

The heat is on: Rising temperatures alter when, how, and where  
large terrestrial mammals move

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# 1 Effects of temperature on movement rates

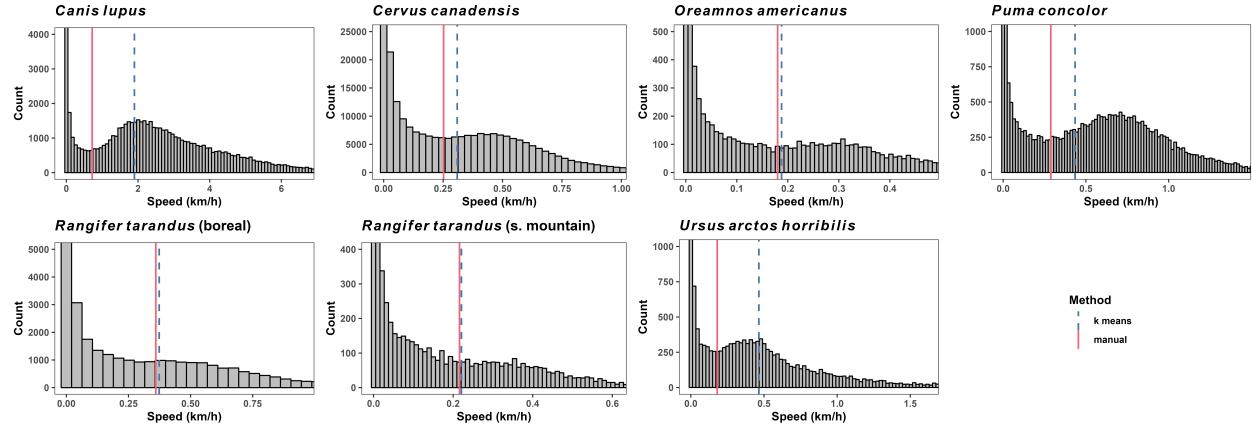


Figure B1: Histograms of each species' estimated speed. Continuous red lines indicate the values used to determine whether an animal was moving or not, which were determined visually using the inflection points of the histograms. Dashed blue lines indicate the minimum speed corresponding to a moving animal as determined by *k*-means algorithms with 2 clusters. For ease of readability, the x axes range from 0 to the 0.99 quantile, while the y axes range from 0 to one fortieth of the total number of estimates.

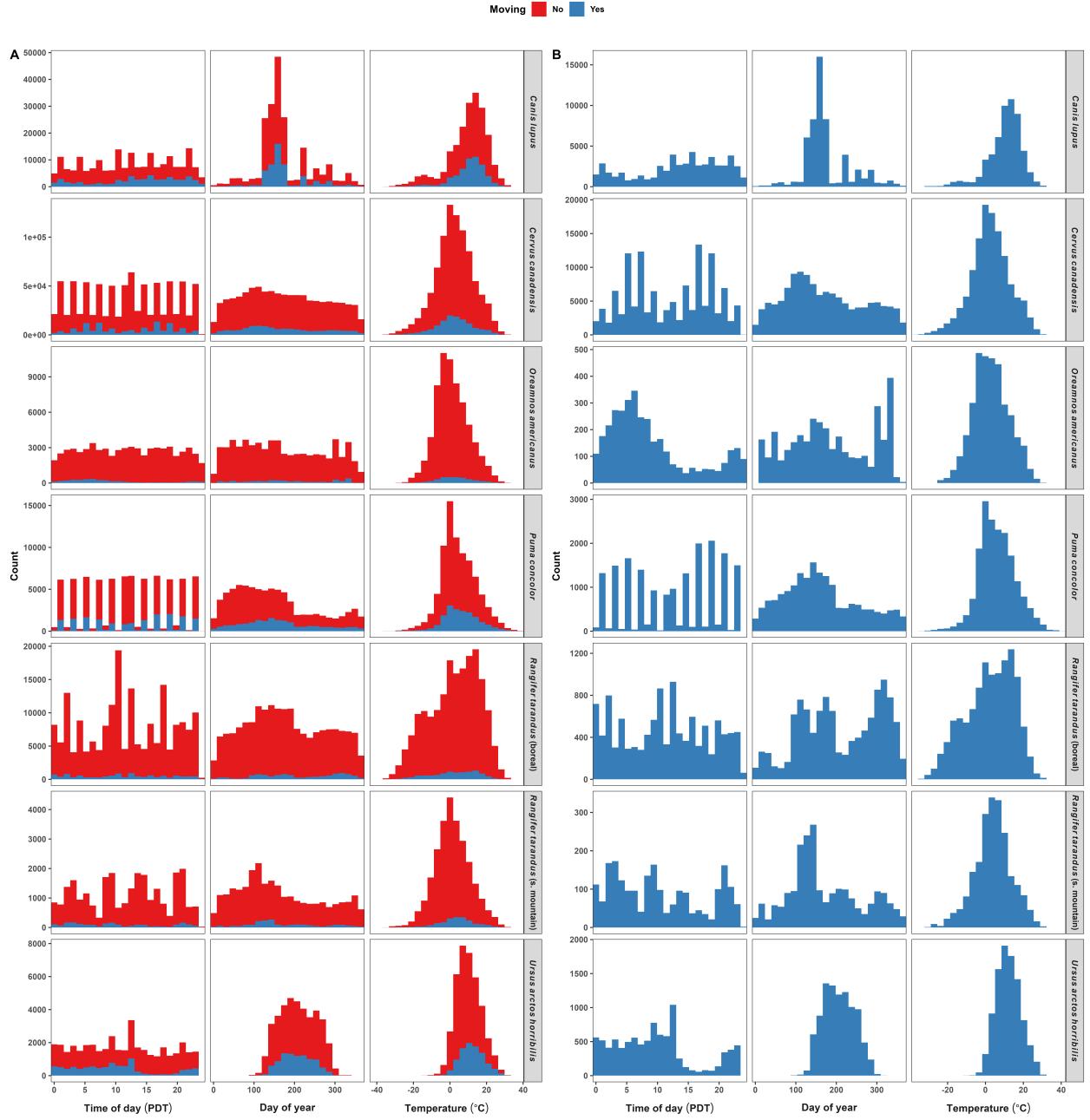


Figure B2: Histograms of the number of states (**A**) and speed estimates (**B**) over time of day (Pacific Daylight Time, PDT), day of year (Julian date), and air temperature.

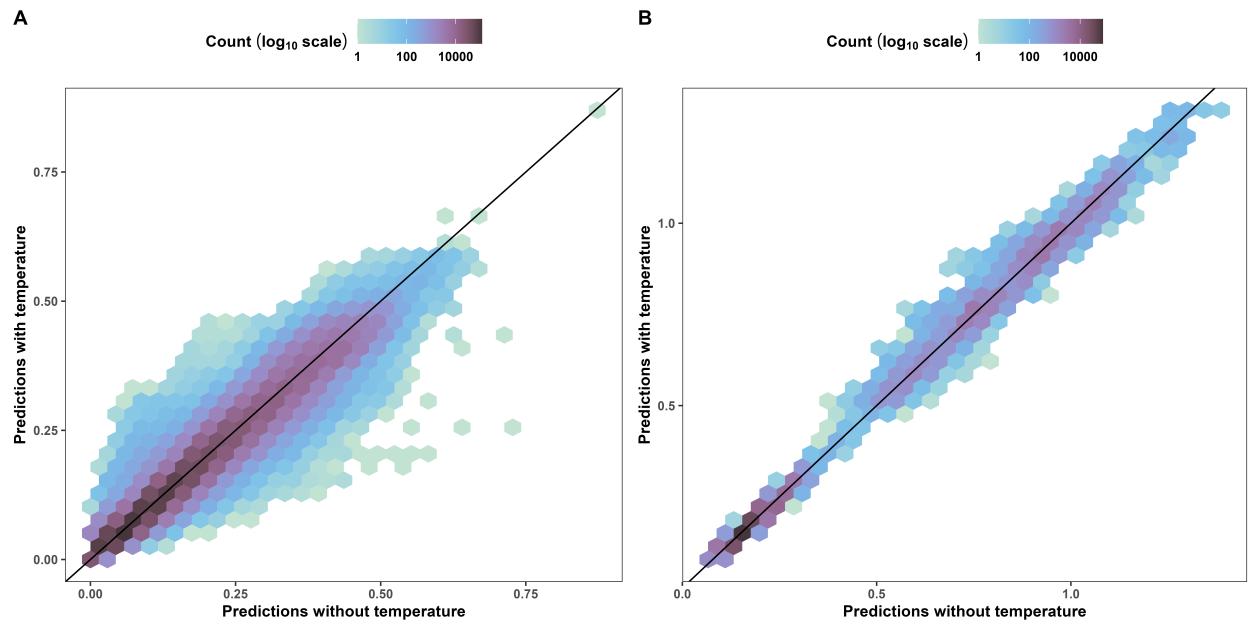


Figure B3: Hexplot of the fitted values from the HGAMs with and without including temperature for the probability of movement (**A**) and speed when moving (**B**).

