Ecoregion 170 – Mackenzie Mountains

Area of ecoregion km² (mi²)	Area of planning region km² (mi²)	Minimum Dynamic Reserve km² (mi²)	Number of protected area benchmarks	Number of new system-level benchmarks	Number of candidate benchmark networks (groups)
87,066 (33,616)	259,592 (100,229)	625 (241)	2	2,050	157 (11)

The identification of candidate benchmark networks is a three-step process. First, potential benchmarks are identified based on size, intactness, and hydrologic connectivity. Next, benchmarks are assembled into candidate networks that are representative of the planning region, where the number of benchmarks required to achieve representation may vary amongst planning regions (e.g., ecoregions). Finally, if multiple benchmark network options exist, to assist with the selection process, the networks are ranked using additional criteria such as benchmark properties, climate change, and focal species, as done below.

Benchmark potential of existing protected areas and regions that support the construction of benchmark networks

The planning region for ecoregion 170 is defined by the ecoregion and intersecting hydrologic units (HUC8s in Alaska and FDAs in Canada). Prior to identifying new benchmarks, existing protected areas (PAs) were clipped to the planning region and evaluated for their potential to serve as system- and subsystem-level benchmarks for the ecoregion (Figure 1). System-level benchmarks are assemblages of intact catchments that are of sufficient size to capture large-scale processes and maintain habitats vulnerable to natural disturbance (*i.e.*, Minimum Dynamic Reserve or MDR). There are two levels of benchmark intactness, I and II, which denote a minimum catchment intactness of 100% and 80%, respectively. Subsystem-level benchmarks do not meet the size and/or intactness criteria for system-level benchmarks. For this study, subsystem benchmarks are no less than 80% MDR in size. Four potential PA system-level II benchmark were identified in ecoregion 170 (Figure 1; Table 1). Two have sufficient overlap with the ecoregion (≥ 80% MDR) to be included as explicit benchmarks in the design of benchmark networks (PA_2 and PA_4). While these two PA benchmarks are sufficient to achieve representation targets, we identified new benchmarks to provide land managers with additional options. Ecoregion 170 has high benchmark potential with benchmarks identified over 99% of the ecoregion (Figure 1), which includes areas with existing protection.

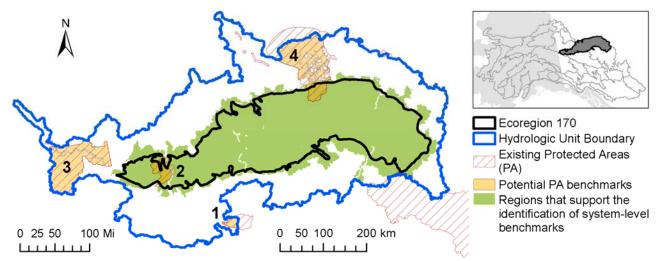


Figure 1: The planning region for ecoregion 170 is defined by the ecoregion (black outline) buffered by HUC8s and FDAs (blue outline). Potential protected area (PA) system-level II benchmarks are shown in orange; labels correspond to the PA ID in Table 1. Regions that support the identification of new system-level benchmarks (≥80% catchment-intactness) are shown in green.

Table 1: Characteristics of the four system-level II protected area (PA) benchmarks identified in Ecoregion 170 (Figure 1). To be included in the design of benchmark networks, the ecoregion portion of the PA benchmark must be ≥ 80% MDR in size.

PA ID	Area km² (mi²)	Benchmark Type	% MDR area intersecting ecoregion
PA_1	557 (215)	System-level II	0
PA_2	1,567 (605)	System-level II	211 (2.1 x MDR)
PA_3	9,555 (3,689)	System-level II	0
PA_4	9,102 (3,514)	System-level II	218 (2.2 x MDR)

