direction	Reference
<i>rbcL</i>	1pIF	ATGTCACCACAAACGGAGACTAA ARCAGGT	Forward	WILSON et al. 2004
	600R	CCAACGCATAAATGGTTGAG	Reverse	GRADSTEIN et al. 2006
	210F	TGGACTACGGTTTGGACTGA	Forward	GRADSTEIN et al. 2006
	1200R	TGYCCYAAAGTTCCACCACC	Reverse	WILSON et al. 2004
	640F	CTCAACCATTATGCGTTGG	Forward	GRADSTEIN et al. 2006
	1390R	CTTTCCAWAYTTCRCAAGCAGCRG	Reverse	WILSON et al. 2004

Table A2S2 Sequences ID from Genbank present in the phylogenetic tree.

Species	Location	Genbank ID
<i>Riccardia vitrea</i>	Japan: Honshu, Tokyo, Ohshima	AB476596.1
<i>R. palmata</i>	USA: Pennsylvania	KC305704.1
<i>R. capillacea</i>	Venezuela: Merida	AY877392.1
<i>R. latifrons</i>	United Kingdom	KC305702.1
<i>R. multifida</i> I	USA: North Carolina	KC305703.1
<i>R. multifida</i> II	Canada: Vancouver Island	DQ268980.1
<i>Lobatiriccardia yakusimensis</i>	Japan: Kagoshima	AB673209.1
<i>Aneura pinguis</i>	Poland	KY705822.1
Unpublished sequences		
<i>R. vitrea</i>	Canada: Quebec	OK302923
<i>R. vitrea</i> I	China: Anhui	OK302922
<i>R. vitrea</i> II	China: Anhui	OK302921

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ANNEXE C

EFFETS HORS SITE DE L'EXPLOITATION MINIÈRE SUR LA FRÉQUENCE
ET L'ABONDANCE DE CINQ ESPÈCES DE PLANTES DE SOUS-BOIS DANS
L'OUEST DU QUÉBEC (CANADA)

OFFSITE EFFECTS OF MINING ON THE FREQUENCY AND ABUNDANCE
OF FIVE UNDERSTOREY PLANT SPECIES IN WESTERN QUÉBEC (CANADA)

Rémi Boisvert, Xiangbo Yin & Nicole J. Fenton

Botany, DOI: 10.1139/cjb-2020-0158

A3.1 Résumé

Bien que l'industrie minière fournisse des minéraux et des métaux pour le marché mondial et représente d'importantes occasions économiques, elle constitue aussi une perturbation anthropique majeure dans les écosystèmes où elle prend place. Toutefois, on sait peu de choses sur ses impacts hors site sur les communautés végétales. Les auteurs ont évalué la fréquence et l'abondance de cinq espèces de plantes de sous-bois boréales communes (*Coptis trifolia* (L.) Salisb., *Cornus canadensis* L., *Linnaea borealis* L., *Lysimachia borealis* (Raf.) U.Manns & Anderb., and *Maianthemum canadense* Desf.) à différentes distances autour de six mines en Abitibi-Témiscamingue et dans le Nord-du-Québec (Québec). Leurs résultats ont montré que *Coptis trifolia* et *M. canadense* étaient affectés négativement par la présence des mines, ce qui suggère qu'ils pourraient être sensibles à la perte du couvert forestier, au dépôt de particules et à la contamination du sol induits par l'exploitation minière. À l'inverse, l'activité minière a favorisé, *Cornus canadensis*, ce qui indique sa résilience au stress causé par les mines et sa capacité à exploiter l'espace de niche laissé par les espèces plus sensibles. Aucun effet sur *Linnaea borealis* et *Lysimachia borealis* n'a été observé. Cette étude confirme l'existence d'impacts hors site des mines sur les communautés végétales. Des études supplémentaires portant sur les impacts non conventionnels devraient être réalisées afin de développer notre compréhension de l'empreinte environnementale plus large de l'activité minière. [Traduit par la Rédaction]

Mots-clés : forêt boréale, impacts non conventionnels, impacts hors site, communautés végétales, stress.

