

Fabronia pusilla

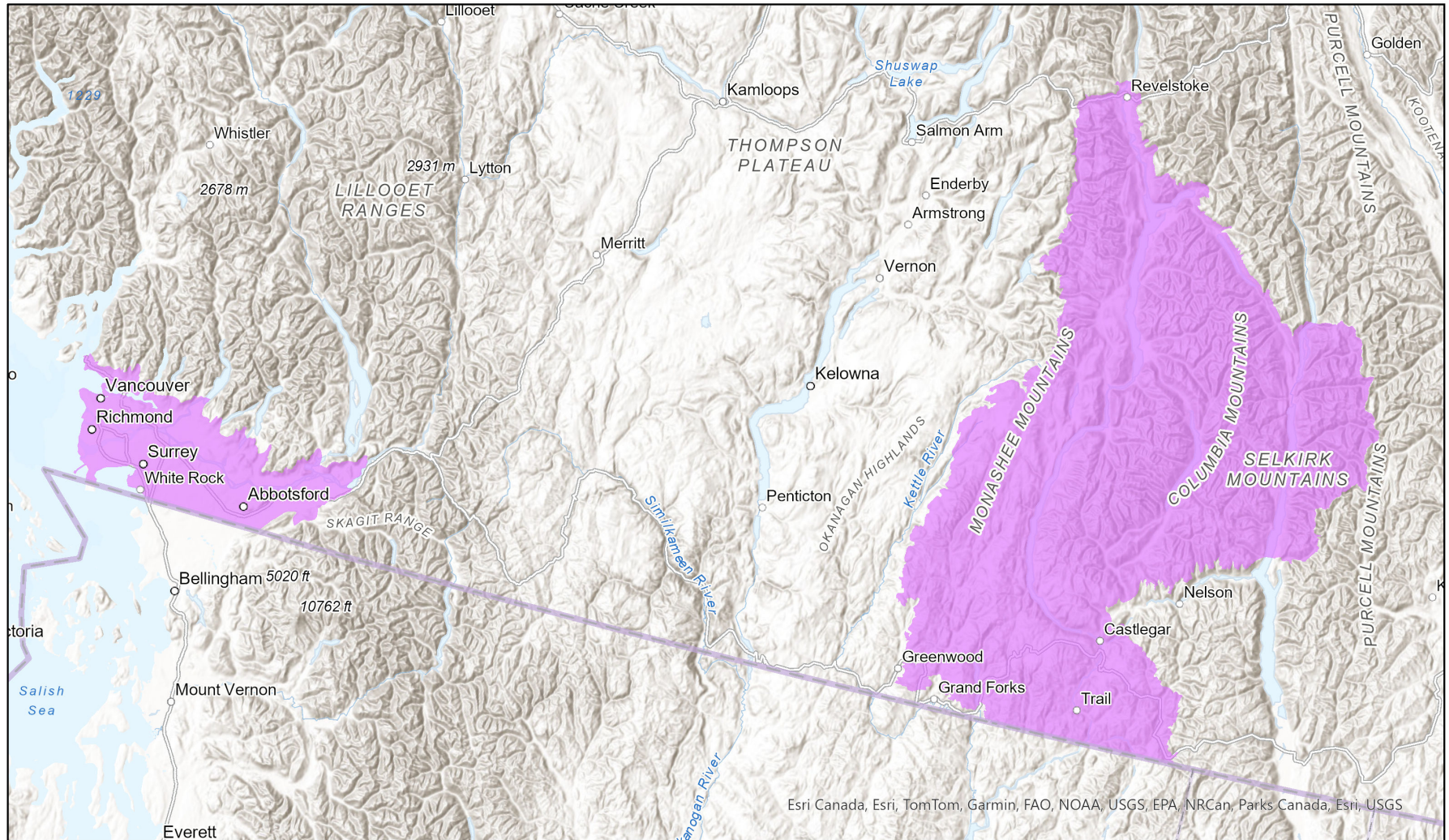


- Present
- Presence Expected
- Historical

Ecosystem-based Automated Range (EBAR)

Date Generated: March 13, 2025; Version: 1.0; Stage: Expert Reviewed (National); Scope: Canadian

Synonyms Used: None



Esri Canada, Esri, TomTom, Garmin, FAO, NOAA, USGS, EPA, NRCAN, Parks Canada, Esri, USGS

Input Records - 3 BC Non-sensitive Element Occurrences, 4 GBIF, 6 iDigBio; Expert Reviews - Ryan Batten

Map centre: 119.8838°W 49.7086°N
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EBAR is relatively coarse scale data and not intended for all applications and analysis. Please see full disclaimer in metadata.

