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Current breeding distributions and predicted range shifts under climate change in two subspecies of Black-tailed Godwits in Asia

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Abstract

Habitat loss and shifts associated with climate change threaten global biodiversity, with impacts likely to be most pronounced at high latitudes. With the disappearance of the tundra breeding habitats, migratory shorebirds that breed at these high latitudes are likely to be even more vulnerable to climate change than those in temperate regions. We examined this idea using new distributional information on two subspecies of Black-tailed Godwits *Limosa limosa* in Asia: the northerly, bog-breeding *L. l. boharii* and the more southerly, steppe-breeding *L. l. melanuroides*. Based on breeding locations of tagged and molecularly assayed birds, we modelled the current breeding distributions of the two subspecies with species distribution models, tested those models for robustness, and then used them to predict climatically suitable breeding ranges in 2070 according to bioclimatic variables and different climate change scenarios. Our models were robust and showed that climate change is expected to push *boharii* into the northern rim of the Eurasian continent. *Melanuroides* is also expected to shift northward, stopping in the Yablonovyy and Stanovoy Ranges, and breeding elevation is expected to increase. Climatically suitable breeding habitat ranges would shrink to 16% and 11% of the currently estimated ranges of *boharii* and *melanuroides*, respectively. Overall, this study provides the first predictions for the future distributions of two little-known Black-tailed Godwit subspecies and highlights the importance of factoring in shifts in bird distribution when designing climate-proof conservation strategies.

Keywords: climate change, East Asian-Australasian Flyway, IPCC, Maxent, *Limosa limosa*, shorebirds, species distribution modelling

Introduction

The global temperature has risen by 0.08°C per decade since the Industrial Revolution in the late 19th century (Lindsey & Dahlman 2020, Root *et al.* 2003). This warming has been particularly severe in the Arctic (Collins *et al.* 2013, Houghton *et al.* 2001, Kåresdotter *et al.* 2021, Serreze & Barry 2011). It has led to advanced snowmelt (Chen *et al.* 2021, Zhong *et al.* 2021) and earlier emergence of insects (Saalfeld *et al.* 2021), forcing animals such as shorebirds to breed and migrate earlier (Lameris *et al.* 2017), with one cost being lower adult survival (Rakhimberdiev *et al.* 2018). In general, many organisms have advanced their timing of reproduction to counter the adverse effects of the warming climate (Jenni & Kéry 2003, Root *et al.* 2003, Walther *et al.* 2002); the taxa that have advanced the least have shown the most substantial population declines (Lameris *et al.* 2018, Ross *et al.* 2018).

To mitigate the detrimental effects of global warming, species are expected to leave their current breeding range and move to more climatically suitable breeding habitats. Indeed, birds that breed at

low latitudes in Eurasia and North America tend to move to the cooler north (Hitch & Leberg 2007, McClure *et al.* 2012, Thomas & Lennon 1999, Virkkala *et al.* 2018, Zuckerberg *et al.* 2009).

However, in the Arctic, northward shifts are constrained by the presence of the Arctic Ocean; this results in a compressed breeding range and smaller population sizes, in addition to longer migrations in terms of distance and time (Parmesan 2006, Rehfish & Crick 2003, Wauchope *et al.* 2017).

Black-tailed Godwits *Limosa limosa* (hereafter “godwits”) are migratory shorebirds that breed across the temperate and boreal zones of the Palearctic (Prater *et al.* 1977, Zhu *et al.* 2020). In Europe, *L. l. limosa* shows little evidence of change in the timing of homeward migration and breeding (Kleijn *et al.* 2010, Schroeder *et al.* 2012), but advancing nesting dates were detected in *L. l. islandica* (Gill *et al.* 2014). With the gradual increase in average April temperatures over the past century, observations have also suggested a northward expansion of *L. l. limosa* in NW Russia (Popov & Starikov 2015). Studies on the East Asian godwits are scarce, with little knowledge on *L. l. melanuroides* in the East Asian-Australasian Flyway and a recent discovery of the hitherto unknown subspecies *L. l. bohaili* in China (Zhu *et al.* 2021a). The previously defined breeding range of *melanuroides*, which included seven fragmented sites from ca. 45 °N to ca. 68 °N (Birdlife International 2018), is consequently being reconsidered because some of those areas are occupied by *bohaili*. Recent tracking revealed two areas in the Arctic region of the Russian Far East where *bohaili* breeds (59 °N to ca. 65 °N). Breeding *melanuroides* are mainly found in the temperate steppe of East Asia (45 °N to ca. 52 °N, Zhu *et al.* 2021b).

The latitudinal contrast between the two East Asian godwit populations provides a suitable system to investigate and compare how climate change may differentially impact the breeding range. On the basis of species distribution models (SDMs, Pacifici *et al.* 2015), we delineate the breeding ranges of the two subspecies *bohaili* and *melanuroides* and examine which environmental factors determine these distributions. We then use this understanding to evaluate how global warming will affect the currently climatically suitable breeding habitats of the two subspecies; to do this, we project two greenhouse emissions scenarios in the models, capturing a range of possibilities by using both a moderate and a pessimistic climate change trend (IPCC, 2013).

