

# Explaining deepSSF habitat selection

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Here we want to associate covariate values with the predicted habitat selection probabilities from the deepSSF model. This helps to understand what the model has learned and can help to infer something about the spatial ecology of the species, particularly in relation to temporal dynamics. We show in the examples below how the selection for covariates such as NDVI changes throughout the day, which also changes through the year. This indicates that the model has represented the multiscale temporal dynamics present in the data, which will come out when simulating from the models.

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## Loading packages

```
library(tidyverse)
packages <- c("terra", "beeswarm", "tictoc", "viridis")
walk(packages, require, character.only = T)
```

## Load the habitat selection selection

The habitat selection probability values are from the `deepSSF_landscape_preds.ipynb` script. The predictions are stored in a CSV file that contains the NDVI, canopy cover, herbaceous vegetation, and slope values for each cell in the landscape. The predictions are stored as columns, one for each hour of the day.

```
# yday 45
habitat_selection_yday45 <-
  read_csv("Python/outputs/landscape_predictions/2005/yday45/id2005_hourly_habitat_suitability")
```

Rows: 707350 Columns: 28  
-- Column specification -----  
Delimiter: ","  
dbl (28): NDVI, Canopy\_cover, Herbaceous\_vegetation, Slope, 1, 2, 3, 4, 5, 6...  
  
i Use `spec()` to retrieve the full column specification for this data.  
i Specify the column types or set `show\_col\_types = FALSE` to quiet this message.

```
# yday 225
habitat_selection_yday225 <-
  read_csv("Python/outputs/landscape_predictions/2005/yday225/id2005_hourly_habitat_suitability")
```

Rows: 707350 Columns: 28  
-- Column specification -----  
Delimiter: ","  
dbl (28): NDVI, Canopy\_cover, Herbaceous\_vegetation, Slope, 1, 2, 3, 4, 5, 6...  
  
i Use `spec()` to retrieve the full column specification for this data.  
i Specify the column types or set `show\_col\_types = FALSE` to quiet this message.

```
# filter out the edge cells that were masked
habitat_selection_yday45 <- habitat_selection_yday45 %>% filter(!NDVI %in% c(Inf, -Inf))
habitat_selection_yday225 <- habitat_selection_yday225 %>% filter(!NDVI %in% c(Inf, -Inf))
```

## Tidy the data frame

Rescale the covariates to their original values and select a subset of the data to speed up the plotting process. As we're taking a subset there will be some random variation in the plots each time the script is run, although there are many cells so the results should be robust.

```
# from the stack of local layers (Python script)
ndvi_max = 0.8220
ndvi_min = -0.2772
canopy_max = 82.5000
canopy_min = 0.0
herby_max = 1.0
herby_min = 0.0
slope_max = 12.2981
slope_min = 0.0006

habitat_selection_yday45_long <- habitat_selection_yday45 %>%
  # take a random sample of the data
  slice_sample(prop = 0.1) %>%
  # rescale the covariates to their original scale
  mutate(NDVI = (NDVI*(ndvi_max - ndvi_min) + ndvi_min),
         Canopy_cover = round((Canopy_cover*(canopy_max - canopy_min) + canopy_min), digits),
         Herbaceous_vegetation = (Herbaceous_vegetation*(herby_max - herby_min) + herby_min),
         Slope = (Slope*(slope_max - slope_min) + slope_min)) %>%
  # pivot the data frame to long format for plotting with ggplot
  pivot_longer(cols = -c(NDVI, Canopy_cover, Herbaceous_vegetation, Slope),
               names_to = "Hour", values_to = "Values") %>%
  # convert the hour column to numeric
  mutate(Hour = as.numeric(Hour))

head(habitat_selection_yday45_long)

# A tibble: 6 x 6
  NDVI Canopy_cover Herbaceous_vegetation Slope Hour     Values
  <dbl>      <dbl>                 <dbl>   <dbl> <dbl>      <dbl>
1 0.206        32.5                  1  0.0898    1  0.000000494
2 0.206        32.5                  1  0.0898    2  0.000000431
3 0.206        32.5                  1  0.0898    3  0.000000377
4 0.206        32.5                  1  0.0898    4  0.000000347
5 0.206        32.5                  1  0.0898    5  0.000000430
6 0.206        32.5                  1  0.0898    6  0.00000117
```

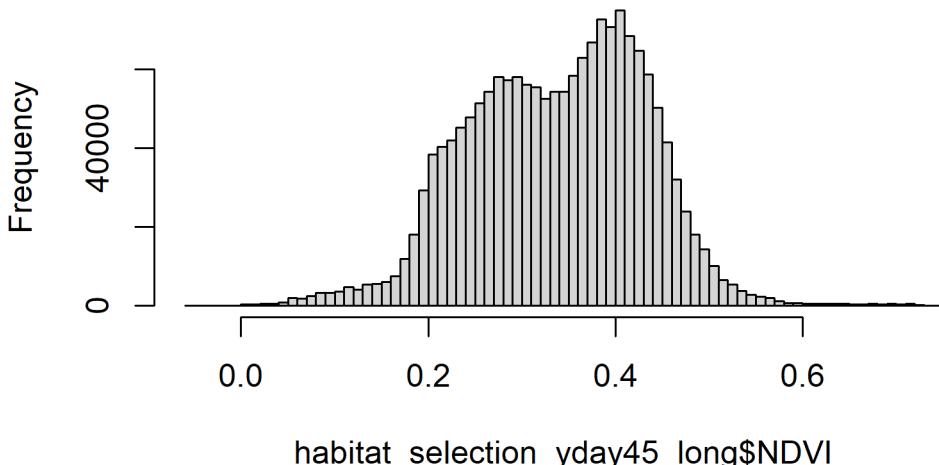
Do the same for the yday 225 predictions.

```
habitat_selection_yday225_long <- habitat_selection_yday225 %>%  
  
# take a random sample of the data  
slice_sample(prop = 0.01) %>%  
  
# rescale the covariates to their original scale  
mutate(NDVI = (NDVI*(ndvi_max - ndvi_min) + ndvi_min),  
       Canopy_cover = round((Canopy_cover*(canopy_max - canopy_min) + canopy_min), digits  
       Herbaceous_vegetation = (Herbaceous_vegetation*(herby_max - herby_min) + herby_min)  
       Slope = (Slope*(slope_max - slope_min) + slope_min)) %>%  
  
# pivot the data frame to long format for plotting with ggplot  
pivot_longer(cols = -c(NDVI, Canopy_cover, Herbaceous_vegetation, Slope),  
             names_to = "Hour", values_to = "Values") %>%  
  
# convert the hour column to numeric  
mutate(Hour = as.numeric(Hour))
```

### Check the distributions of the covariates

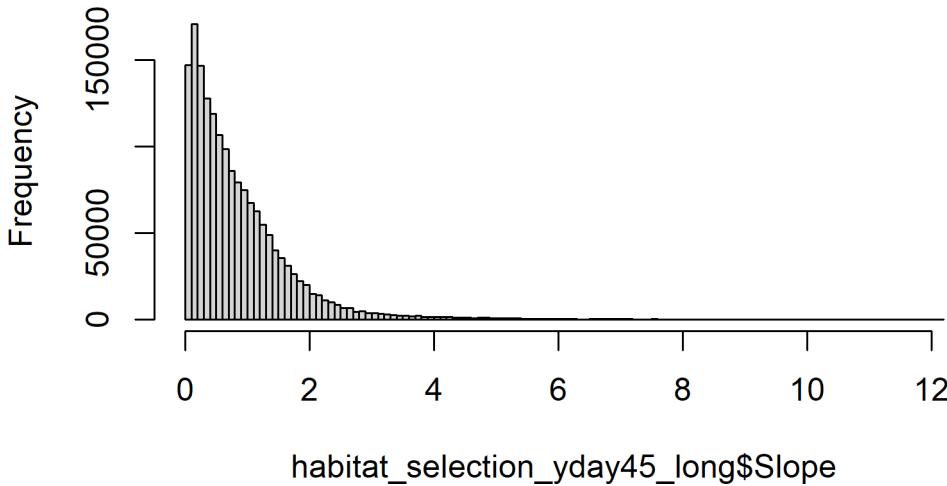
```
hist(habitat_selection_yday45_long$NDVI, breaks = 100, main = "NDVI distribution")
```

**NDVI distribution**



```
hist(habitat_selection_yday45_long$Slope, breaks = 100, main = "Slope distribution")
```

## Slope distribution



There are some extreme values in the NDVI and slope covariates. We will remove these values to get a better idea of the relationship between the covariates and the habitat selection predictions.

```
# NDVI quantiles
NDVI_yday45_quantiles <- quantile(habitat_selection_yday45_long$NDVI, probs = c(0.01, 0.99))
NDVI_yday225_quantiles <- quantile(habitat_selection_yday225_long$NDVI, probs = c(0.01, 0.99))

# Slope quantiles (same throughout the year)
slope_quantiles <- quantile(habitat_selection_yday45_long$Slope, probs = c(0.01, 0.99))

# filter out the extreme values for yday 45
habitat_selection_yday45_long <- habitat_selection_yday45_long %>%
  filter(NDVI >= NDVI_yday45_quantiles[1] &
         NDVI <= NDVI_yday45_quantiles[2] &
         Slope >= slope_quantiles[1] &
         Slope <= slope_quantiles[2])

# filter out the extreme values for yday 225
habitat_selection_yday225_long <- habitat_selection_yday225_long %>%
  filter(NDVI >= NDVI_yday225_quantiles[1] &
         NDVI <= NDVI_yday225_quantiles[2] &
         Slope >= slope_quantiles[1] &
         Slope <= slope_quantiles[2])
```

## Plot the covariate vs prediction values

To try and understand the influence of certain covariates on the habitat selection predictions, we can plot the habitat selection predictions against the covariate values. We will plot the NDVI, canopy cover, herbaceous vegetation and slope covariates against the habitat selection predictions for the yday 45 and yday 225 predictions.

***Note the tabs below for the different covariates***

### NDVI

First we can look at scatterplots for a couple different hours of the day. We can see that the prediction values change quite a bit across the day, indicating that the model has created a dynamic representation of buffalo habitat selection behaviour.

- At hour 1, there are higher selection probabilities associated with higher NDVI values.
- At hour 7, there is a more even distribution of selection probabilities across the NDVI values.
- At hour 12, there are higher selection probabilities associated with lower NDVI values.

