

Ecoregion 31 – Yukon-Old Crow Basin

Area of ecoregion km ² (mi ²)	Area of planning region km ² (mi ²)	Minimum Dynamic Reserve km ² (mi ²)	Number of protected area benchmarks	Number of new system-level benchmarks	Number of candidate benchmark networks (groups)
72,696 (28,068)	231,128 (89,238)	7,161 (2,765)	2	649	61 (9)

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 fundamental 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 31 is defined by the ecoregion and intersecting hydrologic units (HUC8 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 benchmarks were identified in ecoregion 31 (Figure 1; Table 1). However, only two (PA_2 and PA_4) had sufficient overlap with the ecoregion (80% MDR) to be included in the design of benchmark networks. While these two PA benchmarks together achieve representation targets, we identified new benchmarks to provide land managers with additional options. Ecoregion 31 has high benchmark potential with benchmarks identified over 95% of the ecoregion (Figure 1), which includes areas with existing protection.

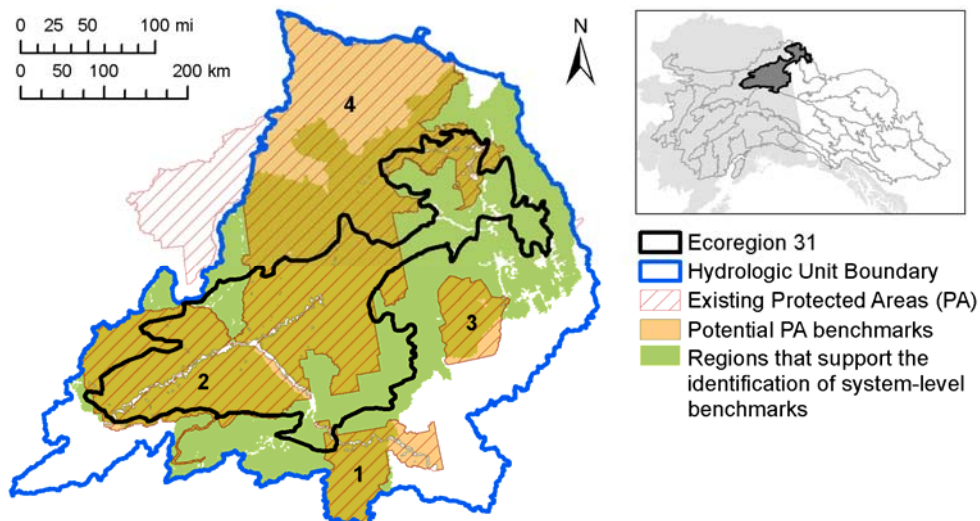


Figure 1: The planning region for ecoregion 31 is defined by the ecoregion (black outline) buffered by HUC8s and FDAs (blue outline). Potential protected area (PA) system-level II benchmarks are shown in orange; labels correspond to the PA ID in Table 1. Regions that support the identification of new 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 31 (Figure 1). To be included in the design of benchmark networks, the ecoregion portion of the PA benchmark must be $\geq 80\%$ MDR in size.

PA ID	Area km ² (mi ²)	Benchmark Type	% MDR area intersecting ecoregion
PA_1	9,687 (3,740)	System-level II	5
PA_2	26,670 (10,297)	System-level II	273 (2.7 x MDR)
PA_3	6,481 (2,502)	System-level II	0
PA_4	76,946 (29,709)	System-level II	373 (3.7 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 $0.1 \times \text{MDR}$. For ecoregion 31, existing protected area benchmarks (PA_2 and PA_4) achieve the representation targets for all indicator classes. Only the ecoregion portions of PA_2 and PA_4 are needed to achieve representation and satisfy size and intactness requirements of system-level benchmarks (N1; Figure 2). As such, only the ecoregion portions of PA_2 and PA_4 were used to assess fundamental benchmark properties and resilience to climate change. However, the full extent of PA_2 and PA_4 were used when evaluating the representation of focal species habitat. Networks designed from three new system-level benchmarks achieve all targets. In total, 61 benchmark networks were identified and assigned to nine spatial groups. For reporting, the set of candidate benchmark networks was reduced to the top network from each spatial group (N1-N9; 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 networks selected from the nine 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 ID	Area km ² (mi ²)	Upstream area km ² (mi ²)	KS & BC Dissimilarity Metrics				
			CMI	GPP	LED	Land Cover	Mean Dissimilarity
N1	46,215 (17,844)	172,507 (66,605)	0.143	0.059	0.015	0.072	0.072
N2	21,421 (8,270)	22,989 (8,876)	0.173	0.065	0.147	0.139	0.131
N3	21,440 (8,278)	46,628 (18,003)	0.224	0.061	0.291	0.145	0.180
N4	21,509 (8,304)	39,362 (15,198)	0.163	0.081	0.173	0.105	0.130
N5	20,907 (8,072)	107,327 (41,439)	0.181	0.113	0.159	0.114	0.142
N6	20,847 (8,049)	39,475 (15,241)	0.128	0.133	0.046	0.118	0.106
N7	21,273 (8,213)	42,504 (16,411)	0.151	0.039	0.146	0.126	0.115
N8	21,485 (8,295)	121,871 (47,054)	0.061	0.023	0.108	0.102	0.074
N9	21,510 (8,305)	130,481 (50,379)	0.367	0.046	0.192	0.104	0.177