Identification of candidate benchmark networks

Candidate benchmark networks for the planning region were identified based on the representation of four indicators of environmental variation: Climate Moisture Index (CMI), Gross Primary Productivity (GPP), Lake-Edge Density (LED), and Land Cover. Representation was assessed using MDR-based representation targets for indicator classes. Representation targets were derived for each class by multiplying the MDR for the ecoregion by the proportion of the class in the ecoregion. For example, if the class makes up 10% of the ecoregion, the target would be 0.1 x MDR. For a representation target to be achieved, it must be fully met within a single benchmark, except when benchmarks overlap. For ecoregion 170, we identified two PA system-level benchmark networks that achieve all MDR-based representation targets, networks N1 and N2 (Table 2, Figure 2). Only the ecoregion portion of N1 and N2 is needed to achieve representation and satisfy the size and intactness requirements of a system-level benchmark. As such, only the ecoregion portion of N1 and N2 was used to assess fundamental benchmark properties and resilience to climate change. However, the full extent of N1 and N2 was used when evaluating the representation of focal species habitat. Networks were also identified from new benchmarks only. Networks designed from two new system-level benchmarks achieve all representation targets. In total, 157 candidate benchmark networks were identified, and assigned to 11 spatial groups (Figure 2). For reporting, the set of candidate benchmark networks was reduced to the top network from each spatial group (N1-N11; Figure 2, Table 2). The top networks were selected using the same criteria and methods as described below for ranking candidate benchmark networks. For a full description of the methods, see the main report.

Table 2: Area and representation characteristics of the top network selected from each of the 11 spatial groups in Figure 2. The networks differ in area due to variable overlap of benchmarks within networks. All networks achieved MDR-based representation targets for CMI, LED, GPP, and land cover, but representation varied amongst networks when evaluated using Kolmogorov-Smirnov (KS; continuous indicators CMI, LED, and GPP) and Bray-Curtis (BC; categorical indicator land cover) dissimilarity metrics (DMs). DMs range from 0 to 1, with values closer to 0 indicating better representation.

Network	Area km² (mi²)	Upstream area	KS & BC Dissimilarity Metrics					
ID	ID Area kiii (iiii)	km² (mi²)	CMI	GPP	LED	Land Cover	Mean Dissimilarity	
N1	1,377 (532)	16 (6)	0.205	0.100	0.098	0.222	0.156	
N2	2,697 (1,041)	315 (122)	0.179	0.106	0.090	0.235	0.152	
N3	1,253 (484)	539 (208)	0.127	0.128	0.049	0.088	0.098	
N4	1,257 (485)	557 (215)	0.141	0.153	0.231	0.190	0.179	
N5	1,255 (485)	1,110 (429)	0.093	0.128	0.077	0.181	0.120	
N6	1,256 (485)	62 (24)	0.056	0.040	0.048	0.104	0.062	
N7	1,253 (484)	2,168 (837)	0.094	0.049	0.095	0.108	0.087	
N8	1,226 (473)	4,702 (1,815)	0.062	0.025	0.102	0.051	0.060	
N9	1,255 (484)	2,256 (871)	0.117	0.117	0.032	0.131	0.099	
N10	1,258 (486)	2,838 (1,096)	0.196	0.065	0.038	0.162	0.115	
N11	1,252 (483)	4,763 (1,839)	0.103	0.078	0.105	0.103	0.097	