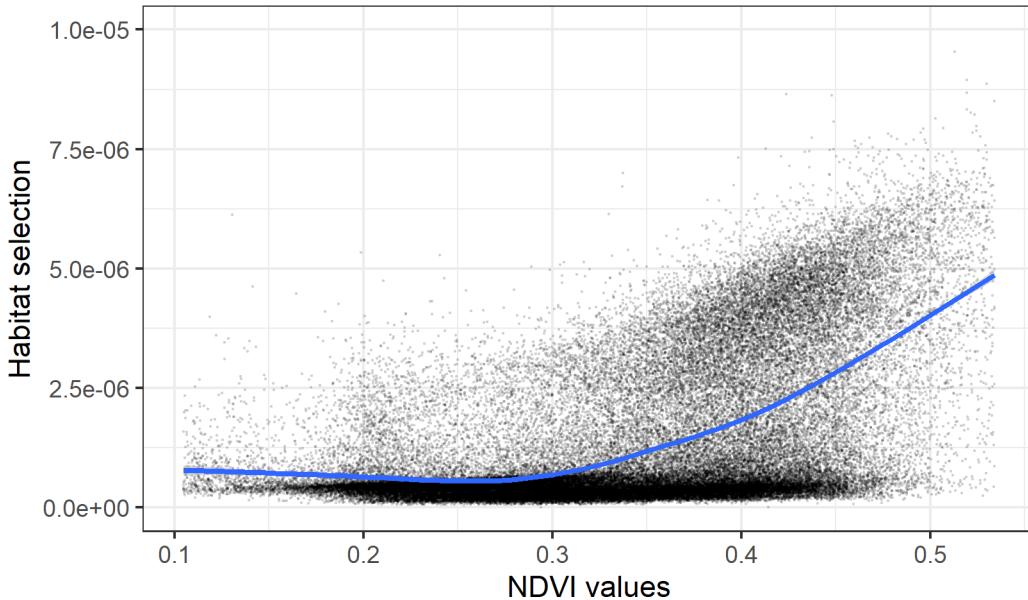
To highlight the relationship we will fit a smoother to the data.

```
habitat_selection_yday45_long %>% filter(Hour == 1) %>%
  ggplot(aes(x = NDVI, y = Values)) +
  geom_point(size = 0.1, alpha = 0.1) +
  geom_smooth(method = "gam") +
  scale_y_continuous(limits = c(0, 1e-5)) +
  theme_bw() +
  theme(legend.position = "bottom") +
  labs(
    title = "NDVI vs Habitat Selection - Hour 1",
    x = "NDVI values",
    y = "Habitat selection"
  )

`geom_smooth()` using formula = 'y ~ s(x, bs = "cs")'

Warning: Removed 907 rows containing non-finite outside the scale range
(`stat_smooth()`).
```

## NDVI vs Habitat Selection - Hour 1



```
ggsave("outputs/ndvi_habitat_selection_yday45_hour1.png", width = 150, height = 90,
       units = "mm", dpi = 300)
```

```
`geom_smooth()` using formula = 'y ~ s(x, bs = "cs")'
```

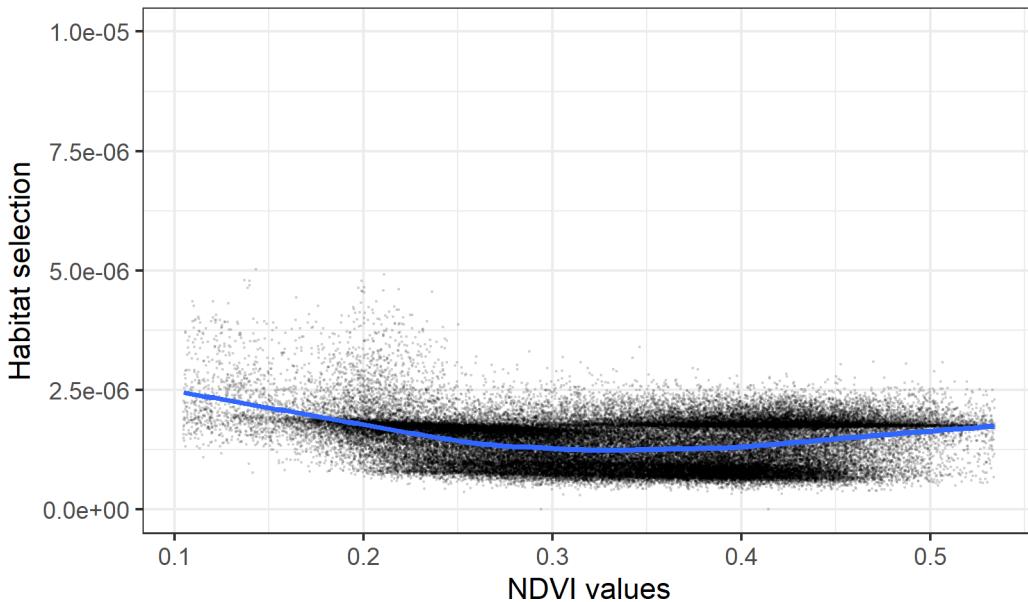
```
Warning: Removed 907 rows containing non-finite outside the scale range
(`stat_smooth()`).
```

```
habitat_selection_yday45_long %>% filter(Hour == 7) %>%
  ggplot(aes(x = NDVI, y = Values)) +
  geom_point(size = 0.1, alpha = 0.1) +
  geom_smooth(method = "gam") +
  scale_y_continuous(limits = c(0, 1e-05)) +
  theme_bw() +
  theme(legend.position = "bottom") +
  labs(
    title = "NDVI vs Habitat Selection - Hour 7",
    x = "NDVI values",
    y = "Habitat selection"
  )
```

```
`geom_smooth()` using formula = 'y ~ s(x, bs = "cs")'
```

```
Warning: Removed 907 rows containing non-finite outside the scale range
(`stat_smooth()`).
```

## NDVI vs Habitat Selection - Hour 7



```
ggsave("outputs/ndvi_habitat_selection_yday45_hour7.png", width = 150, height = 90,  
       units = "mm", dpi = 300)
```

```
`geom_smooth()` using formula = 'y ~ s(x, bs = "cs")'
```

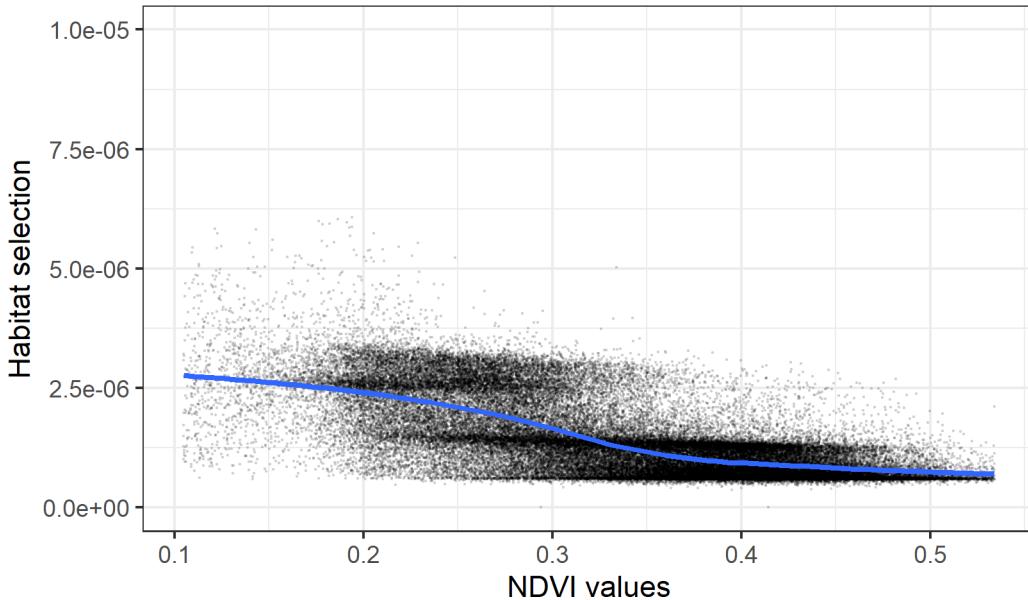
```
Warning: Removed 907 rows containing non-finite outside the scale range  
(`stat_smooth()`).
```

```
habitat_selection_yday45_long %>% filter(Hour == 12) %>%  
  ggplot(aes(x = NDVI, y = Values)) +  
  geom_point(size = 0.1, alpha = 0.1) +  
  geom_smooth(method = "gam") +  
  scale_y_continuous(limits = c(0, 1e-05)) +  
  theme_bw() +  
  theme(legend.position = "bottom") +  
  labs(  
    title = "NDVI vs Habitat Selection - Hour 12",  
    x = "NDVI values",  
    y = "Habitat selection"  
)
```

```
`geom_smooth()` using formula = 'y ~ s(x, bs = "cs")'
```

```
Warning: Removed 907 rows containing non-finite outside the scale range  
(`stat_smooth()`).
```

## NDVI vs Habitat Selection - Hour 12



```
ggsave("outputs/ndvi_habitat_selection_yday45_hour12.png", width = 150, height = 90,
       units = "mm", dpi = 300)
```

```
`geom_smooth()` using formula = 'y ~ s(x, bs = "cs")'
```

```
Warning: Removed 907 rows containing non-finite outside the scale range
(`stat_smooth()`).
```

## All hours

We can also plot all the hours of the day on the same plot to see how the relationship between NDVI and habitat selection changes throughout the day.