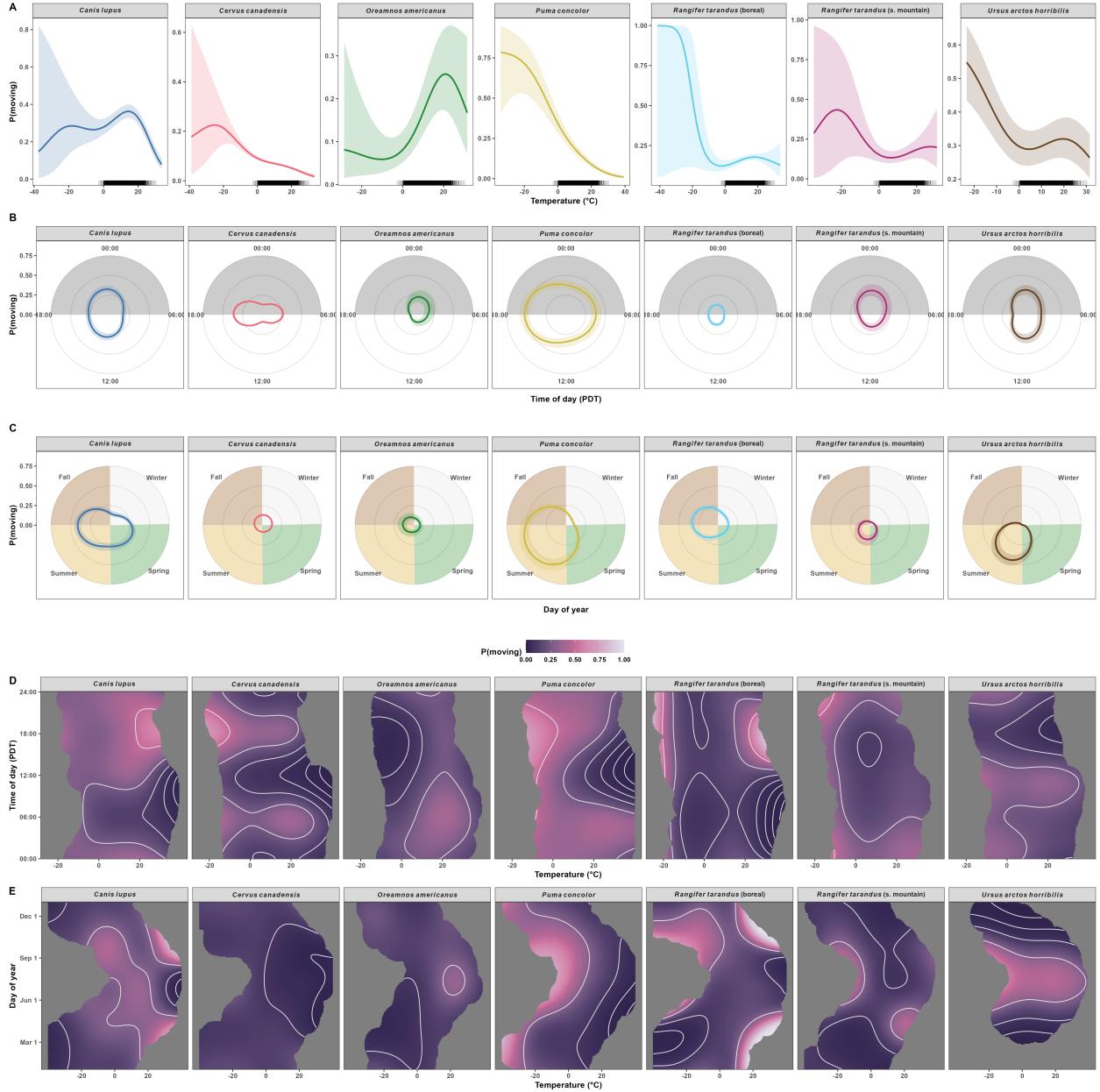


Figure B4: **A.** Estimated effects of temperature on each species' probability of moving on June 1<sup>st</sup> at 12:00, Pacific Daylight Time (PDT). The rug plot indicates each species' data on June 1<sup>st</sup>. **B.** Estimated effects of time of day on each species' probability of moving on June 1<sup>st</sup> at 0°C. The grey area indicates evening and night (hours between 18:00 and 6:00). **C.** Estimated effects of day of year on each species' probability of moving at 12:00 with a temperature of 0°C. The year is divided into the four seasons: winter (white), spring (green), summer (gold), and fall (brown). In panels A-C, ribbons indicate 95% Bayesian Credible Intervals, and the sampling rate was post-stratified to  $\Delta t = 1$  hour for all species. **D.** Effects of time of day and temperature on species' probability of moving on June 1<sup>st</sup> at 12:00 PDT. Surfaces in panels D and E extend to 10% of the range away from each datum.

