

Introduction

Background

Every year since 1995, citizen scientists gather for a weekend in September at the floodplain (the pools area) of the Mana Pools National Park and perform transects, with the goal of counting the number of animals of specific species in this pools area of the park. Counts are done in the mornings and the afternoons of the Saturday and Sunday.

Area

The Mana Pools National Park is located in Northern Zimbabwe against the Zambezi river. The park itself is 219600ha and is home to a large variety of game animals, including the big 5. A 4500ha area of the park is made up of a floodplain (the pools area) which retains water better than the rest of the park and generally has more lush vegetation.

Data available

The data available is present in 3 datasets. The first contains the counts of animal species found as a result of the aforementioned transects. Species counted are: elephant, buffalo, eland, zebra, waterbuck, kudu, impala, nyala, bushbuck, wild dog, lion, leopard, hyena, jackal, warthog and baboon. Counts are separated according to day and time of day. The second and third datasets are for climatic and vegetation conditions for the 219600ha park and the 4500ha pools area, respectively. The frequency of this climatic and vegetation data is monthly.

study objectives

The primary object of this study is to display my proficiency with wrangling and presenting data using R, thus any correlations that are determined are for interests sake alone. That being said, I plan to study the correlations between game animals present and the amount of precipitation which fell during the previous wet season in the park and pools area, and the correlations between game animals present and various park and pools climatic and vegetation and factors present in the dry season leading up to each game count. Considering that the pools area retains water better and produces better quality vegetation than the rest of the park, and that animals will migrate locally to find better feeding and drinking resources, I expect that one will see that animal counts will vary according to yearly climatic and vegetation scores (increasing with better food and water resources at the pools and poorer food and water resources in the rest of the park and vice-versa).

Table importing and cleaning

Installing the tidyverse

```
library(tidyverse)
```

```
## -- Attaching packages ----- tidyverse 1.3.1 --
```

```
## v ggplot2 3.3.5      v purrr   0.3.4
## v tibble  3.1.4      v dplyr   1.0.7
## v tidyr   1.1.3      v stringr 1.4.0
## v readr   2.0.1      v forcats 0.5.1
```

```
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag() masks stats::lag()
```

```
library(lubridate)
```

```
##
## Attaching package: 'lubridate'
```

```
## The following objects are masked from 'package:base':
##
## date, intersect, setdiff, union
```

Importing the game count df (gc), the park climatic data (pad) df, and the pools climatic data (roi) df.

```
gc <- read_csv('C:/Users/chris/Desktop/Coding/R_work/Projects/Game_counts/game_counts.csv')
```

```
## Rows: 80 Columns: 19
```

```
## -- Column specification -----
## Delimiter: ","
## chr (2): Day, Time
## dbl (17): Year, Elephant, Buffalo, Eland, Zebra, Waterbuck, Kudu, Impala, Ny...
```

```
##
## 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.
```

```
pad <- read_csv('C:/Users/chris/Desktop/Coding/R_work/Projects/Game_counts/PARK_additional_data.csv')
```

```
## Rows: 265 Columns: 18
```

```
## -- Column specification -----
## Delimiter: ","
## dbl (17): aet, def, pdsi, pet, pr, ro, soil, srad, swe, tmmn, tmmx, vap, vp...
## date (1): date
```

```
##
## 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.
```

```
roi <- read_csv('C:/Users/chris/Desktop/Coding/R_work/Projects/Game_counts/ROI_additional_data.csv')
```

```
## Rows: 265 Columns: 18
```

```
## -- Column specification -----
## Delimiter: ","
## dbl (17): aet, def, pdsi, pet, pr, ro, soil, srad, swe, tmmn, tmmx, vap, vp...
## date (1): date
```

```
##
## 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.
```

```
gc
```

```
## # A tibble: 80 x 19
##   Year Day      Time Elephant Buffalo Eland Zebra Waterbuck Kudu Impala Nyala
##   <dbl> <chr>   <chr>   <dbl>   <dbl> <dbl> <dbl>   <dbl> <dbl> <dbl> <dbl>
## 1 1995 Saturd~ am      155     381    179    460     234    66   3005    0
## 2 1995 Saturd~ pm      211     634    209    416     248   105   2345    0
## 3 1995 Sunday am      159     465    136    426     238    62   2376    0
## 4 1995 Sunday pm      173     304    171    581     242   140   2131    0
## 5 1996 Saturd~ am      210     376    173    273     350    54   4099    0
## 6 1996 Saturd~ pm      251     330    167    259     305    51   3443    0
## 7 1996 Sunday am      163     431    254    240     404    23   3686    0
## 8 1996 Sunday pm      327     443    233    231     336    47   2893    0
## 9 1997 Saturd~ am       83     516    260    226     393    55   3480    0
## 10 1997 Saturd~ pm      220     639    176    241     354    70   3455    0
## # ... with 70 more rows, and 8 more variables: Bushbuck <dbl>, Wild dog <dbl>,
## #   Lion <dbl>, Leopard <dbl>, Hyena <dbl>, Jackal <dbl>, Warthog <dbl>,
## #   Baboon <dbl>
```