A3.2 Abstract

Although the mining industry provides minerals and metals to the global market and represents important economic opportunities, it also constitutes a major anthropogenic disturbance in the ecosystems where it takes place. However, little is known about its offsite impacts on plant communities. We assessed the frequency and the abundance of five common boreal understorey plant species (*Coptis trifolia* (L.) Salisb., *Cornus canadensis* L., *Linnaea borealis* L., *Lysimachia borealis* (Raf.) U.Manns & Anderb., and *Maianthemum canadense* Desf.) at different distances around six mines in Abitibi-Témiscamingue and Nord-du-Québec (Québec). Frequency and (or) abundance of *Coptis trifolia* and *M. canadense* decreased near mines, suggesting that they might be sensitive to the loss of forest cover, particle deposition, and soil contamination induced by mining. Conversely, the frequency and abundance of *Cornus canadensis* increased near active mines, which indicates its resilience to the stress caused by mines and its ability to exploit the niche space left by more sensitive species. No effects on *Linnaea borealis* and *Lysimachia borealis* were observed. Our study suggests the possible presence of offsite impacts of mines on plant communities. Further studies considering enigmatic impacts should be conducted in order to develop our understanding of the broader environmental footprint of mining activity.

Key words: boreal forest, enigmatic impacts, offsite impacts, plant communities, stress.

A3.3 Introduction

Although the mining industry provides minerals and metals to the global market and represents important economic opportunities, it also constitutes a major anthropogenic disturbance in the ecosystems where it takes place. In fact, mining

development has significant ecological impacts, contributing to the loss, degradation, and conversion of wildlife habitats (Johnson and Skousen 1995; Simmons et al. 2008; Gould 2011). These impacts are partly the result of land-cover change at the mining site and are direct and easily assessable. However, ecological impacts are not restricted to the immediate location of the mines. Offsite impacts are substantial, although little is known about their extent. Raiter et al. (2014) included offsite impacts as one of their four categories of enigmatic impacts, which they define as the “impacts that are easily and often overlooked in impact evaluations”.

Various offsite impacts are associated with mining activity. Loss of canopy cover can be observed around mines due to the edge effects they generate. In fact, anthropogenic development not only destroys vegetation at its immediate location, but it also induces tree mortality in the surrounding area (Nascimento and Laurance 2004). Anthropogenic openings have been shown to provoke more significant edge effects than natural openings in boreal forests (Vepakomma et al. 2018). Generation of dust is another issue related to mining (Ghose and Majee 2000). The transport of dust particles by wind (particles that can contain metals and metalloids) leads to their deposition in a large perimeter around the mine site and can have negative effects on vegetation. (Ghose and Majee 2000; Csavina et al. 2012; Li et al. 2018). Soil contamination is also a major risk generated by the mining industry, principally by acid mine drainage. The acidification of soil causes the solubilization of toxic heavy metals, which then become available for plants (Zhuang et al. 2009). Significant heavy metal concentrations can be observed several kilometers away from the mine sites (Escarré et al. 2011). The combination of these impacts might have an effect on surrounding plant communities by inducing the loss of native species thus creating a favourable environment for the introduction of exotic ruderal species, which then alters the functioning of the ecosystems (Hooper et al. 2005).