We see there is a rhythmic pattern, with selection for lower values during the middle hours of the day.

```
habitat_selection_yday45_long %>%
  ggplot(aes(x = NDVI, y = Values, colour = as.factor(Hour))) +
  geom_point(size = 0.1, alpha = 0.05) +
  geom_smooth(method = "gam") +
  scale_colour_viridis(discrete = T) +
  scale_y_continuous(limits = c(0, 1e-05)) +
  theme_bw() +
  theme(legend.position = "right") +
  labs(
```

```

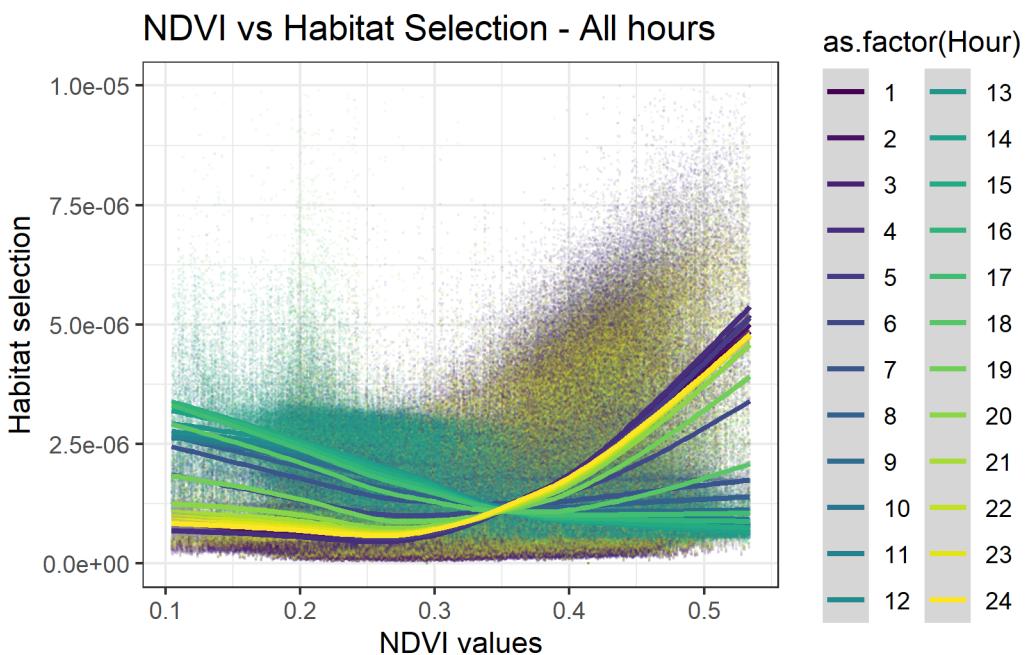
title = "NDVI vs Habitat Selection - All hours",
x = "NDVI values",
y = "Habitat selection"
)

```

```
`geom_smooth()` using formula = 'y ~ s(x, bs = "cs")'
```

Warning: Removed 21911 rows containing non-finite outside the scale range  
(`stat\_smooth()`).

Warning: Removed 143 rows containing missing values or values outside the scale range  
(`geom\_point()`).



```
ggsave("outputs/ndvi_habitat_selection_yday45_all_hours.png", width = 150, height = 90,
       units = "mm", dpi = 300)
```

```
`geom_smooth()` using formula = 'y ~ s(x, bs = "cs")'
```

Warning: Removed 21911 rows containing non-finite outside the scale range  
(`stat\_smooth()`).

Removed 143 rows containing missing values or values outside the scale range  
(`geom\_point()`).

## Selection surface

However, the plot above is fairly messy, so instead we will use the `stat_summary_2d` function to create a 2D histogram of the data. This will allow us to see the average habitat selection values for different NDVI values at different times of the day.

What we are seeing is the *mean habitat selection probability value* for different NDVI values at different times of the day. Higher values indicate a higher mean probability of selecting NDVI values in that range.

This selection surface reflects the plots above, with a pattern of selection for lower NDVI values during the middle hours of the day, and high values in the early morning and evening. This suggests that buffalo are selecting for less vegetated areas during the middle of the day.

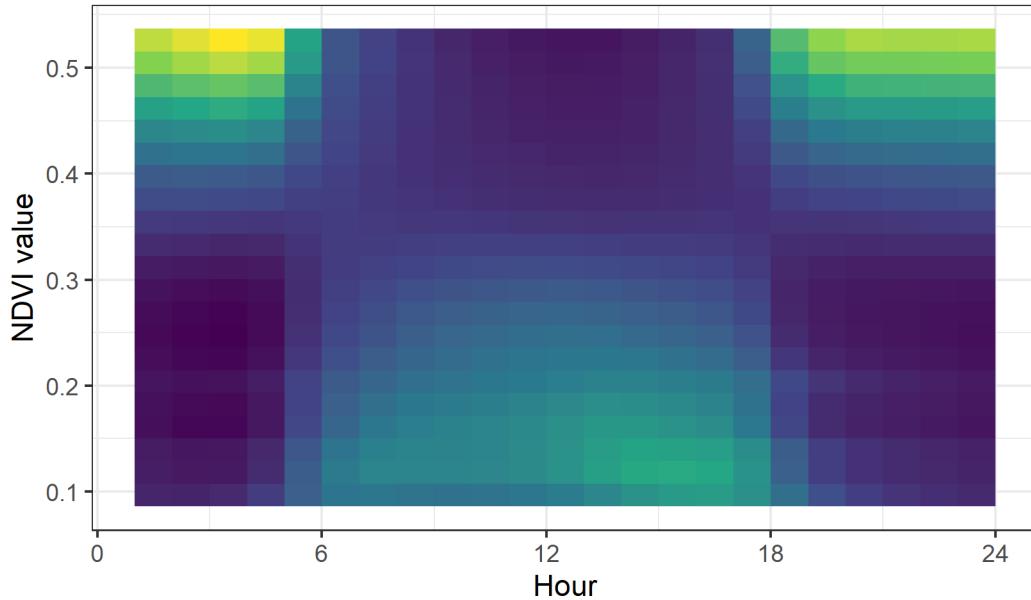
These selection surfaces are also similar to those presented by Forrest et al. (2024), which was achieved by fitting temporal dynamics and quadratic terms to the NDVI covariate. However, this was only fitted to one season of data at a time, whereas the deepSSF model can be fitted with multiscale temporal dynamics.

Day of the year = 45 (mid-February)

```
ggplot(habitat_selection_yday45_long,
       aes(x = Hour, y = NDVI)) +
  stat_summary_2d(aes(z = Values), fun = mean, bins = c(23,20)) +
  scale_fill_viridis_c("Mean probability") +
  scale_x_continuous("Hour", breaks = seq(0,24,6)) +
  scale_y_continuous("NDVI value") +
  ggtitle(paste0("Prediction values: Day 45")) +
  theme_bw() +
  theme(legend.position = "none")
```

Warning: Removed 21768 rows containing non-finite outside the scale range  
(`stat\_summary2d()`).

## Prediction values: Day 45



```
# for manuscript
ggsave("outputs/ndvi_habitat_selection_yday45.png", width = 90, height = 75,
       units = "mm", dpi = 600)
```

Warning: Removed 21768 rows containing non-finite outside the scale range  
(`stat\_summary2d()`).

```
# for website
ggsave("outputs/ndvi_habitat_selection_yday45_website.png", width = 150, height = 90,
       units = "mm", dpi = 300)
```

Warning: Removed 21768 rows containing non-finite outside the scale range  
(`stat\_summary2d()`).

We can illustrate the multitemporal dynamics of the NDVI covariate by plotting the selection surface for different days of the year.

Day of the year = 225 (mid-August)

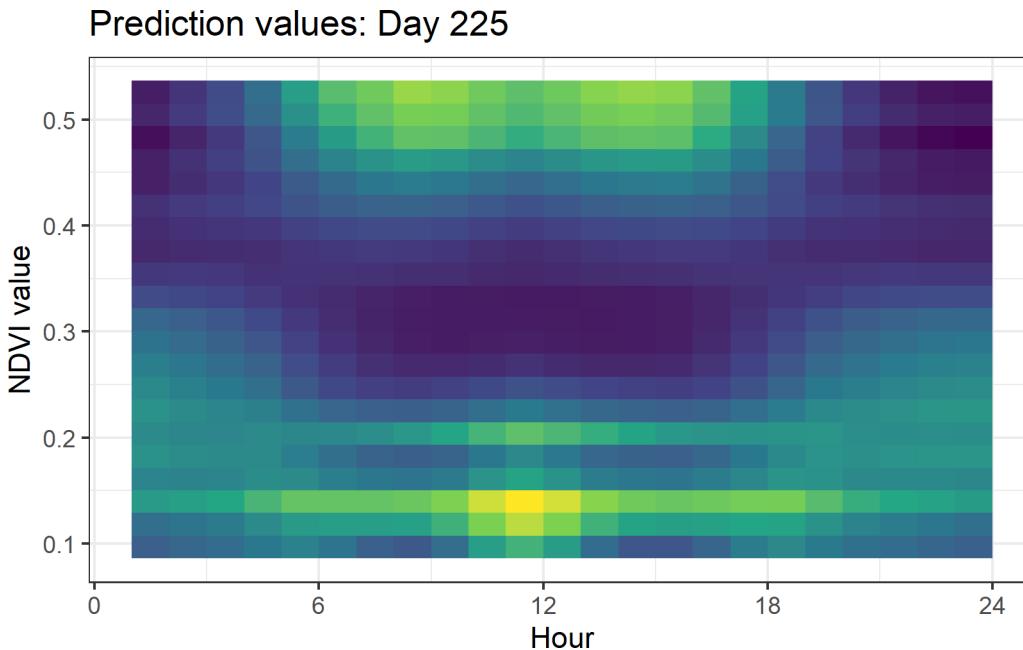
Here we see quite a different response, where there is some bimodal selection during the day, with higher selection probabilities for both low and high NDVI values. This may indicate that there are some features described by low or high values of NDVI that are important for buffalo habitat selection during this time of year, or may just indicate that the selection is more variable in the dry season.

```

ggplot(habitat_selection_yday225_long,
       aes(x = Hour, y = NDVI)) +
  stat_summary_2d(aes(z = Values), fun = mean, bins = c(23,20)) +
  scale_fill_viridis_c("Mean probability") +
  scale_x_continuous("Hour", breaks = seq(0,24,6)) +
  scale_y_continuous("NDVI value") +
  ggtitle(paste0("Prediction values: Day 225")) +
  theme_bw() +
  theme(legend.position = "none")

```

Warning: Removed 2376 rows containing non-finite outside the scale range  
(`stat\_summary2d()`).



```

# for manuscript
ggsave("outputs/ndvi_habitat_selection_yday225.png", width = 90, height = 75,
       units = "mm", dpi = 600)

```

Warning: Removed 2376 rows containing non-finite outside the scale range  
(`stat\_summary2d()`).

```

# for website
ggsave("outputs/ndvi_habitat_selection_yday225_website.png", width = 150, height = 90,
       units = "mm", dpi = 300)

```

```
Warning: Removed 2376 rows containing non-finite outside the scale range
(`stat_summary2d()`).
```