Fortunately gc was small enough to take a quick scan through it. I noticed that in one year the Saturday and Sunday were recorded in shorthand ('Sat' and 'Sun'), so I converted them to Saturday and Sunday.

```
gc <- gc %>% mutate(Day = recode(Day, 'Sun' = 'Sunday', 'Sat' = 'Saturday'))
gc
```

```
## # A tibble: 80 x 19
##   Year Day      Time Elephant Buffalo Eland Zebra Waterbuck Kudu Impala Nyala
##   <dbl> <chr>   <chr>   <dbl>   <dbl> <dbl> <dbl>   <dbl> <dbl> <dbl> <dbl>
## 1 1995 Saturd~ am      155     381    179    460     234    66   3005    0
## 2 1995 Saturd~ pm      211     634    209    416     248   105   2345    0
## 3 1995 Sunday am      159     465    136    426     238    62   2376    0
## 4 1995 Sunday pm      173     304    171    581     242   140   2131    0
## 5 1996 Saturd~ am      210     376    173    273     350    54   4099    0
## 6 1996 Saturd~ pm      251     330    167    259     305    51   3443    0
## 7 1996 Sunday am      163     431    254    240     404    23   3686    0
## 8 1996 Sunday pm      327     443    233    231     336    47   2893    0
## 9 1997 Saturd~ am       83     516    260    226     393    55   3480    0
## 10 1997 Saturd~ pm      220     639    176    241     354    70   3455    0
## # ... with 70 more rows, and 8 more variables: Bushbuck <dbl>, Wild dog <dbl>,
## #   Lion <dbl>, Leopard <dbl>, Hyena <dbl>, Jackal <dbl>, Warthog <dbl>,
## #   Baboon <dbl>
```

As I'm interested in the total counts per year, I could collapse the counts for Saturdays and Sundays, and am and pm into single counts representing each year, this df I called gc_yearly. I then removed all carnivores, warthog and baboon as I'm interested in animals that will locally migrate to and from the pools area, and carnivores, baboons and warthogs generally have well defined territories and so won't migrate locally. I then multiplied the number of animals of a species by the species LSU (large stock unit). This gives a more realistic relative pressure of each species on the environment (a factor to consider when adding species to