Materials and methods

Breeding locations

We obtained breeding ground locations for genetically-confirmed *bohaili* individuals by satellite tracking (Zhu *et al.* 2021b), considering stationary locations recorded during June–August as breeding ground locations (Fig. 1). Stationary locations were identified when a transmitter recorded a speed of ≈ 0 km/h and a local elevation of ≈ 0 m. Breeding ground locations of *melanuroides* were obtained

from our previous genetic work (Zhu *et al.* 2021a). We used known genetic differences between the subspecies to confirm that the individuals breeding in locations identified through satellite tracking versus those breeding in locations identified through published records did indeed belong to the two subspecies (see Zhu *et al.* 2021a). This yielded 60 known breeding ground locations for *bohail* (2015-2018) and 41 for *melanuroides* (1993-2016) to parameterize our species distribution models (SDMs).

Environmental variables

To capture the current breeding habitat in our species distribution models, we obtained (a) 19 bioclimatic variables and (b) surface elevation data from WorldClim (www.worldclim.org, v 2.1, 1970-2000), along with (c) six breeding habitat features variables (e.g. vegetation type, snow cover) from Earthenv (<http://www.earthenv.org/landcover>), (d) the average normalized difference vegetation index (NDVI) for May–July during 2014–2018 from NOAA (<https://www.ncei.noaa.gov/data/avhrr-land-normalized-difference-vegetation-index/access/>), and (e) the Global Human Footprint and Influence indexes which summarize data on human population density, land use, infrastructure and access (<https://sedac.ciesin.columbia.edu/data/collection/wildareas-v2>). See Table S1 for a complete overview and detailed information on all variables. As the distributional record of the *bohail* subspecies was collected more recently than the 1970-2000 bioclimatic data, we verified that the variable that contributed the most to predicting its distribution (Table S2), i.e., temperature seasonality (Bio4), from 1970-2000 to 2010-2018 increased by 1% only.

To predict the breeding range for both subspecies in 2070, we obtained data from the Intergovernmental Panel on Climate Change (IPCC) AR5 from the Global Climate Model (GCM) downscaled data portal (http://www.ccafs-climate.org/data_spatial_downscaling/). We selected two different emissions scenarios to take the uncertainty of the future into account: (1) the Representative Concentration Pathway 4.5 (RCP), a moderate scenario in which greenhouse gas emissions peak around 2040 and then decline, and (2) the more pessimistic RCP 8.5, in which emissions are predicted to continue to grow throughout the 21st century (<https://ar5-syr.ipcc.ch/index.php>). For each of the two emissions scenarios, we selected two different Global Climate Models (GCMs), GFDL-CM3 and MIROC-ESM (Wauchope *et al.* 2017), that predict future bioclimatic conditions, resulting in two sets of predicted bioclimatic conditions for each subspecies (Table S1).

In considering potential godwit breeding habitats, we constrained the range of all environmental variables to above 40°N on the Eurasian continent (Birdlife International 2018). Though there may also be suitable breeding habitats in Alaska or northern Canada, ca. 3000 km to the east of our easternmost records, the likelihood that these areas will become occupied by godwits within the next five decades appears low due to the lack of suitable steppingstone habitats between the current

breeding locations and North America. In Qgis 3.8, all environmental variables were resampled to 2.5 arc-min spatial resolution (ca. 4.5 km at the equator) using the nearest neighbour interpolation. Any missing values were filled with the mean values within five pixels of the missing pixel using the maximum distance method, followed by repeating the smoothing iterations five times after each interpolation. The mean NDVI for a given pixel was calculated using the raster calculator.

Species Distribution Modelling

We constructed current and future SDMs for these two subspecies using maximum entropy modelling (Maxent, version 3.3.4). Maxent is a machine-learning algorithm that uses presence-only data to determine the predicted suitability of local conditions for a given species (Phillips *et al.* 2006; Phillips & Dudik, 2008). Several studies have suggested that Maxent has good predictive power for birds (Hu & Liu, 2014, Wauchope *et al.* 2017). We used occurrence records only when the subspecies present was genetically confirmed, and the breeding status was known because this approach to predicting subspecies distribution has proven more accurate than traditional methods that rely solely on occurrence records (Ikeda *et al.* 2017).

To model the current breeding habitats of the two subspecies, we first tested Pearson's correlations on all 29 environmental variables. Each highly correlated variable pair ($|r| \geq 80\%$) that gave higher values when testing the correlations against the remaining ones were retained. Next, we determined the final selection of the variables by their contributions and jackknife tests. Those variables that contributed less than 1% to the model were removed (Table S2). As a result, 14 variables were selected for the final analyses: annual mean temperature (°C), mean diurnal temperature range (°C), temperature seasonality (SD * 100), mean temperature of the wettest quarter (°C), mean temperature of the driest quarter (°C), mean temperature of the coldest quarter (°C), annual precipitation (mm), precipitation seasonality (variation coefficient), precipitation of the driest quarter (mm), human footprint (range: 1–100, a higher value indicates higher anthropogenic impacts), herbaceous plants coverage (%), shrub coverage (%), elevation (m) and NDVI (see correlation matrix in Table S3). To test the robustness of our models in predicting the current range, we used a model that contained only the western occurrence records of *bohaili* to see whether the model would correctly predict the eastern range more than 1000 km away (Fig 1). We were unable to do the same for *melanuroides* because the subspecies did not have such widely dispersed breeding occurrence records.