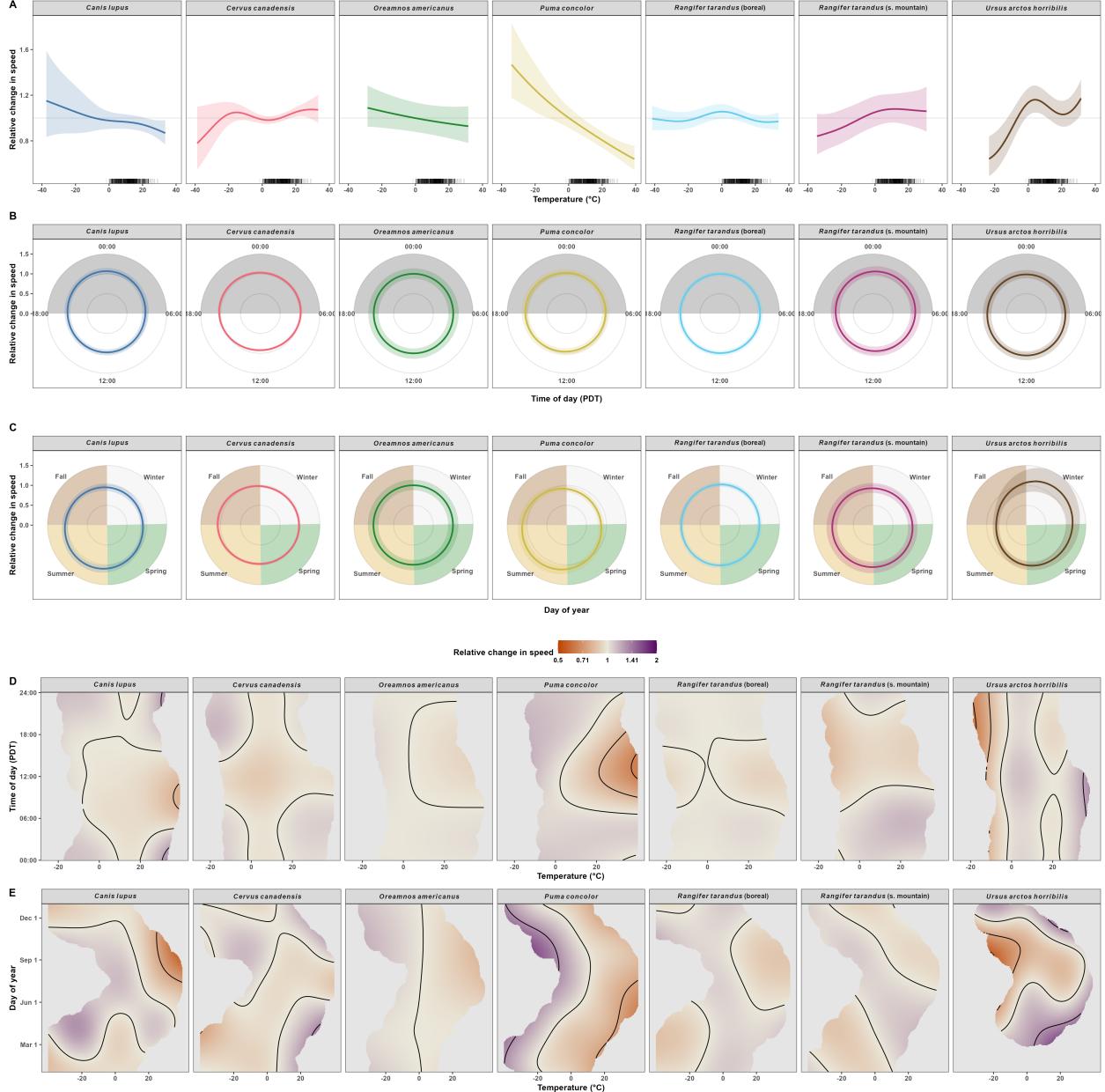


Figure B5: **A.** Estimated effects of temperature on each species' speed when moving on June 1<sup>st</sup> at 12:00, Pacific Daylight Time (PDT). The rug plot indicates each species' data points where an animal was moving on June 1<sup>st</sup>. **B.** Estimated effects of time of day on each species' speed when moving on June 1<sup>st</sup> at 0°C. The grey area indicates evening and night (hours between 18:00 and 6:00). **C.** Estimated effects of day of year on each species' speed when moving at 12:00 with a temperature of 0°C. The year is divided into the four seasons: winter (white), spring (green), summer (gold), and fall (brown). In panels A-C, ribbons indicate 95% Bayesian Credible Intervals, and the sampling rate was post-stratified to  $\Delta t = 1$  hour for all species. **D.** Effects of time of day and temperature on species' speed when moving on June 1<sup>st</sup>. **E.** Effects of day of year and temperature on species' speed when moving, if the animal was moving at 12:00 PM PDT. Surfaces extend to 10% of the range away from each datum. The color bar is on the log<sub>2</sub> scale to help visualize patterns in doubling.

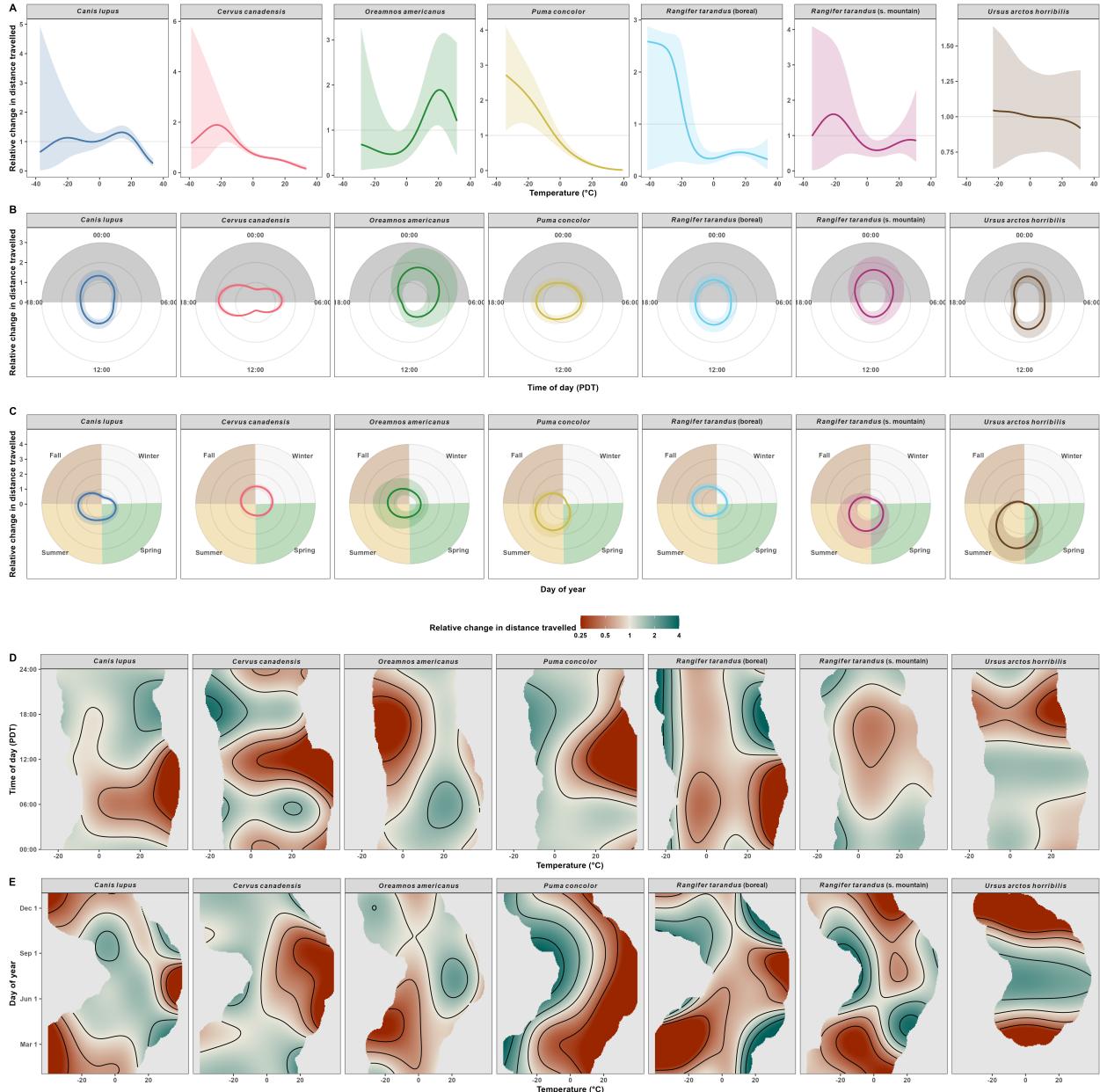


Figure B6: **A.** Estimated effects of temperature on each species' distance traveled on June 1<sup>st</sup> at 12:00, Pacific Daylight Time (PDT). The rug plot indicates each species' data points where an animal was moving on June 1<sup>st</sup>. **B.** Estimated effects of time of day on each species' distance traveled on June 1<sup>st</sup> at 0°C. The grey area indicates evening and night (hours between 18:00 and 6:00). **C.** Estimated effects of day of year on each species' distance traveled at 12:00 with a temperature of 0°C. The year is divided into the four seasons: winter (white), spring (green), summer (gold), and fall (brown). In panels A-C, ribbons indicate 95% Bayesian Credible Intervals, and the sampling rate was post-stratified to  $\Delta t = 1$  hour for all species. **D.** Effects of time of day and temperature on species' distance traveled on June 1<sup>st</sup> at 12:00 PDT. **E.** Effects of day of year and temperature on species' distance traveled at 12:00 PDT. Surfaces extend to 10% of the range away from each datum. The color bar is on the  $\log_2$  scale to help visualize patterns in doubling.