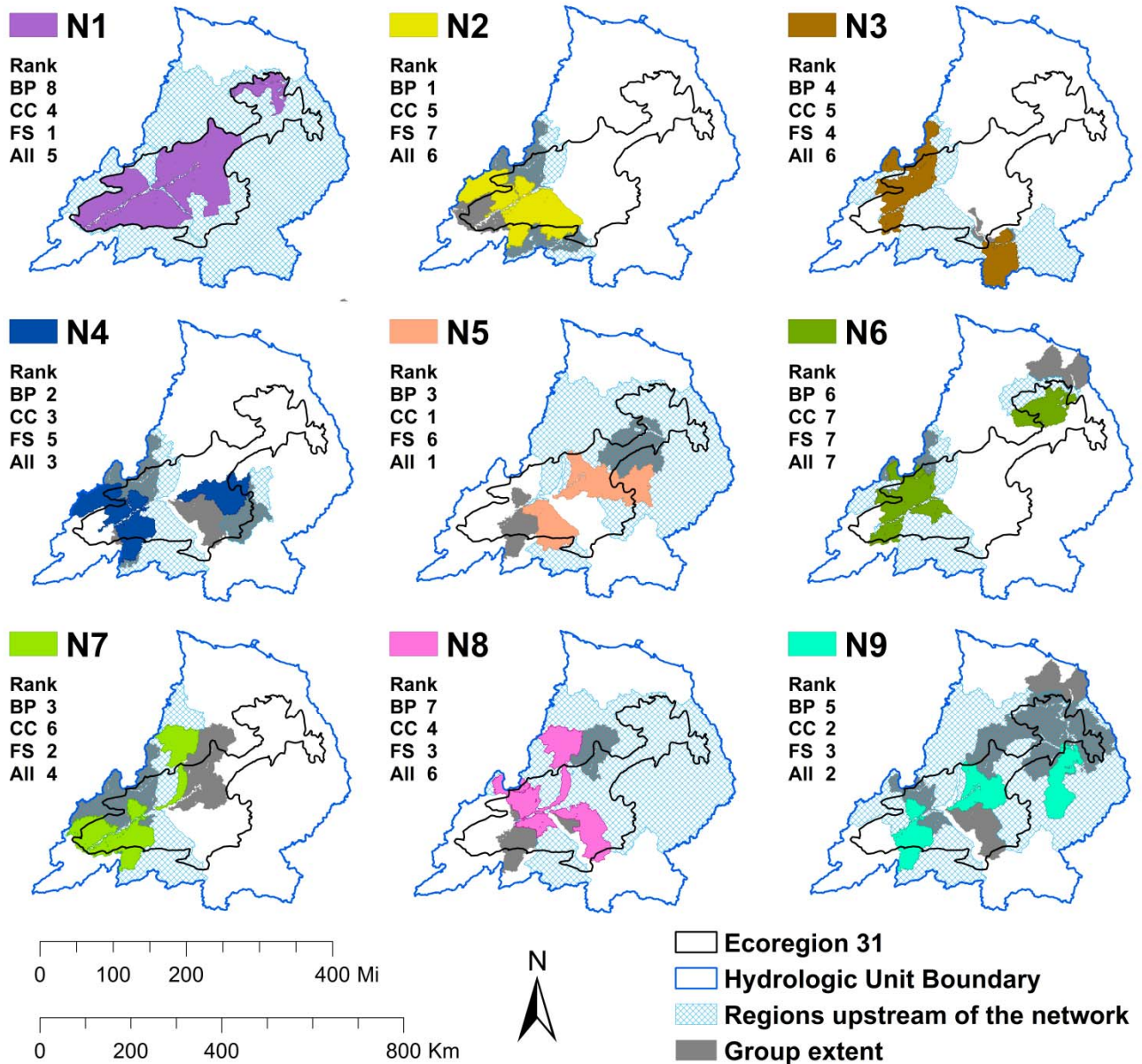


Figure 2: Benchmark networks for ecoregion 31 (N=61) were assigned to nine spatial groups. The top network selected from each group is shown in color. N1 is comprised of protected area system-level II benchmarks PA_2 and PA_4 identified in Figure 1; only the ecoregion portions of PA_2 and PA_4 are required to meet the size, intactness, and representation requirements of a system-level benchmark network, as shown here. N2-N9 are benchmark networks comprised of three new system-level benchmarks. N2-N8 have two overlapping 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).