The godwit breeding habitat in 2070 was estimated by changing the current climatic conditions as predicted in two IPCC greenhouse gas emissions scenarios (RCP 4.5 & 8.5). Since there were no non-climatic variables available in the future scenarios, e.g., landcover, anthropogenic impacts or NDVI, we only used the 19 bioclimatic variables and elevation data for these models. Therefore, the

predictions *only* estimate the future change in climatically suitable breeding habitat ranges and do not deal with other global changes. Consequently, the predictions are relatively conservative.

In all models, 70% of the occurrence records were randomly assigned to the training dataset and 30% to the testing dataset. The “Remove duplicate presence records” option was used to avoid the inclusion of duplicated records in a grid cell. The model performance was evaluated using the mean AUC score (area under the receiving operating curve, mean \pm SD) and threshold-based evaluation methods (Phillips *et al.* 2006). The model output was summarized into a logistic suitability value ranging from 0: unsuitable to 1: suitable. In our case, we defined three suitability levels: low: 0–0.29, medium: 0.3–0.59 and high: 0.6–1. We ran twenty bootstrap replicates for all models and selected the point-wise mean of the 20 output grids to draw the predicted breeding habitat ranges. Results from the two Global Climate Models (GCMs) in each emissions scenario were averaged, yielding two consensus grids for each subspecies. Lastly, we overlaid the predicted current breeding habitat ranges and the predicted climatically suitable breeding habitat for both subspecies in 2070 onto maps in Qgis 3.18 (Fig. 2).

Spatial analysis

The values of the 14 variables that best explained the current range of the breeding distribution were extracted at the occurrence locations using the function “sample raster values” in Qgis 3.18. We used Student t-tests to explore differences between the subspecies for these environmental and climatic variables. To assess the predicted shift in breeding habitat range in 2070, we used Student t-tests to compare mean breeding latitude and elevation in 2020 with those predicted for each subspecies in 2070. Since the highest logistic suitability in 2070 differed between subspecies (see Results), we constructed the polygons with values greater than 0.15 (for *bohail*) and 0.3 (*melanuroides*) in Qgis 3.18. We randomly extracted 200 latitude and elevation datums from each modelled habitat range. We calculated the sizes of the breeding habitats with low, medium and high suitability in current and future scenarios using the “r.report” function. Statistics were performed in R v 3.6.0 (R Core Team 2018) then visualized in the package *ggplot2*.

Results

Current breeding habitat ranges

Confirming the known breeding distribution of *bohail*, the model identified two isolated suitable breeding areas, both located around the Arctic Circle in the Russian Far East (61–68°N; Fig. 1). The western area was in the Sakha Republic, extending from 100°E westwards to the confluence of the

Vilyuy and Lena Rivers and ending in the western foothills of the Verkhoyansk Range (128°E; Fig. 1). The eastern area was between the Momskiy and Kolyma Ranges, in the Kolyma River basin of Magadan Oblast (150–155°E; Fig. 1). For *melanuroides*, the range of suitable breeding habitat was further south (42–52°N). It covered a vast swathe of Asia's temperate interior, stretching from Lake Baikal to the Heilongjiang River and the Sea of Japan (90–140°E; Fig. 1). Some coastal areas in the Russian Far East were also found to be suitable for *melanuroides*, namely Sakhalin Island and an area south of the Kolyma Range (142–160°E; Fig. 1). The AUC scores and threshold statistics suggest that these models provided reasonable predictions for both *bohail* ($AUC_{\text{training}} = 0.996 \pm 0.001$; $AUC_{\text{test}} = 0.993 \pm 0.001$; all threshold statistics $p < 10^{-13}$) and *melanuroides* ($AUC_{\text{training}} = 0.991 \pm 0.007$; $AUC_{\text{test}} = 0.957 \pm 0.049$; all threshold statistics $p < 10^{-5}$). We verified the robustness of our model predictions by testing whether our models would also predict the eastern breeding area of *bohail* by using only the occurrence records from the western part. Encouragingly, the models successfully predicted the eastern area (Fig. 1). The final and test models indicated that the eastern breeding area is suitable for *bohail* (the highest suitabilities were 0.86 and 0.77, respectively), with breeding range sizes of $292 \times 10^3 \text{ km}^2$, $304 \times 10^3 \text{ km}^2$, respectively.