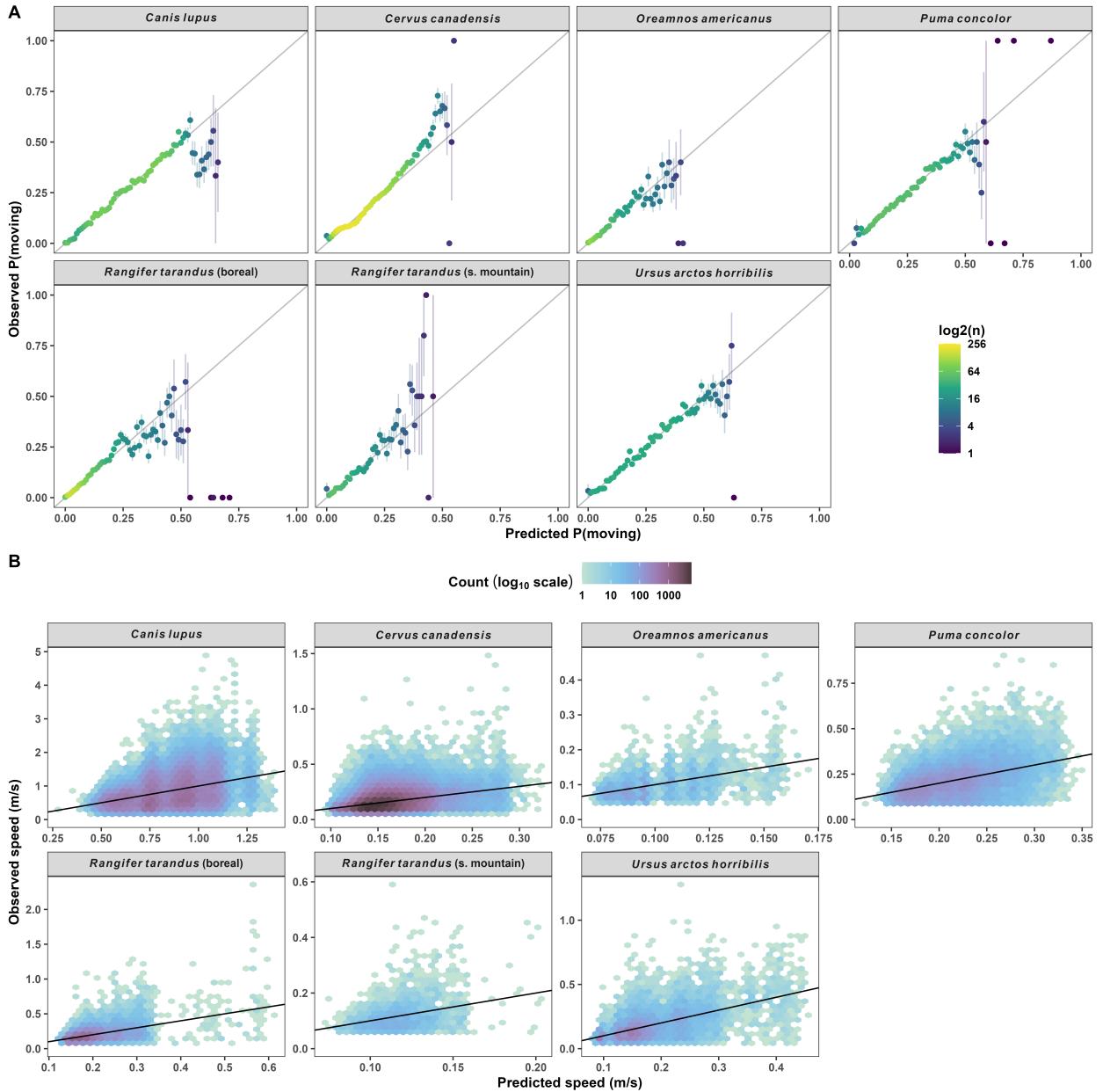


Figure B7: Relationships between the observed and predicted values for probability of movement (**A**) and speed given that an animal was moving (**B**). The color in panel **A** indicates the number of points used to calculate the estimate (on the  $\log_2$  scale), while the grey lines in both panels indicate perfect prediction (i.e., the 1:1 line).

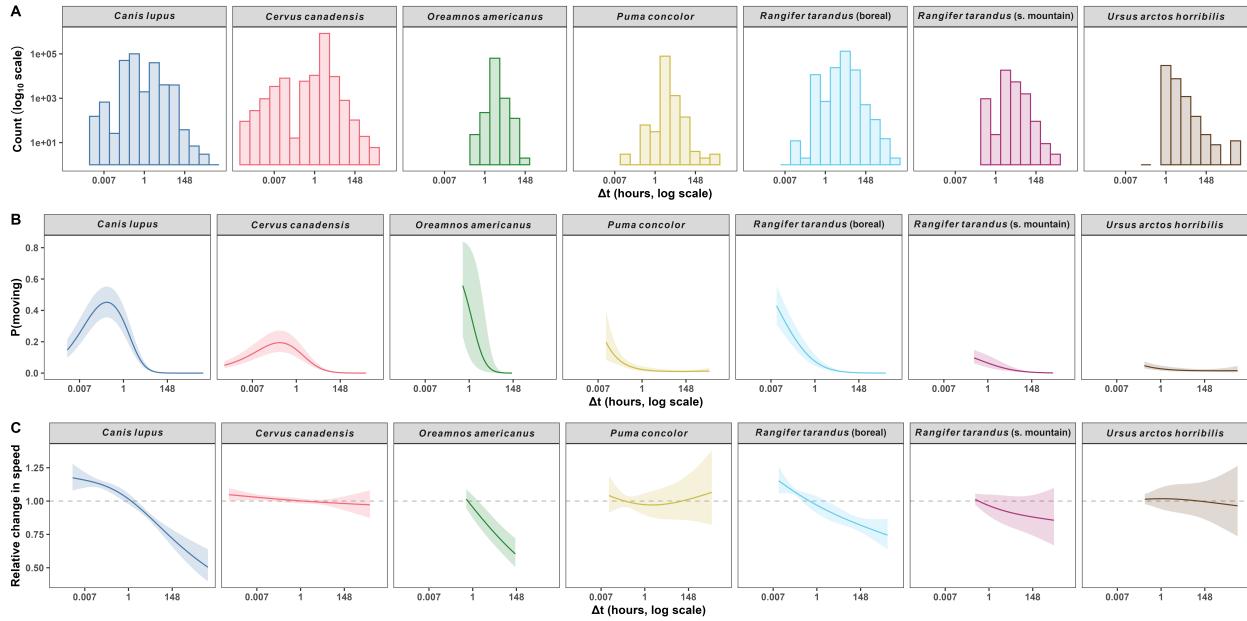


Figure B8: **A.** Histograms of sampling intervals between GPS locations. **B.** Species-level smooth effects of sampling interval on the estimated probability of the animal moving. **C.** Species-level estimated smooth effects of sampling interval on an animal's speed when moving. All x axes are on the natural logarithm scale.

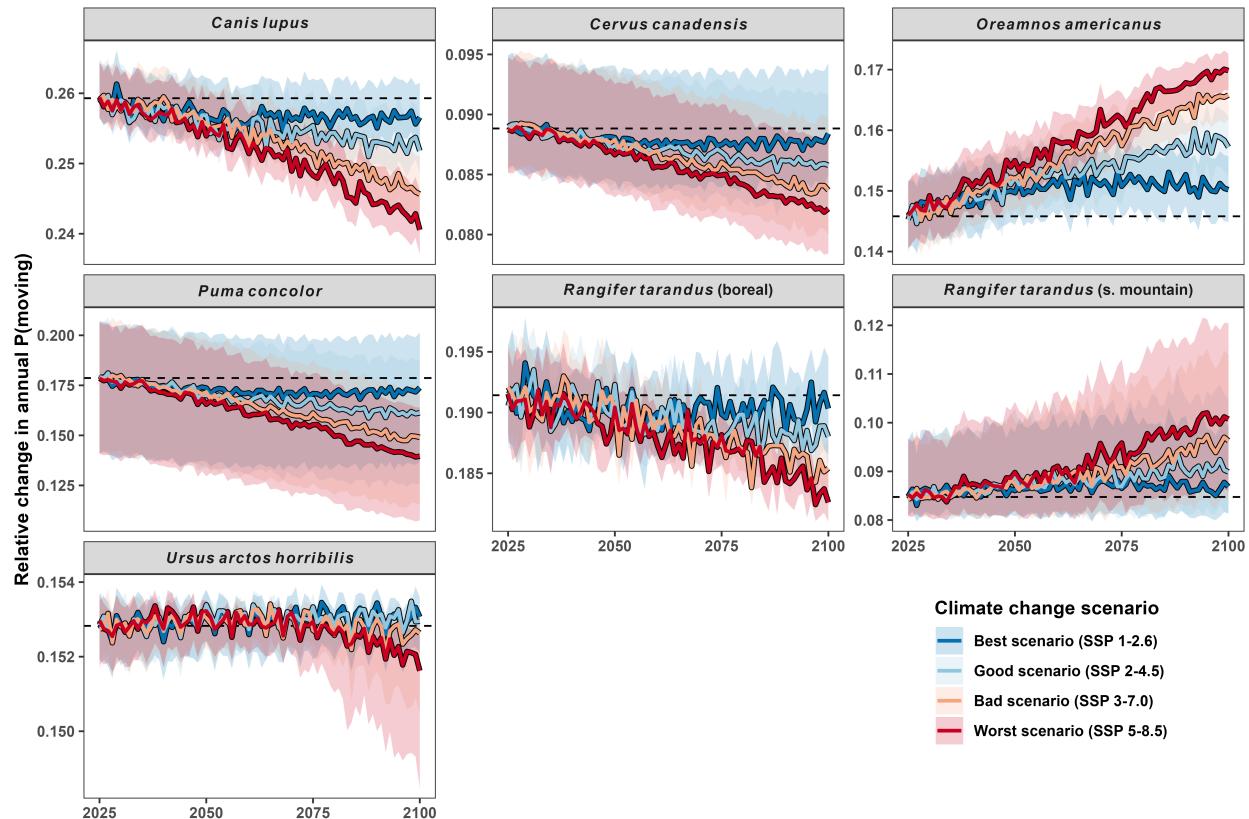


Figure B9: **The direction and magnitude of changes in probability of moving due to climate change varies among species, but worse climate-change scenarios result in the greatest change.** Lines indicate the median projected change in probability of moving due to changes in temperature within the species' current extent. Shaded areas indicate the 90% prediction interval within the range. The dashed black lines indicate the mean probability of movement in 2025 across the four scenarios. The projections only account for changes in movement behavior (i.e., movement frequency and speed) and ignore changes in physiology or movement costs.

