Yuriko Hashimoto 990853225 Penn State MGIS Capstone GEOG 596B

# Evaluating environmental predictors of breeding waterfowl population distribution and abundance in the Central Interior of British Columbia (2007-2017)

#### **ABSTRACT**

Conservation biology has seen an explosion of species distribution models (SDMs) in the recent published literature and growing numbers of governments and organizations responsible for small-scale regional to global conservation activities are actively utilizing them or have embarked on their own predictive studies (Franklin and Miller 2009; Guisan, Wilfried, and Zimmermann 2017). Waterfowl conservation planning in British Columbia (BC) however currently does not have the benefit of breeding waterfowl SDMs. To address this gap I have developed methods to generate SDMs using 11 years of survey data from BC's annual Breeding Waterfowl Survey between 2007 and 2017. Models utilizing a random forest-based approach were used to produce predictive maps for ten species and five species groups within each of the eight ecosections within the central interior representing core provincial breeding waterfowl habitat. The developed methods, techniques and recommendations form a template for future studies in the generation of species distribution maps for guides in conservation planning.

Keywords: waterfowl, random forest, species distribution model, conservation

## **INTRODUCTION**

While the uses and practical applications of species distribution models in conservation decision-making are diverse (Guillera-Arroita et al. 2015; Guisan et al. 2013; Guisan and Thuiller 2005) the overarching aim is to conserve biodiversity and support better land use planning and environmentally sustainable management practices. The breadth of practical applications include predicting climate change adaptation, invasive species management, critical habitat identification, reserve selection, impact assessment and ecological restoration (Elith and Graham 2009; Franklin and Miller 2009).

Recognizing the lack of waterfowl species distribution models (SDMs) to support conservation actions in response to the urgency of conservation challenges in Canada's boreal forests, Barker et al (2014) published the first national-scale waterfowl SDMs in 2014 based on the traditional Waterfowl Breeding Population and Habitat Survey (WBPHS) database. The WBPHS has been described as "arguably the largest and best-designed population survey in the world" (Murray, Anderson, and Steury 2010), continental in scale and running on a continuous annual basis since 1955, the survey captures what is recognized to be "core waterfowl breeding habitat" in North America. Designed primarily to provide annual breeding population estimates and inform hunting regulations in the U.S. and Canada, these data additionally provide a long-term population monitoring dataset that has since informed countless studies on species-habitat relationships in support of waterfowl conservation. While the models performed well over all, the results of extrapolation to out-of-sample areas were variable for a number of reasons as outlined by the authors. British Columbia (BC) is not within the traditional survey and no survey data from BC informed the models which predicted low total densities for the Western Cordillera between the Pacific coast and Rocky Mountains—a result that conflicts with anecdotal expert opinion (A. Breault, personal communication 2018). As such there are currently no adopted waterfowl

breeding population distribution models in use for the province. This study aims to fill the this gap and help support the mandate of the Canadian Wildlife Service (CWS) to conserve biodiversity (Environment and Climate Change Canada 2019; Environment Canada - Biodiversity Convention Office 1995). In the early 2000s after exploratory pilot surveys in BC's central and sub-boreal highlands determined waterfowl population abundances to be significant enough to justify a regional breeding survey program, the British Columbia Breeding Waterfowl Survey began in earnest in 2006. BC's May surveys, run jointly by the CWS and the U.S. Fish and Wildlife Service and conducted in partnership with Ducks Unlimited Canada, inform the annual population status of migratory game birds in the Central Interior Plateau of BC and contributes to adaptive harvest strategies for mallards in the Pacific Flyway (Zimpfer, Breault, and Sanders 2019).

I developed methods to create SDMs for the top ten most abundant species as well as five group classifications based on life history traits and community descriptors (Baldassarre 2014) (Tables 1 and 2). Guisan et al (2013) have observed that despite the oft-cited assumption of the utility of SDMs in conservation decision-making there is little evidence demonstrating their application in real-world conservation management. Following their recommendation that modellers make explicit the objectives of the framework within which models were developed, I provide guidance on how the SDMs may be improved and implemented as a decision-making platform for conservation planning by the Canadian Wildlife Service and partners in conservation.

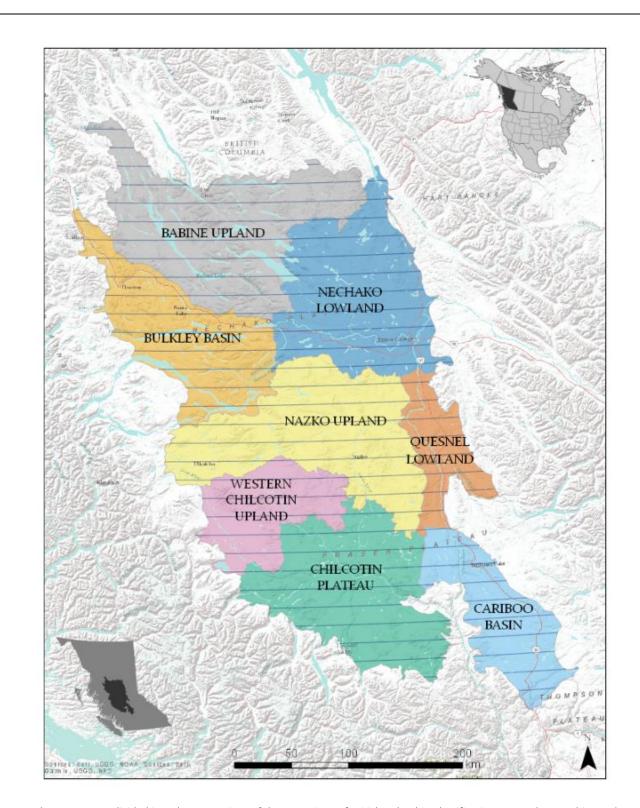
#### **MATERIALS AND METHODS**

#### **Survey Methods**

The study area was designed to capture BC's prime waterfowl breeding habitat of the humid, continental plateaus of the central and sub-boreal Interior between the Coast mountains to the west and Rocky Mountains to the east. The region of interlocking highlands and valleys are mainly forested and sparsely populated. The main industries include forestry, cereal crop agriculture within lowland valleys, ranching and mining (Demarchi 2011).

The survey design consists of latitudinal strip transects 400 metres wide, spaced ten miles apart (16.09 km) and delineated by the boundaries of eight ecosections which represent the finest scale, sub-regional stratum within the Ecoregion Classification System of British Columbia—a small-scale ecosystem management framework stratifying nested regions of "similar climate, physiography, oceanography, hydrology, vegetation and wildlife potential" (Demarchi 2011) (Figure 1). The classification levels ordered from high to low include Ecodomains for global mapping applications, ecodivisions for national-scale, ecoprovinces for provincial scale, ecoregions for regional and ecosections for sub-regional planning. While designed for small-scale mapping ecosection boundaries are drawn at large scale (1:20,000).

The southern two-thirds of the study area falls within the Fraser Plateau Ecoregion. The flat and rolling hills of the region contain numerous meandering streams and low-lying depressions that create an abundance of wetland habitat (Demarchi 2011) while the Central Interior Ecoprovince in which it nests supports 65% of all species known to occur and 61% of all bird species known to breed in BC. with the greatest total abundances of waterfowl occurring in the Cariboo Basin ecosection within the Riske Creek area (Demarchi 2011; Savard, Sean Boyd, and John Smith 1994). The remaining northernmost third of the study area stretches into the Subboreal Interior Ecoprovince which supports 57% of all bird species known to occur and 45% known to breed in the province (Demarchi 2011). Table # provides general descriptions of study area ecosections. The surveys are conducted by helicopter and surveyed by a consistent survey crew of two to three experienced observers (Zimpfer, Breault, and Sanders 2019). The transects when originally conceived were designed within the NAD 1983 geographic coordinate system and were based on the mapped Ecoregion Classification System version 2.0 (1995). The ecosection boundaries have since been updated with finer scale vegetation zonation data in version 2.1 (2006) however the annual surveys continue to follow the extent of the original design. All analyses were based on the areal overlap between the two versions with the boundaries redefined in alignment with version 2.1.



**Figure 1**. The survey area divided into the ecosections of the Ecoregions of British Columbia Classification system (Demarchi, 2011). Latitudinal strip transects (horizontal lines) are spaced ten miles (~16 km) apart within the Central and Sub-boreal plateaus of BC.

Table 1. General summary descriptions of ecosections within study area as described in Demarchi (2011).

| Ecoregion         | Ecosection                  | Topography   | Climate   | Vegetation  | Governance,<br>Protection,<br>Development   |
|-------------------|-----------------------------|--|---|---|---|
| Fraser<br>Basin   | Babine<br>Upland<br>BAU     | Rolling upland with<br>low ridges; many small<br>streams and lakes and<br>several large lakes  | Sub-continental<br>extreme winter<br>cold and snow<br>events, humid and<br>rainy  | Sub-boreal<br>spruce<br>dominated<br>forests  | Sparsely populated,<br>seasonal lodges;<br>many protected<br>areas  |
| Fraser<br>Plateau | Bulkley<br>Basin<br>BUB     | Broad, lowland valleys<br>with many lakes  | Rainshadow effect<br>of Coast<br>mountains  | Dominated by<br>lodgepole pine;<br>trembling aspen<br>in lower south-<br>facing valleys       | Extensive<br>development and<br>agriculture within<br>Yellowhead Hwy<br>corridor and<br>surrounding<br>Francois Lake;<br>protected areas<br>include Francois lake<br>Park |
| Fraser<br>Plateau | Cariboo<br>Basin<br>CAB     | Rolling uplands; many<br>streams, wetlands and<br>lakes  | Warmest<br>temperatures<br>within ecoregion;<br>relatively high<br>summer<br>precipitation  | Vegetation<br>zonation changes  | Small farms, ranches<br>and logging are main<br>industries. Williams<br>Lake is largest city  |
| Fraser<br>Plateau | Chilcotin<br>Plateau<br>CHP | Rolling uplands with<br>many small lakes and<br>wetlands; higher relief<br>in south and<br>northwest abutting<br>Chilcotin Ranges and<br>west Chilcotin; many<br>small streams and<br>rivers | Pronounced rain<br>shadow effect  | Douglas fir near<br>rivers and<br>lodgepole pine at<br>higher elevations                      | Ranching and<br>logging are main<br>industries; Big Creek<br>Park is largest park   |
| Fraser<br>Plateau | Nazko<br>Upland<br>NAU      | Rolling uplands with<br>high relief within<br>north-central region;<br>slow moving streams<br>and wetlands   | Sub-continental climatecold winters, warm summers, maximum precipitation in late spring/early summer; subject to cold balasts of Arctic air | Lodgepole pine<br>in south and<br>white<br>spruce/lodgepole<br>pine/subalpine<br>fir in north | Ranching and First<br>Nations reserves but<br>no large<br>communities   |

| Fraser<br>Basin   | Nechako<br>Lowland<br>NEL             | Flat or gently rolling<br>lowlands  | Sub-boreal<br>climate-humid<br>summers and<br>harsh winters  | Sub-boreal<br>spruce forests<br>and lodgepole<br>pine   | Largest city is Prince<br>George; cereal crop<br>farming in southern<br>lowlying areas and<br>ranching            |  |  |
|-------------------|---------------------------------------|---|--|---|---|--|--|
| Fraser<br>Plateau | Quesnel<br>Lowland<br>QUL             | Lowland trench<br>bissected by the<br>Fraser River; small<br>lakes and wetlands<br>but no large lakes | Subject to cold<br>Arctic air;<br>increased<br>precipitation as air<br>moves east over<br>Columbia<br>Mountains and in<br>summer due to<br>upland wetlands | Douglas-fir on<br>dry south-facing<br>slopes and<br>trembling aspen,<br>lodegpole pine,<br>white spruce to<br>subalpine fir with<br>increasing<br>elevation | Quesnel is the largest and only city but many communities; farming ranching and forestry (with extensive logging) |  |  |
| Fraser<br>Plateau | Western<br>Chilcotin<br>Upland<br>WCU | Rounded uplanded<br>with a large lowland<br>area containing many<br>lakes                             | Sub-continental<br>climate subject to<br>blasts of Arctic<br>winter air  | Mainly lodgepole<br>pine and white<br>spruce in higher<br>elevations  | Anahim Lake is<br>largest community   |  |  |

The survey does not follow the design methodology of the traditional WBPHS, however the survey techniques are consistent with the methods outlined in Smith (1995) and the Standard Operating Procedures (US Fish and Wildlife Service (USFWS) and Canadian Wildlife Service (CWS) 1987). Prior to 2010, the survey technologies employed consisted of paper maps, field notes and GPS units with navigation determined by piloting along bearings to GPS waypoints. In 2010, the mobile GIS software PC-Mapper with Airborne Inspection (version 4.0, Corvallis Microtechnology Inc, 2015) was adopted for both survey navigation and data collection. The software is run on Panasonic Toughbooks (CF-19 and CF-31) with the screen in view of both the pilot and observer. Real-time navigation is guided by the GIS with reference base data containing strip transect boundaries, freshwater polygons and stream segments, ecosection boundaries and fuel waypoints. Digital data collection via georeferenced voice recordings transcribed by the observer post-survey collection have replaced paper analogue methods.