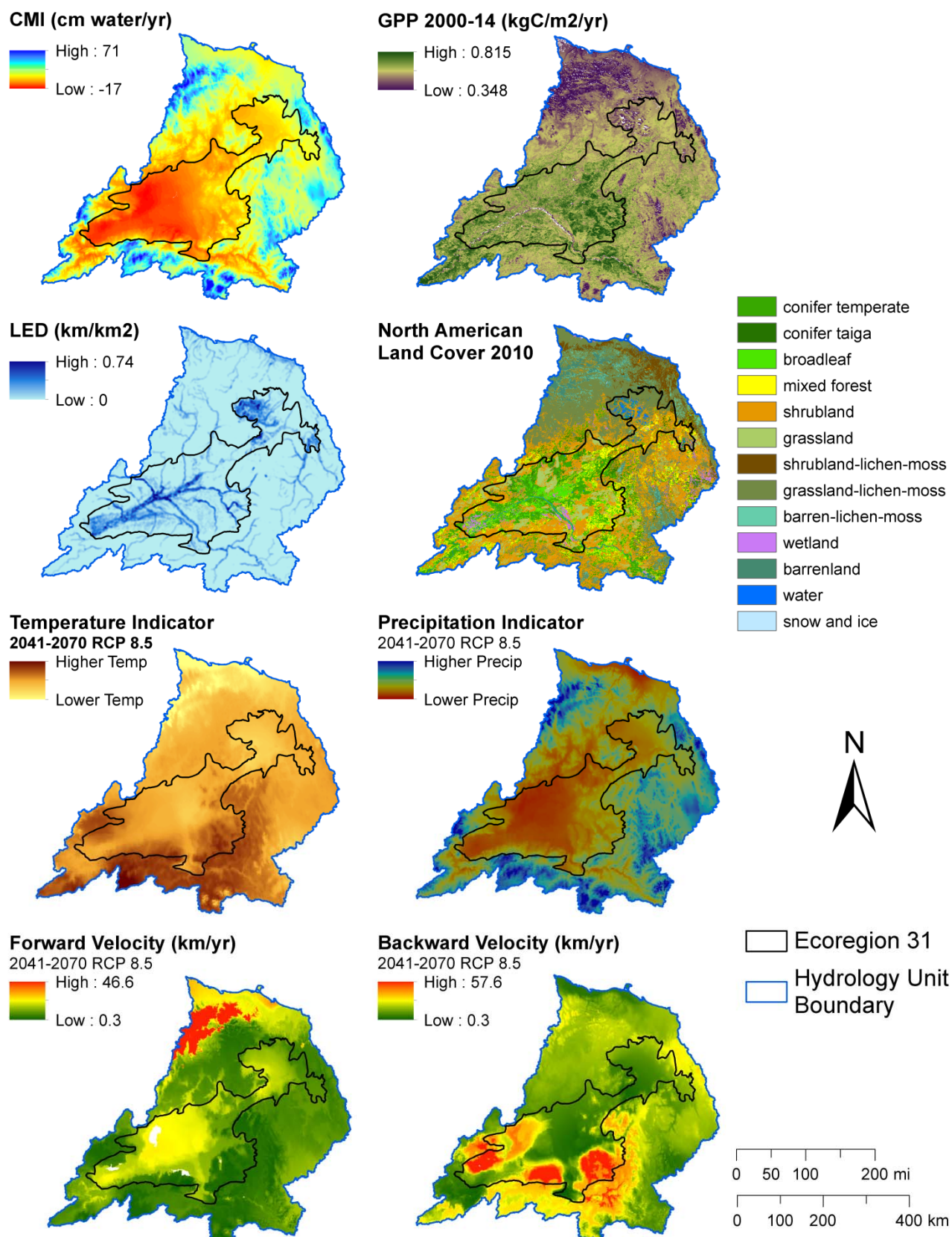


Figure 3: Distribution of the four indicators of environmental variation and four climate change datasets in the ecoregion, which include Climate Moisture Index (CMI, [Wang et al. 2016](#)), Gross Primary Productivity 2000-14 (GPP, [BEACONS 2015](#)), Lake-Edge Density (LED, [BEACONS 2015](#)), and North 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=61) 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 nine candidate networks vary with regards to these properties. While all benchmark networks satisfy MDR-based representation targets, representation varies when measured using dissimilarity metrics (DMs). Mean Dissimilarity ranges from 0.072 to 0.177, with higher values indicating greater dissimilarity and poorer representation (Tables 2 & 3). Seven of the nine networks have moderate to high representation across the four indicators of environmental variation with DM < 0.2 (Table 2). Networks differ widely in the amount of upstream area (22,989 – 172,507 km²), although in all cases the upstream area is largely intact (97-99%; Table 3). All networks are similar with regards to shape (except N1) and internal vulnerability, as well as internal hydrologic connectivity with minimum lwDCI ranging from 0.101 to 0.251 (Table 3), with values closer to 1 indicating greater connectivity.

Table 3: Benchmark networks were ranked using a suite of fundamental 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, which is shown in (). The highest ranked network within each individual benchmark property is highlighted in grey. All metrics are described in the [main report](#). **Overlapping benchmarks within a network were treated as a single benchmark which calculating Shape and lwDCI.**

Network ID	Mean Dissimilarity	Upstream Area km ² (mi ²)	Upstream AWI (%)	Internal Vulnerability	Maximum Shape	Minimum lwDCI	Benchmark Properties Rank
N1	0.072	172,507 (66,605)	97	0.021	5.0	0.101	8 (0.097)
N2	0.131	22,989 (8,876)	98	0.005	3.4	0.251	1 (0.118)
N3	0.180	46,628 (18,003)	98	0.014	3.8	0.145	4 (0.111)
N4	0.130	39,362 (15,198)	99	0.005	3.4	0.251	2 (0.117)
N5	0.142	107,327 (41,439)	97	0.009	3.2	0.138	3 (0.114)