Concerning habitat features, *bohail* bred near rivers in swampy areas with low shrub coverage and low anthropogenic impact (Fig. 3, S1). *Melanuroides* bred mainly in pasture areas close to lakes and with more human activity (Fig. 3, S1). The annual mean temperature and precipitation in the breeding habitats for *bohail* were significantly lower than for *melanuroides* (Fig 3). Within the modelling range where the logistic suitability values were higher than 0, 64% ($8.4 \times 10^5 \text{ km}^2$) and 63% ($4.2 \times 10^6 \text{ km}^2$) of grid cells were predicted to have low breeding suitability for *bohail* and *melanuroides*, respectively. In comparison, 30% ($3.9 \times 10^5 \text{ km}^2$) and 32% ($2.1 \times 10^6 \text{ km}^2$) of grid cells had medium suitability, and 7% ($0.9 \times 10^4 \text{ km}^2$) and 5% ($0.4 \times 10^6 \text{ km}^2$) of grid cells had high suitability (Fig. 4, Table S4).

Climatically suitable breeding habitat ranges in 2070

The models predicted a dramatic range shift and decline of climatically suitable breeding habitats by 2070 for both subspecies under RCP 4.5 and RCP 8.5 scenarios (Fig 2). The mean latitude of *bohail*'s breeding habitat range is predicted to shift northward from the current $64.7 \pm 7.6 \text{ }^\circ\text{N}$ to $71.9 \pm 2.1 \text{ }^\circ\text{N}$ ($p < 0.0001$) in RCP 4.5 and $72.6 \pm 2.3 \text{ }^\circ\text{N}$ ($p < 0.0001$) in RCP 8.5 (Fig 4). In other words, the only remaining climatically suitable breeding habitat would lie at the northern margin of the continent along the Arctic Ocean coast in the Russian Far East (68–70°N, 60–180°E). With this latitudinal shift, the breeding elevation of *bohail* would show a significant decrease from the current $295 \pm 112 \text{ m}$ to $33 \pm 31 \text{ m}$ ($p < 0.0001$) in RCP 4.5 and $40 \pm 28 \text{ m}$ ($p < 0.0001$) in RCP 8.5 (Fig 4). In RCP 4.5, the habitat range would decline to 11% of the current extent ($1.5 \times 10^5 \text{ km}^2$), of which 96% ($1.4 \times 10^5 \text{ km}^2$)

of grid cells were predicted to have low suitability, 5% (6742 km²) of grid cells were predicted to have medium suitability, and no high suitability habitats were expected to remain. In RCP 8.5, the habitat range declined to 9% of the current area, and all areas (1.2*10⁵ km²) were predicted to have low suitability (Fig 4, Table S4).

The climatically suitable breeding habitat range of *melanuroides* is predicted to be split by the Yablonovy and Stanovoy Ranges, with the central part of Mongolia and the south of Lake Baikal no longer suitable for *melanuroides* (Fig. 1, 2). The habitat range would shift to the north of the Heilongjiang River basin, toward the foothills of the Stanovoy Range. The mean latitude of the habitat range is predicted to shift significantly from the current 47.1 ± 3.3 °N to 52.5 ± 2 °N ($p < 0.0001$) in RCP 4.5 and 54.5 ± 2.1 °N ($p < 0.0001$) in RCP 8.5 (Fig 4). The current mean elevation (787 ± 490 m) is predicted to remain the same as in RCP 4.5 (845 ± 304 m, $p = 0.16$), but increased significantly in RCP 8.5 (957 ± 272 m, $p < 0.0001$, Fig 4). Under RCP 4.5, the habitat range declined to 16% of the current area (1.1*10⁶ km²), of which 71% (7.6*10⁵ km²) of grid cells were predicted to have low suitability, 29% (3.2*10⁵ km²) of grid cells were predicted to have medium suitability, and no high suitability habitats were expected to remain. In RCP 8.5, the habitat range declined to 15% of the current area (1*10⁶ km²), with 90% (9.5*10⁵ km²) of the habitat range predicted to have low suitability, 10 % (1*10⁵ km²) of grid cells predicted to have medium suitability, and no high suitability habitats remained (Fig 4, Table S4).

Discussion

The modelling shows that, consistent with common patterns, the breeding ranges of both subspecies of Black-tailed Godwits are projected to shift northward due to a warming climate. More crucial, however, is our finding that both subspecies are also expected to suffer a large decline in climatically suitable breeding habitats because of climate change. The breeding range and climatic suitability for the more northerly breeding subspecies *bohaili* would suffer the most significant decline because there is no land further to the north for the birds to use. In addition, due to the lack of climatically suitable breeding habitat in the north, the southerly subspecies *melanuroides* is expected to be forced onto higher ground to breed. Our results highlight the importance of factoring in future shifts in bird distribution when designing climate-proof conservation strategies.

Current breeding ranges

Currently, *bohaili* godwits are known to breed in Russia along the Arctic circle in two isolated regions east and west of the Verkhoyansk Mountain Range. Our model was able to predict the eastern region based on records from the western region alone (Fig 1.), demonstrating the model's robustness and the similarity of the environmental conditions in both regions. Degtyarev *et al.* (2020) describe a

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population of ca. 11,000 godwits breeding in an area stretching from 62°N to 68°N and from 100°E to 125°E. This area closely matches our prediction of the western breeding range of *bohaili*. The measured body dimensions of members of this breeding population also match those described for *bohaili* (E. Shemyakin & P. Tomkovich, pers. comm., Zhu *et al.* 2021b), which further adds to our confidence that our model correctly predicts the current breeding range for *bohaili*.