The survey is designed to capture the primary breeding period beginning in early May but is weather and climate dependent and has taken place as early as late April (April 28, 2015) and as late as mid-May (May 13, 2011). Heavy snowpack and/or cold spring temperatures can delay accessibility to open water and wetlands. The study area extent is almost 11 million hectares and takes an average 23 days (~105 flying hours) to survey.

**Table 2.** Species included in the study and their nesting and feeding guilds based on Baldasarre (2014) and Cornell (2015). Species-specific models for only the top ten most common species were created however less common species were accounted for in the groups outlined in Table 2.

| Species<br>Code | Common Name         | Scientific Name     | Species Group | Foraging Guild           | Nesting Guild |
|-----------------|---------------------|---------------------|---------------|--------------------------|---------------|
| AMWI            | American Wigeon*    | Mareca americana    | Dabblers      | Dabbler - Plants         | Ground        |
| BAGO            | Barrow's Goldeneye* | Bucephala islandica | Divers        | Surface - Dive - Insects | Cavity        |
| BUFF            | Bufflehead*         | Bucephala albeola   | Divers        | Aerial Dive - Insects    | Floating      |

| BWTE Blue-winged Teal* Spatula discors Dabblers Dabbler - Seeds Ground  CAGO Canada Goose* Branta canadensis Geese Ground Forager - Seeds Ground  CANV Canvasback Aythya valisineria Divers Surface - Dive - Plants Floating  CITE Cinnamon Teal Spatula cyanoptera Dabblers Dabbler - Seeds Ground  COGO Common Goldeneye Bucephala clangula Divers Surface - Dive - Insects Cavity  COME Common Merganser Mergus merganser Mergansers Surface - Dive - Fish Cavity  GADW Gadwall Mareca strepera Dabblers Dabbler - Plants Ground  GWTE Green-winged Teal* Anas crecca Dabblers Dabbler - Seeds Ground  HADU Harlequin Duck Histrionicus histrionicus Divers Surface - Dive - Insects Ground  HOME Hooded Merganser Lophodytes cucullatus Mergansers Surface - Dive - Fish Cavity  LTDU Long-tailed Duck Clangula hyemalis Divers Surface - Dive - Insects Ground |
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| MALL Mallard* Anas platyrhynchos Dabblers Dabbler - Seeds Ground  |
| NOPI Northern Pintail Anas acuta Dabblers Dabbler - Seeds Ground  |
| NSHO Northern Shoveler* Spatula clypeata Dabbler - Plants Ground  |
| RBME Red-breasted Merganser <i>Mergus serrator</i> Mergansers Surface - Dive - Fish Ground  |
| REDH Redhead Aythya americana Divers Surface - Dive - Plants Floating   |
| RNDU Ring-necked Duck* Aythya collaris Divers Surface - Dive - Plants Floating  |
| RUDU Ruddy Duck <i>Oxyura jamaicensis</i> Divers Surface - Dive - Insects Ground  |
| SCAU** Lesser Scaup* Aythya affinis Divers Surface - Dive - Insects Ground  |
| SNGO Snow Goose Anser caerulescens Geese Ground Forager - Plants Ground   |
| TRUS Trumpeter Swan Cygnus buccinator Swans Dabbler - Plants Ground   |
| WFGO White-fronted Goose Anser albifrons Geese Dabbler - Plants Ground  |
| WODU Wood Duck Aix sponsa Dabbler - Plants Ground   |

 $<sup>{}^{*}</sup>$  Species within the top ten most commonly observed for which a species-specific model was created.

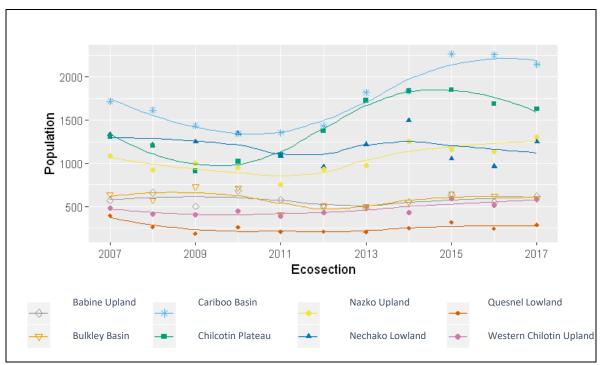
**Table 3.** Group classifications for which group-specific species distribution models were generated (refer to Table 1 for specific species included within the groups).

| Abbreviation | Explanation  |
|--------------|--|
| sp_div       | Index of species richness—number of unique species                     |
| sp_tot       | Count of total indicated breeding population of all waterfowl          |
| dabblers     | Count of total indicated breeding population of dabbling ducks         |
| divers       | Count of total indicated breeding population of diving ducks           |
| cavity       | Count of total indicated breeding population of cavity nesting species |

# **Population estimates**

The surveys are conducted by helicopter along meandering paths at altitudes and speeds lower than fixed-wing aircraft (30-50 m and 40-80 km/h, respectively) therefore no complementary ground surveys are conducted and no visibility correction factor is applied as visibility is assumed to be complete. The total indicated breeding population estimates were derived from raw counts based on species, sex and grouping as outlined in Smith (1995). The temporal subset of the analysis was from 2007 to 2017 inclusive (data from 2006 was excluded due to data inconsistencies and 2018 was excluded due to the absence of interpreted climate data). Survey observation points record the location of the observer within the helicopter and not the bird on the ground. Efforts to correct locations guided by reference data and field observation notes were discontinued from 2012 onward due to resource constraints and inconsistent observational record-keeping. In order to account for this spatial uncertainty, observation location points were collated to the nearest 400m interval along the center line of the strip transect.

<sup>\*\*</sup> The waterfowl population survey does not distinguish scaup species however only lesser scaup are observed in the area.



**Figure 2.** Total indicated breeding population of the top ten most common species observed within the transect (2007-2017).

The Cariboo Basin is by far the most populous ecosection with the greatest total count (Figure 2) and population density (Figure 3) followed by the Chilcotin Plateau, Nechako Lowland and Nazko Upland. The remaining ecosections—Bulkley Basin, Quesnel Lowland, Western Chilcotin Upland and Babine Upland—have similar population densities with the lattermost consistently the least populous. These ecosections share similarly stable year-to-year trends while the densities in the two most populous ecosections, Cariboo Basin and Chilcotin Plateau, display concordant fluctuations and appear to be increasing.

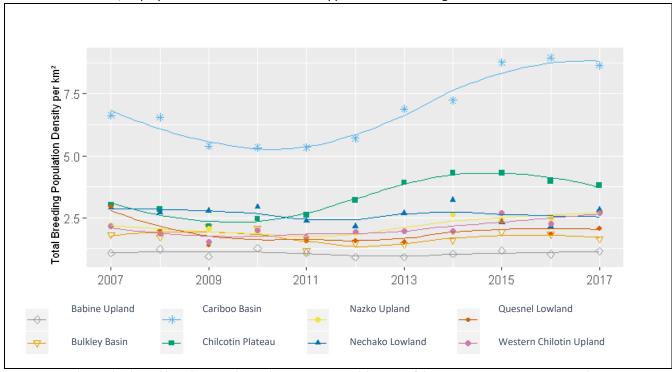
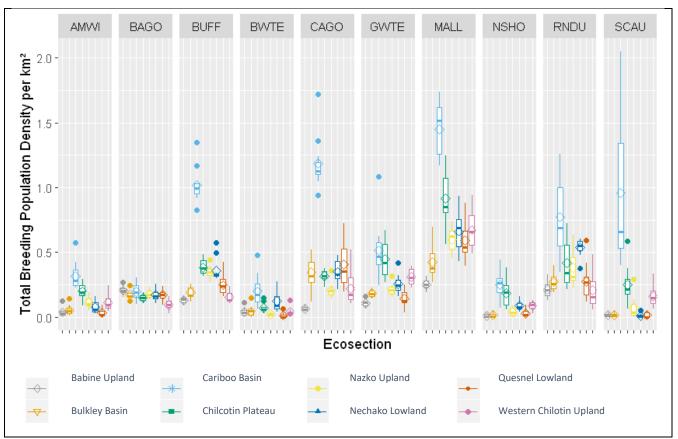


Figure 3. Total annual indicated breeding population density per square kilometre of the top ten most common species by ecosection observed within the transect.



**Figure 4**. Boxplots of total annual indicated breeding population density per square kilometre of the top ten most common species by ecosection observed within the transect.

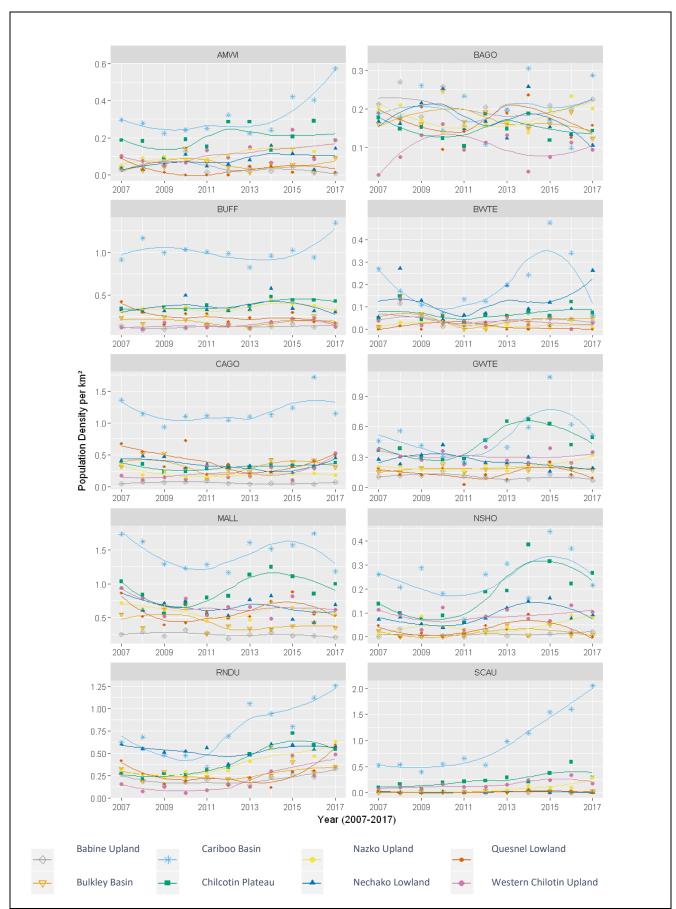


Figure 5. Total indicated breeding population of the top ten most common species observed within the transect (2007-2017).

Of the top ten most commonly observed species, the species demonstrating the greatest year-to-year variability in abundance include Lesser Scaup\* (LESC), Ring-necked Duck (RNDU), Canada Goose (CAGO), and Mallard (MALL) (Figure 5). The distributions suggest it is LESC and RNDU species driving the increasing population trends in the Cariboo Basin and the Chilcotin Plateau. Mallard (MALL), Canada Goose (CAGO), Green-Winged Teal (GWTE), Blue-Winged Teal (BWTE), and Northern Shoveler (NOSH) reflect concordant annual fluctuations—decreasing from 2007 to 2010, increasing between 2010 and 2015, and decreasing again to 2017. American Wigeon (AMWI) to a lesser degree follows the main trend but appears to be slightly increasing in the Cariboo Basin. Barrow's Goldeneye (BAGO) distributions remained steady throughout their distributions in all ecosections during the study period.

#### **Environmental data**

Environmental predictor variables were pre-selected based on ecological theory of life history traits and physiological processes: hydrological features associated with wetlands, and land use practices and disturbances that can potentially influence waterfowl habitat (Table 2). Where possible direct measures of predictor variables rather than proxy data was selected. For example, measures of temperature and precipitation rather than the indirect measure of elevation were included in the analyses. A number of datasets were collated and evaluated but the final data inputs were constrained by availability, resolution, coverage, and currency, and complicated by interpretation and relevance (Appendix 1). Technical constraints—processing hardware and time—necessitated limiting the number of candidate predictors to improve both interpretation and efficiency however more accurate predictions may be generated with fewer restrictions.

Annual, seasonal, monthly and 30-year normal climatic variables were extracted from ClimateBC software (version 6.10, Wang, Hamann, Spittlehouse, & Carroll, 2016) based on the coordinates and elevation at 400m interval points along the transect center line. Growing degree days and average April temperature were included to capture primary productivity. To represent and characterize hydrological regimes which are driven mainly by snowpack accumation in winter within the study area (Demarchi 2011; Islam et al. 2017; R. Pike et al. 2010) precipitation as snow and the climatic moisture deficit (precipitation minus potential evapotranspiration) were selected. The Biogeoclimatic Zones of British Columbia (BEC) is a standardized provincial dataset that classifies ecosystems within nested classes of regional, site and chronological levels of apex vegetaion (Ministry of Forests, Lands, Natural Resource Operations and Rural Development (FLNRO), 2018) and was included to represent the combined influence of soil chemistry, vegetation, topography and climate. Additionally, the BEC Zone classifications provide an alternative regional classification system to the Ecoregions of BC ecosections for supplemental model development. Moreover, predicted changes to BEC Zone boundaries due to predicted climate change scenarios can be extracted from ClimateBC for future climate impact studies.