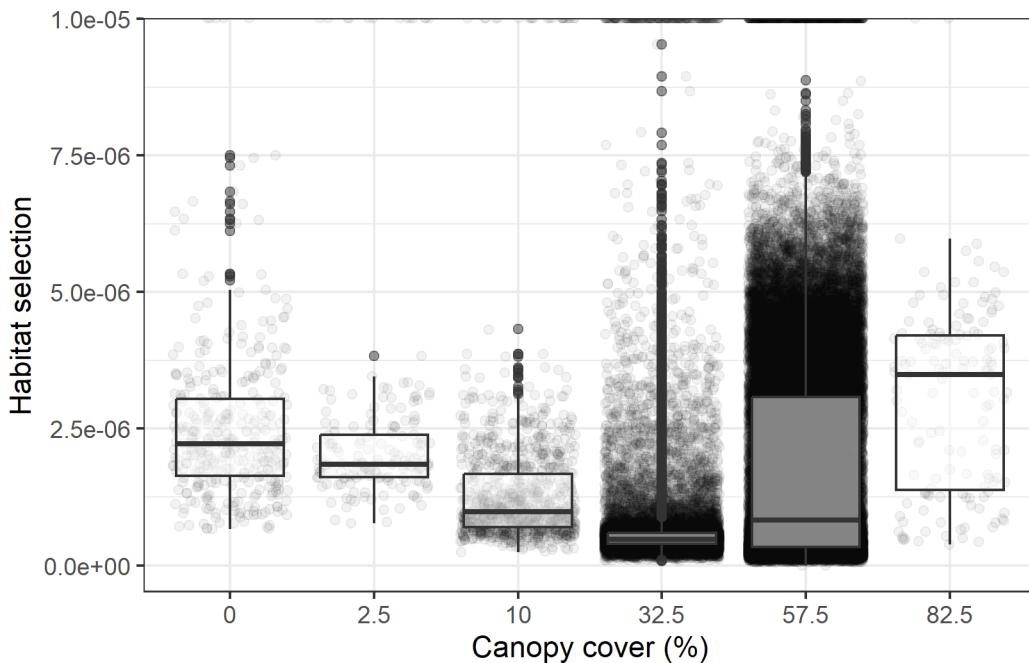
## Canopy cover

We can do the same for canopy cover.

Here we show these as boxplots (keep in mind that the canopy cover covariate is continuous, and the canopy classifications are not equally spaced, but we are treating it as a categorical variable for the purposes of this plot).

```
habitat_selection_yday45_long %>% filter(Hour == 1) %>%
  ggplot(aes(x = as.factor(Canopy_cover), y = Values)) +
  geom_jitter(alpha = 0.05) +
  geom_boxplot(alpha = 0.5) +
  theme_bw() +
  theme(legend.position = "bottom") +
  labs(
    x = "Canopy cover (%)",
    y = "Habitat selection"
  )
```

```
Warning: Removed 907 rows containing non-finite outside the scale range
(`stat_boxplot()`).
```

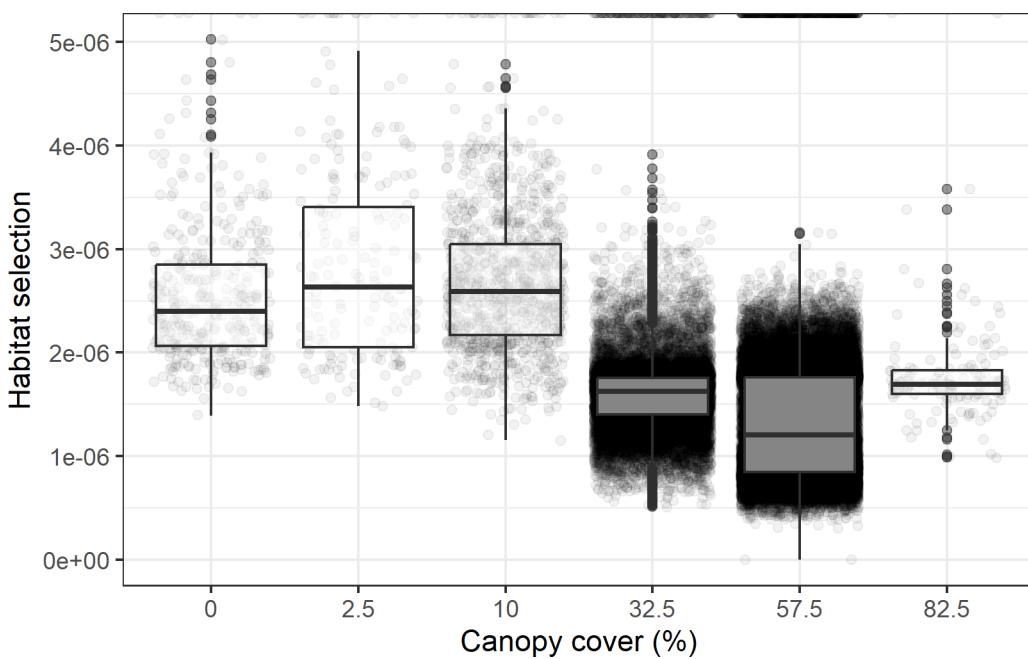


```

habitat_selection_yday45_long %>% filter(Hour == 7) %>%
  ggplot(aes(x = as.factor(Canopy_cover), y = Values)) +
  geom_jitter(alpha = 0.05) +
  geom_boxplot(alpha = 0.5) +
  theme_bw() +
  theme(legend.position = "bottom") +
  labs(
    x = "Canopy cover (%)",
    y = "Habitat selection"
  )

```

Warning: Removed 907 rows containing non-finite outside the scale range (`stat\_boxplot()`).

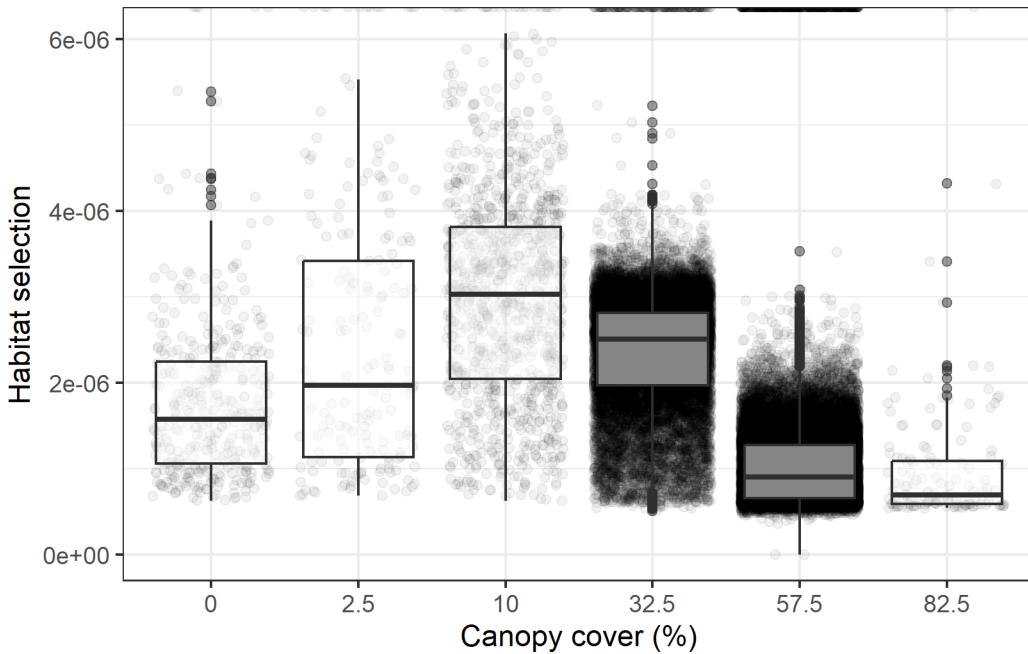


```

habitat_selection_yday45_long %>% filter(Hour == 12) %>%
  ggplot(aes(x = as.factor(Canopy_cover), y = Values)) +
  geom_jitter(alpha = 0.05) +
  geom_boxplot(alpha = 0.5) +
  theme_bw() +
  theme(legend.position = "bottom") +
  labs(
    x = "Canopy cover (%)",
    y = "Habitat selection"
  )

```

Warning: Removed 907 rows containing non-finite outside the scale range  
(`stat\_boxplot()`).



### All hours

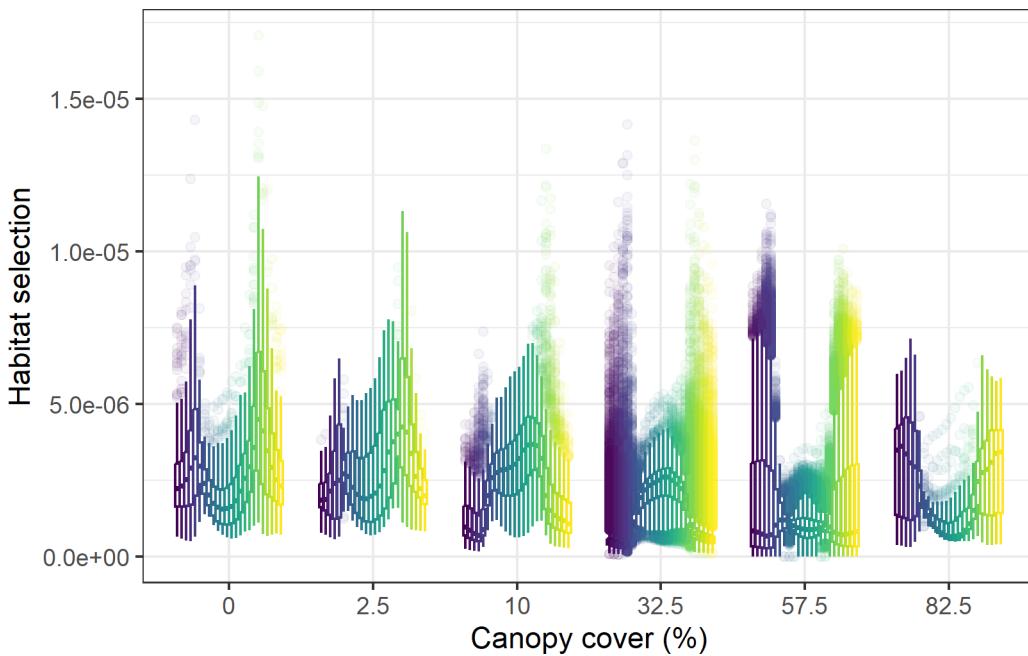
Again, this is a slightly confusing plot, but it shows how the habitat selection probabilities for each canopy cover ‘category’ change across the day.