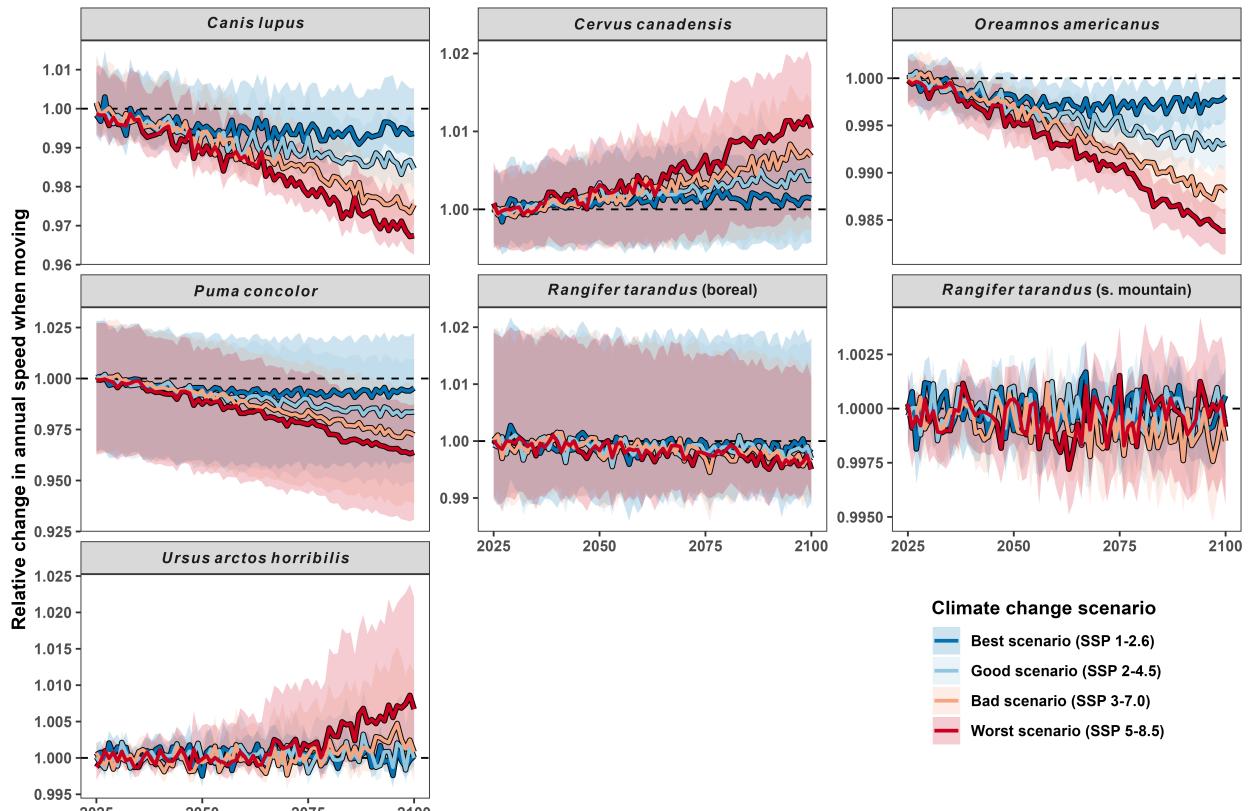


Figure B10: **The direction and magnitude of changes in speed due to climate change varies among species, but worse climate-change scenarios result in the greatest change.** Lines indicate the median projected change in speed due to changes in temperature within the species' current extent. Shaded areas indicate the 90% prediction interval within the range. The dashed black lines indicate the mean speed in 2025 across the four scenarios. The projections only account for changes movement behavior (i.e., movement frequency and speed) and ignore changes in physiology or movement costs.

## 2 Effects of temperature on habitat selection

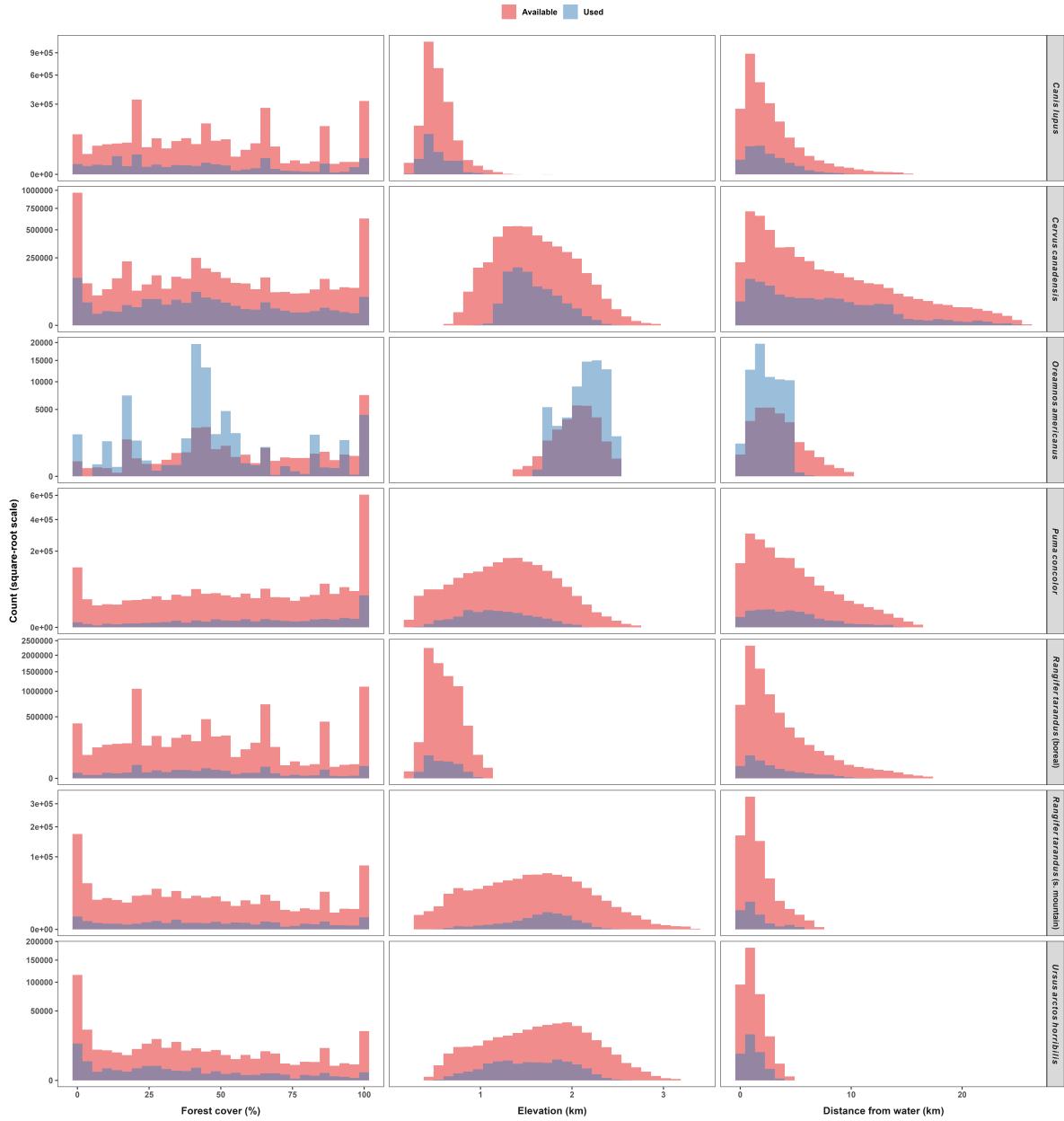


Figure B11: Histograms of available (red) and used (blue) resources. Each GPS fix counts as one observation. The y axis is on the square-root scale to help visualize values with low counts.