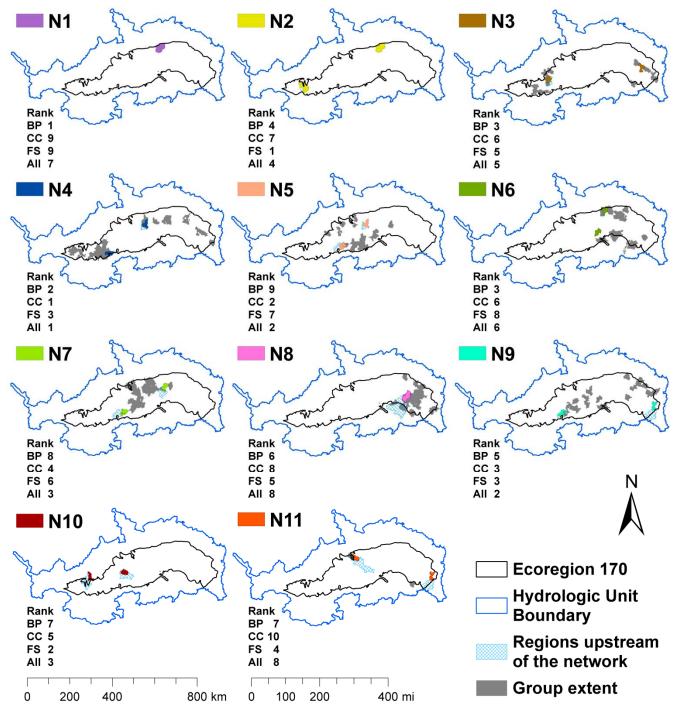


Figure 2: Benchmark networks for ecoregion 170 (N=157) were assigned to 11 spatial groups. The top network selected from each group is shown in color. N1-N2 networks are comprised of protected area system-level II benchmarks. N3-N11 benchmark networks are comprised of two new system-level benchmarks. Catchments upstream of networks are shown in blue cross-hatching. The group extent (grey) is the area covered by all networks in the group. Ranks based on fundamental benchmark properties (BP), resilience to climate change (CC), amount of focal species habitat (FS), and overall rank (ALL) are reported (Tables 3-6).

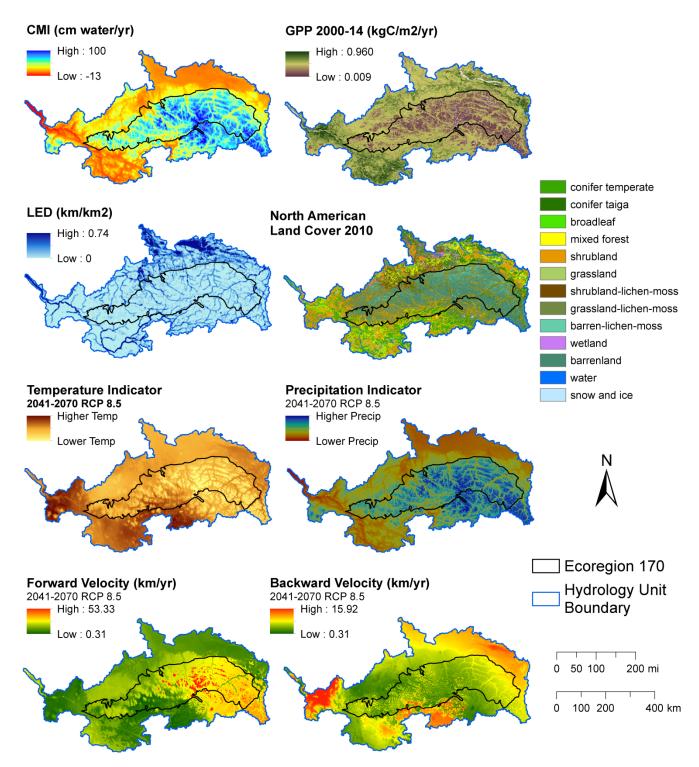


Figure 3: Distribution of the four indicators of environmental variation and four climate change datasets in the ecoregion, which include Climate Moisture Index (CMI, Wang *et al.* 2016), Gross Primary Productivity 2000-14 (GPP, BEACONS 2015), Lake-Edge Density (LED, BEACONS 2015), and North American Land Cover 2010 (CEC 2013), and the climate-projected datasets: Temperature and Precipitation Indicators, and Forward and Backward Velocity (AdaptWest Project 2015).

Benchmark network ranking

Candidate benchmark networks were ranked using three sets of criteria: fundamental benchmark properties (Table 3), resilience to climate change (Table 4), and the amount of focal species habitat (Table 5). Ranks were determined using weighted-rank methods as described in the main report. The ranks across the three sets of criteria were combined to provide an overall rank (Table 6). The results for all benchmark networks (N=157) are available at www.beaconsproject.ca/nwb.