make additional categories, such as browsers). I then added columns which summate the grazers, browsers and angostic herbivores, as these herbivore types use different vegetation types (grass, trees and shrubs, or both, respectively).

```
gc_yearly <- gc %>% group_by(Year) %>% summarize(
  Elephant=sum(Elephant, na.rm=TRUE), Buffalo=sum(Buffalo, na.rm=TRUE),
  Eland=sum(Eland, na.rm=TRUE), Zebra=sum(Zebra, na.rm=TRUE),
  Waterbuck=sum(Waterbuck, na.rm=TRUE), Kudu=sum(Kudu, na.rm=TRUE),
  Nyala=sum(Nyala, na.rm=TRUE), Bushbuck=sum(Bushbuck, na.rm=TRUE),
  Impala=sum(Impala, na.rm=TRUE)
)
gc_yearly <- gc_yearly %>% select(Year, Elephant, Buffalo, Eland, Zebra, Waterbuck, Kudu, Nyala, Bushbuck, Impala)
gc_yearly <- gc_yearly %>% mutate(
  Elephant=Elephant*10, Buffalo=Buffalo*1.07, Eland=Eland*1.08, Zebra=Zebra*1.09,
  Waterbuck=Waterbuck*1.1, Kudu=Kudu*1.11, Nyala=Nyala*1.12, Bushbuck=Bushbuck*1.13, Impala=Impala*1.14
)
gc_yearly <- gc_yearly %>% mutate(
  Grazers = Buffalo + Zebra, Browsers = Kudu + Bushbuck, Agnostic = Elephant + Waterbuck + Nyala + Bushbuck + Impala
)
gc_yearly
```

```
## # A tibble: 20 x 13
##   Year Elephant Buffalo Eland Zebra Waterbuck Kudu Nyala Bushbuck Impala
##   <dbl>     <dbl>   <dbl> <dbl> <dbl>     <dbl> <dbl> <dbl>   <dbl> <dbl>
## 1 1995      6980    1909.  751. 1243.     481  201.  0      1.3  1873.
## 2 1996      9510    1691.  893.  662.     698.  94.5  0      0.91 2683.
## 3 1997      5620    2181.  989.  599.     692  111.  0      0.65 2555.
## 4 1998      5870    3797.  977.  896.     630  119.  0      0.26 3124.
## 5 1999      8220    2223. 1161.  967.     717   68.6  0      0.39 2611.
## 6 2000      7700    5231. 1216. 1063.     863  197.  0      0.13 2690.
## 7 2001      6490    2985. 1112.  756.     834   57.2  0      0.26 2387.
## 8 2002      6390    1509. 1352.  707.     792  163.  0      0.52 3419.
## 9 2003      5140    2472. 1147.  591.     852   71.8  0      0.26 2396.
## 10 2004      5370    4014. 1064.  535.     863  145.  0.23  0     2481.
## 11 2005      7930    1559. 1264.  785.     934  270.  0.23  0.78 3326.
## 12 2006      9660    2391. 1053.  658.     642  126.  0      0.13 3082.
## 13 2007      9930    2400. 1356.  960.     648  172.  0.69  0.13 3933.
## 14 2008     10950    3122. 1232.  698.    1120  153.  0      0.26 3460.
## 15 2009     10870     681.  956.  323.     804.  94.5  0      0.78 3375.
## 16 2010     15300    2227.  927.  327.     840.  217.  0.23  0.52 3594.
## 17 2011     10690    1727.  807.  586.     718.  244.  0      0.13 4344.
## 18 2012      9900     745.  816.  500.     694.  216.  2.07  3.38 4418.
## 19 2013      9470     622.  671.  521.     574  186.  1.38  0.39 4777.
## 20 2014     13050    1655.  566.  530.     522  205.  0      0.26 4422.
## # ... with 3 more variables: Grazers <dbl>, Browsers <dbl>, Agnostic <dbl>
```

I then looked at the pad df.

```
pad
```

```
## # A tibble: 265 x 18
##   date      aet    def pdsi  pet    pr    ro  soil  srad  swe  tmmn
##   <date>    <dbl>  <dbl> <dbl> <dbl>  <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
## 1 1994-01-01 1267.    0    -172. 1267. 230.    11.5 1180. 2017.    0  210.
## 2 1994-02-01 1162.    4.02 -223. 1166. 130.    6.57 1255. 2122.    0  201.
## 3 1994-03-01 706.    862. -330. 1569. 27.2    1.32 807. 2700.    0  190.
## 4 1994-04-01 428.    963. -399. 1390. 23.1    1    598. 2355.    0  173.
## 5 1994-05-01 119.   1087. -458. 1206.  0      0    480. 2153.    0  139.
## 6 1994-06-01 77.6   903. -471.  981.  0      0    402. 1903.    0  112.
## 7 1994-07-01 55.3  1054. -468. 1109.  0      0    347. 1994.    0  117.
```

```
## 8 1994-08-01 41.5 1339. -447. 1381. 0 0 306. 2176. 0 138.
## 9 1994-09-01 34.4 1811. -420. 1846. 0.0519 0 273. 2604. 0 181.
## 10 1994-10-01 284. 1707. -326. 1990. 27.1 1.19 247. 2454. 0 212.
## # ... with 255 more rows, and 7 more variables: tmmx <dbl>, vap <dbl>,
## # vpd <dbl>, vs <dbl>, NDVI <dbl>, EVI <dbl>, water <dbl>
```