The current breeding range of *melanuroides* is not well-known, but the latest summary assessments (BirdLife International 2018) assume that *melanuroides* breeds relatively far north (60–65°N) and thus overlaps with *bohaili*. However, this is not consistent with either our or previous results. Our tracking and modelling efforts do not support the idea that *melanuroides* breeds north of 60°N, and neither do the findings of Lappo *et al.* (2012), while the modelled breeding range for *melanuroides* is significantly further to the south (42–52°N) and therefore does not overlap with *bohaili* at all. Instead, the identified breeding ranges of the two subspecies are separate from one another and differ substantially in landscape, environment and climate: while *bohaili* breeds in swampy natural lowland with lower annual mean temperatures and precipitation, *melanuroides* mainly inhabits warmer and wetter semi-natural pastures near lakes (Fig. 1). Intriguingly, *bohaili*'s current medium and highly suitable breeding habitat range is less than one-fifth of *melanuroides*' (Table S4). A potential explanation for this difference is that there is less open area for *bohaili* to breed due to the presence of Boreal (Taiga) forest and mountain ranges (Fig. 5, S1). In contrast, the much larger breeding range of *melanuroides* is not restricted in this way, since it does not include forests and mountains (Fig. S1).

Projected breeding ranges under climate change scenarios

The modelling results indicate that the climatically suitable breeding ranges of *bohaili* and *melanuroides* may shrink by 89% and 84% while also shifting northward, assuming no adaptive change in habitat preferences. Under both climate change scenarios, the modelled climatically suitable breeding habitat range of *bohaili* is concentrated in the estuaries along the Arctic coast. Here, further northward expansion is constrained by the absence of land (Fig. 2). Meanwhile, *melanuroides* is projected to leave the temperate pastures due to significant loss of climatically suitable habitat in central-eastern Mongolia and NE China (45–50°N). We observed a projected increase in their breeding elevation, with the southern foothills of the Stanovoy Range becoming suitable for *melanuroides* (Fig. 2). However, the latitudinal shift is not as drastic as predicted for *bohaili* because, climatically, most areas to the north of the Stanovoy Range will remain too cold and dry to suit *melanuroides*. Moreover, the northward shift required to reach the next potentially suitable breeding habitat would be nearly impossible to make in only five decades; there are ca. 1900 km of continuous boreal forest separating the current breeding area from the basins of the Vilyuy and Lena rivers in the

north (Fig. 1, Fig. 5). It is important to reemphasize that these modelled ranges take only projected future climate into account. If other changes (e.g., climate-driven changes in land or vegetation cover, water bodies) were also factored in, the actual future distribution of these two subspecies would probably be even smaller than we have projected.

Consistent with the predictions given here, continental godwits (*L. l. limosa*) in NW Russia have already been observed to shift their breeding range northward from 60 °N in 1970 to 64 °N in 2012 (Popov & Starikov 2015). In addition, godwits started colonizing the basins of the Vilyuy and Lena rivers from the south in the early 1990s (Degtyarev *et al.* 2020). Meanwhile, godwits have been observed to vacate southerly regions like the Selenga Delta near Lake Baikal, after a period of continuous population decline, which might be due to the prolonged periods of low water since the late 2000s (I. Fefelov, pers. comm).

The bigger picture

While these two East Asian godwit subspecies do not suffer from as much direct anthropogenic impact on their breeding grounds as the Continental European subspecies (Kruk *et al.* 1997, Kleijn *et al.* 2010), they appear to be threatened by hunting (Zhu *et al.* 2021b) and deteriorating staging sites in the Yellow Sea (Chan *et al.* 2019, Piersma *et al.* 2016, Yang *et al.* 2011). We now show that the two subspecies will also likely have to cope with climate change (1) heavily diminishing the suitability of their current breeding habitat and (2) shifting and considerably shrinking the climatically suitable breeding habitat range. As a result, we expect a decline in the adult survival and reproductive success of those godwits that remain in their current breeding range (Jetz & Dobson 2007, Saalfeld *et al.* 2021).

This expected reduction in suitable breeding habitat contrasts interestingly with observations made for a different subspecies, the Icelandic godwit *Limosa limosa islandica*. At the start of the 20th century, the breeding area for *islandica* was constrained to Iceland's relatively warm southwest corner, and the population numbered only a few thousand (Gill *et al.* 2007). As the climate warmed and many areas were converted to farmland, large parts of the island became more suitable for breeding, and the *islandica* population consequently grew to ca. 47,000 (Gunnarsson *et al.* 2005, Gunnarsson *et al.* 2006). Nevertheless, the impact of a changing climate has also been negative; the original primary breeding area in the southwest of Iceland now appears to have become too warm, leading to reduced reproductive success in that area (Alves *et al.* 2019). These *islandica* observations neatly illustrate that climate change can benefit a godwit population by increasing the suitable breeding habitat, and it can negatively impact a godwit population by decreasing the suitable breeding habitat. Though both effects have been detected in the case of *islandica*, our results show that only the negative effect is expected for *bohii* and *melanuroides*.