N6	0.106	39,475 (15,241)	98	0.015	4.3	0.128	6 (0.108)
N7	0.115	42,504 (16,411)	98	0.016	3.6	0.164	3 (0.114)
N8	0.074	121,871 (47,054)	97	0.004	4.3	0.244	7 (0.105)
N9	0.177	130,481 (50,379)	97	0.012	3.4	0.138	5 (0.110)

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 two climate-projected multivariate indicators of climatic conditions (2041-2070, RCP 8.5)¹, which we refer to as Temperature and Precipitation Indicators given the explanatory power of temperature and precipitation variables in each indicator, respectively. Three networks (N1, N6, N8) maintain representation with moderate to high values for both indicators (DM < 0.2; Table 4), while two additional networks (N7, N9) maintain representation of one indicator. 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 8.5)¹, respectively. Higher velocities indicate greater vulnerability to species loss. Across networks, mean forward and backward velocities range from 3.8 to 7.0 km/yr and 3.4 to 6.6 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 (except N2-N4), mean backward velocity (bv) is lower than mean forward velocity (fv). All networks, except N9, have mean backward velocities higher than the ecoregion-level mean of 3.5 km/yr, whereas six of nine networks have mean forward velocities less than the ecoregion-level mean of 5.7 km/yr. This suggests that while most networks identified for ecoregion 15 favour colonization potential over refugia potential (bv < fv), the networks tend to intersect areas of the ecoregion with greater refugia potential. 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 ID	KS Dissimilarity Metric		Mean Forward Climate Velocity km/yr (mi/yr)	Mean Backward Climate Velocity km/yr (mi/yr)	Climate Change Rank
	Temperature Indicator	Precipitation Indicator			
N1	0.054	0.128	6.3 (3.9)	3.7 (2.3)	4 (0.108)
N2	0.440	0.234	3.8 (2.4)	6.6 (4.1)	5 (0.104)
N3	0.294	0.255	4.1 (2.5)	5.8 (3.6)	5 (0.104)
N4	0.419	0.290	3.8 (2.3)	4.9 (3.1)	3 (0.116)
N5	0.231	0.204	4.0 (2.5)	3.7 (2.3)	1 (0.132)
N6	0.103	0.046	7.0 (4.4)	4.1 (2.5)	7 (0.098)
N7	0.205	0.170	5.3 (3.3)	4.5 (2.8)	6 (0.102)
N8	0.091	0.123	5.8 (3.6)	3.9 (2.4)	4 (0.108)
N9	0.156	0.371	4.9 (3.0)	3.4 (2.1)	2 (0.123)