Land cover variables to characterize habitat were derived from 2010 Landsat imagery published by the North American Land Change Monitoring Sytem (CCRS/CCMEO/NRCan 2017). Cover classes were aggregated and reclassified to account for imbalanced classes (Appendix #). Topographic data included slope and aspect derived from 1:20,000 DEM (FLNRO, 2014).

Hydrological variables were derived from the provincial reference dataset for standardized hydrological features, and included polygons of lakes, rivers, man-made waterbodies and wetlands as well as stream lines (FLRNO, 2011). Lakes were classified by size class and streams were aggregated by stream order values (Appendix 2). To account for shoreline complexity and avoid correlation a lake perimeter to area index was derived.

The line network of unpaved roads was included to represent anthropogenic disturbance of resource extraction activities (Ministry of Forests, Lands, Natural Resource Operations and Rural Development, 2013).

Land management practices were represented by the Agricultural Land Reserve (Agricultural Land Commission, 2018)—provincial designation designed to identify and conserve agricultural productivity, and by the Protected Areas Database of lands under federal, provincial and municipal protection and land conservancy trusts (CWS, 2018).

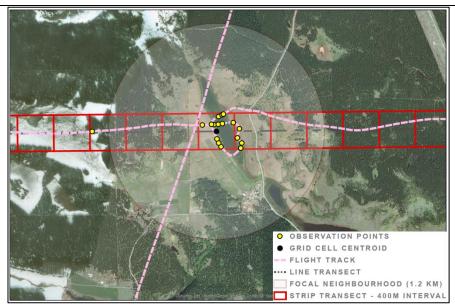
**Table 4.** Environmental predictor variables selected for model inputs.

| Predictor -<br>abbreviation | Predictor - description                                   | Resolution | Time Period       | Source  | Calculation  |
|-----------------------------|---|------------|-------------------|---------|--|
| fwa_1                       | Freshwater Atlas - Lakes<br>< 1 ha                        | 1:20,000   | Static (Variable) | Data BC | Total area within 1.2 km<br>circular radius of centroid point<br>along 400m interval of transect                                       |
| fwa_2                       | Freshwater Atlas - Lakes<br>1-2 ha                        | 1:20,000   | Static (variable) | Data BC | Total area within 1.2 km circular radius of centroid point along 400m interval of transect   |
| fwa_3                       | Freshwater Atlas - Lakes<br>3-5 ha                        | 1:20,000   | Static (variable) | Data BC | Total area within 1.2 km circular radius of centroid point along 400m interval of transect   |
| fwa_4                       | Freshwater Atlas - Lakes<br>5-10 ha                       | 1:20,000   | Static (variable) | Data BC | Total area within 1.2 km circular radius of centroid point along 400m interval of transect   |
| fwa_5                       | Freshwater Atlas - Lakes<br>10-20 ha                      | 1:20,000   | Static (variable) | Data BC | Total area within 1.2 km circular radius of centroid point along 400m interval of transect   |
| fwa_6                       | Freshwater Atlas - Lakes<br>20-50 ha                      | 1:20,000   | Static (variable) | Data BC | Total area within 1.2 km circular radius of centroid point along 400m interval of transect   |
| fwa_7                       | Freshwater Atlas - Lakes<br>50-100 ha                     | 1:20,000   | Static (variable) | Data BC | Total area within 1.2 km circular radius of centroid point along 400m interval of transect   |
| fwa_8                       | Freshwater Atlas - Lakes > 100 ha                         | 1:20,000   | Static (variable) | Data BC | Total area within 1.2 km circular radius of centroid point along 400m interval of transect   |
| fwa_10ha_plus               | Freshwater Atlas - Lakes > 10 ha                          | 1:20,000   | Static (variable) | Data BC | Total area within 1.2 km circular radius of centroid point along 400m interval of transect   |
| shorecx_10plus              | Freshwater Atlas -<br>Shoreline index of lakes<br>< 10 ha | 1:20,000   | Static (variable) | Data BC | Ratio of total perimeter to area of lakes, multiplied by 100,000   |
| shorecx_10plus              | Freshwater Atlas - Shoreline index of lakes > 10 ha       | 1:20,000   | Static (variable) | Data BC | Ratio of total perimeter to area of lakes, multiplied by 100,000   |
| str_s                       | Freshwater Atlas -<br>Stream order 1-3                    | 1:20,000   | Static (variable) | Data BC | Total length of streams of stream orders 1 to 3  |
| str_m                       | Freshwater Atlas -<br>Stream orders 4-6                   | 1:20,000   | Static (variable) | Data BC | Total length of streams of<br>stream orders 4 to 6   |
| fwa_w                       | Freshwater Atlas -<br>Wetlands                            | 1:20,000   | Static (variable) | Data BC | Total area of wetlands within 1.2 km circular radius of centroid point along 400m interval of transect Total area of rivers within 1.2 |
| fwa_r                       | Freshwater Atlas - River                                  | 1:20,000   | Static (variable) | Data BC | km circular radius of centroid<br>point along 400m interval of<br>transect   |
| aspect                      | Compass direction   | 1:20,000   | Static (2011)     | Data BC | Average aspect within 400 x<br>400 m areal interval of strip<br>transect   |
| slope                       | Topographic slope   | 1:20,000   | Static (2011)     | Data BC | Average slope within 400 x 400 m areal interval of strip transect  |
| alr                         | Agricultural Land<br>Reserve                              | 1:20,000   | Static (2014)     | Data BC | Total area within 1.2 km circular radius of centroid point along 400m interval of transect   |

| pa               | Protected Areas                             | 1:20,000            | Static (2018) | CWS<br>Conservation<br>Areas DB                   | Total area within 1.2 km circular radius of centroid point along 400m interval of transect                        |
|------------------|---|---------------------|---------------|---|---|
| dra_u            | Digital Road Atlas,<br>Unpaved Roads        | 1:20,000            | Static (2014) | Data BC   | Total density within 1.2 km circular radius of centroid point along 400m interval of transect                     |
| cec_urban        | Land cover - urban                          | 30 m                | Static (2010) | Commission<br>for<br>Environmental<br>Cooperation | Total area within 1.2 km circular radius of centroid point along 400m interval of transect                        |
| cec_shrubland    | Land cover - shrubland                      | 30 m                | Static (2010) | Commission<br>for<br>Environmental<br>Cooperation | Total area within 1.2 km circular radius of centroid point along 400m interval of transect                        |
| cec_mixed_forest | Land cover - mixed forest                   | 30 m                | Static (2010) | Commission<br>for<br>Environmental<br>Cooperation | Total area within 1.2 km circular radius of centroid point along 400m interval of transect                        |
| cec_broadleaf    | Land cover - broadleaf<br>(deciduous)       | 30 m                | Static (2010) | Commission<br>for<br>Environmental<br>Cooperation | Total area within 1.2 km circular radius of centroid point along 400m interval of transect                        |
| cec_needleleaf   | Land cover - needleleaf<br>(coniferous)     | 30 m                | Static (2010) | Commission<br>for<br>Environmental<br>Cooperation | Total area within 1.2 km circular radius of centroid point along 400m interval of transect                        |
| bec_zn           | Biogeoclimatic Zone                         | 1:20,000            | Static (2018) | Data BC   | Majority area within 400 x 400 m areal interval of strip transect   |
| norm_dd5         | 30 yr normal Growing<br>Degree Days > 5°C   | Scale-free 400<br>m | 1980-2010     | ClimateBC   | Climatic measure at coordinate position and elevation along 400m interval of transect                             |
| norm_cmd         | 30 yr normal Cumulative<br>Moisture Deficit | Scale-free 400<br>m | 1980-2010     | ClimateBC   | Climatic measure at coordinate position and elevation along 400m interval of transect                             |
| pas_wt           | Mean Precipitation as<br>Snow in winter     | Scale-free 400<br>m | Annual        | ClimateBC   | Climatic measure at coordinate position and elevation along 400m interval of transect averaged over survey period |
| tave04           | Average April<br>Temperature                | Scale-free 400<br>m | Annual        | ClimateBC   | Climatic measure at coordinate position and elevation along 400m interval of transect averaged over survey period |

# **Geoprocessing Methods**

Spatial analyses were performed and mapping products were produced in ArcGIS Pro (versions 2.2 to 2.4, ESRI, 2019) and Python (version 3.6.5, Python Software Foundation, 2018). Predictor variables were extracted, projected, rasterized and generalized as required in BC Albers equal area coordinate system within a 1 km buffer of the study area to eliminate edge effects. The location uncertainty of point observations determined the finest scale of the analysis to be a resolution of 400m. Conceptually, survey strip transects were segmented into 400m intervals forming 400m x 400m (16 ha) grid cells—16, 540 in total. Topographic values of slope and aspect were generalized to the mean average within each grid cell. BEC zone classifications were generalized by majority area within the cell and remained categorical. All remaining predictors were represented by continuous values and generalized to a 1.2 km radius of the interval centroid to capture landscape level effects (Figure 6). Predicted response values were projected to a fishnet grid of 400m cell centroid points with attributed predictor values and rasterized for display. Many of the steps required in the pre-processing of spatial data were automated in a Python Script Toolbox (Appendix 3).



**Figure 6.** Representative schematic of the survey methods incorporating survey data from 2017. Observation points (yellow dots) are collected along the meandering flight tracklog (dashed pink line) within the transect. The strip transect was spliced into 400m wide intervals (red squares) with a circular focal neighbourhood of 1.2 km (pale circle) of the 400x400m grid cell centroid (black dot) used to characterize landscape level effects of environmental variables.

## **Statistical Methods**

Random forest is an ensemble machine learning algorithm that creates a series of decision trees based on the principles of bagging (or bootstrapped aggregation—the drawing of a large number of data samples by random sampling with replacement) and random permutation. Each individual decision tree is based on a random subset of the data and at each node of the tree the data is partitioned into two branches based on a randomly selected but predetermined number of predictor variables evaluated by the algorithm to produce the best split. As individual trees are highly susceptible to noise in the data and sensitive to local optima, each tree is considered a 'weak learner' (Guisan, Wilfried, and Zimmermann 2017). But by combining and averaging the results of several regression trees, a 'strong learner' is created in the final ensembled prediction.

A random forest-based approach was selected for its predictive accuracy, resistance to overfitting, ability to account for imbalanced classes, ability to handle both continuous and categorical variables, and its independence from requirements of feature scaling and centering as well as assumptions of normality (Cutler, Cutler, and Stevens 2012; Guisan, Wilfried, and Zimmermann 2017). The observation frequency distribution of bird counts reflected a zero-inflated negative binomial distribution typical of ecological count data (Qian 2010) but not amenable to standard regression-based approaches. Preliminary explorations of zero-inflated generalized mixed models indicated it is a promising approach for future explanatory model building.

All statistical modeling and data manipulation was performed in R (version 3.5.3, R Core Team, 2018). The R 'party' package (version 1.3-3) was selected for its implementation of the random forest and bagging algorithm 'cforest\_unbiased' function which utilizes conditional inference trees that account for correlation structures between variables in the permutation scheme of variable selection; additionally, the function implements a permutation importance measure, function 'varimp', that is immune to erroneous calculations due to correlated responses and is not biased towards continuous data or categorical variables with many classifications unlike the Gini accuracy of traditional random forest implementations (Strobl et al. 2008; Strobl, Hothorn, and Zeileis 2009). Slight differences in selected predictor variables between ecosection models were due to zero or near-zero variance, correlated data structures and/or mismatched factor variables in ecosection-based data inputs for model training and prediction.

Preliminary trial results of model inputs of annual survey data indicated low variable importance rankings for the dynamic climate variables. As all other environmental data was static the model input values were based on mean averaged counts and climate measures. Due to high zero-inflation (average of 93.6% frequency of zeroes for the top ten most common species) the data was not subset into training, validation, and test sets to evaluate model performance, instead the internal out-of-bag (OOB) error measures were determined. As each tree is based on a random subset of the training data, a collection of datasets which do not contain a particular record can be ensembled for each record. This collection forms the out-of-bag examples which are used as an unbiased test set to assess prediction accuracy by averaging the error rate. The predetermined number of candidate variables for each node split ('mtry' parameter of 'cforest') was left at the default value of 5 which resulted in better OOB estimates than 8 which was tested using Briemann's rule of thumb to divide the number of candidate variables by three for regression trees (Guisan, Wilfried, and Zimmermann 2017). The stability of 'varimp' ranks helped to inform the number of trees generated for each model which was set to 5,000.

Separate models for each species or species group were developed for each ecosection. Preliminary trials of generalized models on the entire study area indicated significant ecosection differences in predictor variables. Additionally, the computational intensity of the algorithm required for the entire study area was greater than available resources.