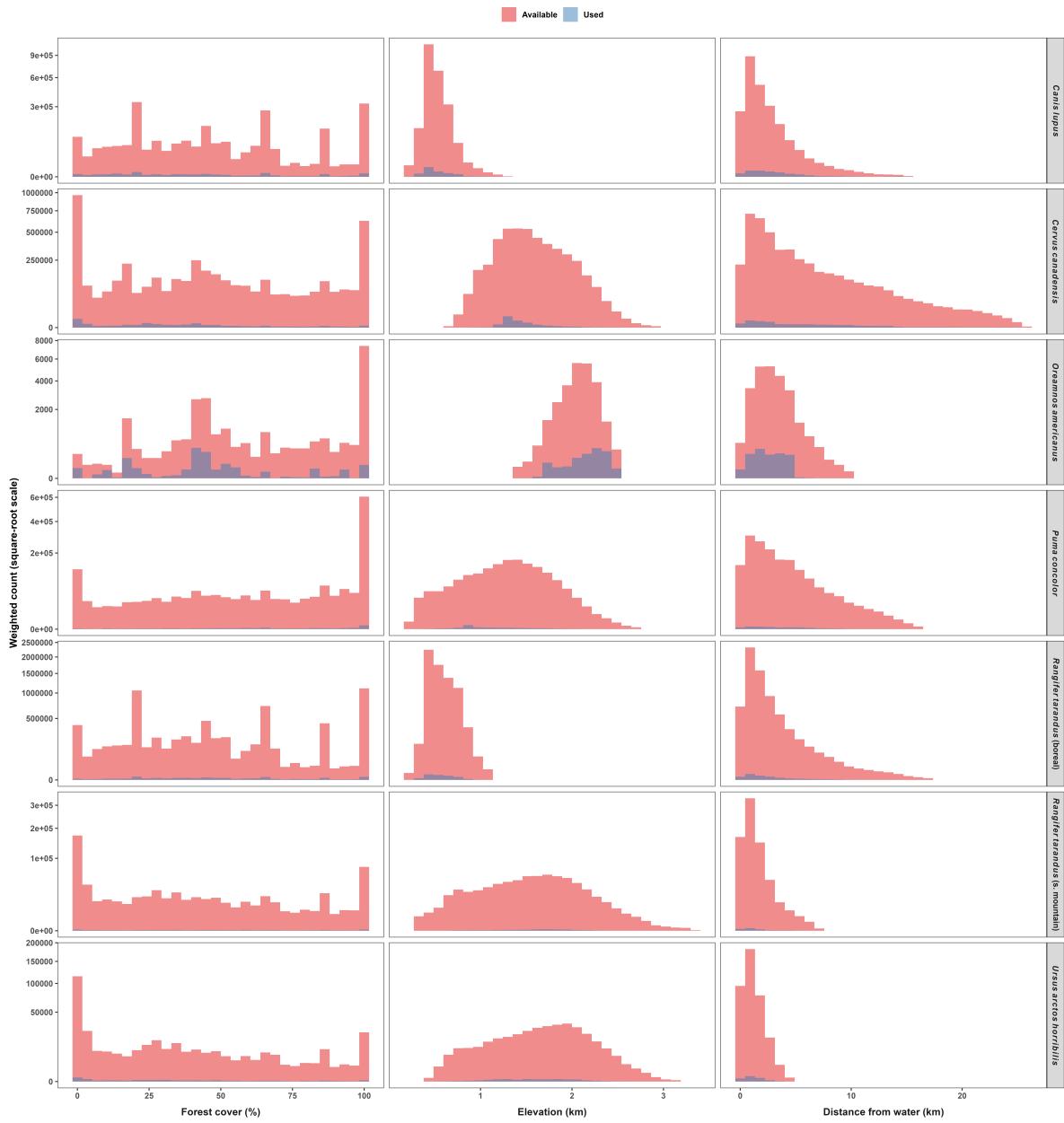


Figure B12: Histograms of available (red) and used (blue) resources. All GPS fixes are weighted according to their independence relative to the respective animals' range crossing time and Autocorrelated Kernel Density Estimate (i.e., the weights returned by AKDE weights multiplied by the number of home-range-crossing degrees of freedom). The y axis is on the square-root scale to help visualize values with low counts.

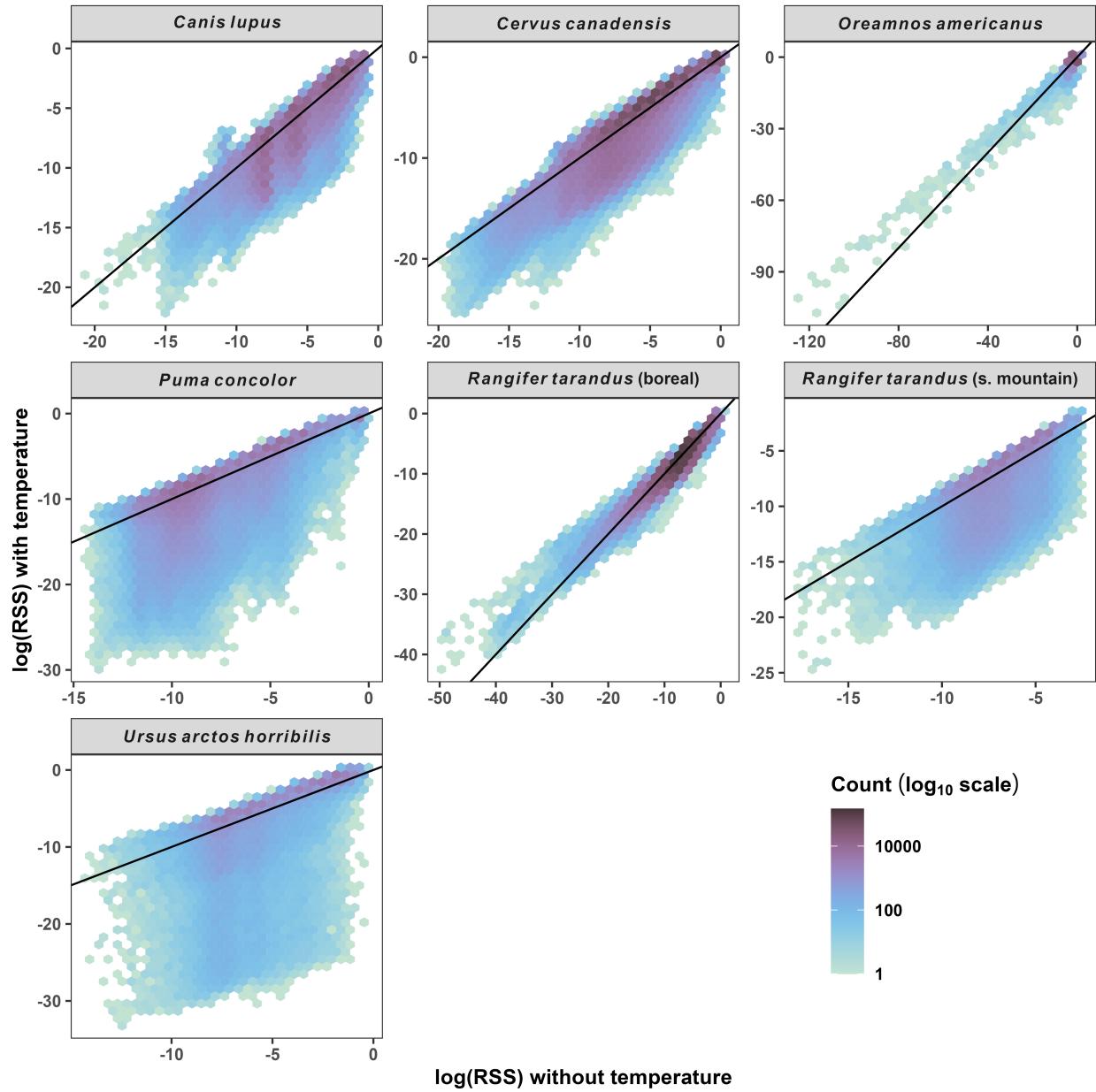


Figure B13: Hexplot of the fitted values from the HRSFs with and without including temperature.