I renamed the columns to their full names and dropped the columns which are irrelevant to the study. I then used the date column to create a Year column and a Month column. As the game counts occurred in the beginning of September every year, I added two columns which gave an altered Year and Month value such that the Year_altered and Month_altered columns ran from September to August. This would make it easier to group climatic data for the periods before game counts were done.

```
pad <- pad %>% rename(Actual_evapotranspiration=aet,
                      Climate_water_deficit=def,
                      Palmer_drought_severity_index=pdsi,
                      Predicted_evapotranspiration=pet,
                      Precipitation=pr,
                      Runoff=ro,
                      Soil_moisture=soil,
                      Surface_radiation=srad,
                      Snow_water_Equivalent=swe,
                      Min_temp=tmmn,
                      Max_temp=tmmx,
                      Vapor_pressure=vap,
                      Vapor_pressure_deficit=vpd,
                      Wind_speed=vs,
                      Normalized_diff_veg_index=NDVI,
                      Enhanced_veg_index=EVI,
                      Water=water)
pad <- pad %>% select(c(1,4,6,8,16,17,18))
pad <- pad %>% mutate(Year=year(date),Month=month(date))
pad <- pad %>% select(c(2,3,4,5,6,7,8,9))
pad <- pad %>% mutate(Year_altered=lead(Year, n=4))
pad <- pad %>% mutate(Month_altered=lead(Month,n=4))
pad
```

```
## # A tibble: 265 x 10
##   Palmer_drought_severity_index Precipitation Soil_moisture Normalized_diff_ve-
##                                <dbl>          <dbl>          <dbl>          <dbl>
## 1                               -172.         230.         1180.           0
## 2                               -223.         130.         1255.           0
## 3                               -330.          27.2          807.           0
## 4                               -399.          23.1          598.           0
## 5                               -458.           0          480.           0
## 6                               -471.           0          402.           0
## 7                               -468.           0          347.           0
## 8                               -447.           0          306.           0
## 9                               -420.          0.0519          273.           0
## 10                              -326.          27.1          247.           0
## # ... with 255 more rows, and 6 more variables: Enhanced_veg_index <dbl>,
## #   Water <dbl>, Year <dbl>, Month <dbl>, Year_altered <dbl>,
## #   Month_altered <dbl>
```

I then applied to roi what I had applied to pad.

```

roi <- roi %>% rename(Actual_evapotranspiration=aet,
                     Climate_water_deficit=def,
                     Palmer_drought_severity_index=pdsi,
                     Predicted_evapotranspiration=pet,
                     Precipitation=pr,
                     Runoff=ro,
                     Soil_moisture=soil,
                     Surface_radiation=srad,
                     Snow_water_Equivalent=swe,
                     Min_temp=tmmn,
                     Max_temp=tmmx,
                     Vapor_pressure=vap,
                     Vapor_pressure_deficit=vpd,
                     Wind_speed=vs,
                     Normalized_diff_veg_index=NDVI,
                     Enhanced_veg_index=EVI,
                     Water=water)
roi <- roi %>% select(c(1,4,6,8,16,17,18))
roi <- roi %>% mutate(Year=year(date),Month=month(date))
roi <- roi %>% select(c(2,3,4,5,6,7,8,9))
roi <- roi %>% mutate(Year_altered=lead(Year, n=4))
roi <- roi %>% mutate(Month_altered=lead(Month,n=4))
roi

```

```

## # A tibble: 265 x 10
##   Palmer_drought_severity_index Precipitation Soil_moisture Normalized_diff_ve-
##   <dbl> <dbl> <dbl> <dbl>
## 1 -176. 225. 1147. 0
## 2 -220. 125. 1151. 0
## 3 -328. 20.1 799. 0
## 4 -397. 20.2 620. 0
## 5 -447. 0 509. 0
## 6 -457. 0 433. 0
## 7 -454. 0 377. 0
## 8 -437. 0 335. 0
## 9 -411. 0 301. 0
## 10 -322. 25.8 274. 0
## # ... with 255 more rows, and 6 more variables: Enhanced_veg_index <dbl>,
## #   Water <dbl>, Year <dbl>, Month <dbl>, Year_altered <dbl>,
## #   Month_altered <dbl>