The conditional inference trees utilized by 'party' package are insensitive to highly correlated data structures but in order to reduce processing time, data preparation included addressing correlation and multicollinearity by step-wise elimination of correlated variables with threshold measures of r = 0.8 followed by elimination of variance inflation factors of 5 and greater corresponding to general thresholds recommended by Guisan et al (2017). The 'varimp' function to derive variable importance can be parameterized to produced unbiased importance measures of highly correlated data however preliminary trial results in Cariboo Basin reflected exaggerated processing times that restricted the application in this study. Neither correlation nor multicollinearity in predictor variables reduces predictive accuracy of random forests however the candidate predictors removed in the preprocessing step for the generation of 'varimp' values were not re-incorporated in the final prediction models. It is recommended these be included in future modeling exercises. The predictive models were re-run setting different seed values in order to confirm the stability of importance measures.

Random forest methods are widely recognized as fast and able to handle large amounts of data and model variables but the unbiased algorithm of the 'cforest' implementation is more computationally intensive than standard approaches. For example model training took approximately 10 minutes to process ~3,000 records and 27 variable inputs for mallards in Babine Upland, while model forecasting to the ecosection, an area ~ 40 times greater, took over 18 hours on a 64-bit OS workstation with 48.0 GB RAM and an Intel Xeon(R) CPU 3.60GHz. The R package 'caret' is designed specifically for data preprocessing and model generation and contains a number of machine learning methods including a more traditional random forest implementation and future studies are recommended these be explored.

The R scripts developed for statistical analyses as well as links to a Github repository of latest project updates and an interactive Jupyter Notebook documenting major processing steps are provided in Appendix 4.

#### **RESULTS AND DISCUSSION**

#### Variable Importance

Predictive modelling algorithms like random forest forsake the theoretical hypotheses of explanatory causal models for accurate forecasting (Shmueli 2011). The variable importance measures derived from recursive partitioning methods are unlike regression coefficients in that they do not provide a linear measure of the relationship between the predictor and response variables, rather the reported measure reflects the drop in prediction accuracy of the model by random permutation of the variable (Guisan, Wilfried, and Zimmermann 2017; Strobl et al. 2008). The most unambiguous application of these values is in variable selection for model

building. In summary, variable importance values are not by themselves interpretive however they may reveal causal mechanisms to inform the development of explanatory models (Shmueli, 2011). Predictors were ranked by their variable important measures and summaries are provided in Table # and Table #, respectively (see Appendix # for variable importance plots of each unique species within each ecosection). Note that the study is informed by a management framework and explores a place-based approach focusing on regional ecosections rather than the species perspective (e.g. Variable importance plots are grouped by ecosection not species).

#### **Model Performance**

Model performance was assessed with the out-of-bag (OOB) estimate. The coefficient of determination ( $R^2$ ) and mean absolute error (MAE) are reported in Table #.  $R^2$  represents the measure of variance in response values that is explained by the model and corresponds to the correlation between the observed and expected values. The MAE provides a measure of the difference between the observed and expected values and is less sensitive to outliers than the root mean square error (RMSE). In general terms, the lower the MAE and higher the  $R^2$  the better the performance.

Table 2. Model performance of each species-ecosection model estimated by the coefficient of determination ( $R^2$ ) and mean absolute error (MAE) derived from out-of-bag (OOB) estimates.

|                               | Upla           | ine<br>and | Bulkley | Basin | Cari<br>Ba     | boo<br>sin | Child<br>Plat  |      | Na:<br>Upl | -    | Nech<br>Low    |      | Que<br>Low     | snel<br>land | Child          | tern<br>cotin<br>land |
|-------------------------------|----------------|------------|---------|-------|----------------|------------|----------------|------|------------|------|----------------|------|----------------|--------------|----------------|-----------------------|
| Species                       | R <sup>2</sup> | MAE        | $R^2$   | MAE   | R <sup>2</sup> | MAE        | R <sup>2</sup> | MAE  | $R^2$      | MAE  | R <sup>2</sup> | MAE  | R <sup>2</sup> | MAE          | R <sup>2</sup> | MAE                   |
| AMWI                          | 0.10           | 0.01       | 0.07    | 0.02  | 0.09           | 0.08       | 0.18           | 0.05 | 0.21       | 0.03 | 0.07           | 0.02 | 0.02           | 0.01         | 0.26           | 0.02                  |
| BAGO                          | 0.26           | 0.05       | 0.20    | 0.04  | 0.17           | 0.05       | 0.11           | 0.04 | 0.27       | 0.04 | 0.13           | 0.03 | 0.09           | 0.05         | 0.13           | 0.03                  |
| BUFF                          | 0.18           | 0.03       | 0.24    | 0.04  | 0.30           | 0.20       | 0.23           | 0.09 | 0.24       | 0.08 | 0.16           | 0.09 | 0.10           | 0.06         | 0.23           | 0.04                  |
| BWTE                          | 0.04           | 0.01       | 0.05    | 0.01  | 0.09           | 0.05       | 0.14           | 0.02 | 0.13       | 0.01 | 0.13           | 0.03 | 0.02           | 0.00         | 0.14           | 0.01                  |
| CAGO                          | 0.08           | 0.02       | 0.02    | 0.00  | 0.15           | 0.26       | 0.18           | 0.07 | 0.13       | 0.05 | 0.18           | 0.09 | 0.05           | 0.11         | 0.05           | 0.06                  |
| GWTE                          | 0.11           | 0.03       | 0.16    | 0.05  | 0.17           | 0.11       | 0.19           | 0.10 | 0.19       | 0.05 | 0.13           | 0.07 | 0.04           | 0.04         | 0.21           | 0.07                  |
| MALL                          | 0.15           | 0.06       | 0.15    | 0.10  | 0.24           | 0.26       | 0.26           | 0.17 | 0.24       | 0.12 | 0.20           | 0.14 | 0.10           | 0.14         | 0.28           | 0.13                  |
| NOSH                          | 0.01           | 0.00       | 0.02    | 0.00  | 0.11           | 0.07       | 0.19           | 0.05 | 0.18       | 0.01 | 0.10           | 0.02 | 0.01           | 0.01         | 0.22           | 0.02                  |
| RNDU                          | 0.18           | 0.05       | 0.20    | 0.06  | 0.20           | 0.16       | 0.23           | 0.09 | 0.20       | 0.09 | 0.21           | 0.12 | 0.06           | 0.07         | 0.12           | 0.05                  |
| SCAU                          | 0.01           | 0.00       | 0.07    | 0.00  | 0.20           | 0.22       | 0.21           | 0.06 | 0.15       | 0.02 | 0.05           | 0.00 | 0.01           | 0.01         | 0.21           | 0.04                  |
| Dabblers                      | 0.16           | 0.10       | 0.16    | 0.16  | 0.21           | 0.61       | 0.29           | 0.36 | 0.26       | 0.21 | 0.19           | 0.27 | 0.08           | 0.19         | 0.36           | 0.25                  |
| Divers                        | 0.30           | 0.16       | 0.31    | 0.17  | 0.29           | 0.64       | 0.28           | 0.25 | 0.30       | 0.23 | 0.25           | 0.28 | 0.12           | 0.20         | 0.28           | 0.15                  |
| Cavity-<br>Nesters<br>Species | 0.26           | 0.07       | 0.23    | 0.06  | 0.18           | 0.07       | 0.11           | 0.05 | 0.27       | 0.06 | 0.22           | 0.07 | 0.09           | 0.07         | 0.15           | 0.03                  |
| Diversity                     | 0.29           | 0.09       | 0.35    | 0.12  | 0.37           | 0.34       | 0.37           | 0.20 | 0.39       | 0.14 | 0.28           | 0.17 | 0.18           | 0.15         | 0.43           | 0.11                  |
| All Species Average           | 0.27           | 0.24       | 0.22    | 0.36  | 0.27           | 0.30       | 0.33           | 0.63 | 0.33       | 0.43 | 0.24           | 0.55 | 0.10           | 0.43         | 0.36           | 0.40                  |

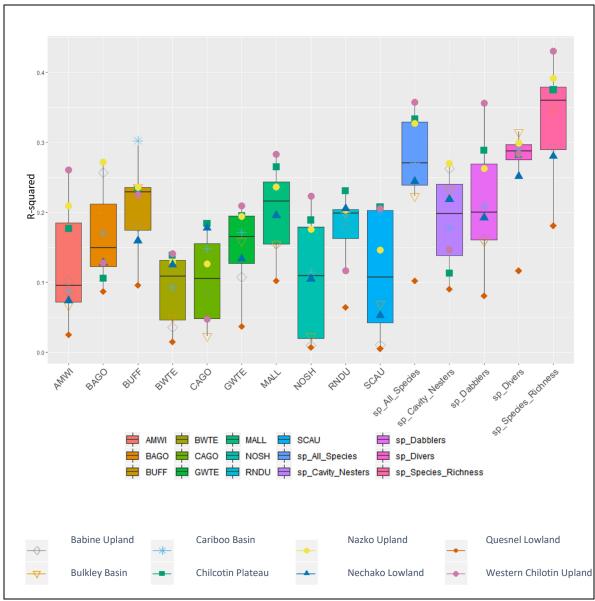


Figure 7. Boxplots of the coefficient of determination (R2) of each species-ecosection specific model.

Overall, the models on average explained 18% of the variation in species distribution (Figure 7). The generalized groups representing all species ("sp\_All\_species") and species richness ("sp\_Species\_Richness") performed best followed by divers, dabblers and cavity nesters. The least variability in R² values was for divers. Diving ducks tend to prefer larger, deeper lakes that are likely to be better represented in the data than smaller wetlands. Models for Western Chilcotin Upland, Chilcotin Plateau and Nazko Upland performed best while Quesnel Lowland consistently scored poorly. Quesnel Lowland is the smallest ecosection in the study area and the predictors selected do not appear to have sufficient explanatory variability. While the models for Western Chilcotin Upland and Chilcotin Plateau performed best overall their performance for cavity-nesting species, Barrow's goldeneyes and Bufflehead, were relatively poor. Cavity-nesting duck species do not excavate their own nests and rely on abandoned (or natural) cavities within aspen or mixed aspen/coniferous forests (Baldasarre, 2014). Buffleheads as North America's smallest diving duck, prefer entrance holes excavated by northern flickers which are smaller than those made by pileated woodpeckers to which goldeneyes are restricted (Baldasarre, 2014). Model parameters for cavity-nesting species may be improved with species distributions of these nest cavity excavating land birds.

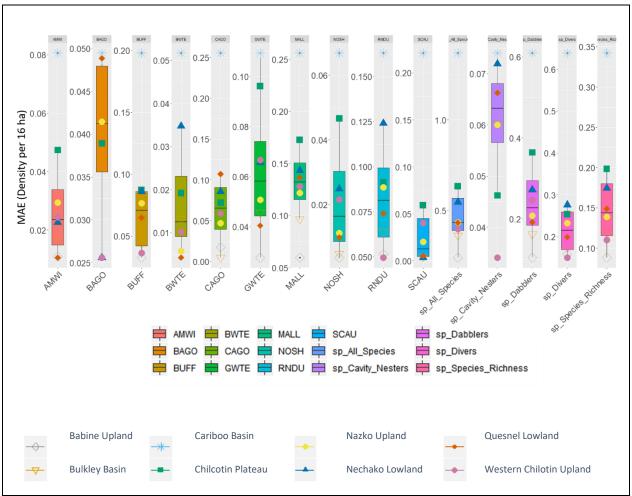


Figure 7. Boxplots of the mean absolute error (MAE) of each species-ecosection specific model (note the dynamic scale range values).

The overall average MAE was 13%, but this varied widely between species. The generalized group classifications excluding cavity-nesters had the largest values and greatest range in variability between ecosections. Cariboo Basin, consistently scored the highest error rate. The high error rates are likely due to the zero-inflation of observation frequencies—birds are patchily distributed throughput the landscape with zero observations in most cells punctuated by hotspots of high abundance.

## **Model Limitations**

We recognize "[d]ucks like water" (Pimm 1994), but these waters have yet to be mapped in detail: Canada, unlike most industrialized nations, does not have a national wetland dataset (Canadian Wetland Inventory — Ducks Unlimited Canada, 2019). The delineation of highly productive wetlands, such as ephemeral wetlands due to increased nutrient mineralization by aerobic microbes (Schlesinger and Bernhardt 2013) and smaller wetlands due to increased light penetration at shallower depths and greater shoreline emergent vegetation, are difficult to capture with freely available, coarser-scale remote sensing imagery (e.g. Landsat imagery at 30m resolution). Techniques for finer-scale wetland feature extraction based on Landsat have been developed for open wetlands but have not yet been fully developed and tested on forested wetlands (Halabisky et al. 2018).

Relative to other provinces, BC has an abundance of fine-scale (1:20,000) base reference spatial data however the creation of much of these data has been driven by the forestry sector which is a significant component of the BC economy. Consequently, much of the mapping of environmental features has been focused on supporting forestry management practices and the delineation of wetlands (areas unsuited to logging) especially at higher elevations is poor (D. Filatow, personal communication April 28, 2018). Moreover, despite

mapping standards, BC TRIM data was produced on a mapsheet by mapsheet basis and variability exists between areas. There are ongoing efforts to develop methodologies for wetland mapping coincident with the May surveys utilizing Radarsat-2 technologies in a collaborative effort between the CWS and the Science and Technology branches of Environment and Climate Change Canada which are promising but results are still preliminary and the project is far from complete (K. Moore, personal communication, September 30, 2018). Additionally, the Canadian Wetland Inventory project spearheaded by Ducks Unlimited Canada aims to provide a comprehensive national wetland inventory but BC has yet to be mapped (Canadian Wetland Inventory — Ducks Unlimited Canada, 2019). Global wetland mapping projects include the European Space Agency's Climate Change Initiative which has a number of exciting projects and remote sensing products of varying temporal and spatial resolution and should be monitored for updates (European Space Agency, 2019). Model performance and predictive accuracy can be expected to improve with the incorporation of improved wetland cover products, but proactive management measures should not be delayed while waiting for these data.