Looking at 32.5% canopy cover for instance, there is lower selection probability in the early morning and evening, and higher selection probability in the middle of the day. This suggests that buffalo are selecting for areas with 32.5% canopy cover during the middle of the day.

For open canopy (0%), selection is highest around dawn and dusk, which correlates with the high movement periods of buffalo.

```
habitat_selection_yday45_long %>%
  ggplot(aes(x = as.factor(Canopy_cover), y = Values, colour = as.factor(Hour))) +
  geom_boxplot(alpha = 0.05) +
  scale_colour_viridis(discrete = T) +
  theme_bw() +
  theme(legend.position = "none") +
  labs(
    x = "Canopy cover (%)",
    y = "Habitat selection"
  )
```

```
Warning: Removed 21768 rows containing non-finite outside the scale range
(`stat_boxplot()`).
```



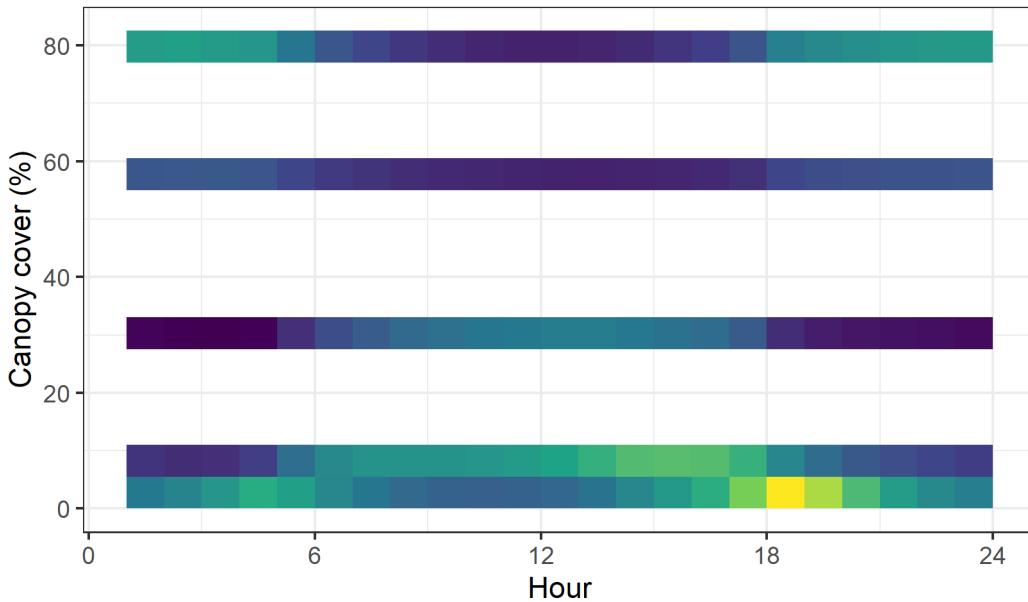
### Selection surface

Day of the year = 45 (mid-February)

```
ggplot(habitat_selection_yday45_long,
       aes(x = Hour, y = Canopy_cover)) +
  stat_summary_2d(aes(z = Values), fun = mean, bins = c(23,15)) +
  scale_fill_viridis_c("Mean probability") +
  scale_x_continuous("Hour", breaks = seq(0,24,6)) +
  scale_y_continuous("Canopy cover (%)") +
  ggtitle(paste0("Prediction values: Day 45")) +
  theme_bw() +
  theme(legend.position = "none")
```

```
Warning: Removed 21768 rows containing non-finite outside the scale range
(`stat_summary2d()`).
```

## Prediction values: Day 45



```
ggsave("outputs/canopy_habitat_selection_yday45.png", width = 90, height = 75,
       units = "mm", dpi = 600)
```

Warning: Removed 21768 rows containing non-finite outside the scale range  
(`stat\_summary2d()`).

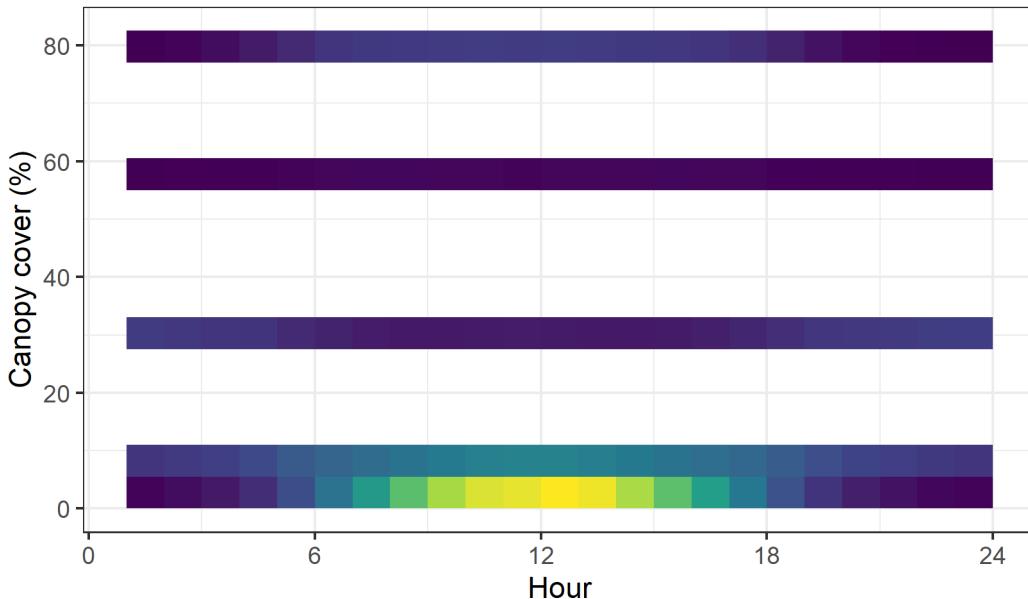
Day of the year = 225 (mid-August)

There is clearly high selection for open canopy in the middle of the day during the dry season.

```
ggplot(habitat_selection_yday225_long,
       aes(x = Hour, y = Canopy_cover)) +
  stat_summary_2d(aes(z = Values), fun = mean, bins = c(23,15)) +
  scale_fill_viridis_c("Mean probability") +
  scale_x_continuous("Hour", breaks = seq(0,24,6)) +
  scale_y_continuous("Canopy cover (%)") +
  ggtitle(paste0("Prediction values: Day 225")) +
  theme_bw() +
  theme(legend.position = "none")
```

Warning: Removed 2376 rows containing non-finite outside the scale range  
(`stat\_summary2d()`).

## Prediction values: Day 225



```
ggsave("outputs/canopy_habitat_selection_yday225.png", width = 90, height = 75,  
       units = "mm", dpi = 600)
```

Warning: Removed 2376 rows containing non-finite outside the scale range  
(`stat\_summary2d()`).

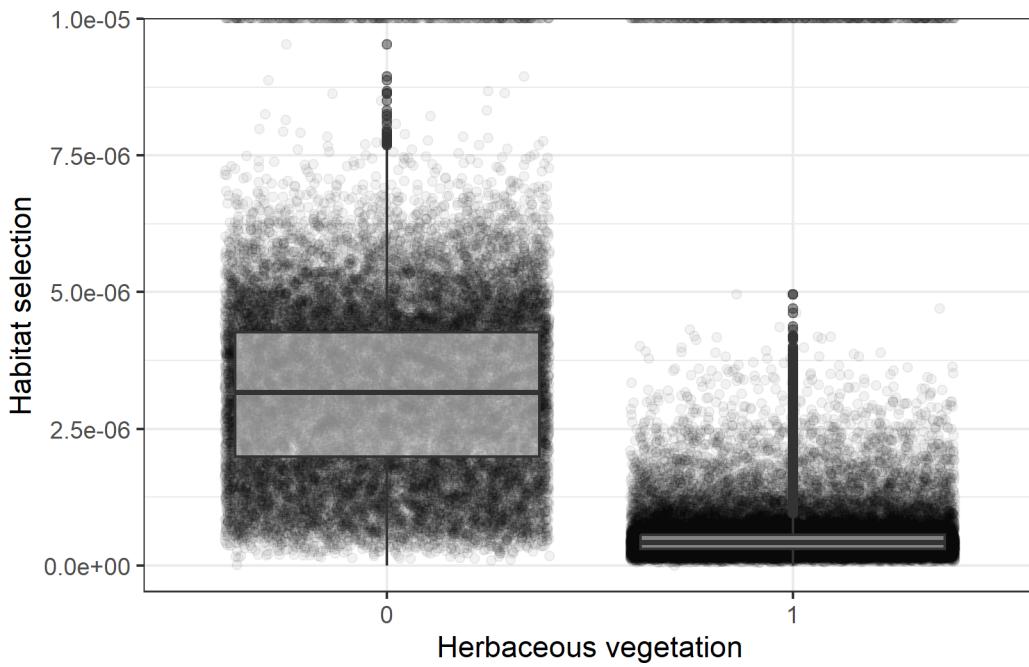
## Herbaceous vegetation

As herbaceous vegetation is binary, the plotting is a bit different to interpret, but the plots do show a relationship between habitat selection and herbaceous vegetation.

Recall that 0 is ‘woody vegetation’ and 1 is ‘herbaceous vegetation’.