Ecosystem-based Automated Range (EBAR) Metadata

Species

National Scientific Name:	<i>Fabronia pusilla</i> Raddi
Scientific Name Reference:	Flora of North America Editorial Committee (FNA). 2014a. Flora of North America north of Mexico. Vol. 28. Bryophytes: Mosses, part 2. Oxford Univ. Press, New York. vii + 702 pp.
National English Name:	Silver Hair Moss
National French Name:	Fabronie naine
Element National ID:	187709
Element Global ID:	122148 (go to NatureServe Explorer)
Element Code:	NBMUS2V030
Endemism Type:	N
Canadian COSEWIC Name:	
Canadian COSEWIC ID:	741

Rank/Status

Global Rank:	G4G5 (reviewed March 31, 2005)
National Rank (Canada):	N1N3 (reviewed 2024)
Subnational Ranks (Canada):	BC=S1S3
National Rank (United States):	NNR
Subnational Ranks (United States):	AZ=SNR, CA=SNR, CO=SNR, ID=SNR, MT=S1, OR=S3, WA=SNR
National Rank (Mexico):	None
Subnational Ranks (Mexico):	None
Canadian SARA Status:	Endangered/En voie de disparition (January 12, 2005)
Canadian COSEWIC Status:	Endangered (May 04, 2012)
US ESA Status:	None

Range Map

Date Generated:	March 13, 2025
Version:	1.0
Stage:	Expert Reviewed (National)
Scope:	Canadian
Metadata:	Primary Species - <i>Fabronia pusilla</i> Raddi Input Records - 3 BC Non-sensitive Element Occurrences, 4 GBIF, 6 iDigBio; Expert Reviews - Ryan Batten
Comments:	None Please see spatial data for Ecoshape-level reviewer comments.
Disclaimer:	Please review our methods document before using EBAR. EBAR range data are relatively coarse scale and appropriate for screening and education purposes, but are not intended for all types of applications and analysis. The absence of data in any geographic areas does not necessarily mean that a species is not present. An ecoshape with a presence value does not necessarily mean that a species is present throughout the entire geographic area.
Presence Definitions:	(Please see Comments above for any exceptions) Present - the species is found within the ecoshape based on species observation data, Element Occurrences, Source Features, Canadian Federal Critical Habitat, or expert opinion. Presence Expected - expert opinion the species may be present, or the ecoshape overlapped with a range estimate or a habitat suitability model. Historical - all species occurrence data within the ecoshape contains observation data greater than 40 years old or an Element Occurrence (EO) that was ranked as Extirpated or Historical (EO Rank of H, H?, X or X?), or expert opinion that the species is extirpated or historical.
Usage Type Definitions:	(Please see Comments above for any exceptions) Breeding - the species is thought to breed within the ecoshape based on eBird Breeding and Behaviour Codes or expert opinion. Possible Breeding - the species is probably or possibly breeding within the ecoshape based on eBird, BBA or jurisdiction Breeding and Behaviour Codes, or on expert opinion.
Map Projection:	North America Albers Equal Area Conic (WKID 4269)

Credits

Suggested Citation:	NatureServe Canada, 2020. Ecosystem-based Automated Range (EBAR) for <i>Fabronia pusilla</i> , Version 1.0, Expert Reviewed (National) (Canadian Scope). Ottawa, Canada. Retrieved from [insert url] on [insert date]
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Project Website:	www.natureserve.org/canada/ebars
Contact:	ebars-kba@natureserve.ca
Input References:	BC Non-sensitive Element Occurrences - British Columbia Conservation Data Centre GBIF - Global Biodiversity Information Facility GBIF - GBIF Occurrence Download https://doi.org/10.15468/dl.e3ax32 Accessed from R via rgbif (https://github.com/ropensci/rgbif) on 2024-06-21 iDigBio - Integrated Digital Biocollection
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