```

I then found the total precipitation for the prior wet season (Months `altered` 2 to 9, annually) and the Palmer_drought_severity_index, Soil_moisture, Enhanced_veg_index and Water for the 3 months prior to the game count (Months `altered` 10 to 12, annually). These new dfs were named `pad_wet_precip` and `pad_dry_clim`, respectively for the park data and `roi_wet_precip` and `roi_dry_clim`, respectively for the pools data. To these dfs I added a column to represent the area the df was referring to to (i.e. pools or park).

```

pad_wet_precip <- pad %>% filter(Month_altered >= 2 & Month_altered <= 9) %>% group_by(Year_altered) %>%
pad_dry_clim <- pad %>% filter(Month_altered==10|Month_altered==11|Month_altered==12) %>% group_by(Year_altered) %>%

roi_wet_precip <- roi %>% filter(Month_altered >= 2 & Month_altered <= 9) %>% group_by(Year_altered) %>%
roi_dry_clim <- roi %>% filter(Month_altered==10|Month_altered==11|Month_altered==12) %>% group_by(Year_altered) %>%

```

```
pad_wet_precip
```

```
## # A tibble: 22 x 3
##   Year Precipitation Area
##   <dbl>         <dbl> <chr>
## 1 1994           411. park
## 2 1995           512. park
## 3 1996           683. park
## 4 1997          1036. park
## 5 1998           678. park
## 6 1999           950. park
## 7 2000           824. park
## 8 2001          1028. park
## 9 2002           651. park
## 10 2003           912. park
## # ... with 12 more rows
```

```
pad_dry_clim
```

```
## # A tibble: 22 x 6
##   Year Palmer_drought_severity_in~ Soil_moisture Enhanced_veg_ind~ Water Area
##   <dbl>         <dbl>         <dbl>         <dbl> <dbl> <chr>
## 1 1994          -1386.         1055.           0 3.02 park
## 2 1995          -2338.           514.           0 2.02 park
## 3 1996          -1322.         1361.           0 2.01 park
## 4 1997          1424.         2886.           0 2.11 park
## 5 1998          -911.         1261.           0 2.02 park
## 6 1999           703.         2167.           0 1.01 park
## 7 2000           879.         2461.        6662. 0.846 park
## 8 2001          1489.         2556.        7333. 2.02 park
## 9 2002         -1106.         1055.        6086. 1.01 park
## 10 2003           703.         2345.        7121. 1.01 park
## # ... with 12 more rows
```

```
roi_wet_precip
```

```
## # A tibble: 22 x 3
##   Year Precipitation Area
##   <dbl>         <dbl> <chr>
## 1 1994           390. pools
## 2 1995           515. pools
## 3 1996           661. pools
## 4 1997           976. pools
## 5 1998           637. pools
## 6 1999           894. pools
## 7 2000           781. pools
## 8 2001           990. pools
## 9 2002           620. pools
## 10 2003           899. pools
## # ... with 12 more rows
```

```
roi_dry_clim
```

```
## # A tibble: 22 x 6
##   Year Palmer_drought_severity_in~ Soil_moisture Enhanced_veg_ind~ Water Area
##   <dbl>           <dbl>           <dbl>           <dbl> <dbl> <chr>
## 1 1994          -1348.           1146.             0   3.33 pools
## 2 1995          -2185.             485.             0   2.23 pools
## 3 1996          -1323.           1331.             0   2.20 pools
## 4 1997           1472.           3435.             0   2.21 pools
## 5 1998          -937.           1329.             0   2.21 pools
## 6 1999           895.           2761.             0   1.11 pools
## 7 2000           633.           2285.          5011.   1.02 pools
## 8 2001          1632.           3261.          5826.   2.27 pools
## 9 2002         -1104.           1111.          4941.   1.13 pools
## 10 2003           812.           2561.          5715.   1.15 pools
## # ... with 12 more rows
```