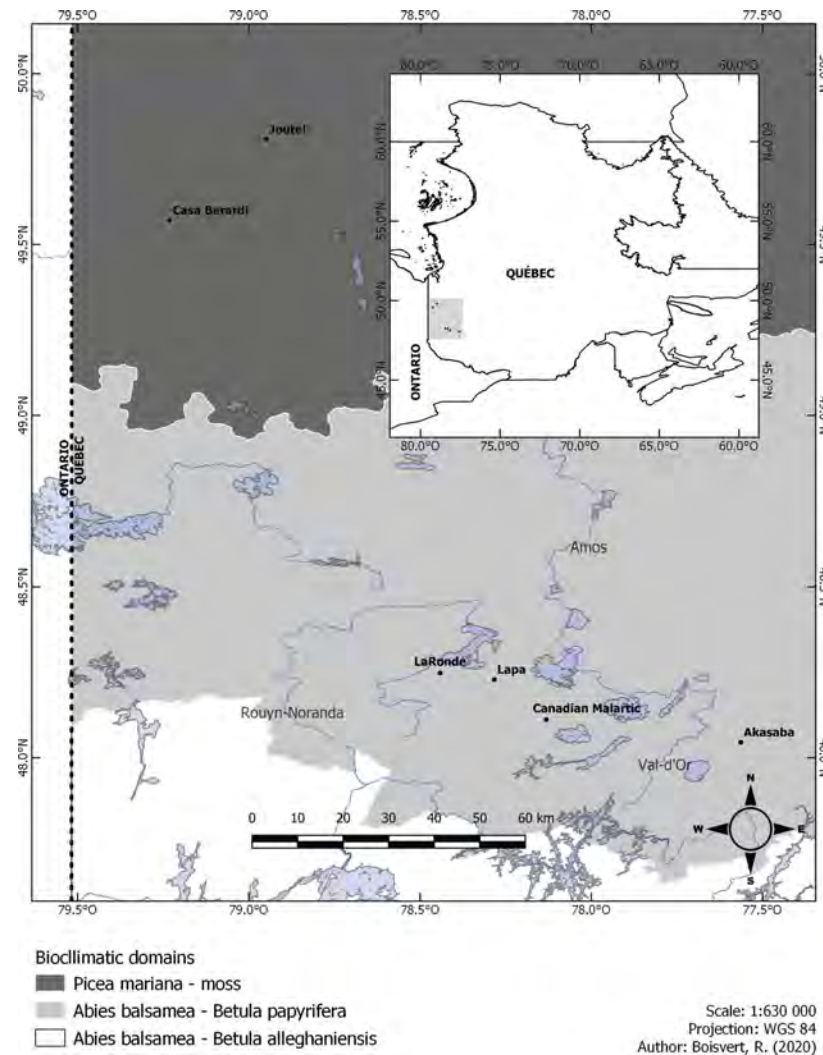


Figure A3.1 Geographic position of the six mines sampled: Casa Berardi (Hecla Québec Inc.), Joutel (Agnico Eagle Mines Ltd.), Akasaba (Agnico Eagle Mines Ltd.), Canadian Malartic (Canadian Malartic Corp.), Lapa (Agnico Eagle Mines Ltd.) and LaRonde (Agnico Eagle Mines Ltd.). Province boundary data from Statistics Canada (2011). Municipality data from Ministère de l'Énergie et des Ressources Naturelles (2018). Water system data from Ministère de l'Énergie et des Ressources Naturelles (2019). Bioclimatic domain data from Ministère des Forêts, de la Faune et des Parcs (2016). Themap was created in QGIS (QGIS Development Team 2020). [Colour online.]

Table A3.1: Description of the sampling design, including the number of transects sampled, the number of plots of each ecosystem type encountered around the mine and the number total of plots sampled for each mine.

Mines	Akasaba	Canadian Malaric	Lapa	LaRonde	Joutel	Casa Berardi	Total
Number of transects	6	8	7	8	7	8	44
Number of plots per ecosystem type	22 coniferous	17 coniferous	21 coniferous	9 coniferous	22 coniferous	32 coniferous	123 coniferous
	1 deciduous	9 deciduous	5 deciduous	27 deciduous	9 deciduous	2 deciduous	53 deciduous
	6 mixedwood	24 mixedwood	19 mixedwood	17 mixedwood	12 mixedwood	6 mixedwood	84 mixedwood
	13 wetland	6 wetland	4 wetland	4 wetland	6 wetland	16 wetland	49 wetlands
Total number of plots	42	56	49	56	49	56	308

Table A3.2 Fixed effect variables and interactions included in each model after the forward selection method.

Species	Frequency models	Abundance models
<i>Coptis trifolia</i>	Variables: Distance from mine Ecosystem type Interaction: Ecosystem type: distance	Variables: Distance from mine Ecosystem type Mine stage Interaction: -
<i>Cornus canadensis</i>	Variables: Distance from mine Ecosystem type Mine stage Interaction: Mine stage: distance	Variables: Distance from mine Ecosystem type Mine stage Interactions: Mine stage: distance Ecosystem type: mine stage
<i>Linnaea borealis</i>	Variables: Distance from mine Ecosystem type Mine stage Interactions: Ecosystem type: distance Mine stage: distance	Variable: Distance from mine Interaction: -
<i>Lysimachia borealis</i>	Variables: Distance from mine Ecosystem type Mine stage Interaction: Ecosystem type: distance	Variables: Distance from mine Ecosystem type Mine stage Interaction: Ecosystem type: mine stage
<i>Maianthemum canadense</i>	Variables: Distance from mine Ecosystem type Mine stage Interactions: Ecosystem type: mine stage	Variables: Distance from mine Ecosystem type Mine stage Interactions: Mine stage: distance Ecosystem type: distance

This study aimed to improve our understanding of the offsite impacts of mines on understorey plant communities in the boreal forest. Specifically, our objective was to investigate the offsite impacts of mining on five common understorey plant species: *Coptis trifolia* (L.) Salisb., *Cornus canadensis* L., *Linnaea borealis* L., *Lysimachia borealis* (Raf.) U.Manns & Anderb., and *Maianthemum canadense* Desf. To do so,

we assessed the frequency and the abundance of these species surrounding six mines in Abitibi-Témiscamingue and Nord-du-Québec (Québec). Because the loss of canopy cover, soil contamination, and dust deposition generated by the mines are expected to be greater at shorter distances, we expected that both frequency and abundance of all five species would increase with distance from the mine sites.