Hydrological regimes that create and maintain wetland ecosystems are influenced by a host of factors such as the depth of snowpack, the rate of snowmelt, levels of groundwater, precipitation, glacial retreat, land management practices, and anthropogenic and natural disturbance (Adamus 2014; R. G. Pike et al. 2010). Such complexities are challenging to represent but future studies may be improved by incorporating snow basin, drainage and/or watershed data. Additionally, BC's forested ecosystems have been impacted by the mountain pine beetle outbreak from 1999 to 2015 followed by the extreme fire seasons of 2017 and 2018. The short and longer-term effects of these regional disturbances on forest hydrology are difficult to predict (R. Pike et al. 2010; R. G. Pike et al. 2010) but are worthy of monitoring and future investigation (Bunnell, Fred L, Wells, Ralph 2010; MacKenzie and Moran 2004; Snauffer, Hsieh, and Cannon 2016).

The population estimates reflect sampled observations over a limited period of time and therefore reflect only a snapshot of the species-habitat relationship assumed to be in pseudo-equilibrium (Guisan, Wilfried, and Zimmermann 2017). Moreover, the models did not account for density-dependence, breeding philopatry, site fidelity, territoriality, influence of breeding phenology, species nesting chronologies or lagged response to environmental shifts. A recent study on the breeding phenology of cavity-nesting birds identified observable impacts on nesting activities with critical temperature periods of local temperature as short as 4 days (Drake and Martin 2018). Daily temperature datasets from Natural Resources Canada at 1 km resolution (McKenney et al. 2011) were collated and extracted but not included in the generation of the models as these data were considered more useful as explanatory variables than predictive determinants. Preliminary trials indicated significant variable importance estimates for averaged weekly temperatures and future efforts are encouraged to explore the incorporation of these and other datasets as identified in Appendix 1.

#### **Species Abundance and Distribution**

# <insert discussion of predicted distributions and abundance of mapped outputs>

# **Model Applications**

This modeling study had two main, related objectives: first, develop the methods and techniques required to standardize the BC May Survey dataset for analysis and second, create SDMs of relative abundance and distribution of breeding waterfowl to support conservation planning and habitat management. The predicted distribution maps can be used to highlight hotspot areas of high abundance and diversity to evaluate and assess protection or mitigation measures, provide finer-scale population estimates for environmental impact assessments, and help guide adaptive management measures to prepare for climate change. Additionally, the results can inform future studies of ecological relationships, and serve as guidance further model development that addresses the models' limitations.

Changes in BC's hydrological regimes due to increased warming and drying trends are predicted to lead to wetland losses at low to mid-elevations. As higher elevations are less sensitive to temperature changes that

can affect snowpack accumulation, BC's higher elevation wetlands may buffer the impacts for waterfowl and other wetland species (Bunnell, Fred L, Wells, Ralph 2010; R. Pike et al. 2010; R. G. Pike et al. 2010). The breeding habitat within BC's Central Interior is by no means as productive as the Prairie Pothole Region (PPR), However the conservation value of our wetlands, especially at elevations greater than 1200 metres (Bunnell, Fred L, Wells, Ralph 2010), may rise with the present and predicted climate-related shifts in wetland productivity of the PPR (N. D. Niemuth, Fleming, and Reynolds 2014; N. Niemuth, Wangler, and Reynolds 2010; Zhao et al. 2016). Future climate impact studies should take into account the wide range in variability of projections between the global models available and employ ensemble methods that encompass this range of variability to account for uncertainty in impact assessments and adaptive management strategies (R. G. Pike et al. 2010; Spittlehouse and Wang 2016).

#### **CONCLUSION**

"The single story creates stereotypes, and the problem with stereotypes is not that they are untrue, but that they are incomplete. They make one story become the only story" (Adichie 2009).

With the quote above, the Nigerian author Chimamanda Ngozi Adichie was referring to how a limited perspective can lead to flawed conclusions and racial stereotyping; how any single observation however true is not representative. Models, like stories, must be developed and communicated in meaningful ways. A species distribution model is a single story: a simplified human construct of a complex phenomenon shared to represent and reveal processes and patterns through a single lens. Future studies are encouraged to validate and build upon these results by integrating expert opinion, employing best available methods and data, addressing uncertainty, and continuing to support systematic monitoring and ecological research. These models were created to assist the CWS, governmental and other partners in conservation to assess our efforts, guide action, and inspire us to ask more questions to better our understanding of fundamental biological and ecological theory (Sinclair et al, 2010) to inform planning and decision-making. This study is a single story but it is a story worth telling in support of waterfowl conservation in British Columbia.

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# **APPENDIX 1 – Dataset Sources**

Environmental datasets collated and evaluated for the modeling study. The table describes each dataset and provides a brief description of variables and their assessment.

# **APPENDIX 2—Reclassification**

<reclass tables for land cover and FWA>

# **APPENDIX 3 – Python Script Toolbox**

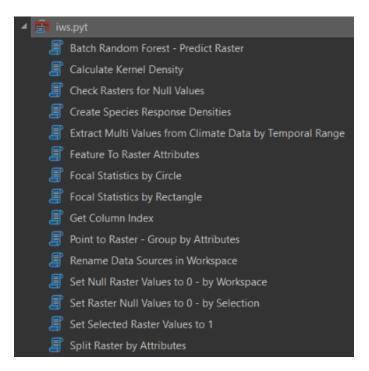


Figure 3. Python Script Toolbox as displayed in ArcGis Pro.