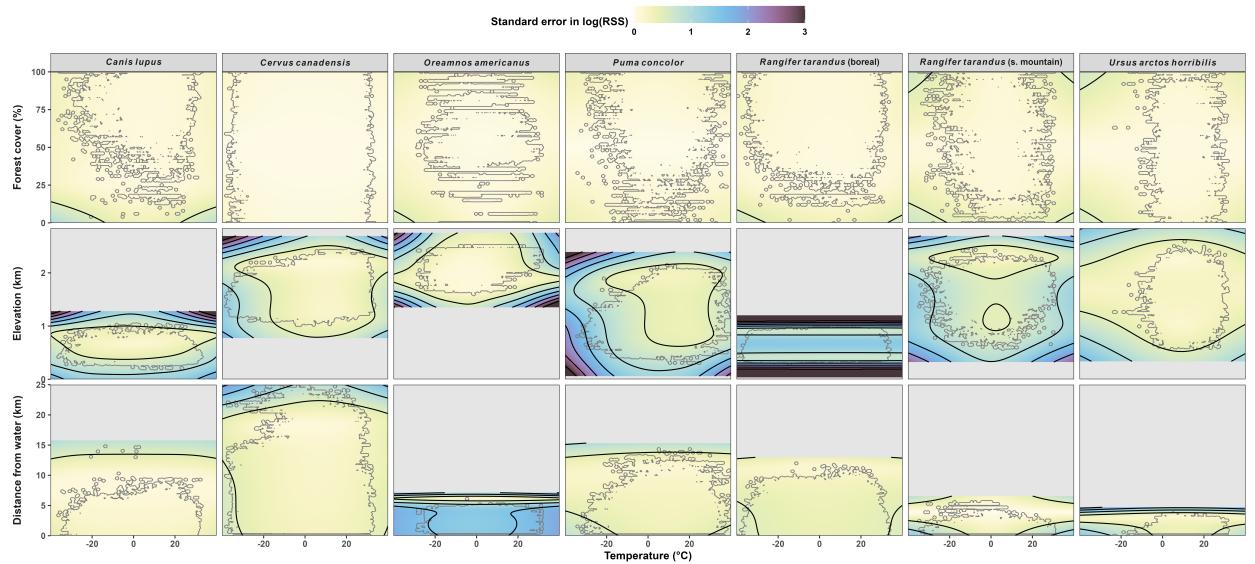


Figure B14: Estimated standard error in the relative selection strength (RSS) for forest cover (%), elevation (km), and distance from water (km) as a function of temperature (see Fig. 3 in the main text). The grey contours indicate the extent of each species' observed data. For ease of readability, the color bar values were capped to a maximum of 3 (original maximum was 7.00).

### 3 Model selection

Table B1: Changes in deviance explained ( $\Delta DE$ ) and Akaike Information Criterion ( $\Delta AIC$ ) from including temperature in the models estimating probability of movement ( $P(M)$ ) and speed when moving (Speed) as well as in species-level Hierarchical Resource Selection Functions (HRSF). Unlike conventional marginal AIC (Akaike, 1974), The AIC values provided by the `mgcv` package for R are calculated using the models' estimated effective degrees of freedom and the Bayesian variance-covariance matrices, which avoids issues related to random effects or Restricted Maximum Likelihood (REML).

Model	Species	$\Delta DE$	$\Delta AIC$
$P(M)$	All	0.5%	5927
Speed	All	0.2%	1682
HRSF	<i>Canis lupus</i>	9.2%	6566
HRSF	<i>Cervus canadensis</i>	4.6%	4448
HRSF	<i>Oreamnos americanus</i>	18.6%	813
HRSF	<i>Puma concolor</i>	8%	966
HRSF	<i>Rangifer tarandus</i> (boreal)	5.3%	5637
HRSF	<i>Rangifer tarandus</i> (s. mountain)	12%	342
HRSF	<i>Ursus arctos horribilis</i>	17.7%	914

### 3.1 Analyses of Deviance

Here we produce approximate p-values for analyses of deviance (a form of generalized likelihood ratio tests) following the methods of Section 3.3 in Wood (2017). We use an  $F$  test rather than a  $\chi^2$  test for the Gamma models because the scale parameter for the Gamma distribution is estimated rather than known or fixed (as in the case of binomial or Poisson distributions – see table 3.1 on page 104 of Wood, 2017).

```
# P(moving)
anova(readRDS('../models/binomial-gam-without-temperature.rds'),
      readRDS('../models/binomial-gam.rds'),
      test = 'Chisq')

## Analysis of Deviance Table
##
## Model 1: moving ~ s(animal, bs = "re") + species + s(tod_pdt, by = species,
##           k = 5, bs = "cc") + s(doy, by = species, k = 5, bs = "cc") +
##           ti(doy, tod_pdt, by = species, k = 5, bs = c("cc", "cc")) +
##           s(log(dt), k = 3) + s(log(dt), species, k = 3, bs = "fs")
## Model 2: moving ~ s(animal, bs = "re") + species + s(tod_pdt, by = species,
##           k = 5, bs = "cc") + s(doy, by = species, k = 5, bs = "cc") +
##           s(temp_c, by = species, k = 5, bs = "tp") + ti(doy, tod_pdt,
##           by = species, k = 5, bs = c("cc", "cc")) + ti(temp_c, tod_pdt,
##           by = species, k = 5, bs = c("tp", "cc")) + ti(temp_c, doy,
##           by = species, k = 5, bs = c("tp", "cc")) + s(log(dt), k = 3) +
##           s(log(dt), species, k = 3, bs = "fs")
##   Resid. Df Resid. Dev      Df Deviance  Pr(>Chi)
## 1    1476577    1129510
## 2    1476436    1123326 141.59      6184 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```

# speed when moving
anova(readRDS('../models/gamma-gam-without-temperature.rds'),
      readRDS('../models/gamma-gam.rds'),
      test = 'F')

## Analysis of Deviance Table
##
## Model 1: speed_est ~ s(animal, bs = "re") + species + s(tod_pdt, by = species,
##                 k = 5, bs = "cc") + s(doy, by = species, k = 5, bs = "cc") +
##                 ti(doy, tod_pdt, by = species, k = 5, bs = c("cc", "cc")) +
##                 s(log(dt), k = 3) + s(log(dt), species, k = 3, bs = "fs")
## Model 2: speed_est ~ s(animal, bs = "re") + species + s(tod_pdt, by = species,
##                 k = 5, bs = "cc") + s(doy, by = species, k = 5, bs = "cc") +
##                 s(temp_c, by = species, k = 5, bs = "tp") + ti(doy, tod_pdt,
##                 by = species, k = 5, bs = c("cc", "cc")) + ti(temp_c, tod_pdt,
##                 by = species, k = 5, bs = c("tp", "cc")) + ti(temp_c, doy,
##                 by = species, k = 5, bs = c("tp", "cc")) + s(log(dt), k = 3) +
##                 s(log(dt), species, k = 3, bs = "fs")
##   Resid. Df Resid. Dev      Df Deviance      F      Pr(>F)
## 1     223798    37934
## 2     223710    37649  87.751   285.11 17.456 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```

# HRSFs
anova(readRDS('../models/rsf-Canis_lupus-2025-01-20.rds'),
      readRDS('../models/rsf-Canis_lupus-no-temperature-2025-01-20.rds'),
      test = 'Chisq')