Fundamental Benchmark Properties

In addition to being large and intact, benchmarks are designed to have high internal hydrologic connectivity (e.g., DCI), minimal vulnerability to external and internal disturbances (e.g., Upstream Area), and a compact shape (e.g., Shape Index), and selected to be representative of environmental variation. The 11 candidate networks vary with regards to these properties. While all benchmark networks satisfy MDR-based representation targets, representation varies widely when measured using dissimilarity metrics (DMs). Mean Dissimilarity ranges from 0.060 to 0.179, with higher values indicating greater dissimilarity and poorer representation (Tables 2 & 3). Eight of the 11 networks have moderate to high representation with DM < 0.2 across the four indicators of environmental variation (Table 2). Networks also differ widely in the amount of upstream area (16 – 4,763 km²), although in all cases the upstream area is largely intact (94-100%; Table 3). Networks are similar with regards to internal vulnerability, but differ with regards to shape (1.3-3.3), and internal hydrologic connectivity, with minimum lwDCI ranging from 0.148 to 0.923 (Table 3), with values closer to 1 indicating greater connectivity.

Table 3: Benchmark networks were ranked using a suite of benchmark properties. Mean Dissimilarity is the mean dissimilarity metric for the four indicators of environmental variation, and ranges from 0 to 1, with values closer to 0 indicating better representation, and ranking higher. Upstream Area is a measure of vulnerability to external influences via the stream network; lower values rank higher. Upstream AWI is the mean area-weighted intactness of catchments upstream of the network; higher values rank higher. Internal Vulnerability is the proportion of low (<80%) intact areas within the network; lower values rank higher. Maximum Shape is the shape index for the benchmark in the network that most deviates from a circle (shape index = 1); lower values rank higher. Minimum lwDCI is the mean length-weighted Dendritic Connectivity Index (0-1; low to high connectivity) for the benchmark with the lowest internal hydrologic connectivity in the network; higher values rank higher. Benchmark Properties Rank is based on the network-level mean weighted rank across all properties, shown in (). The highest ranked network within each individual benchmark property is highlighted in grey. All metrics are described in the main report.

Network ID	Mean Dissimilarity	Upstream Area km² (mi²)	Upstream AWI (%)	Internal Vulnerability	Maximum Shape	Minimum lwDCl	Benchmark Properties Rank
N1	0.156	16 (6)	100	0.000	1.3	0.173	1 (0.100)
N2	0.152	315 (122)	100	0.000	2.1	0.173	4 (0.092)
N3	0.098	539 (208)	100	0.000	2.1	0.577	3 (0.095)
N4	0.179	557 (215)	100	0.000	1.9	0.923	2 (0.099)
N5	0.120	1,110 (429)	97	0.003	3.3	0.148	9 (0.083)
N6	0.062	62 (24)	98	0.002	2.0	0.459	3 (0.095)
N7	0.087	2,168 (837)	98	0.003	2.9	0.251	8 (0.084)
N8	0.060	4,702 (1,815)	100	0.000	2.3	0.701	6 (0.088)
N9	0.099	2,256 (871)	100	0.000	2.4	0.488	5 (0.089)
N10	0.115	2,838 (1,096)	94	0.000	2.4	0.187	7 (0.085)
N11	0.097	4,763 (1,839)	100	0.000	2.4	0.488	7 (0.085)

Climate Change Resilience

Changes in patterns of environmental variation are expected under climate change. To address this, we ranked benchmark networks based on their ability to maintain representation, as measured by dissimilarity metrics (DM), using climate-projected multivariate indicators of climatic conditions (2041-2070, RCP 8.5¹; Figure 3), which we refer to as Temperature and Precipitation Indicators given the explanatory power of temperature and precipitation variables in each indicator, respectively. Five of the eleven networks maintain representation with moderate values for both indicators (DM < 0.2; Table 4). With the exception of N1 and N2, the remaining networks have one indicator with moderate to good representation. To address the vulnerability of benchmark networks and their support of biodiversity under climate change, we evaluated the ability of species to persist within and colonize benchmark networks, using forward and backward climate velocity (2041-2070, RCP 8.5¹; Figure 3), respectively. Higher velocities indicate greater vulnerability to species loss. Across networks, mean forward and backward velocities range from 4.3 to 12.1 km/yr and 2.3 to 4.3 km/yr, respectively (Table 4). Lower forward velocities indicate higher refugia potential for species, whereas lower backward velocities indicate higher colonization potential. In all networks, backward velocity is lower than forward velocity, which suggests that the networks favour colonization potential. However, while most networks have backward velocities (bv) similar to the ecoregion-level mean bv of 2.6 km/yr, five networks favour areas with higher refugia potential with forward velocities (fv) 3-4 km/yr lower than the ecoregion-level mean fv of 8.2 km/yr. The output is available such that users have the flexibility to select a subset of climate datasets to rank networks.