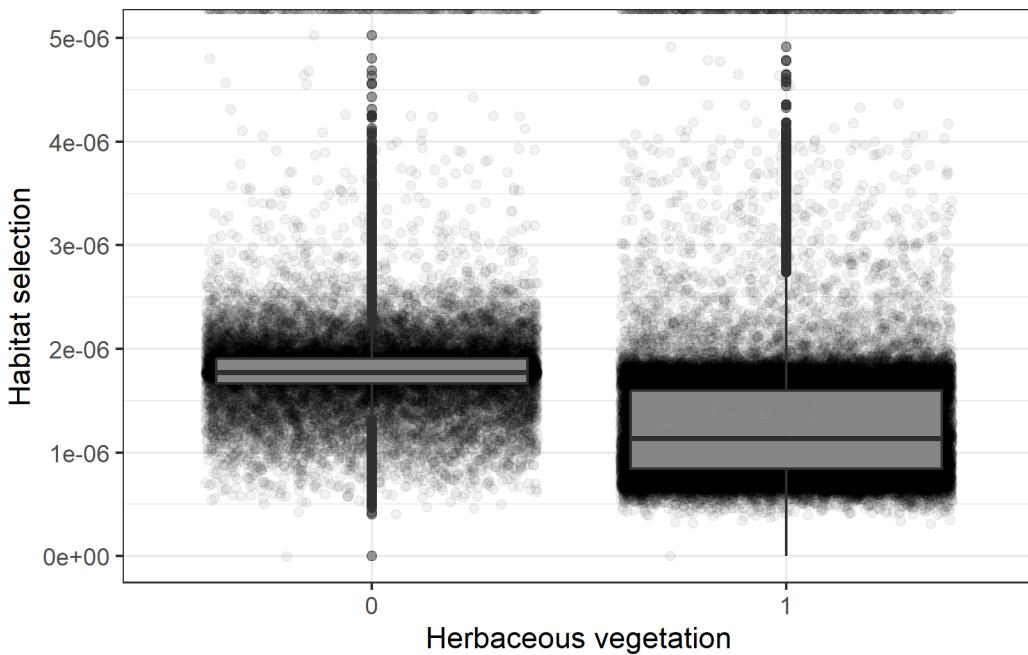
```
habitat_selection_yday45_long %>% filter(Hour == 1) %>%  
  ggplot(aes(x = factor(Herbaceous_vegetation), y = Values)) +  
  geom_jitter(alpha = 0.05) +  
  geom_boxplot(alpha = 0.5) +  
  theme_bw() +  
  theme(legend.position = "bottom") +  
  labs(  
    x = "Herbaceous vegetation",  
    y = "Habitat selection"  
)
```

Warning: Removed 907 rows containing non-finite outside the scale range  
(`stat\_boxplot()`).



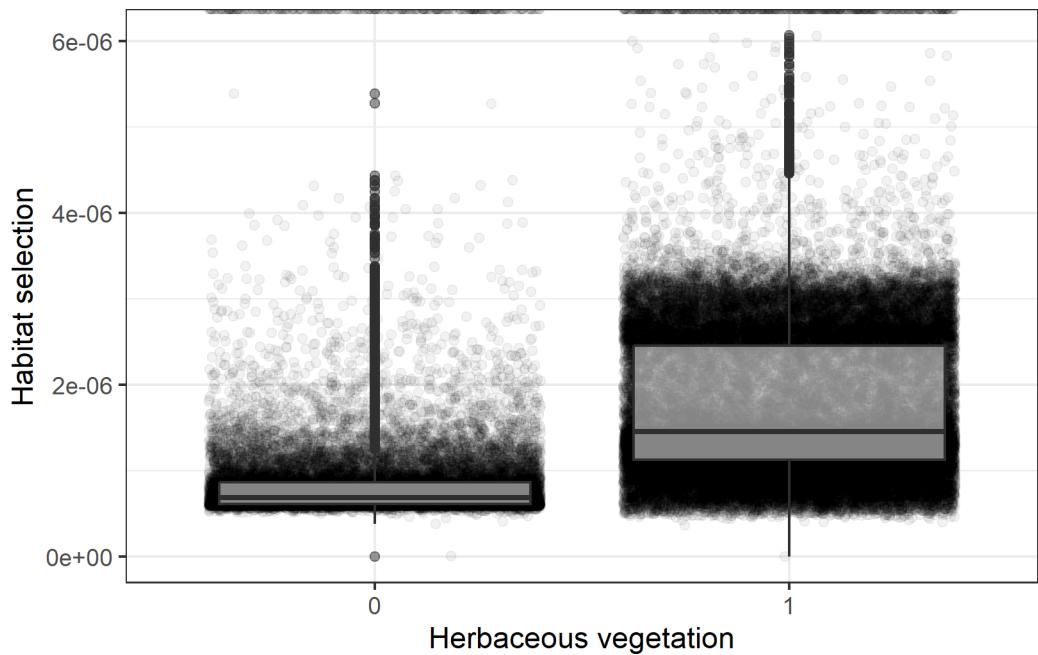
```
habitat_selection_yday45_long %>% filter(Hour == 7) %>%
  ggplot(aes(x = factor(Herbaceous_vegetation), y = Values)) +
  geom_jitter(alpha = 0.05) +
  geom_boxplot(alpha = 0.5) +
  theme_bw() +
  theme(legend.position = "bottom") +
  labs(
    x = "Herbaceous vegetation",
    y = "Habitat selection"
  )
```

Warning: Removed 907 rows containing non-finite outside the scale range  
(`stat\_boxplot()`).



```
habitat_selection_yday45_long %>% filter(Hour == 12) %>%
  ggplot(aes(x = factor(Herbaceous_vegetation), y = Values)) +
  geom_jitter(alpha = 0.05) +
  geom_boxplot(alpha = 0.5) +
  theme_bw() +
  theme(legend.position = "bottom") +
  labs(
    x = "Herbaceous vegetation",
    y = "Habitat selection"
  )
```

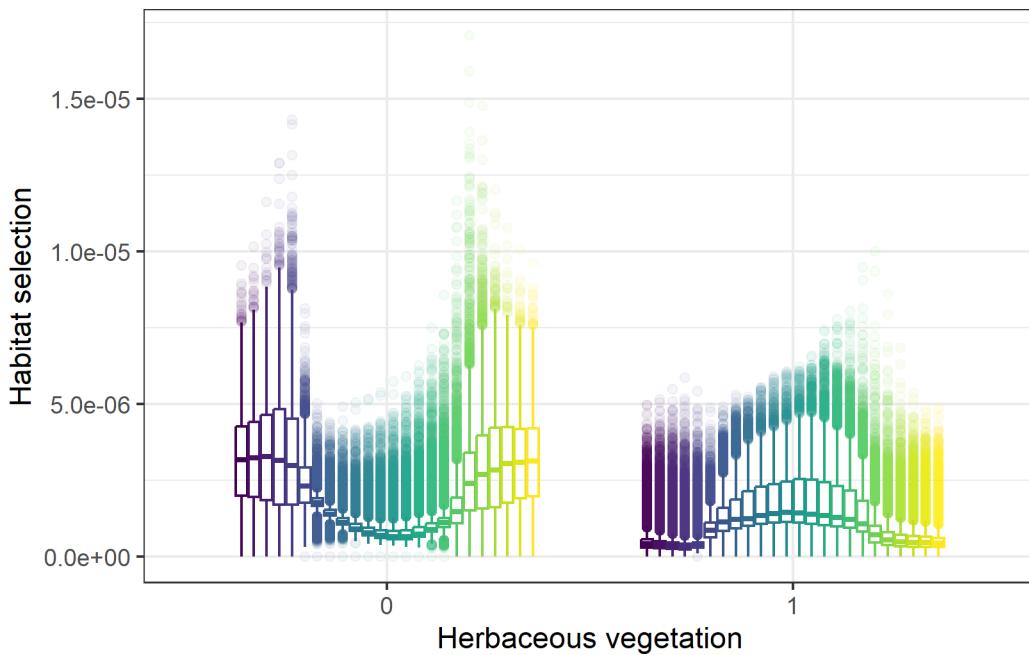
Warning: Removed 907 rows containing non-finite outside the scale range  
(`stat\_boxplot()`).



### All hours

```
habitat_selection_yday45_long %>%
  ggplot(aes(x = as.factor(Herbaceous_vegetation), y = Values, colour = as.factor(Hour))) +
  geom_boxplot(alpha = 0.05) +
  scale_colour_viridis(discrete = T) +
  theme_bw() +
  theme(legend.position = "none") +
  labs(
    x = "Herbaceous vegetation",
    y = "Habitat selection"
  )
```

Warning: Removed 21768 rows containing non-finite outside the scale range  
(`stat\_boxplot()`).

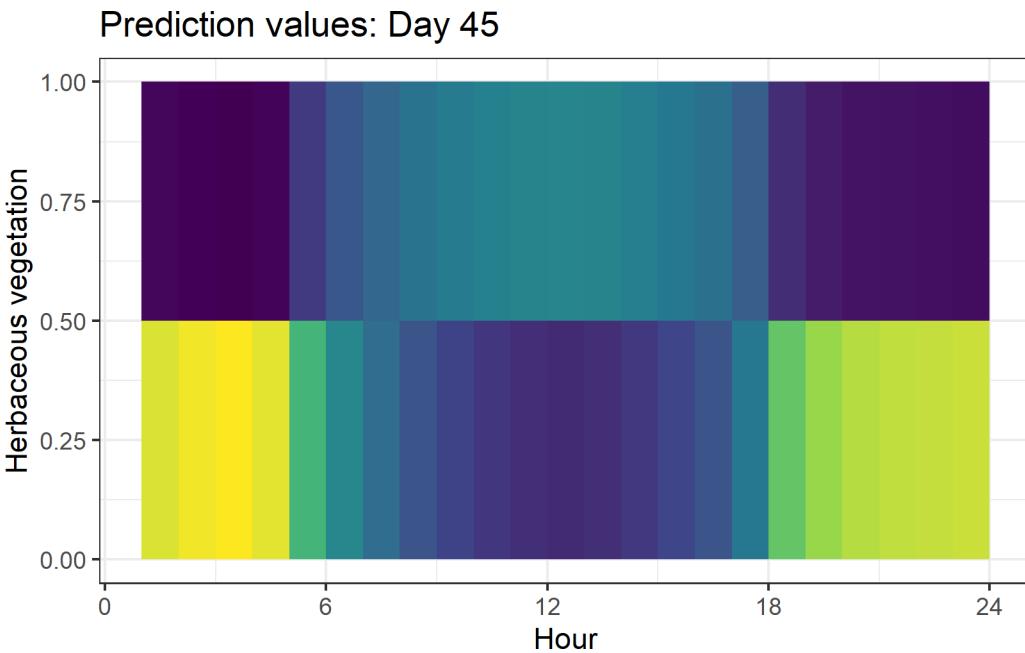


### Selection surface

Day of the year = 45 (mid-February)

```
ggplot(habitat_selection_yday45_long,
       aes(x = Hour, y = Herbaceous_vegetation)) +
  stat_summary_2d(aes(z = Values), fun = mean, bins = c(23, 2)) +
  scale_fill_viridis_c("Mean probability") +
  scale_x_continuous("Hour", breaks = seq(0, 24, 6)) +
  scale_y_continuous("Herbaceous vegetation") +
  ggtitle(paste0("Prediction values: Day 45")) +
  theme_bw() +
  theme(legend.position = "none")
```