```

```

## Analysis of Deviance Table
##
## Model 1: detected ~ s(forest_perc, k = 4, bs = "cr") + s(elevation_m,
##                 k = 4, bs = "cr") + s(dist_water_m, k = 4, bs = "cr") + s(animal,
##                 bs = "re") + s(forest_perc, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##                 s(elevation_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##                 s(dist_water_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##                 ti(forest_perc, temperature_C, k = 4, bs = "cr") + ti(elevation_m,
##                 temperature_C, k = 4, bs = "cr") + ti(dist_water_m, temperature_C,
##                 k = 4, bs = "cr") + s(temperature_C, k = 4, bs = "cr") +
##                 s(temperature_C, animal, k = 4, bs = "fs", xt = list(bc = "cr"))
## Model 2: detected ~ s(forest_perc, k = 4, bs = "cr") + s(elevation_m,
##                 k = 4, bs = "cr") + s(dist_water_m, k = 4, bs = "cr") + s(animal,
##                 bs = "re") + s(forest_perc, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##                 s(elevation_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##                 s(dist_water_m, animal, k = 4, bs = "fs", xt = list(bc = "cr"))
##   Resid. Df Resid. Dev      Df Deviance  Pr(>Chi)
## 1     639545    39183
## 2     639643    45920 -97.944   -6736.8 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```

anova(readRDS('../models/rsf-Cervus canadensis-2025-01-20.rds'),
      readRDS('../models/rsf-Cervus canadensis-no-temperature-2025-01-20.rds'),
      test = 'Chisq')

## Analysis of Deviance Table
##
## Model 1: detected ~ s(forest_perc, k = 4, bs = "cr") + s(elevation_m,
##   k = 4, bs = "cr") + s(dist_water_m, k = 4, bs = "cr") + s(animal,
##   bs = "re") + s(forest_perc, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(elevation_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(dist_water_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   ti(forest_perc, temperature_C, k = 4, bs = "cr") + ti(elevation_m,
##   temperature_C, k = 4, bs = "cr") + ti(dist_water_m, temperature_C,
##   k = 4, bs = "cr") + s(temperature_C, k = 4, bs = "cr") +
##   s(temperature_C, animal, k = 4, bs = "fs", xt = list(bc = "cr"))
## Model 2: detected ~ s(forest_perc, k = 4, bs = "cr") + s(elevation_m,
##   k = 4, bs = "cr") + s(dist_water_m, k = 4, bs = "cr") + s(animal,
##   bs = "re") + s(forest_perc, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(elevation_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(dist_water_m, animal, k = 4, bs = "fs", xt = list(bc = "cr"))
##   Resid. Df Resid. Dev      Df Deviance Pr(>Chi)
## 1    1436458     19199
## 2    1436594    23816 -136.27   -4617.1 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

anova(readRDS('../models/rsf-Oreamnos americanus-2025-01-20.rds'),
      readRDS('../models/rsf-Oreamnos americanus-no-temperature-2025-01-20.rds'),
      test = 'Chisq')

## Analysis of Deviance Table
##
## Model 1: detected ~ s(forest_perc, k = 4, bs = "cr") + s(elevation_m,
##   k = 4, bs = "cr") + s(dist_water_m, k = 4, bs = "cr") + s(animal,
##   bs = "re") + s(forest_perc, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(elevation_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(dist_water_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   ti(forest_perc, temperature_C, k = 4, bs = "cr") + ti(elevation_m,
##   temperature_C, k = 4, bs = "cr") + ti(dist_water_m, temperature_C,
##   k = 4, bs = "cr") + s(temperature_C, k = 4, bs = "cr") +
##   s(temperature_C, animal, k = 4, bs = "fs", xt = list(bc = "cr"))
## Model 2: detected ~ s(forest_perc, k = 4, bs = "cr") + s(elevation_m,
##   k = 4, bs = "cr") + s(dist_water_m, k = 4, bs = "cr") + s(animal,
##   bs = "re") + s(forest_perc, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(elevation_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(dist_water_m, animal, k = 4, bs = "fs", xt = list(bc = "cr"))
##   Resid. Df Resid. Dev      Df Deviance Pr(>Chi)
## 1    70696     1576.8
## 2    70694    2387.3 2.1921   -810.56

```

```

anova(readRDS('../models/rsf-Puma concolor-2025-01-20.rds'),
      readRDS('../models/rsf-Puma concolor-no-temperature-2025-01-20.rds'),
      test = 'Chisq')

## Analysis of Deviance Table
##
## Model 1: detected ~ s(forest_perc, k = 4, bs = "cr") + s(elevation_m,
##                      k = 4, bs = "cr") + s(dist_water_m, k = 4, bs = "cr") + s(animal,
##                      bs = "re") + s(forest_perc, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##                      s(elevation_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##                      s(dist_water_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##                      ti(forest_perc, temperature_C, k = 4, bs = "cr") + ti(elevation_m,
##                      temperature_C, k = 4, bs = "cr") + ti(dist_water_m, temperature_C,
##                      k = 4, bs = "cr") + s(temperature_C, k = 4, bs = "cr") +
##                      s(temperature_C, animal, k = 4, bs = "fs", xt = list(bc = "cr"))
## Model 2: detected ~ s(forest_perc, k = 4, bs = "cr") + s(elevation_m,
##                      k = 4, bs = "cr") + s(dist_water_m, k = 4, bs = "cr") + s(animal,
##                      bs = "re") + s(forest_perc, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##                      s(elevation_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##                      s(dist_water_m, animal, k = 4, bs = "fs", xt = list(bc = "cr"))
##   Resid. Df Resid. Dev      Df Deviance Pr(>Chi)
## 1     324734    4076.5
## 2     324772    5099.7 -38.033  -1023.3 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```

anova(
  readRDS('../models/rsf-Rangifer tarandus boreal-2025-01-21.rds'),
  readRDS('../models/rsf-Rangifer tarandus boreal-no-temperature-2025-01-20.rds'),
  test = 'Chisq')

## Analysis of Deviance Table
##
## Model 1: detected ~ s(forest_perc, k = 4, bs = "cr") + s(elevation_m,
##                      k = 4, bs = "cr") + s(dist_water_m, k = 4, bs = "cr") + s(animal,
##                      bs = "re") + s(forest_perc, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##                      s(elevation_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##                      s(dist_water_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##                      ti(forest_perc, temperature_C, k = 4, bs = "cr") + ti(elevation_m,
##                      temperature_C, k = 4, bs = "cr") + ti(dist_water_m, temperature_C,
##                      k = 4, bs = "cr") + s(temperature_C, k = 4, bs = "cr") +
##                      s(temperature_C, animal, k = 4, bs = "fs", xt = list(bc = "cr"))
## Model 2: detected ~ s(forest_perc, k = 4, bs = "cr") + s(elevation_m,
##                      k = 4, bs = "cr") + s(dist_water_m, k = 4, bs = "cr") + s(animal,
##                      bs = "re") + s(forest_perc, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##                      s(elevation_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##                      s(dist_water_m, animal, k = 4, bs = "fs", xt = list(bc = "cr"))
##   Resid. Df Resid. Dev      Df Deviance Pr(>Chi)
## 1     1452896     48216
## 2     1452959     53927 -62.606  -5711.3 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```

anova(readRDS('../models/rsf-Rangifer tarandus southern mountain-2025-01-20.rds'),
      readRDS('../models/rsf-Rangifer tarandus southern mountain-no-temperature-2025-01-20.rds'),
      test = 'Chisq')

```

```

## Analysis of Deviance Table
##
## Model 1: detected ~ s(forest_perc, k = 4, bs = "cr") + s(elevation_m,
##   k = 4, bs = "cr") + s(dist_water_m, k = 4, bs = "cr") + s(animal,
##   bs = "re") + s(forest_perc, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(elevation_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(dist_water_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   ti(forest_perc, temperature_C, k = 4, bs = "cr") + ti(elevation_m,
##   temperature_C, k = 4, bs = "cr") + ti(dist_water_m, temperature_C,
##   k = 4, bs = "cr") + s(temperature_C, k = 4, bs = "cr") +
##   s(temperature_C, animal, k = 4, bs = "fs", xt = list(bc = "cr"))
## Model 2: detected ~ s(forest_perc, k = 4, bs = "cr") + s(elevation_m,
##   k = 4, bs = "cr") + s(dist_water_m, k = 4, bs = "cr") + s(animal,
##   bs = "re") + s(forest_perc, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(elevation_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(dist_water_m, animal, k = 4, bs = "fs", xt = list(bc = "cr"))
##   Resid. Df Resid. Dev      Df Deviance  Pr(>Chi)
## 1     132580    2026.1
## 2     132589  2384.5 -8.8792  -358.34 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```

anova(readRDS('../models/rsf-Ursus arctos horribilis-2025-01-20.rds'),
      readRDS('../models/rsf-Ursus arctos horribilis-no-temperature-2025-01-20.rds'),
      test = 'Chisq')

```

```

## Analysis of Deviance Table
##
## Model 1: detected ~ s(forest_perc, k = 4, bs = "cr") + s(elevation_m,
##   k = 4, bs = "cr") + s(dist_water_m, k = 4, bs = "cr") + s(animal,
##   bs = "re") + s(forest_perc, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(elevation_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(dist_water_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   ti(forest_perc, temperature_C, k = 4, bs = "cr") + ti(elevation_m,
##   temperature_C, k = 4, bs = "cr") + ti(dist_water_m, temperature_C,
##   k = 4, bs = "cr") + s(temperature_C, k = 4, bs = "cr") +
##   s(temperature_C, animal, k = 4, bs = "fs", xt = list(bc = "cr"))
## Model 2: detected ~ s(forest_perc, k = 4, bs = "cr") + s(elevation_m,
##   k = 4, bs = "cr") + s(dist_water_m, k = 4, bs = "cr") + s(animal,
##   bs = "re") + s(forest_perc, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(elevation_m, animal, k = 4, bs = "fs", xt = list(bc = "cr")) +
##   s(dist_water_m, animal, k = 4, bs = "fs", xt = list(bc = "cr"))
##   Resid. Df Resid. Dev      Df Deviance  Pr(>Chi)
## 1     99164    2356.7
## 2     99165  3275.0 -1.1525  -918.26 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

## References

- Akaike H (1974) A new look at the statistical model identification. *IEEE Transactions on Automatic Control*, **19**, 716–723.
- Wood SN (2017) Generalized additive models: An introduction with R, Second edition edn. CRC Press/Taylor & Francis Group, Boca Raton.