A3.4 Materials and methods

A3.4.1 Study area

The study was conducted on six gold mines in western Québec (Figure A3.1): Casa Berardi (Hecla Québec Inc.) and Joutel (Agnico Eagle Mines Ltd.), both located in the *Picea mariana*–moss bioclimatic domain and in the Nord-du-Québec administrative region; Akasaba (Agnico Eagle Mines Ltd.), Canadian Malartic (Canadian Malartic Corp.), Lapa (Agnico Eagle Mines Ltd.), and LaRonde (Agnico Eagle Mines Ltd.), all located in the *Abies balsamea*–*Betula papyrifera* bioclimatic domain and in the Abitibi-Témiscamingue administrative region (Saucier et al. 1998). At the time of the study, three of the mines were operating (Canadian Malartic, since 2005; Casa Berardi, since 1988; LaRonde, since 1988), and the three other mines were non-operating (Akasaba, establishing; Lapa, closed since 2018; Joutel, closed since 1998). Most of the mines extracted other commodities in addition to the gold, such as copper (in Akasaba, LaRonde, and Joutel), silver (in Canadian Malartic, Casa Berardi, and LaRonde), and zinc (in LaRonde).

A3.4.2 Plant species

Plant species were selected based on the fact they are all widespread in the boreal forest understorey. All five species are common in coniferous, mixedwood, and deciduous stands, except for *Coptis trifolia*, which is mostly associated with

coniferous stands (Carleton and Maycock 1980; Howard 1993). All of the species also occasionally occur in wetlands (Taft and Solecki 1990; Sullivan 1992; Howard 1993; Pavék 1993; Gucker 2012). *Coptis trifolia* and *Lysimachia borealis* are mainly associated with low light conditions and mature forests, while *Cornus canadensis*, *Linnaea borealis*, and *M. canadense* tolerate a wide spectrum of light intensity and can be found in every successional stage, although *M. canadense* is predominantly found in late successional forests (Howard 1993; Pavék 1993; Howard and Lee 2002; Moola and Vasseur 2004; Gucker 2012). All species occurred around every mine studied.

A3.4.3 Sampling design

Six to eight transects were established perpendicular to each mine's periphery through different types of ecosystems (coniferous, mixedwood, deciduous stands, and wetlands). Along each transect, seven plots, each 3 m in radius, were positioned at different distances from the mine's edge (0, 20, 50, 100, 200, 500 and 1000m) and three 1 m² quadrats were placed randomly within each plot. In total, 123 plots were in coniferous stands, 84 were in mixed wood stands, 53 were in deciduous stands, and 49 were in wetlands (Table A3.1). The presence or absence of each species was recorded in every 3 m radius plot. A presence record in any one of those plots was considered as one event of frequency. Abundances of the five species were assessed by visually estimating their percent cover in each 1 m² quadrat. Sampling around all six mines was completed between June and August 2019.

Table A3.3 Effects of distance from mine, mine stage (non-operating compared to operating) and interactions between distance from mine and mine stage on the frequency and abundance of *Coptis trifolia*, *Cornus canadensis*, *Linnaea borealis*, *Lysimachia borealis* and *Maianthemum canadense*.

Specie		Effect of distance from the mine	Effect of mine stage (non-operating)	Effect of the interactions between distance from mine and mine stage (non-operating)
<i>Coptis trifolia</i>	Frequency ^a	1.705	-	-
	Abundance ^b	(p=0.008*) 0.110 (p=0.06)	-0.088 (p=0.463)	-
<i>Cornus canadensis</i>	Frequency	-0.364 (p=0.47)	-	2.705 (p=0.0008*)
	Abundance	0.057 (p=0.78)	1.174(p=0.004*) -0.340 (p=0.26)	0.625 (p=0.049*)
<i>Linnaea borealis</i>	Frequency	0.618 (p=0.48)	-0.514 (p=0.42)	-0.693 (p=0.51)
	Abundance	-0.113 (p=0.13)	-	-
<i>Lysimachia borealis</i>	Frequency	-1.372 (p=0.06)	-0.704 (p=0.32)	-
	Abundance	0.073 (p=0.29)	-0.072 (p=0.66)	-
<i>Maianthemum canadense</i>	Frequency	0.803	-0.989 (p=0.36)	-
	Abundance	(p=0.04*) 0.202 (p=0.50)	-0.071 (p=0.87)	-0.163 (p=0.58)