Table 4: Benchmark networks were ranked based on their capacity to represent future climatic conditions (temperature and precipitation indicators) and vulnerability to changing climatic conditions (forward and backward velocity). **Temperature and Precipitation Indicators** were assessed using the Kolmogorov-Smirnov (KS) dissimilarity metric, which ranges from 0 to 1; lower values indicate better representation, and rank higher. **Climate Velocities** are calculated as the geometric mean across all benchmarks from each network; lower values rank higher. **Climate Change Rank** is based on the network-level mean weighted rank across the four climatic measures, shown in (). The highest ranked network within each indicator/velocity is highlighted in grey. For dataset details, see the main report.

Network -	KS Dissimila	rity Metrics	Mean Forward	Mean Backward	Climate Change	
ID	Temperature Indicator	'		Climate Velocity km/yr (mi/yr)	Rank	
N1	0.417	0.182	10.9 (6.7)	2.8 (1.7)	9 (0.067)	
N2	0.180	0.206	8.5 (5.3)	2.9 (1.8)	7 (0.074)	
N3	0.114	0.150	8.0 (5.0)	2.9 (1.8)	6 (0.076)	
N4	0.156	0.102	4.3 (2.6)	2.7 (1.7)	1 (0.126)	
N5	0.189	0.092	4.3 (2.6)	2.9 (1.8)	2 (0.123)	
N6	0.331	0.040	12.1 (7.5)	2.3 (1.4)	6 (0.076)	
N7	0.212	0.069	5.0 (3.1)	2.8 (1.7)	4 (0.107)	
N8	0.130	0.054	11.7 (7.2)	3.1 (1.9)	8 (0.068)	
N9	0.228	0.140	4.4 (2.7)	4.0 (2.5)	3 (0.109)	
N10	0.201	0.179	5.6 (3.5)	2.3 (1.4)	5 (0.102)	
N11	0.155	0.088	10.6 (6.6)	3.3 (2.1)	10 (0.066)	

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¹ All climate-projected datasets used to rank networks were for the period 2041-2070 and were created using RCP 8.5, the Representative Concentration Pathway with the highest greenhouse gas emissions from IPCC (2014). Additional rankings based on 2011-2040 and 2071-2100 and RCP 4.5 are available at www.beaconsproject.ca/nwb.

Focal Species

An extensive review of management plans for the NWBLCC did not reveal specific conservation targets for focal species (see focal species report). As such, the objective for all focal species was to maximize the protection of current and future habitat when ranking benchmark networks. For some species, there are multiple datasets (N=3-15). When multiple datasets were used, the network rank for the species (or guild) is a mean of the weighted ranks generated for each dataset. Within each network, ranks vary across species. For this evaluation, Networks N1 & N2 include the full extent of the PA benchmarks PA_1 and PA_2 (Figure 1), not just the ecoregion portion (Figure 2). For example, the ranks of individual species within network N1 ranges from 4 to 10 (Table 5a). The high focal species rank of the protected area benchmark network (N2) may in part be due to network size, which is 8.5 times larger than networks N3-11). To remove the potential influence of the larger networks N1 and N2 on the weighted rank values, the ranking process with focal species was repeated with N1 and N2 excluded (Table 5b). The relative ranks of networks N3-N11 may differ from Table 5a, as differences masked by the larger networks are revealed. The output is available such that users have the flexibility to select a subset of species datasets to rank networks.

Table 5a: Benchmark networks were ranked based on the amount of focal species habitat they capture. Data were not available for Beaver, Broad Whitefish, Chinook Salmon and Chum Salmon. Values in () are weighted ranks. When multiple datasets were used for a species (e.g., Rusty Blackbird N=3), networks were ranked using the mean of weighted ranks from across datasets. The highest ranked network within each individual species is highlighted in grey. **Focal Species Rank** is based on the network-level mean weighted rank across all species. For further details on the datasets and methods see main and focal species reports. Additional information on each focal species and their datasets is available at www.beaconsproject.ca/nwb.