#### The code for the Python script toolbox is provided below.

```
import arcpy, os
from arcpy import env
arcpy.env.overwriteOutput = True
arcpy.env.voverw rteouppu = 17ue
from arcpy.sa import *
arcpy.CheckOutExtension("Spatial")
arcpy.env.qualifiedFieldMames = False
# scratch_ws = arcpy.CreateScratchName(workspace=arcpy.env.scratchGDB)
arcpy.env.snapRaster = r"C:\Users\hashimotoy\Desktop\ws\base_alb.gdb\dem"
arcpy.env.mask = r"C:\Users\hashimotoy\Desktop\ws\base_alb.gdb\survey_mask"
scratch_ws = r"C:\Users\hashimotoy\Desktop\ws_prep\scratch.gdb"
  def checkFieldExists(in_tbl, field_nm, field_type):
    fields = arcpy.ListFields(in_tbl)
    x = False
    for field in fields:
        if field.name == field_nm:
        x = Talse
 def getBaseName(in_fc):
    desc = arcpy.Describe(in_fc)
    basename = desc.baseName
    return basename
   def unique_values(table, field):
    with arcpy.da.SearchCursor(table, [field]) as cursor:
        return sorted({row[0] for row in cursor})
   class Toolbox(object):

def __init_(self):

"""befine the toolbox (the name of the toolbox is the name of the
.pyt file)."""

self.label = "IWS Data Tools"

self.alias = ""
                           # List of tool classes associated with this toolbox
self.tools = [featureToRasterAttributes,
                                                                          [featureIoRasterAttributes,
setRasterNullValuesToO,
setRasterNullValuesToOWorkspace,
renameDataInWorkspace,
kernelDensityCalculations,
splitRasterByAttributes,
focalStatsByCircle,
focalStatsByRectangle,
setPasterValuesTo1
                                                                           rocalstatsbykectangle,
setRasterValuesTo1,
pointToRasterGroupByAttributes,
getColumnIndex,
extractClimateValuesForRange,
                                                                           checkRasterForNullValues,
batchRandomForest,
createPredictedDensitySurfaces]
```

```
ss Tool(object):
def __init__(self):
   """Define the tool (tool name is the name of the class)."""
   self.label = "Tool"
   self.description = ""
                               self.canRunInBackground = False
           def getParameterInfo(self):
    """Define parameter definitions"""
    # First parameter
    param0 = arcpy.Parameter(
                                            displayName="Input Features",
name="in_features",
datatype="GPFeatureLayer",
parameterType="Required",
direction="Input")
                            # Second parameter
param1 = arcpy.Parameter(
    displayName="Sinuosity Field",
    name="sinuosity_field",
    datatype="Field",
    parameterType="Optional",
    direction="Input")
                              param1.value = "sinuosity"
                             # Third parameter
param2 = arcpy.Parameter(
param2 = arcpy.Parameter()
                             param2.parameterDependencies = [param0.name]
param2.schema.clone = True
           def isLicensed(self):
    """Set whether tool is licensed to execute."""
    return True
           def updateParameters(self, parameters):
    """Modify the values and properties of parameters before internal
    validation is performed. This method is called whenever a parameter
    has been changed."""
    return
            def updateMessages(self, parameters):
    """Modify the messages created by internal validation for each tool
    parameter. This method is called after internal validation."""
    return
           def execute(self, parameters, messages):
    """The source code of the tool."""
    return
lass batchRandomForest(object):
    def    init (self):
                            atchkandominerset(object):
__init__(self):
__i
          def getParameterInfo(self):
    """Define parameter definitions"""
                              param2 = arcpy.Parameter(
                                             displayName="Output Workspace",
name="out_ws",
datatype="DEWorkspace",
parameterType="Required",
direction="Input")
```

```
name="n_tree",
datatype="GPLong",
parameterType="Required",
direction="Input")
           param6 = arcpy.Parameter(
    displayName="Compensate for Sparse Categories",
    name="sparse",
    datatype="GPBoolean",
    parameterType="Required",
    direction="Input")
param6.value = True
           param7 = arcpy.Parameter(
    displayName="Percent Excluded for Validation",
    name="pct_train",
    datatype="GPlong",
    parameterType="Required",
    direction="Input")
param7.filter.type = "ValueList"
param7.filter.list = [10, 20, 30]
           # Fifth parameter
param9 = arcpy.Parameter(
    displayName="Select Explanatory Rasters",
    name="in rasters",
    datatype="DERasterDataset",
    parameterType="Required",
    direction="Input",
    multiValue=True)
            def isLicensed(self):
    """Set whether tool is licensed to execute."""
    return True
def updateParameters(self, parameters):
    """Modify the values and properties of parameters before internal
    validation is performed. This method is called whenever a parameter
    has been changed."""
    return
def updateMessages(self, parameters):
def execute(self, parameters, messages):
    ws_fishnet = r"C:\Users\hashimotoy\Desktop\ws_prep\fishnet.gdb"
           in_fc = parameters[0].valueAsText
lst_sp = parameters[1].valueAsText
out_ws = parameters[2].valueAsText
prefix = parameters[3].valueAsText
n_tree = parameters[4].valueAsText
m_try = parameters[6].valueAsText
sparse = parameters[6].valueAsText
pret_train = parameters[7].valueAsText
n_validation = parameters[8].valueAsText
in_rasters = parameters[9].valueAsText
            prediction_type = "PREDICT_RASTER"
nm = getBaseName(in_fc)
in_features = in_fc
arcpy.env.mask = r"C:\Users\hashimotoy\Desktop\ws_prep\raster_attributes.gdb\\" + "eco_" + nm
             lst_rasters = []
lst_matching = [] # Explanatory raster matching
for raster in in_rasters.split(';'):
   lst_rasters.append(raster)
   sub_lst = [raster, raster]
   lst_matching.append(sub_lst)
             treat_variable_as_categorical = None
explanatory_variables = None
distance_features = None
explanatory_rasters = lst_rasters
            explanatory_rasters = ist_rasters
features_to_predict = None
explanatory_variable_matching = None
explanatory_distance_matching = None
explanatory_rasters_matching = lst_matching
use_raster_values = True
number_of_trees = n_tree
minimum_leaf_size = None
maximum_level = None
sample_size = None
```

```
percentage_for_training = pct_train
output_classification_table = None
compensate_sparse_categories = True
number_validation_runs = n_validation
                       for sp in lst_sp.split(';'):
    arcpy.AddMessage("Running forest for " + sp)
    variable_predict = sp
    file_prefix = prefix + "." + nm + "." + sp + "."
    output_features = os.path.join(out_ws, file_prefix + "_output_features")
    output_trained_features = os.path.join(out_ws, file_prefix + "_trained")
    output_trained_features = os.path.join(out_ws, file_prefix + "_trained")
    output_importance_table = os.path.join(out_ws, file_prefix + "_varimp")
    output_validation_table = os.path.join(out_ws, file_prefix + "_validation")
                                    arcpy.stats.Forest(prediction_type,
in_features,
variable_predict,
treat_variable_as_categorical,
explanatory_variables,
distance_features,
                                                                                                     distance_features,
explanatory_rasters,
features_to_predict,
output_features, output_raster
explanatory_distance_matching,
explanatory_rasters_matching,
output_trained_features,
output_importance_table,
use_raster_values,
number_of_trees,
minimum leaf size.
                                                                                                     minimum_leaf_size,
maximum_level,
sample_size,
                                                                                                      random_sample,
percentage_for_training,
output_classification_table
output_validation_table,
                                                                                                      compensate_sparse_categories
number_validation_runs)
                       __init__(self):
"""Define the tool (tool name is the name of the class)."""
self.label = "Check Rasters for Null Values"
self.description = "Interrogates selected rasters for NULL values"
self.canRunInBackground = True
         def getParameterInfo(self):
    """Define parameter definitions"""
    # First parameter
                      # First parameter
param0 = arcpy.Parameter(
    displayName="Select Rasters",
    name="in_rasters",
    datatype="DERasterDataset",
    parameterType="Required",
    direction="Input",
    multiValue=True)
                       parameters = [param0, param1]
                       """Set whether tool is licensed to execute."""
return True
         def updateParameters(self, parameters):
    """Modify the values and properties of parameters before internal
    validation is performed. This method is called whenever a parameter
    has been changed."""
    return
          def updateMessages(self, parameters):
    """Modify the messages created by internal validation for each tool
    parameter. This method is called after internal validation."""
    return
          def execute(self, parameters, messages):
   lst_rasters = parameters[0].valueAsText
   out_ws = parameters[1].valueAsText
                       for raster in lst_rasters.split(';'):
    nm = getBaseName(raster)
    arcpy.AddMessage("Reading {0}".format(raster))
    outras = IsNull(raster)
    outras.save("{0}/{1}".format(out_ws, nm))
    arcpy.AddMessage("Saving {0}".format(raster))
return
class setRasterNullValuesToOWorkspace(object):
    def init (self):
                                        t_(self):
fine the tool (tool name is the name of the class)."""
label = "Set Null Raster Values to 0 _______
```

```
self.canRunInBackground = True
def getParameterInfo(self):
    """Define parameter definitions"""
          # First parameter
param0 = arcpy.Parameter(
                   displayName="Kaster Works|
name="in_rasters",
datatype="DEWorkspace",
parameterType="Required",
direction="Input")
         # Second parameter
param1 = arcpy.Parameter(
    displayName="Output Workspace",
    name="Out_ws",
    datatype="DEWorkspace",
    parameterType="Required",
    direction="Input")
          return parameters
def isLicensed(self):
    """Set whether tool is licensed to execute."""
    return True
def updateParameters(self, parameters):
    """Modify the values and properties of parameters before internal
    validation is performed. This method is called whenever a parameter
    has been changed."""
 def updateMessages(self, parameters):
    """Modify the messages created by internal validation for each tool
    parameter. This method is called after internal validation."""
def execute(self, parameters, messages):
   in_ws = parameters[0].valueAsText
   out_ws = parameters[1].valueAsText
   arcpy.env.workspace = in_ws
          arcpy.env.workspace = in_ws
rasters = arcpy.ListRasters()
          ror raster in rasters:
    nm = getBaseName(raster)
    arcpy.AddMessage("Reading {0}".format(raster))
    outras = Con(IsNull(raster), 0, raster)
    arcpy.AddMessage("Setting null for {0}".format(raster))
    outras.save("{0}/{1}".format(out_ws, nm))
    arcpy.AddMessage("Saving {0}".format(raster))
return
          __init__(self):
__init__(self):
__init__(self):
"""Define the tool (tool name is the name of the class)."""
self.label = "Set Raster Null Values to 0 - by Selection"
self.description = "Sets all null raster values to 0 in new raster"
self.canRunInBackground = True
parameters = [param0, param1]
def isLicensed(self):
    """Set whether tool is licensed to execute."""
    return True
def updateParameters(self, parameters):
    """Modify the values and properties of parameters before internal
    validation is performed. This method is called whenever a parameter
    has been changed."""
    return
def updateMessages(self, parameters):
    """Modify the messages created by internal validation for each tool
    parameter. This method is called after internal validation."""
def execute(self, parameters, messages):
   lst_rasters = parameters[0].valueAsText
   out_ws = parameters[1].valueAsText
          for raster in lst_rasters.split(';'):
    nm = getBaseName(raster)
    arcpy.AddMessage("Reading {0}".format(raster))
```

```
outras = Con(IsNull(raster), 0, raster)
arcpy.AddMessage("Setting null for {0}".format(raster))
outras.save("{0}/{1}".format(out_ws, nm))
arcpy.AddMessage("Saving {0}".format(raster))
                              __init_(self):
__init
            def getParameterInfo(self):
    # Define parameter definitions
                                               displayName="Input Featur
name="in_features",
tatatype="DEFeatureClass"
parameterType="Required",
direction="Input")
                              # Second parameter
param1 = arcpy.Parameter(
    displayName="Attribute Field",
    name="field",
    datatype="Field",
    parameterType="Required",
    direction="Input")
                              param1.filter.list = ['TEXT', 'LONG']
param1.parameterDependencies = [param0.name]
                               parameters = [param0, param1, param2]
return parameters
            def isLicensed(self):
    """Set whether tool is licensed to execute."""
    return True
             def updateParameters(self, parameters):
             def updateMessages(self, parameters):
    """Modify the messages created by internal validation for each tool
    parameter. This method is called after internal validation."""
    return
                               in_fc = parameters[0].valueAsText
in_field = parameters[1].valueAsText
out_ws = r"C:\Users\hashimotoy\Desktop\ws_prep\raster.gdb"
attributes_ws = r"C:\Users\hashimotoy\Desktop\ws_prep\raster.gdb"
                              arcpy.AddMessage("Converting to raster: " + out_raster + "...")
raster = arcpy.FeatureToRaster_conversion(in_fc, in_field, os.path.join(out_ws, out_raster))
# raster = r"C:\Users\hashimotoy\Desktop\ws_prep\raster.gdb\vri"
values = unique_values(raster, in_field)
# Extract By Attributes each unique value
for value in values:
    if value = "":
        pass
                                                                  e:
    print(" Extracting " + value)
    where_clause = in_field + " = '" + value + "'"
    arcpy.AddMessage(" " + where_clause)
    extract = ExtractByAttributes(raster, where_clause)
    out_nm = (in_field + "_" + value).lower()
    extract.save(os.path.join(attributes_ws, out_nm))
lass renameDataInWorkspace(object):
    def         init (self):
                             def getParameterInfo(self):
    """Define parameter definitions"""
# First parameter
                               param0 = arcpy.Parameter(
displayName="Input Workspace",
```

```
displayName="Select String Parsing Functio
name="parse_function",
datatype="GPString",
parameterType="Required",
direction="Input")
                 # Second parameter
param2 = arcpy.Parameter(
                         displayName="Input string",
name="in_string",
datatype="GPString",
parameterType="Required",
direction="Input")
                          displayName="Output strin
name="out_string",
datatype="GPString",
parameterType="Optional",
direction="Input")
                 parameters = [param0, param1, param2, param3]
       def isLicensed(self):
    """Set whether tool is licensed to execute."""
    return True
       def updateParameters(self, parameters):
   parameters[3].enabled = False
   if parameters[1].value:
      if parameters[1].valueASTEXT == "REPLACE SUBSTRING":
            parameters[3].enabled = True
      else:
       def updateMessages(self, parameters):
    """Modify the messages created by internal validation for each tool
    parameter. This method is called after internal validation."""
       def execute(self, parameters, messages):
    in_ws = parameters[0].valueAsText
    parse_function = parameters[1].valueAsText
    in_string = parameters[2].valueAsText
    out_string = parameters[3].valueAsText
                 env.workspace = in_ws
fcs = arcpy.ListFeatureClasses()
rasters = arcpy.ListRasters()
                 for data in lst:
    nm = getBaseName(data)
    if in_string in nm:
                          arcpy.Rename_management(data, nm.replace(in_string, out_string))
arcpy.AddMessage("\nRenamed " + nm + " to " + nm.replace(in_string, out_string))
arcpy.Rename_management(data, out_string)
arcpy.AddMessage("\nRenamed " + nm + " to " + out_string)