=====Note: An “*” indicates a significant effect ($p < 0.05$). A “-” means that the variable was not included in the model. (a) Frequencies were analyzed with generalised linear mixed-effects models with a binomial distribution (0 corresponding to absence of the species and 1 to its presence). (b) Abundances were log-transformed and analyzed with linear mixed-effects models.

A3.5 Statistical analysis

The frequency of each species was analyzed with a generalised linear mixed-effects models with a binomial distribution using the `glmer` function of the `lme4` package in R version 3.5.1 (Bates et al. 2015; RCoreTeam2018). Abundances were log-transformed and analyzed for each species with a linear mixed-effects model using the `lme` function of the `nlme` package in R version 3.5.1 (R Core Team 2018; Pinheiro et al. 2019). Mine sites were included as a random effect variable for both frequency

and abundances models. Fixed-effect variables were selected using the forward selection method with the following variables and their interactions: distance from mine, ecosystem type, and mine stage (operating or non-operating). Distance from mine was always included. Only variables and interactions that had a statistically significant effect or lowered the p value of the effect of distance from mine were conserved. Table A3.2 shows variables and interactions included in the models following this method. The ecosystem type variable acted as a control variable in the models since the study focused on the effects of mining. Therefore, only the effect of distance from mine, mine stage, and the interaction between distance from mine and mine stage were analyzed.

A3.6 Results and discussion

Species response to mining activity

Of the five species, *Coptis trifolia* and *M. canadense* appear to have been negatively influenced by mines. Their frequency increased with distance from the mines ($p = 0.008$ and $p = 0.040$, respectively) (see Table A3.3). This pattern was observed around both operating and nonoperating mines. The effect was more significant for *Coptis trifolia*, which was observed in less than 16% of the plots between 0m and 50 m of distance, while at distances of 500 m and 1000 m, frequencies were 30% and 27%, respectively (see Figure A3.2). For *M. canadense*, the effect seemed more significant at the edge of mine, where the species was observed in 27% of the plots, while it was observed in 50% of the plots at a distance of 1000 m. From 50 m to 500 m away from the mines, frequencies fluctuated between 34% and 39% (see Figure A3.2). Other studies have also shown that these species are strongly affected by disturbance, although they focused on the effects of clear-cut harvesting (Moola and Vasseur 2004; Roberts and Gilliam 1995; Aikens et al. 2007; Zenner and Berger 2008). *Coptis trifolia* and *M. canadense* might have been influenced by the loss of

canopy cover induced by the mines' edge effect because both species are mostly associated with mid-late to late-seral forests (Howard and Lee 2002; Moola and Vasseur 2004). Another potential cause of the decrease of *Coptis trifolia* and *M. canadense* next to mining sites might have been the stress induced by dust deposition and the subsequent soil contamination generated by the mines. Although, to our knowledge, no study has specifically investigated the effects of mines or metal contamination on these species, it is known that dust deposition can reduce plant photosynthetic rate, stomatal conductance, and transpiration rate (Li et al. 2018), and that metal contamination can extend several kilometers away from its source and modify vegetal community composition (Escarré et al. 2011). *Coptis trifolia* and *M. canadense* could have been disadvantaged by those stresses, and other species not studied in this research could have been similarly affected.