I then merged pad_wet_precip and roi_wet_precip, and pad_dry_clim and roi_dry_clim. These dfs were called wet_precip and dry_clim, respectively.

```
wet_precip <- pad_wet_precip %>% full_join(roi_wet_precip) %>% arrange(Year)
```

```
## Joining, by = c("Year", "Precipitation", "Area")
```

```
wet_precip
```

```
## # A tibble: 44 x 3
##   Year Precipitation Area
##   <dbl>           <dbl> <chr>
## 1 1994           411. park
## 2 1994           390. pools
## 3 1995           512. park
## 4 1995           515. pools
## 5 1996           683. park
## 6 1996           661. pools
## 7 1997          1036. park
## 8 1997           976. pools
## 9 1998           678. park
## 10 1998           637. pools
## # ... with 34 more rows
```

```
dry_clim <- pad_dry_clim %>% full_join(roi_dry_clim) %>% arrange(Year)
```

```
## Joining, by = c("Year", "Palmer_drought_severity_index", "Soil_moisture", "Enhanced_veg_index", "Wat
```

```
dry_clim
```

```
## # A tibble: 44 x 6
##   Year Palmer_drought_severity_in~ Soil_moisture Enhanced_veg_ind~ Water Area
##   <dbl>           <dbl>           <dbl>           <dbl> <dbl> <chr>
```



```
## 1 1994 -1386. 1055. 0 3.02 park
## 2 1994 -1348. 1146. 0 3.33 pools
## 3 1995 -2338. 514. 0 2.02 park
## 4 1995 -2185. 485. 0 2.23 pools
## 5 1996 -1322. 1361. 0 2.01 park
## 6 1996 -1323. 1331. 0 2.20 pools
## 7 1997 1424. 2886. 0 2.11 park
## 8 1997 1472. 3435. 0 2.21 pools
## 9 1998 -911. 1261. 0 2.02 park
## 10 1998 -937. 1329. 0 2.21 pools
## # ... with 34 more rows
```

I then merged gc_yearly and wet_precip to make gc_precip and gc_yearly and dry_clim to make gc_clim

```
gc_precip <- gc_yearly %>% full_join(wet_precip)
```

```
## Joining, by = "Year"
```

```
gc_clim <- gc_yearly %>% full_join(dry_clim)
```

```
## Joining, by = "Year"
```

```
gc_precip
```

```
## # A tibble: 44 x 15
##   Year Elephant Buffalo Eland Zebra Waterbuck Kudu Nyala Bushbuck Impala
##   <dbl>   <dbl>   <dbl> <dbl> <dbl>   <dbl> <dbl> <dbl>   <dbl> <dbl>
## 1 1995     6980    1909.  751. 1243.    481  201.    0     1.3  1873.
## 2 1995     6980    1909.  751. 1243.    481  201.    0     1.3  1873.
## 3 1996     9510    1691.  893.  662.    698.  94.5    0     0.91 2683.
## 4 1996     9510    1691.  893.  662.    698.  94.5    0     0.91 2683.
## 5 1997     5620    2181.  989.  599.    692  111.    0     0.65 2555.
## 6 1997     5620    2181.  989.  599.    692  111.    0     0.65 2555.
## 7 1998     5870    3797.  977.  896.    630  119.    0     0.26 3124.
## 8 1998     5870    3797.  977.  896.    630  119.    0     0.26 3124.
## 9 1999     8220    2223. 1161  967.    717  68.6    0     0.39 2611.
## 10 1999     8220    2223. 1161  967.    717  68.6    0     0.39 2611.
## # ... with 34 more rows, and 5 more variables: Grazers <dbl>, Browsers <dbl>,
## #   Agnostic <dbl>, Precipitation <dbl>, Area <chr>
```

```
gc_clim
```

```
## # A tibble: 44 x 18
##   Year Elephant Buffalo Eland Zebra Waterbuck Kudu Nyala Bushbuck Impala
##   <dbl>   <dbl>   <dbl> <dbl> <dbl>   <dbl> <dbl> <dbl>   <dbl> <dbl>
## 1 1995     6980    1909.  751. 1243.    481  201.    0     1.3  1873.
## 2 1995     6980    1909.  751. 1243.    481  201.    0     1.3  1873.
## 3 1996     9510    1691.  893.  662.    698.  94.5    0     0.91 2683.
## 4 1996     9510    1691.  893.  662.    698.  94.5    0     0.91 2683.
## 5 1997     5620    2181.  989.  599.    692  111.    0     0.65 2555.
## 6 1997     5620    2181.  989.  599.    692  111.    0     0.65 2555.
```

```
## 7 1998      5870    3797.  977.  896.      630 119.      0    0.26 3124.
## 8 1998      5870    3797.  977.  896.      630 119.      0    0.26 3124.
## 9 1999      8220    2223. 1161  967.      717  68.6     0    0.39 2611.
## 10 1999     8220    2223. 1161  967.      717  68.6     0    0.39 2611.
## # ... with 34 more rows, and 8 more variables: Grazers <dbl>, Browsers <dbl>,
## #   Agnostic <dbl>, Palmer_drought_severity_index <dbl>, Soil_moisture <dbl>,
## #   Enhanced_veg_index <dbl>, Water <dbl>, Area <chr>
```