Warning: Removed 21768 rows containing non-finite outside the scale range  
(`stat\_summary2d()`).



```
ggsave("outputs/herby_habitat_selection_yday45.png", width = 90, height = 75,
       units = "mm", dpi = 600)
```

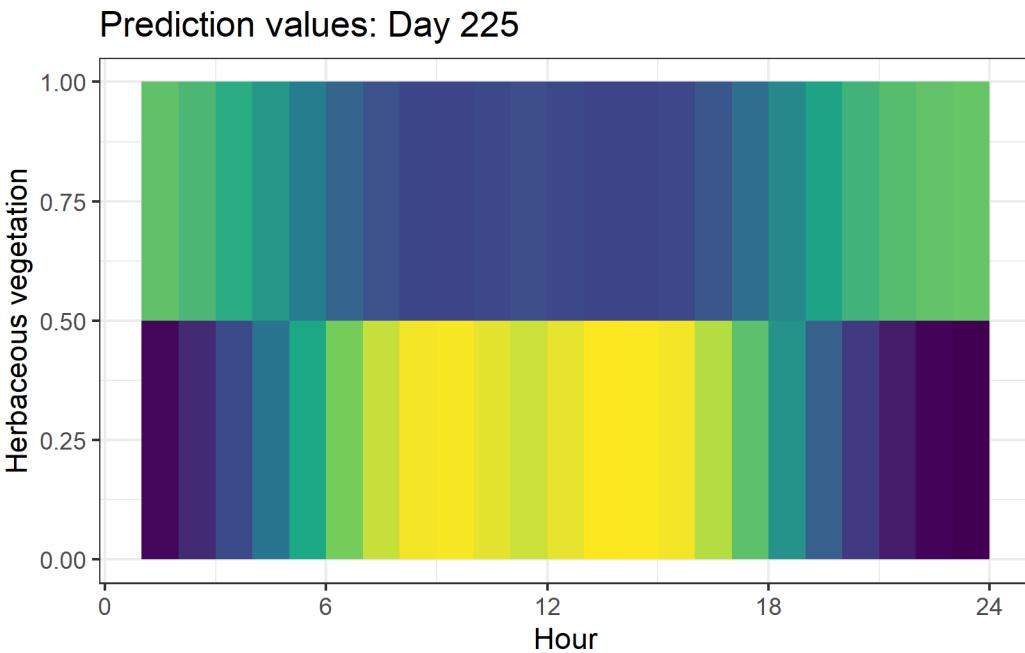
Warning: Removed 21768 rows containing non-finite outside the scale range  
(`stat\_summary2d()`).

Again there are seemingly opposing selection patterns between the wet and dry seasons.

Day of the year = 225 (mid-August)

```
ggplot(habitat_selection_yday225_long,
       aes(x = Hour, y = Herbaceous_vegetation)) +
  stat_summary_2d(aes(z = Values), fun = mean, bins = c(23, 2)) +
  scale_fill_viridis_c("Mean probability") +
  scale_x_continuous("Hour", breaks = seq(0,24,6)) +
  scale_y_continuous("Herbaceous vegetation") +
  ggtitle(paste0("Prediction values: Day 225")) +
  theme_bw() +
  theme(legend.position = "none")
```

Warning: Removed 2376 rows containing non-finite outside the scale range  
(`stat\_summary2d()`).



```
ggsave("outputs/herby_habitat_selection_yday225.png", width = 90, height = 75,
       units = "mm", dpi = 600)
```

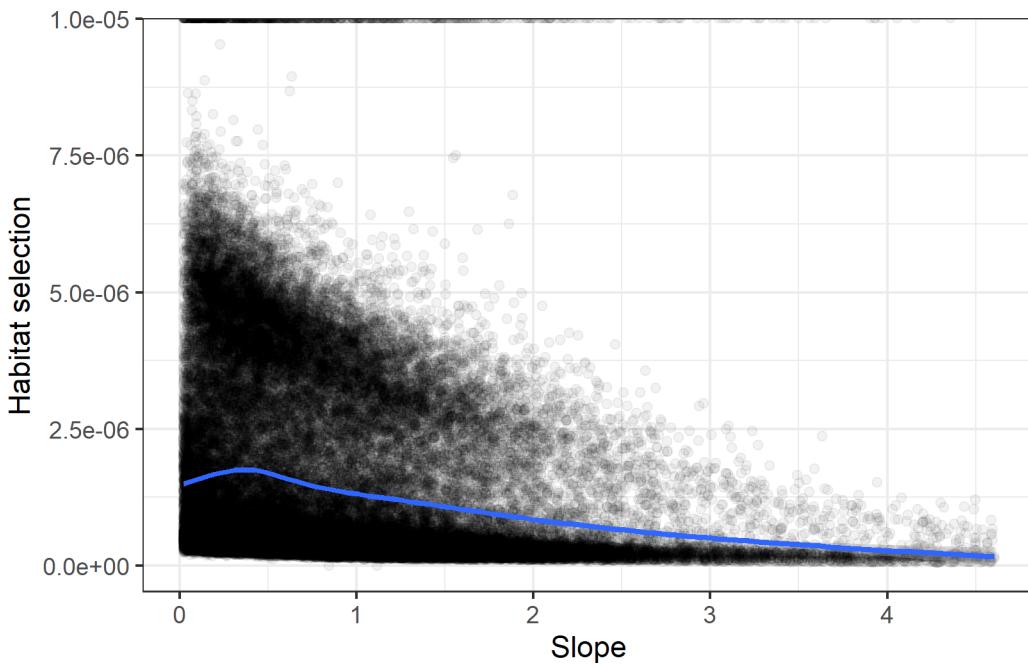
Warning: Removed 2376 rows containing non-finite outside the scale range  
`(`stat_summary2d()`).`

## Slope

```
habitat_selection_yday45_long %>% filter(Hour == 1) %>%
  ggplot(aes(x = Slope, y = Values)) +
  geom_point(alpha = 0.05) +
  geom_smooth(method = "gam") +
  theme_bw() +
  theme(legend.position = "bottom") +
  labs(
    x = "Slope",
    y = "Habitat selection"
  )
```

```
`geom_smooth()` using formula = 'y ~ s(x, bs = "cs")'
```

Warning: Removed 907 rows containing non-finite outside the scale range  
`(`stat_smooth()`).`

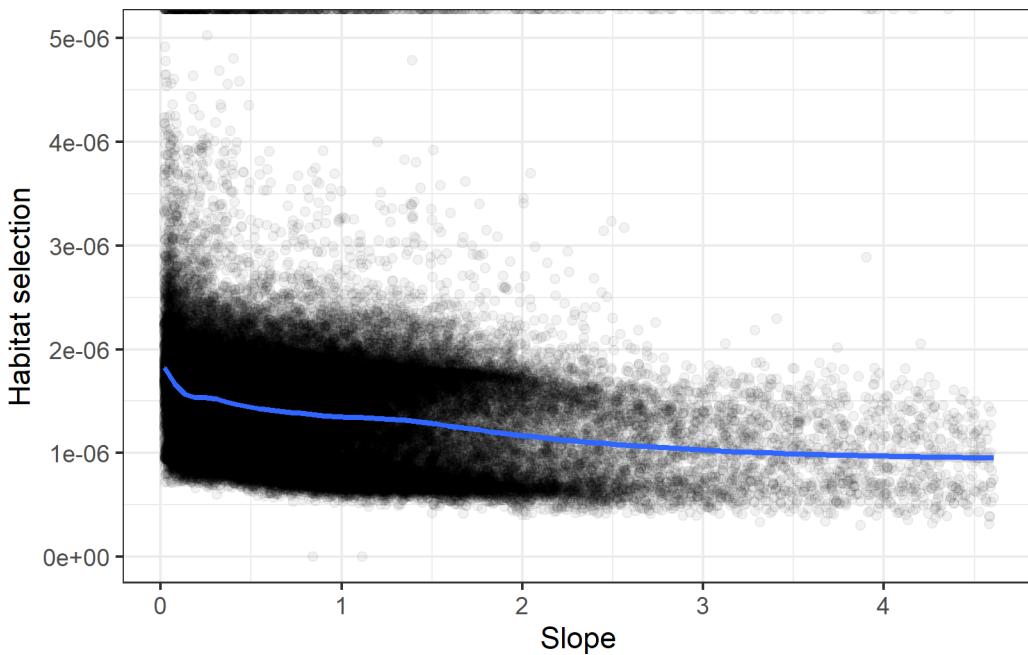


```

habitat_selection_yday45_long %>% filter(Hour == 7) %>%
  ggplot(aes(x = Slope, y = Values)) +
  geom_point(alpha = 0.05) +
  geom_smooth(method = "gam") +
  theme_bw() +
  theme(legend.position = "bottom") +
  labs(
    x = "Slope",
    y = "Habitat selection"
  )
`geom_smooth()` using formula = 'y ~ s(x, bs = "cs")'

```

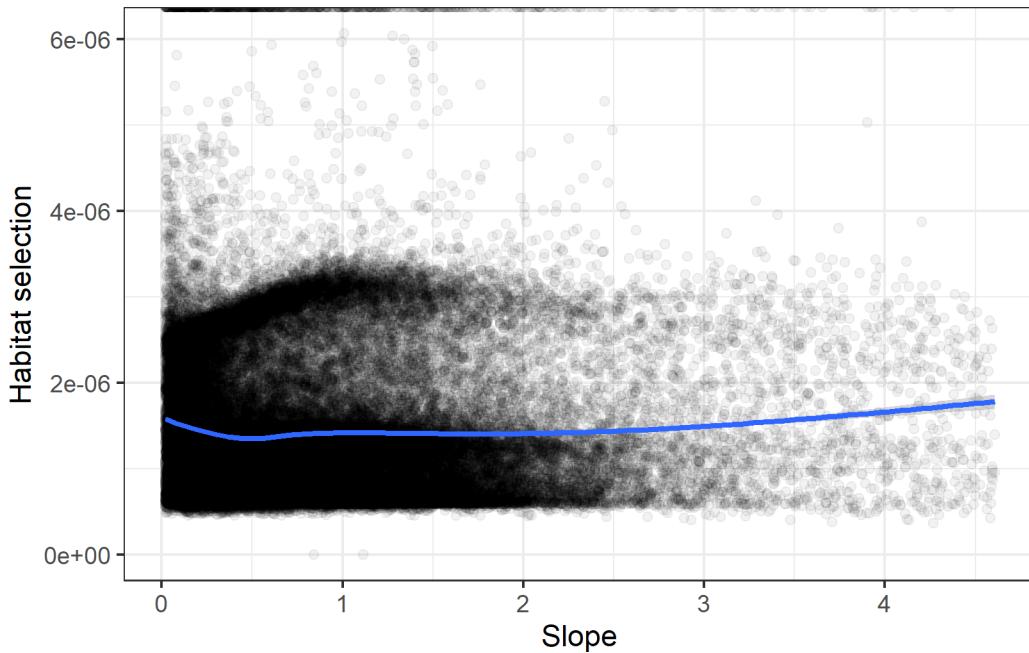
Warning: Removed 907 rows containing non-finite outside the scale range  
(`stat\_smooth()`).



```
habitat_selection_yday45_long %>% filter(Hour == 12) %>%
  ggplot(aes(x = Slope, y = Values)) +
  geom_point(alpha = 0.05) +
  geom_smooth(method = "gam") +
  theme_bw() +
  theme(legend.position = "bottom") +
  labs(
    x = "Slope",
    y = "Habitat selection"
  )
```

```
`geom_smooth()` using formula = 'y ~ s(x, bs = "cs")'
```

```
Warning: Removed 907 rows containing non-finite outside the scale range
(`stat_smooth()`).
```