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 NWB LCC 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. This ecoregion crosses the Yukon-Alaska border, and there are transboundary species datasets as well as datasets restricted to Alaska and Yukon. To prevent bias in network rankings given inconsistent spatial extents for datasets, all networks were ranked using transboundary species datasets only (Table 5a, 5b), and networks restricted to Alaska were ranked using transboundary and Alaska datasets (Table 5c). There are no networks restricted to the Yukon. For some species, there are multiple datasets (N=2-15). When multiple datasets were used, the rank of a network for each species (or guild) is a mean of the weighted ranks generated for each dataset. Within each network, ranks vary across species. **For this evaluation, N1 includes the full extent of benchmarks PA_2 and PA_4 (Figure 1), not just the ecoregion portion (Figure 2).** The top ranks of the protected area benchmark network (N1) are likely due to network size, which is 4.8 to 4.9 times larger than the other networks (N2-9). To remove the influence of the large network N1 on the weighted rank values, the ranking process with focal species was repeated with N1 excluded (Table 5b). The relative ranks of networks N2-N9 may differ from Table 5a, as differences masked by the large network N1 are revealed. The output is available such that users have the flexibility to select a subset of species datasets to rank networks.

Table 5a: All benchmark networks were ranked based on the amount of focal species habitat captured. **Only transboundary datasets were used.** Data were not available for Broad Whitefish, and only Alaska and/or Yukon datasets were available for Beaver, Salmon species, and Waterfowl. 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, shown in (). For further details on 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.

Network ID	Area km ² (mi ²)	Rank (mean weighted rank)					Focal Species Rank
		Caribou (N=1)	Dall Sheep (N=1)	Moose (N=1)	Old-Forest Birds (N=15) ¹	Rusty Blackbird (N=3)	
N1	103,616 (40,006) ²	1 (0.150)	5 (0.103)	1 (0.147)	7 (0.104)	2 (0.117)	1 (0.124)
N2	21,421 (8,270)	9 (0.091)	7 (0.092)	2 (0.120)	4 (0.111)	7 (0.104)	7 (0.103)
N3	21,440 (8,278)	3 (0.114)	4 (0.112)	7 (0.100)	2 (0.117)	5 (0.108)	4 (0.110)
N4	21,509 (8,304)	6 (0.105)	7 (0.092)	3 (0.114)	3 (0.114)	4 (0.109)	5 (0.106)
N5	20,907 (8,072)	4 (0.113)	8 (0.090)	4 (0.109)	5 (0.110)	6 (0.106)	6 (0.105)
N6	20,847 (8,049)	5 (0.106)	3 (0.118)	8 (0.089)	8 (0.097)	6 (0.106)	7 (0.103)
N7	21,273 (8,213)	8 (0.098)	2 (0.146)	6 (0.104)	3 (0.114)	3 (0.114)	2 (0.115)
N8	21,485 (8,295)	7 (0.103)	1 (0.149)	6 (0.104)	6 (0.105)	5 (0.108)	3 (0.113)
N9	21,510 (8,305)	2 (0.118)	6 (0.095)	5 (0.108)	1 (0.121)	1 (0.124)	3 (0.113)

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 The area reported for N1 differs from Table 2 because for the focal species analysis N1 includes the full extent of PA_2 and PA_4 (Figure 1), not just the ecoregion portion (Figure 2).

Table 5b. Benchmark networks designed from **new system-level benchmarks only** were ranked based on the amount of focal species habitat they capture. **Only transboundary datasets were used.** Data were not available for Broad Whitefish, and only Alaska and/or Yukon datasets were available for Beaver, Salmon species and Waterfowl. 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 the network-level mean weighted rank across all species, shown in (). 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.