Incase the y axis scales needed for graphs resulted in difficulty seeing the trends in some species/groups while encompassing other species/groups. I also created a table with species/group column normalized as a percentage of the maximum value for the species/group seeing in a single year.

```
gc_max <- gc_yearly %>% summarize(across(2:13, max)) %>% mutate(Year=100) %>% relocate(Year)
gc_max <- gc_max %>% slice(rep(1:n(), each=20))
gc_yearly_norm <- gc_yearly/gc_max*100

gc_norm_precip <- gc_yearly_norm %>% full_join(wet_precip)
```

```
## Joining, by = "Year"
```

```
gc_norm_clim <- gc_yearly_norm %>% full_join(dry_clim)
```

```
## Joining, by = "Year"
```

Data analysis

Correlation between precipitation and animal counts

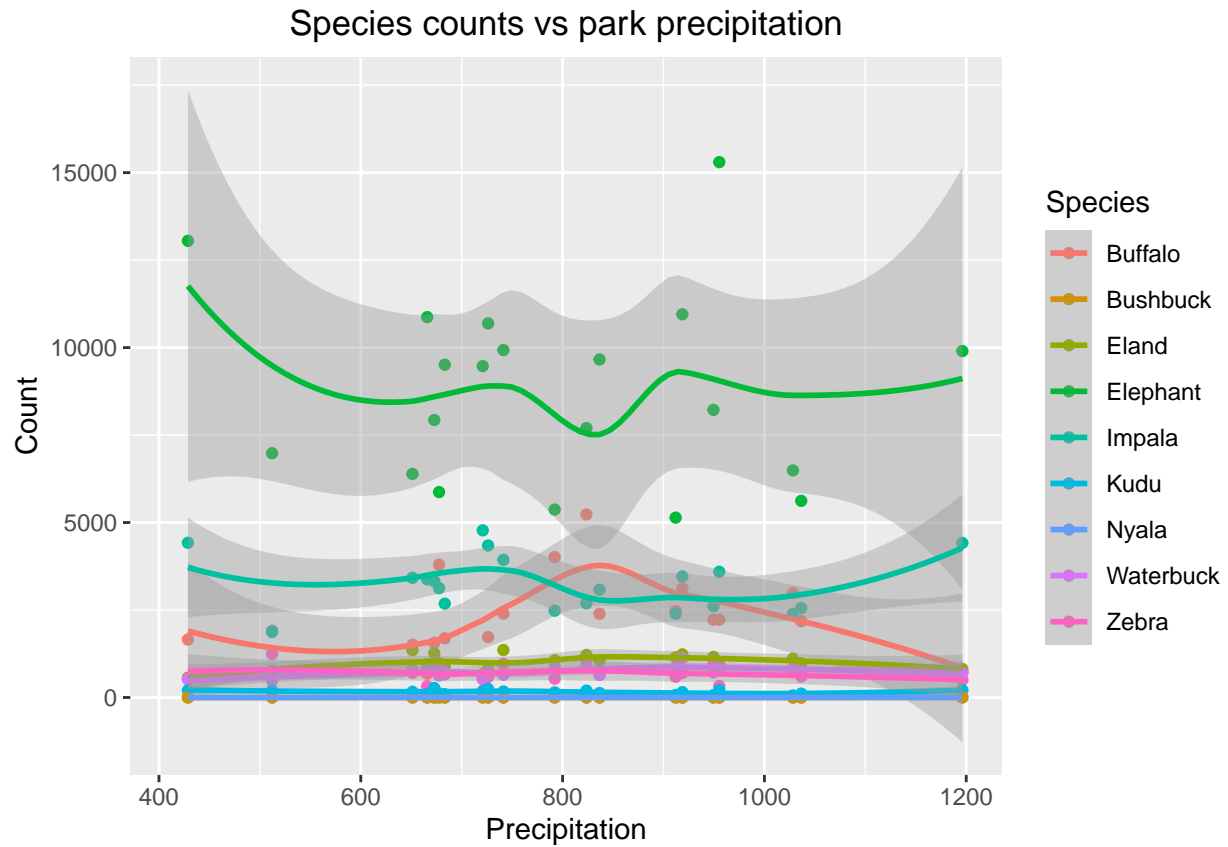
I graphed park and pools precipitations against LSU values for each species, and park and pools precipitations against LSU values for each herbivore type.

```
gc_precip_for_spp_plot <- gc_precip %>% gather(c(2,3,4,5,6,7,8,9,10), key='Species', value='Count')
ggplot(data=filter(gc_precip_for_spp_plot, Area=="park")) + geom_point(mapping=aes(x=Precipitation, y=C

## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'

## Warning: Removed 18 rows containing non-finite values (stat_smooth).

## Warning: Removed 18 rows containing missing values (geom_point).
```