		Rank (mean weighted rank)						
Network ID	Area km² (mi²)	Caribou (N=1)	Dall Sheep (N=1)	Moose (N=1)	Old-Forest Birds (N=15) ¹	Rusty Blackbird (N=3)	Waterfowl (N=3) ²	Focal Species Rank
N1	9,102 (3,514)	9 (0.068)	9 (0.076)	9 (0.079)	10 (0.082)	3 (0.096)	9 (0.083)	9 (0.080)
N2	10,669 (4,119)	1 (0.112)	1 (0.124)	4 (0.090)	8 (0.087)	1 (0.103)	1 (0.105)	1 (0.103)
N3	1,253 (484)	4 (0.094)	5 (0.088)	3 (0.094)	7 (0.088)	7 (0.086)	6 (0.089)	5 (0.089)
N4	1,257 (485)	7 (0.076)	3 (0.096)	1 (0.120)	5 (0.093)	9 (0.084)	3 (0.093)	3 (0.093)
N5	1,255 (485)	8 (0.069)	4 (0.092)	7 (0.084)	2 (0.098)	10 (0.083)	2 (0.095)	7 (0.086)
N6	1,256 (485)	9 (0.068)	7 (0.083)	6 (0.085)	11 (0.079)	4 (0.094)	7 (0.085)	8 (0.082)
N7	1,253 (484)	5 (0.092)	5 (0.088)	8 (0.080)	4 (0.095)	11 (0.081)	4 (0.092)	6 (0.088)
N8	1,226 (473)	3 (0.108)	10 (0.075)	3 (0.094)	9 (0.084)	6 (0.091)	8 (0.084)	5 (0.089)
N9	1,255 (484)	6 (0.089)	8 (0.082)	2 (0.108)	1 (0.101)	8 (0.085)	3 (0.093)	3 (0.093)
N10	1,258 (486)	2 (0.109)	2 (0.105)	10 (0.072)	3 (0.096)	5 (0.093)	5 (0.091)	2 (0.094)
N11	1,252 (483)	2 (0.109)	6 (0.086)	5 (0.087)	6 (0.092)	2 (0.098)	7 (0.085)	4 (0.092)

¹ Guild composed of Boreal Chickadee, Brown Creeper, Pine Grosbeak, Swainson's Thrush, and White-Winged Crossbill. In this case, the weighted rank shown in parenthesis is the mean across 15 datasets (3 per species).

² Guild composed of Lesser Scaup (1 dataset), White-Winged Scoter (1 dataset), and Trumpeter Swan (1 dataset). The weighted rank shown in parenthesis is the mean across all species.

Table 6b: Benchmark networks were ranked based on the amount of focal species habitat they capture. Data were not available for Beaver, Broad Whitefish, and Chinook and Chum Salmon. Values in () are weighted ranks. When multiple datasets were used for a species (*e.g.*, Rusty Blackbird N=3), networks were ranked using the mean of weighted ranks from across datasets. The highest ranked network within each individual species is highlighted in grey. **Focal Species Rank** is a network-level mean weighted rank across all species. For further details on the datasets and methods see main and focal species reports. Additional information on each focal species and their datasets is available at www.beaconsproject.ca/nwb.