Network ID	Rank (mean weighted rank)					Focal Species Rank
	Caribou (N=1)	Dall Sheep (N=1)	Moose (N=1)	Old-Forest Birds (N=15) ¹	Rusty Blackbird (N=3)	
N2	8 (0.092)	7 (0.102)	1 (0.153)	4 (0.125)	7 (0.118)	7 (0.118)
N3	2 (0.142)	4 (0.125)	5 (0.113)	2 (0.132)	4 (0.122)	4 (0.126)
N4	5 (0.122)	7 (0.102)	2 (0.141)	3 (0.128)	3 (0.123)	5 (0.123)
N5	3 (0.140)	8 (0.101)	3 (0.129)	5 (0.124)	6 (0.120)	6 (0.122)
N6	4 (0.124)	3 (0.131)	6 (0.092)	7 (0.107)	5 (0.121)	8 (0.115)
N7	7 (0.106)	2 (0.162)	4 (0.120)	3 (0.128)	2 (0.129)	2 (0.129)
N8	6 (0.118)	1 (0.166)	4 (0.120)	6 (0.118)	4 (0.122)	3 (0.128)
N9	1 (0.152)	6 (0.106)	3 (0.129)	1 (0.135)	1 (0.140)	1 (0.132)

Table 5c. Benchmark networks restricted to **Alaska** and designed from **new system-level benchmarks only** were ranked based on the amount of focal species habitat they capture. **Only Alaska datasets were used.** Data were not available for Broad Whitefish. Values in () are weighted ranks. When multiple datasets were used for a species (e.g., Caribou N=5), 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 the network-level mean weighted rank across all species, shown in (). 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.

Network ID	Rank (mean weighted rank)									Focal Species Rank
	Beaver (N=1)	Caribou (N=5)	Chinook Salmon (N=3)	Chum Salmon (N=2)	Dall Sheep (N=2)	Moose (N=2)	Old-Growth Forest Birds (N=15) ¹	Rusty Blackbird (N=3)	Waterfowl (N=3) ²	
N2	1 (0.256)	5 (0.169)	2 (0.211)	2 (0.200)	4 (0.174)	1 (0.265)	4 (0.191)	4 (0.192)	3 (0.202)	1 (0.206)
N3	4 (0.165)	1 (0.227)	3 (0.193)	5 (0.179)	1 (0.232)	3 (0.174)	1 (0.208)	2 (0.205)	5 (0.173)	4 (0.195)
N4	5 (0.155)	4 (0.180)	1 (0.238)	1 (0.231)	5 (0.164)	2 (0.218)	2 (0.206)	3 (0.203)	4 (0.183)	3 (0.197)
N7	3 (0.188)	3 (0.208)	4 (0.189)	4 (0.191)	2 (0.218)	5 (0.169)	3 (0.204)	1 (0.207)	1 (0.234)	2 (0.200)
N8	2 (0.233)	2 (0.214)	5 (0.166)	3 (0.197)	3 (0.210)	4 (0.170)	5 (0.188)	5 (0.191)	2 (0.205)	3 (0.197)

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 42-100% and 64-100% 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 6: 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%	0.072	8 (0.097)	4 (0.108)	1 (0.124)	5 (0.109)
N2	62.9%	84.0%	0.131	1 (0.118)	5 (0.104)	7 (0.103)	6 (0.108)
N3	42.8%	69.5%	0.180	4 (0.111)	5 (0.104)	4 (0.110)	6 (0.108)
N4	64.3%	81.6%	0.130	2 (0.117)	3 (0.116)	5 (0.106)	3 (0.113)
N5	69.2%	64.2%	0.142	3 (0.114)	1 (0.132)	6 (0.105)	1 (0.117)
N6	78.1%	70.4%	0.106	6 (0.108)	7 (0.098)	7 (0.103)	7 (0.103)
N7	54.6%	91.3%	0.115	3 (0.114)	6 (0.102)	2 (0.115)	4 (0.110)
N8	65.9%	71.2%	0.074	7 (0.105)	4 (0.108)	3 (0.113)	6 (0.108)
N9	57.2%	79.4%	0.177	5 (0.110)	2 (0.123)	3 (0.113)	2 (0.115)

¹ Focal species rank from Table 5a