### All hours

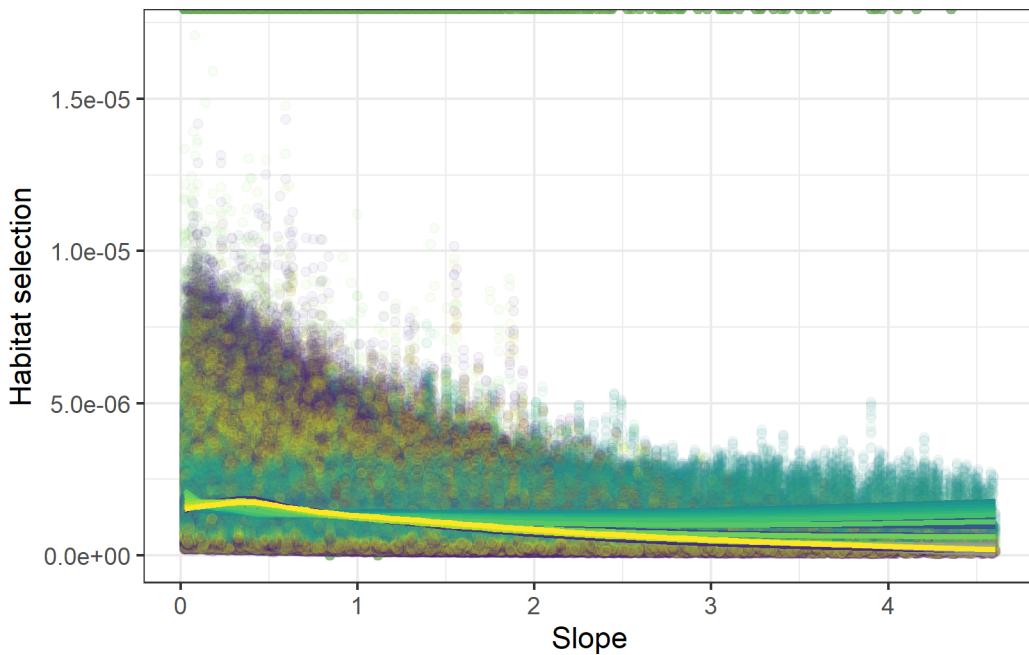
```

habitat_selection_yday45_long %>%
  ggplot(aes(x = Slope, y = Values, colour = factor(Hour))) +
  geom_jitter(alpha = 0.05) +
  geom_smooth(method = "gam") +
  scale_colour_viridis(discrete = T) +
  theme_bw() +
  theme(legend.position = "none") +
  labs(
    x = "Slope",
    y = "Habitat selection"
  )

`geom_smooth()` using formula = 'y ~ s(x, bs = "cs")'

Warning: Removed 21768 rows containing non-finite outside the scale range
(`stat_smooth()`).

```



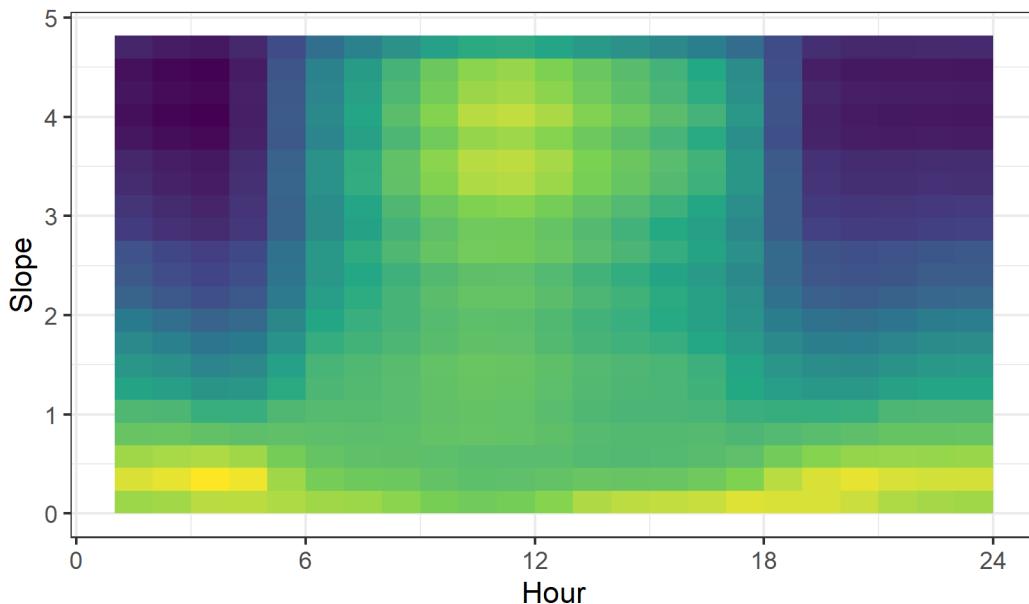
### Selection surface

Day of the year = 45 (mid-February)

```
ggplot(habitat_selection_yday45_long,
       aes(x = Hour, y = Slope)) +
  stat_summary_2d(aes(z = Values), fun = mean, bins = c(23,20)) +
  scale_fill_viridis_c("Mean probability") +
  scale_x_continuous("Hour", breaks = seq(0,24,6)) +
  scale_y_continuous("Slope") +
  ggtitle(paste0("Prediction values: Day 45")) +
  theme_bw() +
  theme(legend.position = "none")
```

Warning: Removed 21768 rows containing non-finite outside the scale range  
(`stat\_summary2d()`).

## Prediction values: Day 45



```
ggsave("outputs/slope_habitat_selection_yday45.png", width = 90, height = 75,  
       units = "mm", dpi = 600)
```

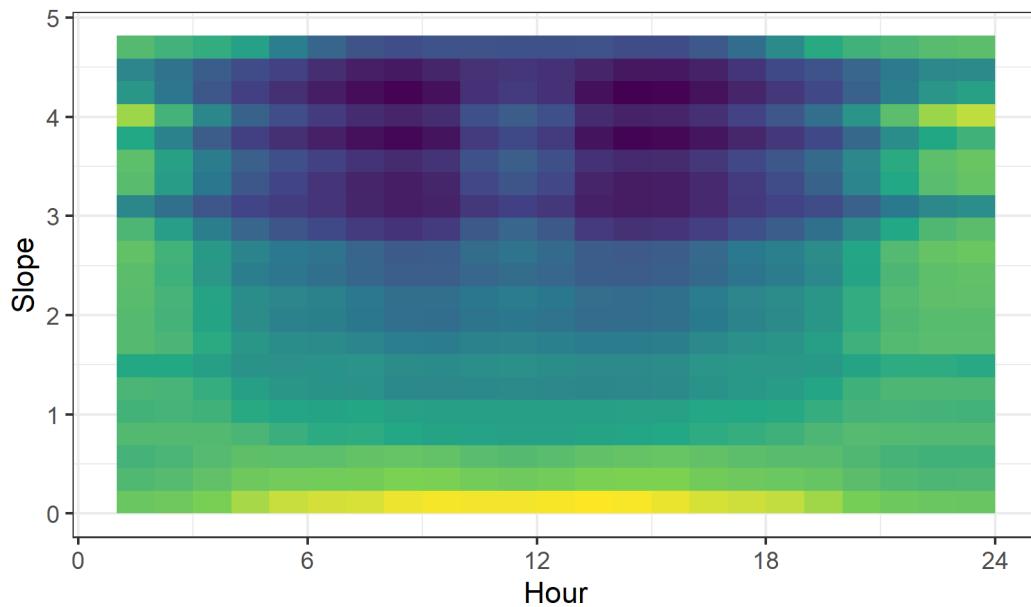
Warning: Removed 21768 rows containing non-finite outside the scale range  
(`stat\_summary2d()`).

Day of the year = 225 (mid-August)

```
ggplot(habitat_selection_yday225_long,  
       aes(x = Hour, y = Slope)) +  
  stat_summary_2d(aes(z = Values), fun = mean, bins = c(23,20)) +  
  scale_fill_viridis_c("Mean probability") +  
  scale_x_continuous("Hour", breaks = seq(0,24,6)) +  
  scale_y_continuous("Slope") +  
  ggtitle(paste0("Prediction values: Day 225")) +  
  theme_bw() +  
  theme(legend.position = "none")
```

Warning: Removed 2376 rows containing non-finite outside the scale range  
(`stat\_summary2d()`).

## Prediction values: Day 225



```
ggsave("outputs/slope_habitat_selection_yday225.png", width = 90, height = 75,  
       units = "mm", dpi = 600)
```

Warning: Removed 2376 rows containing non-finite outside the scale range  
(`stat\_summary2d()`).

Forrest, Scott W, Dan Pagendam, Michael Bode, Christopher Drovandi, Jonathan R Potts, Justin Perry, Eric Vanderduys, and Andrew J Hoskins. 2024. “Predicting Fine-scale Distributions and Emergent Spatiotemporal Patterns from Temporally Dynamic Step Selection Simulations.” *Ecography*, December. <https://doi.org/10.1111/ecog.07421>.