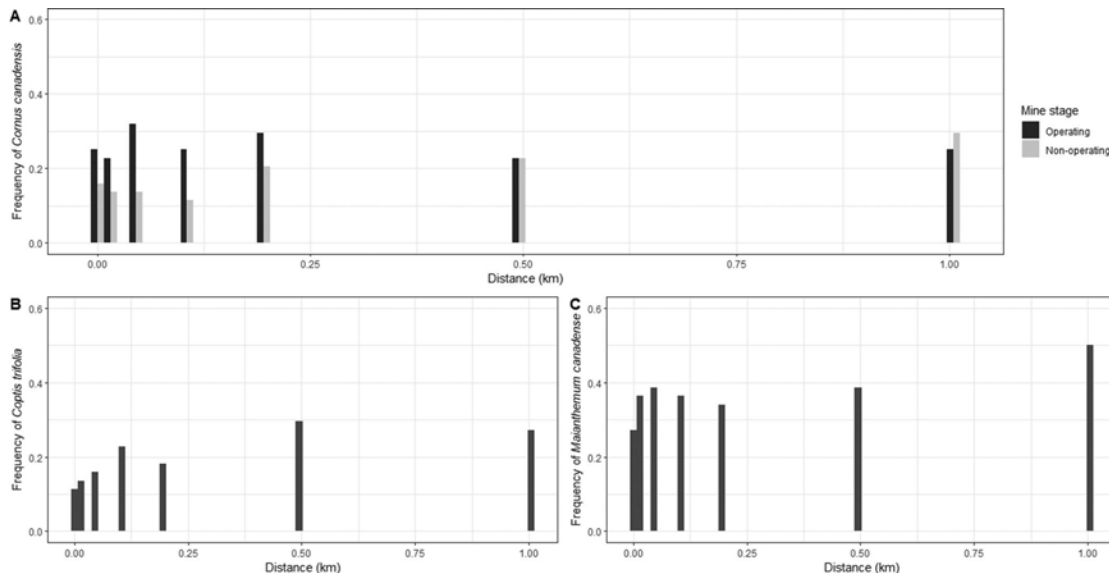


Figure A3.2. Normalized frequencies at 0 km, 0.02 km, 0.05 km, 0.1 km, 0.2 km, 0.5 km, and 1 km from mines for (A) *Cornus canadensis*, according to mine stage, (B) *Coptis trifolia*, and (C) *Maianthemum canadense*.

Conversely, *Cornus canadensis* appeared to favour mining activity. The species was significantly more frequent around operating mines than around non-operating mines ($p = 0.004$). Moreover, significant differences in the effect of distance on frequency

($p = 0.0008$) and abundance ($p = 0.049$) were observed between operating and nonoperating mines (see Table A3.3). In fact, *Cornus canadensis* was negatively influenced around non-operating mines, while it was not influenced around operating mines (see Figure A3.2). These results suggest that *Cornus canadensis* is tolerant to the stress induced by mining activity. Similarly, previous studies have shown that this species persisted following a disturbance (Moola and Vasseur 2004) and some research even showed that its abundance increased (Chávez and Macdonald 2010). *Cornus canadensis*' wide plasticity and its ability to adapt to different soil and light conditions might allow it to exploit the niche left by species that are more sensitive to environmental changes (Kumordzi et al. 2019). The reduced frequency and abundance observed next to non-operating mines compared to operating ones suggests that it may have been shaded out by vegetation regeneration after mine closure. Canopy cover loss would therefore be an important variable advantaging *Cornus canadensis*. The combination of soil contamination and shade might explain the low frequencies observed near the non-operating mines. With respect to *Linnaea borealis* and *Lysimachia canadensis*, specimen frequencies and abundances did not have a statistically significant relationship with distance from both operating and non-operating mines.

A3.7 Extent of the observed effects

Effects of mining activity on vegetation were observed from 0.1 km to 0.2 km away from the mine sites, although this differed slightly between species. Considering frequencies of *Coptis trifolia*, offsite effects of mining seemed to extend 0.2 km away from the mine sites. In fact, *Coptis trifolia* reached its highest frequencies at 0.5 km and 1.0 km from the mines, while its lowest frequencies were observed at 0, 0.02, and 0.05 km from the mines (see Figure A3.2). A similar pattern was observed with *Cornus canadensis* around non-operating mines, where frequencies were considerably lower at 0, 0.02, 0.05, and 0.1 km, and slightly lower at 0.2 km from the

mines. For *M. canadense*, no clear threshold distance was detected, although minimum frequency was observed at the edge of the mine and maximum frequency was observed at 1.0 km away. Further studies should be conducted to evaluate the extent of soil contamination, dust deposition, and loss of canopy cover around mines in boreal forests. Our study suggests the possible presence of offsite impacts of mines on plant communities. The stresses and changes commonly induced by mining activity may lead to a decrease of sensitive species and to an increase of stress tolerant species in the surrounding communities. Further studies considering enigmatic impacts should be made to develop our understanding of the environmental footprint of mining activity.

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