In the longer term, we expect the populations of *bohaili* and *melanuroides* to become smaller than at present. We also expect the adverse effects of climate change to be greater for *bohaili* than for *melanuroides*, because *bohaili* has a substantially smaller range and potentially small population size, precisely the two factors identified as affecting the susceptibility and extinction risk of populations in response to climate change (Beyer & Manica 2020, Gaston & Blackburn 1996). In addition, *bohaili* would face more extreme weather events (e.g., drought and wildfires, see Karuk *et al.* 2021, Yasunari *et al.* 2021) while experiencing both higher predation rates and increased competition for breeding space as more species and populations become concentrated in the Arctic rim (Killengreen *et al.* 2007, Layton-Matthews *et al.* 2020, Vallejos *et al.* 2020). These factors might partially explain the decline of wintering waterbirds breeding in Siberia compared to temperate Asia (Sung *et al.* 2021). Our results highlight the importance of thinking ahead and factoring in shifts in bird distribution when designing climate-proof conservation strategies to salvage our heritage of migratory birds and, ultimately, the ecosystems we all belong to.

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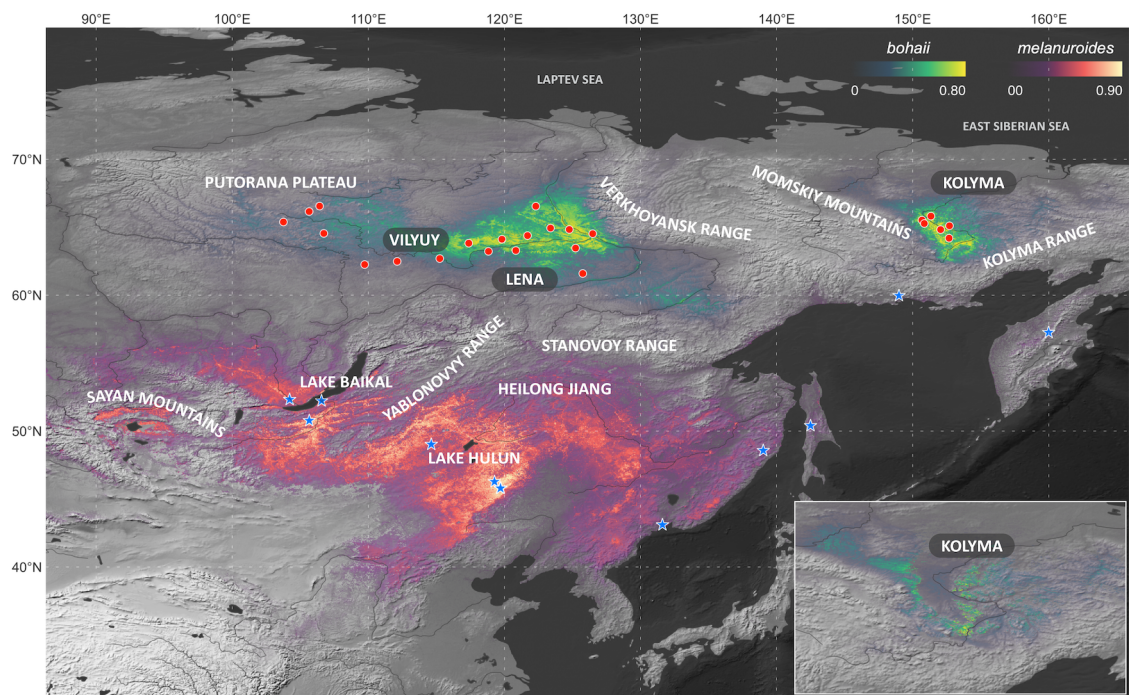
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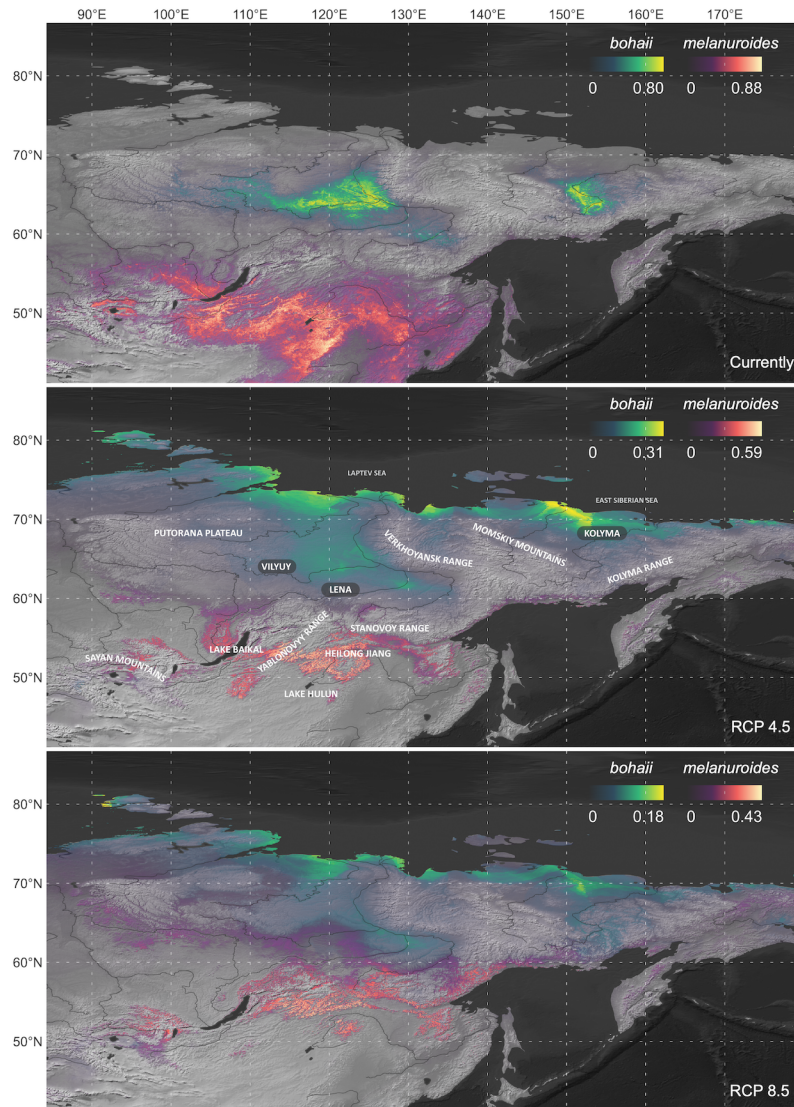
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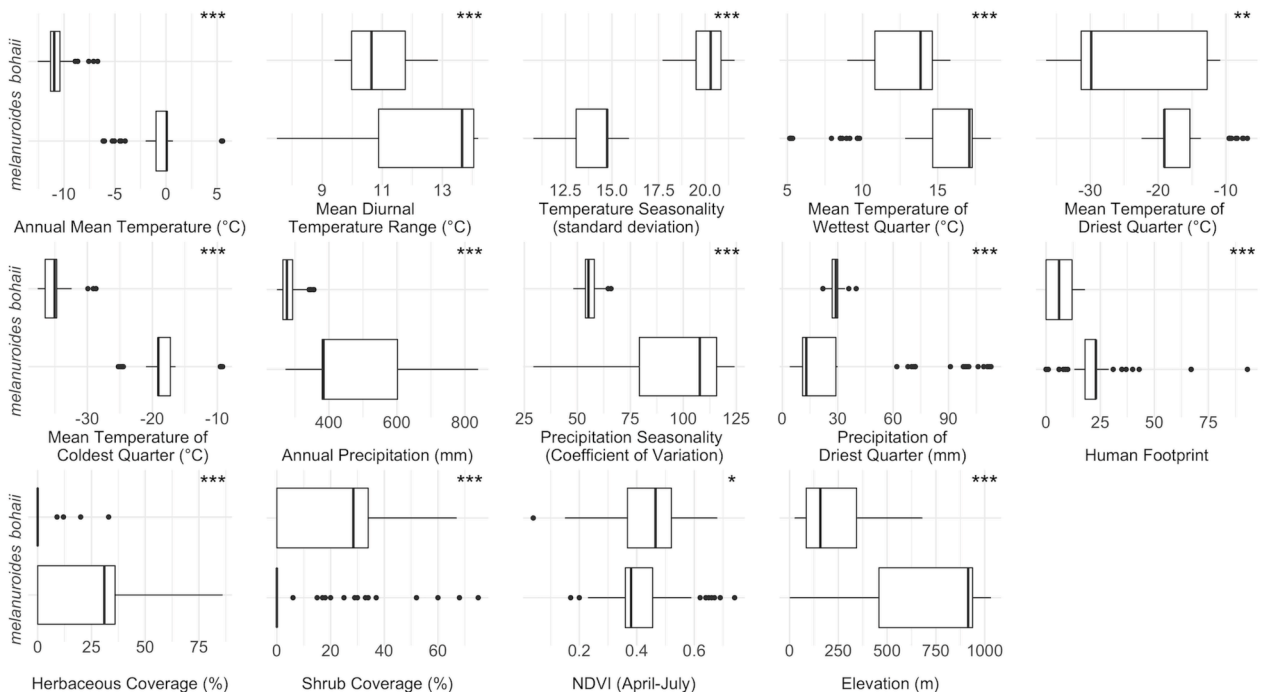
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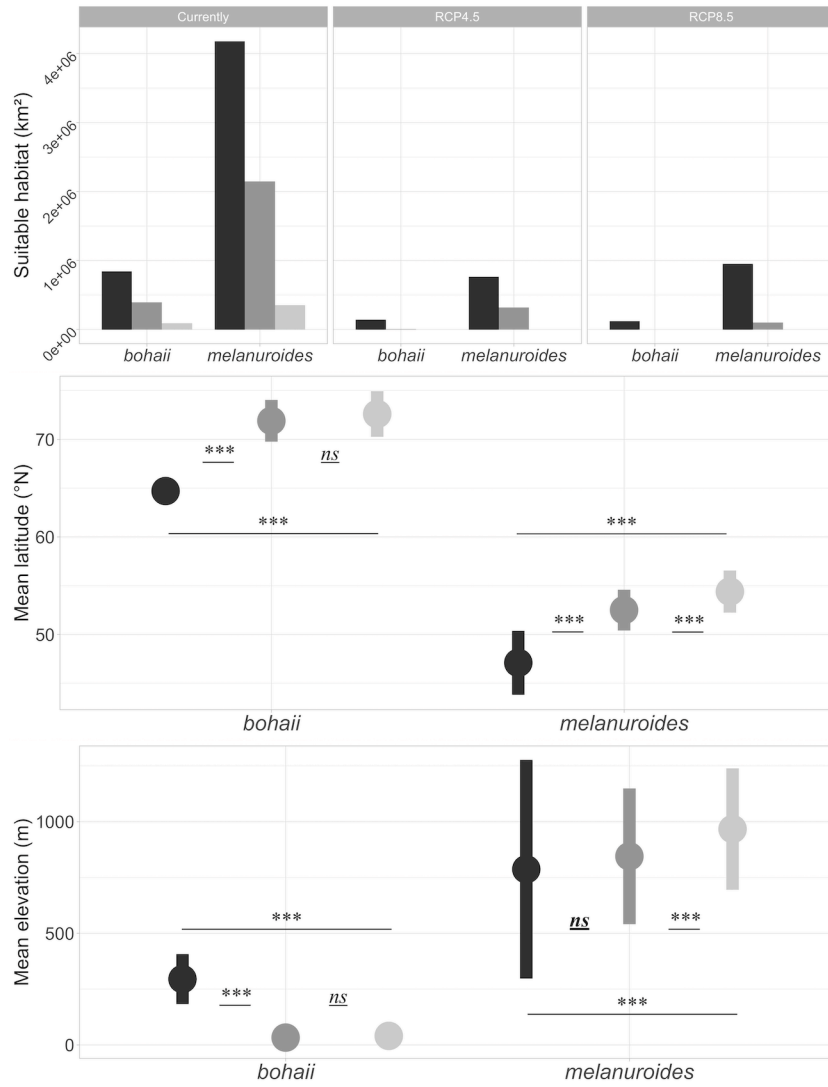
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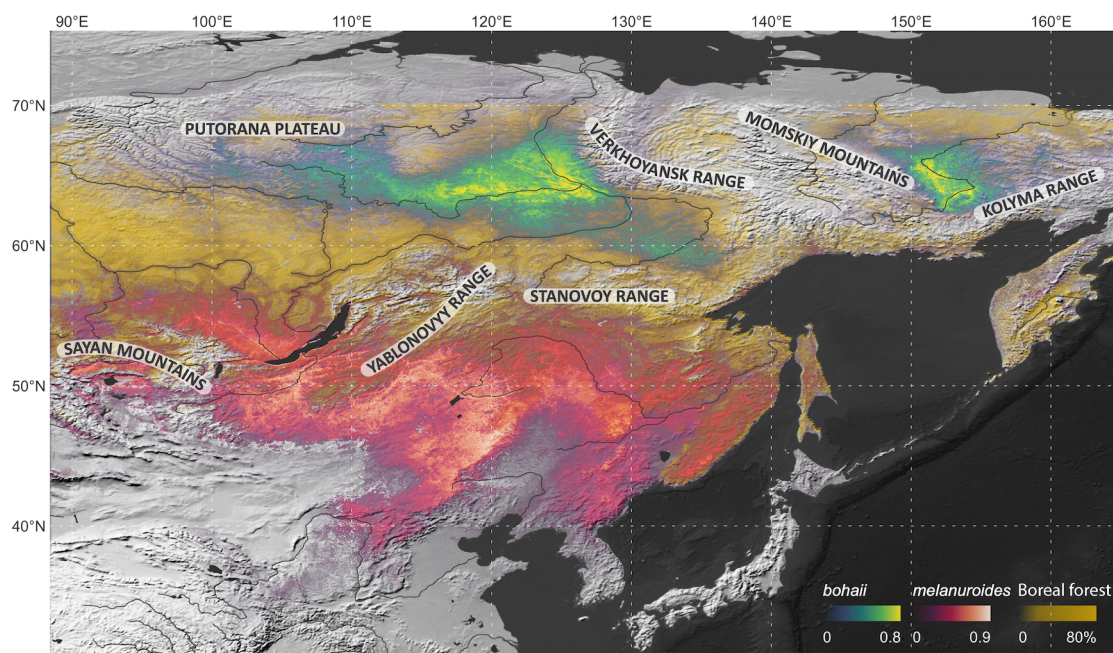
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GCB_16308_6.Fig3_Zhu et al_GCB_resized.tiff



GCB_16308_7.Fig4_Zhu et al_GCB_resized.tiff



GCB_16308_8.Fig5_Zhu et al_GCB.tiff