return
class kernelDensityCalculations(object):
    def    init (self):
                 rmeTubensityCartCriatrons(object).
__init_(self):
__init_(self):
"""Define the tool (tool name is the name of the class)."""
self.label = "Calculate Kernel Density"
self.description = "Tool to generalize vector point and line inputs (use the Focal Statistics tool for raster inputs)"
self.canRunInBackground = True
       def getParameterInfo(self):
    """Define parameter definitions"""
    """Define parameter definitions"""
    # First parameter
                          displayName="Input Feature Classes",
name="lst_in_features",
datatype="DEFeatureClass",
parameterType="Required",
direction="Input",
multiValue=True)
                param1 = arcpy.Parameter(
    displayName="Search radius",
    name="search_radius",
    datatype="GPLong",
    parameterType="Required",
    direction="Input")
param1.value = 564 # pi 'r' squared ~ 1sqkm
```

```
parameters = [param0, param1, param2, param3, param4]
# parameters = [param0, param1, param2, param3]
return parameters
def isLicensed(self):
    """Set whether tool is licensed to execute."""
def updateParameters(self, parameters):
    """Modify the values and properties of parameters before internal
    validation is performed. This method is called whenever a parameter
    has been changed."""
    return
def updateMessages(self, parameters):
    """Modify the messages created by internal validation for each tool
    parameter. This method is called after internal validation."""
    return
         lst_inputs = parameters[0].valueAsText
search_radius = parameters[1].valueAsText
cell_size = parameters[2].valueAsText
area_unit_scale_factor = parameters[3].valueAsText
out_ws = parameters[4].valueAsText
         e=cell_size,
                                                                                unit_scale_factor=area_unit_scale_factor,
all_values="DENSITIES",
!="PLANAR")
                                                                                                  =search_radius.
          __init__(self):
"""Define the tool (tool name is the name of the class)."""
self.label = "Split Raster by Attributes"
self.description = "Output rasters are named according to the attribute field and value"
          self.canRunInBackground = False
def getParameterInfo(self):
    """Define parameter definitions"""
    # First parameter
    param0 = arcpy.Parameter(
                  displayName="Input Raster",
name="in_raster",
datatype=["DERasterDataset", "GPRasterLayer"],
parameterType="Required",
direction="Input")
         # Second parameter
param1 = arcpy.Parameter(
    displayName="Attribute Field",
    name="field",
    datatype="Field",
    parameterType="Required",
    direction="Input")
         param1.filter.list = ['TEXT', 'LONG']
param1.parameterDependencies = [param0.name]
                  displayName="Output workspace",
name="out_ws",
datatype="DEWorkspace",
parameterType="Required",
direction="Input"
         parameters = [param0, param1, param2]
return parameters
```

```
def updateParameters(self, parameters):
    """Modify the values and properties of parameters before internal
    validation is performed. This method is called whenever a parameter
    has been changed."""
    return
def updateMessages(self, parameters):
    """Modify the messages created by internal validation for each tool
    parameter. This method is called after internal validation."""
def execute(self, parameters, messages):
   in_raster = parameters[0].valueAsText
   in_field = parameters[1].valueAsText
   out_ws = parameters[2].valueAsText
         values = unique_values(in_raster, in_field)
# Extract By Attributes each unique value
for value in values:
   if value == "":
                              print(" Extracting " + value)
where_clause = in_field + " = '" + value + "'"
arcpy.AddMessage(" " + where_clause)
extract = ExtractByAttributes(in_raster, where_clause)
out_nm = (in_field + "_" + value).lower()
extract.save(os.path.join(out_ws, out_nm))
          __init__(self):

self.label = "Focal Statistics by Circle"

self.description = "Tool to generalize raster inputs (use the Kernel Density tool for vector points and lines)"

self.canRunInBackground = True
def getParameterInfo(self):
         param0 = arcpy.Parameter(
         param1 = arcpy.rarameter(
    displayName="Neighbourhood in map units",
    name="neighbourhood",
    datatype="GPLong",
    parameterType="Required",
    direction="Input")
param1.value = 1200 # Circular neighbourhood of 1.2km to account for center of 400m cell grid
         parameters = [param0, param1, param2, param3]
return parameters
           """Set whether tool is licensed to execute."""
return True
def updateParameters(self, parameters):
    """Modify the values and properties of parameters before internal
    validation is performed. This method is called whenever a parameter
    has been changed."""
def updateMessages(self, parameters):
    """Modify the messages created by internal validation for each tool
    parameter. This method is called after internal validation."""
    return
          rasters = parameters[0].valueAsText
neighbourhood = parameters[1].valueAsText
stats_type = parameters[2].valueAsText
out_ws = parameters[3].valueAsText
           for in_raster in rasters.split(';'):
    arcpy.AddMessage(in_raster)
    nm = getBaseName(in_raster)
                    arcpy.AddMessage("\nProcessing focal statistics for " + nm)
fs = FocalStatistics(in_raster, NbrCircle(neighbourhood, "M
```

```
stats_type,"DATA
fs.save(os.path.join(out_ws, nm))
lass focalStatsByRectangle(object):
                          __init__(self):
self.label = "Focal Statistics by Rectangle"
self.label = "Focal Statistics by Rectangle"
self.description = "Tool to generalize raster inputs (use the Kernel Density tool for vector points and lines)"
self.canRunInBackground = True
          def getParameterInfo(self):
                         parameters = [param0, param1, param2, param3]
return parameters
          def isLicensed(self):
    """Set whether tool is licensed to execute."""
    return True
           def updateParameters(self, parameters):
"""Modify the values and properties of parameters before internal
validation is performed. This method is called whenever a parameter
has been changed."""
return
           def updateMessages(self, parameters):
    """Modify the messages created by internal validation for each tool
    parameter. This method is called after internal validation."""
                         rasters = parameters[0].valueAsText
neighbourhood = parameters[1].valueAsText
stats_type = parameters[2].valueAsText
out_ws = parameters[3].valueAsText
                          for in_raster in rasters.split(';'):
    arcpy.AddMessage(in_raster)
    nm = getBaseName(in_raster)
                                        nm=- getoasename()ncraster)
arcpy.AddMessage("\nProcessing focal statistics for " + nm)
fs = FocalStatistics(in_raster, NbrRectangle(neighbourhood, neighbourhood, "MAP"),
stats_type, "DATA") #
fs.save(os.path.join(out_ws, nm))
lass setRasterValuesTo1(object):
    def    init (self):
                         init_(self):
__init__(self):
_
         def getParameterInfo(self):
    """Define parameter definitions"""
```

```
parameters = [param0, param1]
return parameters
 def isLicensed(self):
    """Set whether tool is licensed to execute."""
    return True
 def updateParameters(self, parameters):
    """Modify the values and properties of parameters before internal
    validation is performed. This method is called whenever a parameter
    has been changed."""
    return
 def updateMessages(self, parameters):
    """Modify the messages created by internal validation for each tool
    parameter. This method is called after internal validation."""
 def execute(self, parameters, messages):
   lst_rasters = parameters[0].valueAsText
   out_ws = parameters[1].valueAsText
           for in_raster in lst_rasters.split(';'):
    nm = getBaseName(in_raster)
    arcpy.AddMessage("\n" + nm)
    out_con = Con(~IsNull(in_raster), 1, 0)
    out_con.save(os.path.join(out_ws, nm))
          ointlokasteruroupbyAttributes(object):

"""Define the tool (tool name is the name of the class)."""

self.label = "Point to Raster - Group by Attributes"

self.label = "Point to Raster - Group by Attributes"

self.description = "Create raster surfaces from one or more attributes in an input point feature class." \

" Output rasters will be prefixed by user defined string" \

" and appended with the field value. The optional 'group by' parameter will output individual rasters" \
" based on attribute."

self.canRunInBackground = True
# Second parameter
param1 = arcpy.Parameter(
            param1.parameterDependencies = [param0.name]
           displayName="Output Workspace",
    name="out_ws",
    datatype="DEWorkspace",
    parameterType="Required",
    direction="Input")
param2.defaultEnvironmentName = "workspace"
# param2.filter.list = ["Local Database"]
           # Fourth parameter
param3 = arcpy.Parameter(
    displayName="Prefix for out raster name",
    name="prefix",
    datatype="(PString",
    parameterType="Required",
    direction="Input")
           # Sixth parameter
param5 = arcpy.Parameter(
    displayName="Cell Size",
    name="cell_size",
    datatype="CPLong",
    parameterType="Input",
    direction="Input")
param5.value = 400
           # Seventh Optional parameter
param6 = arcpy.Parameter(
__displayName="Optional - Group By attribute",
```

```
datatype="Field",
parameterType="Optional",
direction="Input",
multiValue = False)
             param6.parameterDependencies = [param0.name]
    def isLicensed(self):
    """Set whether tool is licensed to execute."""
     def updateParameters(self, parameters):
    """Modify the values and properties of parameters before internal
    validation is performed. This method is called whenever a parameter
    has been changed."""
    return
     def updateMessages(self, parameters):
    """Modify the messages created by internal validation for each tool
    parameter. This method is called after internal validation."""
      def execute(self, parameters, messages):
    # Set environment settings
             in_pts = parameters[0].valueAsText
fields = parameters[1].valueAsText
out_ws = parameters[2].valueAsText
prefix = parameters[3].valueAsText
assignment_type = parameters[4].valueAsText
assignment_type = parameters[5].valueAsText
             cell_size = parameters[5].valueAsText
groupby = parameters[6].valueAsText
             env.workspace = out_ws
             for field in fields.split(';'):
    valField = field
                    lass getColumnIndex(object):
    def __init (self):
             __init__(self):
"""Define the tool (tool name is the name of the class)."""
self.label = "Get Column Index"
self.description = "Using pandas module, read a csv file and get the index of selected columns"
self.canRunInBackground = True
     def getParameterInfo(self):
    """Define parameter definitions"""
    # First parameter
    param0 = arcpy.Parameter(
             # Second parameter
param1 = arcpy.Parameter(
     def isLicensed(self):
    """Set whether tool is licensed to execute."""
    return True
      def updateParameters(self, parameters):
    """Modify the values and properties of parameters before internal
    validation is performed. This method is called whenever a parameter
    has been changed."""
```

```
def updateMessages(self, parameters):
    """Modify the messages created by internal validation for each tool
    parameter. This method is called after internal validation."""
    return
def execute(self, parameters, messages):
   import pandas as pd
        in_csv = parameters[0].valueAsText
fields = parameters[1].valueAsText
        arcpy.AddMessage("\n\nGetting indices for selected columns...\n\n") \\ csv = pd.read\_csv(in\_csv)
        for field in fields.split(';'):
   index = csv.columns.get_loc(field)
   arcpy.AddMessage("\n " + field + " : " + str(index) + "\n")
        def getParameterInfo(self):
    """Define parameter definitions"""
        # First parameter defin:
# First parameter
param0 = arcpy.Parameter(
                displayName="Input Featur
name="in_features",
datatype="GPFeatureLayer"
parameterType="Required",
direction="Input")
        # Second parameter
param1 = arcpy.Parameter(
        # Third parameter
param2 = arcpy.Parameter(
       return parameters
def isLicensed(self):
    """Set whether tool is licensed to execute."""
    return True
def updateParameters(self, parameters):
    """Modify the values and properties of parameters before internal
    validation is performed. This method is called whenever a parameter
    has been changed."""
def updateMessages(self, parameters):
    """Modify the messages created by internal validation for each tool
    parameter. This method is called after internal validation."""
    return
def execute(self, parameters, messages):
   pts = parameters[0].valueAsText
   out_ws = parameters[1].valueAsText
   clim_var = parameters[2].valueAsText
   ws_clim = parameters[3].valueAsText
        base_nm = getBaseName(pts)
         lst_merge = []
               _merge = []
i in range(2007,2018):
out_fc = os.path.join(out_ws,base_nm + "clim_" + str(i))
arcpy.AddMessage("\nRunning value extraction for " + out_fc)
in_pts = arcpy.CopyFeatures_management(pts, out_fc)
checkFieldExists(in_pts, "year_txt", "TEXT")
lst_clim = []
for var in clim_var.split(';'):
    clim_raster = os.path.join(ws_clim,"clim_" + var + "_" + str(i))
    arcpy.AddMessage(" " + clim_raster)
    lst_clim.append([clim_raster, var])
```

```
arcpy.CalculateField_management(out_fc, "year_txt", "'" + str(i) + "'")
ExtractMultiValuesToPoints(out_fc, lst_clim)
Ist_merge.append(out_fc)
arcpy.AddMessage("\n\nCompleted extractions by year. \n\nMerging into final dataset...")
arcpy.Merge_management(lst_merge, os.path.join(out_ws, base_nm + "_annual_clim"))
                      altereditectoristy)Surfaces(go.)
__init__(self):
__init__
def getParameterInfo(self):
    """Define parameter definitions"""
                      """Define parame
# First parameter
param0 = arcpy.Parameter(
    displayName="Input Directory of CSV response files",
                                            displayName="Input Direct
name="in_csv",
datatype="DEWorkspace",
parameterType="Required",
direction="Input")
                      # Second parameter
param1 = arcpy.Parameter(
    displayName="Geodatabase of fishnets for each ecosection",
    name="ws_fishnet",
    datatype="DEWorkspace",
    parameterType="Required",
    direction="Input")
                      # Third parameter
param2 = arcpy.Parameter(
param2 = arcyName="Qutput Workspace",
                        # Fourth parameter
param3 = arcpy.Parameter(
                                           displayName="Snap Raster",
name="snap_raster",
datatype="DERasterDataset",
parameterType="Required",
direction="Input")
                        return parameters
 def isLicensed(self):
    """Set whether tool is licensed to execute."""
    return True
 def updateParameters(self, parameters):
    """Modify the values and properties of parameters before internal
    validation is performed. This method is called whenever a parameter
has been changed."""
  def updateMessages(self, parameters):
    """Modify the messages created by internal validation for each tool
    parameter. This method is called after internal validation."""
    return
 def execute(self, parameters, messages):
    ws_response = parameters[0].valueAsText
    ws_fishnet = parameters[1].valueAsText
    ws_output = parameters[2].valueAsText
    snap_raster = parameters[3].valueAsText
                     # Loop through the CSV files of predicted densities
# For each CSV file, get the basename, join it to the appropriate ecosection fishnet
# Output a raster
env.workspace = ws_response
arcpy.env.snapRaster = snap_raster
files = arcpy.ListFiles("".csv")
for f in files:
    f_nm = getBaseName(f)
    arcpy.AddMessage(f_nm)
    eco = f_nm[7:1.0]
    sp = f_nm[11:len(f_nm)]
    arcpy.AddMessage("Preparing " + sp.upper() + " for " + eco + "...")
    fishnet_lyr = arcpy.MakeFeatureLayer_management(os.path.join(ws_fishnet,eco), "fishnet_lyr")
    joined = arcpy.AddMossage("n\nagmanagement(fishnet_lyr, "id_fishnet", f, "id_fishnet")
    arcpy.FeatureToRaster_conversion(joined, sp, os.path.join(ws_output, eco + "_" + sp), cell_size=400)
arcpy.AddMessage("\n\nCompleted rasterized densities. Check results in " + ws_output)
return
```