			Rank (mean	ank (mean weighted rank)					
Network ID	Caribou (N=1)	Dall Sheep (N=1)	Moose (N=1)	Old-Forest Birds (N=15) ¹	Rusty Blackbird (N=3)	Waterfowl (N=3) ²	Focal Species Rank		
N3	3 (0.115)	5 (0.109)	3 (0.114)	7 (0.106)	6 (0.106)	6 (0.105)	4 (0.109)		
N4	6 (0.092)	2 (0.124)	1 (0.144)	5 (0.112)	7 (0.104)	4 (0.115)	2 (0.115)		
N5	7 (0.083)	3 (0.117)	6 (0.101)	2 (0.118)	7 (0.104)	1 (0.126)	5 (0.108)		
N6	8 (0.082)	7 (0.100)	5 (0.103)	9 (0.095)	3 (0.118)	7 (0.101)	7 (0.099)		
N7	4 (0.112)	4 (0.110)	7 (0.097)	4 (0.114)	8 (0.098)	2 (0.121)	5 (0.108)		
N8	2 (0.133)	9 (0.086)	3 (0.114)	8 (0.101)	4 (0.111)	8 (0.100)	6 (0.107)		
N9	5 (0.108)	8 (0.098)	2 (0.130)	1 (0.122)	5 (0.107)	3 (0.120)	3 (0.114)		
N10	1 (0.135)	1 (0.143)	8 (0.087)	3 (0.116)	2 (0.120)	5 (0.107)	1 (0.118)		
N11	1 (0.135)	6 (0.107)	4 (0.105)	6 (0.111)	1 (0.126)	8 (0.100)	3 (0.114)		

¹ Guild composed of Boreal Chickadee, Brown Creeper, Pine Grosbeak, Swainson's Thrush, and White-Winged Crossbill. In this case, the weighted rank shown in parenthesis is the mean across 15 datasets (3 per species).

² Guild composed of Lesser Scaup (1 dataset), White-Winged Scoter (1 dataset), and Trumpeter Swan (1 dataset). The weighted rank shown in parenthesis is the mean across all species.

Overall Rank

Candidate benchmark networks were assigned an overall rank based on fundamental benchmark properties (Table 3), resilience to climate change (Table 4) and the amount of focal species habitat (Table 5a). Attributes were given equal weighting. However, users may wish to prioritize some attributes over others. The results are available in a format that gives users the flexibility to modify and re-rank networks. Additional attributes can also be considered. For example, if the conservation priority is the protection of focal species habitat within the ecoregion, networks with greater overlap with the ecoregion are more likely to achieve this objective. Greater overlap with protected areas may facilitate implementation, given existing protection. Overlap with the ecoregion and existing protected areas ranges from 58-100% and 0-100% across benchmark networks, respectively (Table 6). While benchmark networks that most overlap with the ecoregion may best reflect the environmental variation of the ecoregion, networks that extend beyond the ecoregion boundary may contribute to the benchmark networks of neighbouring ecoregions, leading to greater efficiency in the design of a protected areas network for the NWBLCC planning region.

Table 7: Overall Rank is based on the network-level mean weighted rank for fundamental benchmark properties (Table 3), climate change (Table 4) and focal species (Table 5a). Values in () are weighted ranks. **Overlap with ecoregion** and **Overlap with existing PAs** with high levels of protection may be used as additional ranking criteria.

Network ID	Overlap with Ecoregion	Overlap with Existing PAs	Mean Dissimilarity	Benchmark Properties Rank	Climate Change Rank	Focal Species Rank ¹	Overall Rank
N1	100.0%	100.0%	0.156	1 (0.100)	9 (0.067)	9 (0.080)	7 (0.082)
N2	100.0%	100.0%	0.152	4 (0.092)	7 (0.074)	1 (0.103)	4 (0.089)
N3	100.0%	0.0%	0.098	3 (0.095)	6 (0.076)	5 (0.089)	5 (0.086)
N4	88.0%	0.0%	0.179	2 (0.099)	1 (0.126)	3 (0.093)	1 (0.106)
N5	100.0%	0.0%	0.120	9 (0.083)	2 (0.123)	7 (0.086)	2 (0.097)
N6	95.1%	49.6%	0.062	3 (0.095)	6 (0.076)	8 (0.082)	6 (0.084)
N7	100.0%	0.0%	0.087	8 (0.084)	4 (0.107)	6 (0.088)	3 (0.093)
N8	100.0%	0.0%	0.060	6 (0.088)	8 (0.068)	5 (0.089)	8 (0.081)
N9	97.0%	0.0%	0.099	5 (0.089)	3 (0.109)	3 (0.093)	2 (0.097)
N10	58.4%	2.1%	0.115	7 (0.085)	5 (0.102)	2 (0.094)	3 (0.093)
N11	90.2%	0.0%	0.097	7 (0.085)	10 (0.066)	4 (0.092)	8 (0.081)

¹ Focal Species Rank is from Table 5a.