#### APPENDIX 4 - R Scripts - Project Repository

Latest updates to the project will be uploaded to <a href="https://github.com/Yuhash/iws">https://github.com/Yuhash/iws</a>. The repository contains the base survey data inputs as well as an interactive <a href="Jupyter notebook">Jupyter notebook</a> running the R-kernel that documents the major script processes in the workflow. For optimal viewing (without interactive code functionality) in Notebook Viewer go to <a href="https://nbviewer.jupyter.org/github/Yuhash/iws/blob/master/IWS">https://nbviewer.jupyter.org/github/Yuhash/iws/blob/master/IWS</a> Notebook.jpynb.

#### **Workflow Model**

```
# For data pre-processing steps please refer to the Github repository and the Jupyter Notebook
# https://journal.r-project.org/archive/2009/RJ-2009-013/RJ-2009-013.pdf
# cforest model script loops through ecosections and species/species group
# mclapply is a parallelizing function available only on MAC OS. On Windows change 'mclapply' to 'lapply'
useMaxCoreCount <- 2
nTree <- 5000
mTry <- 5
seed <- 42
PREDICT <- TRUE
gitHubRoot <- "/Volumes/Black/Yuri/iws/inputs/by_location/"</pre>
outputRoot <- "/Users/hashimotoy/Desktop/iws-results/"</pre>
fgcRequire <- function(pkg) {</pre>
  if(pkg %in% rownames(installed.packages()) == FALSE) {
    install.packages(pkg)
    library(pkg)
fgcRequire("ggplot2")
fgcRequire("icesTAF")
fgcRequire("party")
fgcRequire("Hmisc")
fgcRequire("lattice")
fgcRequire("beepr")
#install.packages("parallel") # is base, should not be updated (MacOS / R 3.6.0 / RStudio Version
1.2.1335 )
#install.packages("tools") # is base, should not be updated (MacOS / R 3.6.0 / RStudio Version 1.2.1335
library(parallel)
library(lattice)
library(Hmisc)
library(party)
library(tools)
library(icesTAF)
library(beepr)
wd <- gitHubRoot
outwd <- paste0(outputRoot,paste0("results-seed,"-year-studyarea/"))</pre>
set.seed(seed)
mkdir(outwd)
setwd(wd)
getwd()
locList <- c("BAU", "NAU", "NEL", "CHP", "BUB", "CAB", "WCU", "QUL")
.c("amwi","buff","bago","bwte","cago","gwte","mall","nsho","rndu","scau","dabblers","divers","cavity"," sp_div","sp_tot")
mclapply(spList, function(sp) {
  lapply(locList, function(loc) {
    setwd(wd)
    mydata <- read.csv(paste0(wd,"id1_years_",loc,".csv"))</pre>
```

```
mydata[ 2:ncol(mydata) ] <- lapply(mydata[2:ncol(mydata)], as.numeric)</pre>
    #sum(is.na(mydata))
    fishnet <- read.csv(paste0(gitHubRoot, "fishnet_", loc, ".csv"))</pre>
    fishnet[ 2:ncol(fishnet) ] <- lapply(fishnet[2:ncol(fishnet)], as.numeric)</pre>
    setwd(outwd) # DOING OUTPUT NOW
    rdsFilename <- paste0("cf_", sp, "_", loc, ".rds")
vi_fn <- paste0("VI-",sp,"_",loc,".csv")</pre>
    print(paste("START Processing cforest for:", sp))
    cf <- if (
      !file.exists(paste0(outwd, rdsFilename))
      #colnames(mydata)[colnames(mydata)==sp] <- "sp" RENAME HACK</pre>
      cf <- cforest(
        eval(parse(text = paste0(sp,
                                  "~ aspect + slope + alr + pa + cec_urban + cec_shrubland + ",
                                  " cec_mixed_forest + cec_grassland + cec_broadleaf + cec_needleleaf +
                                  " fwa_1 + fwa_2 + fwa_3 + fwa_4 + fwa_10ha_more + str_s + str_m + ",
                                     fwa r + fwa_w + dra_u + ",
                                  # bec_zn + # removed becase: model used for CAB on the eco section
basis
                                  #eco + # removed because we have bined by eco
                                  "norm_dd5 + norm_cmd + pas_wt + tave04 +'
                                  "shorecx_10plus_100000 + shorecx_10less_100000"
        )))
        data = mydata,
        controls = cforest_unbiased(ntree = nTree, mtry = mTry)
      print(paste("cforest done:", sp))
      saveRDS(cf, paste0(outwd, rdsFilename))
      cf
    } else {
      readRDS(paste0(outwd, rdsFilename))
    if (!file.exists(paste0(outwd, vi_fn))
    ) {
      vi <- varimp(cf, conditional = FALSE, OOB = TRUE)</pre>
      # write.csv(vi, paste(fn, ".csv", sep = ""), row.names = FALSE)
      df_vi <- as.data.frame(vi)</pre>
      write.csv(df_vi,vi_fn);
    #### Not incorporated in batch: deriving OOB, Response, Probability
    if (PREDICT) {
      ofResp <- paste0(outwd,sp,"-response_",loc,".csv")
cf_response <- predict(cf, newdata = fishnet, OOB = TRUE, type = 'response')
      df_result <- data.frame(fishnet$id_fishnet, cf_response)</pre>
      write.csv(df_result, ofResp)
    print(paste("COMPLETED Anaylsis of:", sp))
 })
}
 mc.cores = min(useMaxCoreCount,detectCores())
#-----
# Ranking and Plots for varimp
# Written if RDS has to be loaded
setwd(wd)
rdss <- list.files(wd, pattern ="*rds$")</pre>
lapply(rdss, function(f){
 cf <- readRDS(f)
```

```
fn <- substr(basename(f), 4, nchar(basename(f)) - 4)</pre>
 eco <- substr(fn, nchar(fn) - 2, nchar(fn))
sp <- sub(paste0("_", eco), "", fn)
vi_fn <- paste0(wd, "vi_", fn, ".csv")
if (!file.exists(vi_fn)) {</pre>
    vi <- varimp(cf, conditional = FALSE, OOB = TRUE)
df_vi <- as.data.frame(vi)</pre>
    v <- as.vector(df_vi$vi)</pre>
    vi <- varimp(cf, conditional = FALSE, OOB = TRUE)</pre>
    df_vi <- as.data.frame(vi)</pre>
    df_vi$X <- names(vi)</pre>
    df_vi$eco <- eco
    df_vi$sp <- sp
    # Add metadata fields
    df_vi <- as.data.frame(vi)</pre>
    df_vi$X <- names(vi)</pre>
    df_vi$eco <- eco
    df_vi$sp <- sp
    write.csv(df_vi, vi_fn, row.names = FALSE)
    # Rank
    df vi <- df vi %>%
      mutate(rank = dense_rank(desc(df_vi$vi)))
    colnames(df_vi)[colnames(df_vi) == "rank"] <-paste(eco, sp, sep = "_")</pre>
    rank_vi <- subset(df_vi, select = -c(vi, eco, sp))</pre>
    write.csv(df_vi, vi_fn, row.names = FALSE)
})
## Create plots
library(tools)
setwd(wd)
varimpFiles <- list.files(wd, pattern = "^vi.*csv$")</pre>
lapply(varimpFiles, function(f) {
 fn <- substr(basename(f),4,nchar(basename(f))-4)</pre>
 print(fn)
 eco <- substr(fn,nchar(fn)-2,nchar(fn))</pre>
 sp <- sub(paste0("_",eco),"", fn)</pre>
 print(paste("Starting plot", f))
 fn <- file_path_sans_ext(basename(f))</pre>
 title <- paste(toupper(sp),eco)
 print(title)
 df_vi <- read.csv(f)</pre>
 v <- as.vector(df_vi$vi)</pre>
 names(v) <- as.vector(df_vi$X)</pre>
 print(names(v))
 xlab = "varImp", main = title)
 dev.copy(png, paste(wd, paste("varimp_", title, ".png", sep = ""), sep = "/"))
 dev.off()
 print(paste("plot done:", fn))
print("ALL DONE")
R Snippets - Data Manipulation
# Create a data frame from a list of csv files converted to a list of data frames merged by common column
lst <- list.files(wd, pattern ="^rank.*csv$" )</pre>
merged <- do.call("cbind", lapply(ranked, read.csv))</pre>
#-----
# Flat correlation matrix with P and r values
library(tidyr)
library(tibble)
library(Hmisc)
```

```
flat_cor_mat <- function(cor_r, cor_p){</pre>
 #This function provides a simple formatting of a correlation matrix
 #into a table with 4 columns containing :
 # Column 1 : row names (variable 1 for the correlation test)
 # Column 2 : column names (variable 2 for the correlation test)
 # Column 3 : the correlation coefficients
 # Column 4 : the p-values of the correlations
 library(tidyr)
 library(tibble)
 cor_r <- rownames_to_column(as.data.frame(cor_r), var = "row")</pre>
 cor_r <- gather(cor_r, column, cor, -1)</pre>
 cor_p <- rownames_to_column(as.data.frame(cor_p), var = "row")</pre>
 cor_p <- gather(cor_p, column, p, -1)</pre>
 cor_p_matrix <- left_join(cor_r, cor_p, by = c("row", "column"))</pre>
 cor_p_matrix
library(corrplot)
cor <- rcorr(as.matrix(mtcars[, 1:7]))</pre>
my_cor_matrix <- flat_cor_mat(cor$r, cor$P)</pre>
head(my_cor_matrix)
corr <- rcorr(as.matrix(df))</pre>
corrplot(corr$r, method='number', number.cex= 12/ncol(df_sub))#To alter font size
# Check for NA values
df %>%
 select if(function(x) any(is.na(x))) %>%
 summarise_each(funs(sum(is.na(.)))) -> extra_NA
# Set NA values to 0
df[is.na(df)] <- 0</pre>
# The function complete.cases() returns a logical vector indicating which cases are complete.
# list rows of data that have missing values
df[!complete.cases(df),]
# ------
# Check which version of R is running
Sys.getenv("R_ARCH")
# "/i386" or "/x64"
# UPDATE PACKAGES
update.packages(checkBuilt=TRUE, ask=FALSE)
# -----
mypath <- path
multmerge = function(mypath){
 filenames=list.files(path=mypath, full.names=TRUE)
 \label{lapsequation} {\tt datalist = lapply(filenames, function(x)\{read.csv(file=x,header=T)\})}
 Reduce(function(x,y) {merge(x,y)}, datalist)}
mymergeddata = multmerge(path)
# UPDATE R Version
# installing/loading the package:
if(!require(installr)) {
 install.packages("installr"); require(installr)} #load / install+load installr
library(installr)
# using the package:
updateR()
```

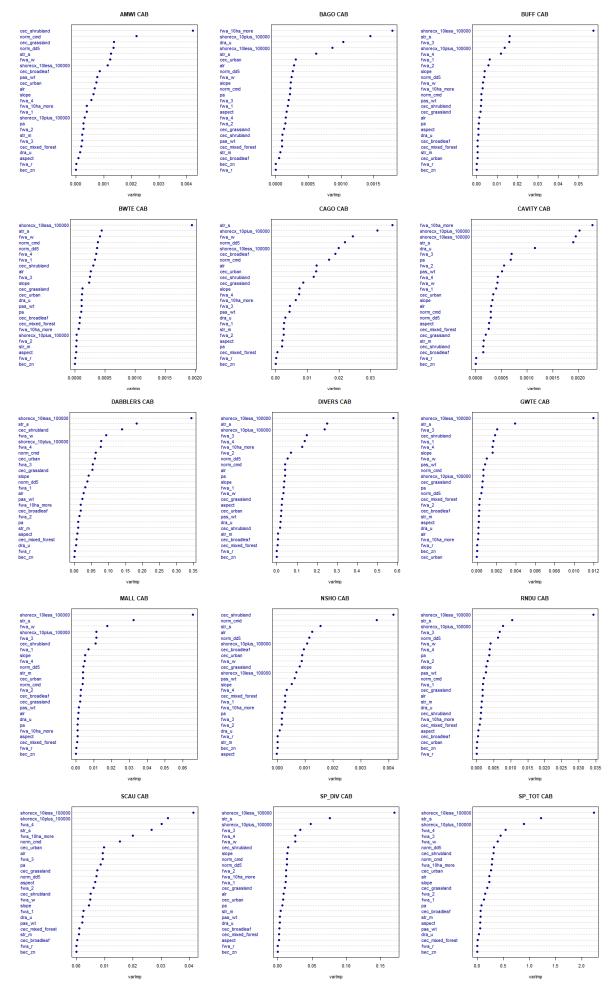
```
# # Rename a column in R
colnames(data)[colnames(data)=="old_name"] <- "new_name</pre>
sp_start <- grep("dabblers", colnames(df))
sp_end <- grep("tot_sp", colnames(df))</pre>
print(sp_start)
print(sp_end)
sp <- colnames(df)[sp_start:sp_end]</pre>
var_start <- sp_end + 1</pre>
var_end <- ncol(df)
print(var_start)</pre>
print(var_end)
int_eco <- grep("eco", colnames(df))</pre>
int_trans <- grep("trans_id", colnames(df))
int_bec_zn <- grep("bec_zone", colnames(df))
int_bec_sz <- grep("bec_sz", colnames(df))</pre>
for (i in 1:length(sp)){
  nm <- sp[i]
  print(nm)
  int = grep(nm, colnames(df))
  data <- df[c(int,int_eco,int_trans,int_bec_zn, int_bec_sz,var_start:var_end)]</pre>
  colnames(data)[1]<- "sp"</pre>
  write.csv(data, paste(outwd, paste("rf_",scale, "_",nm,".csv", sep = ""), sep = "/"), row.names =
FALSE)
}
```

# **APPENDIX 5 – Variable Importance Plots**

**Babine Upland** 

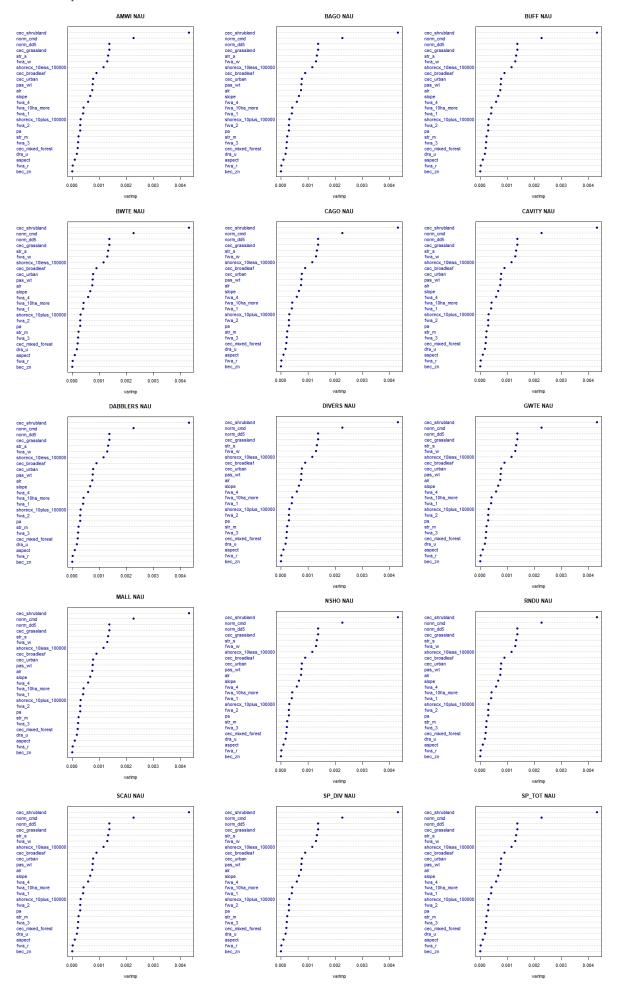
# **Bulkley Basin**

#### **Cariboo Basin**



## **Chilcotin Plateau**

#### **Nazko Upland**



## **Nechako Lowland**

**Quesnel Lowland** 

Western Chilcotin Uupland

# **APPENDIX 6 – Species Distribution Maps**

**American Wigeon** 

**Barrow's Goldeneye** 

**Bufflehead** 

**Canda Goose** 

**Green-winged Teal** 

Pennsylvania State University, MGIS Program