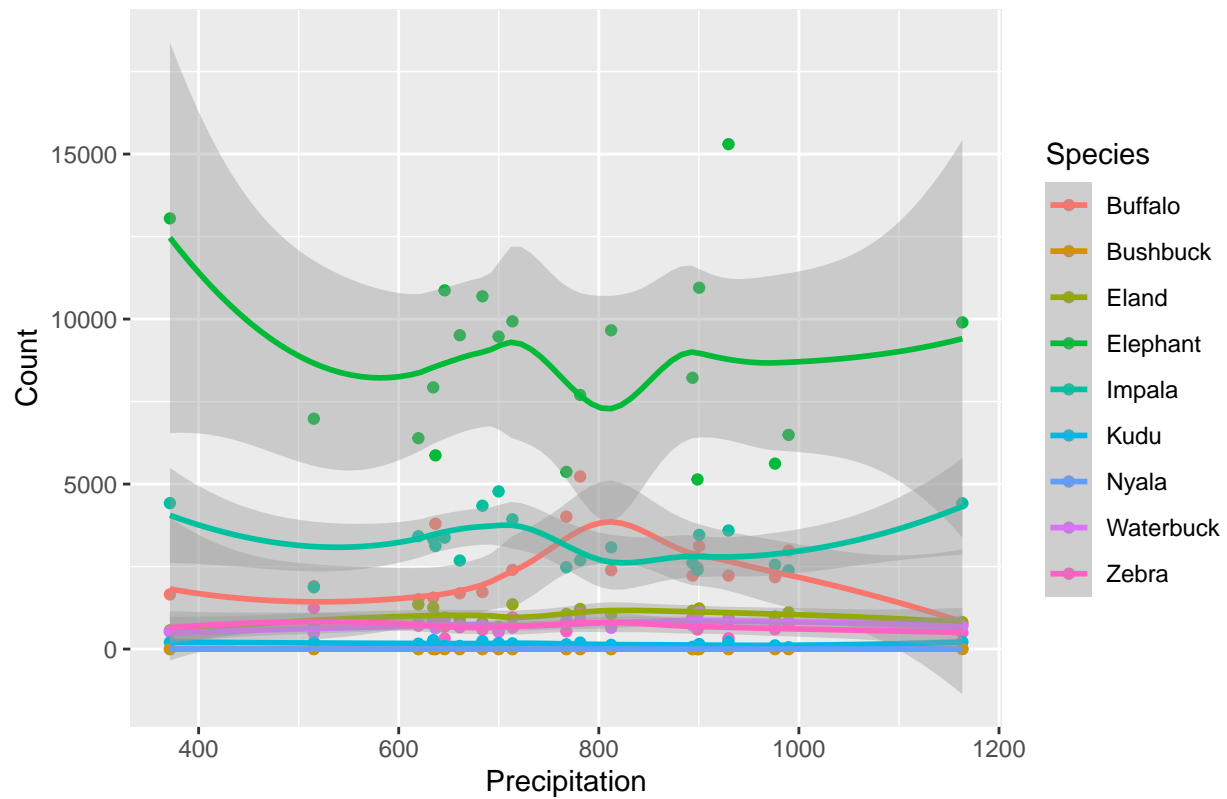
```
ggplot(data=filter(gc_precip_for_spp_plot, Area=="pools")) + geom_point(mapping=aes(x=Precipitation, y=
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 18 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 18 rows containing missing values (geom_point).
```

Species counts vs pools precipitation



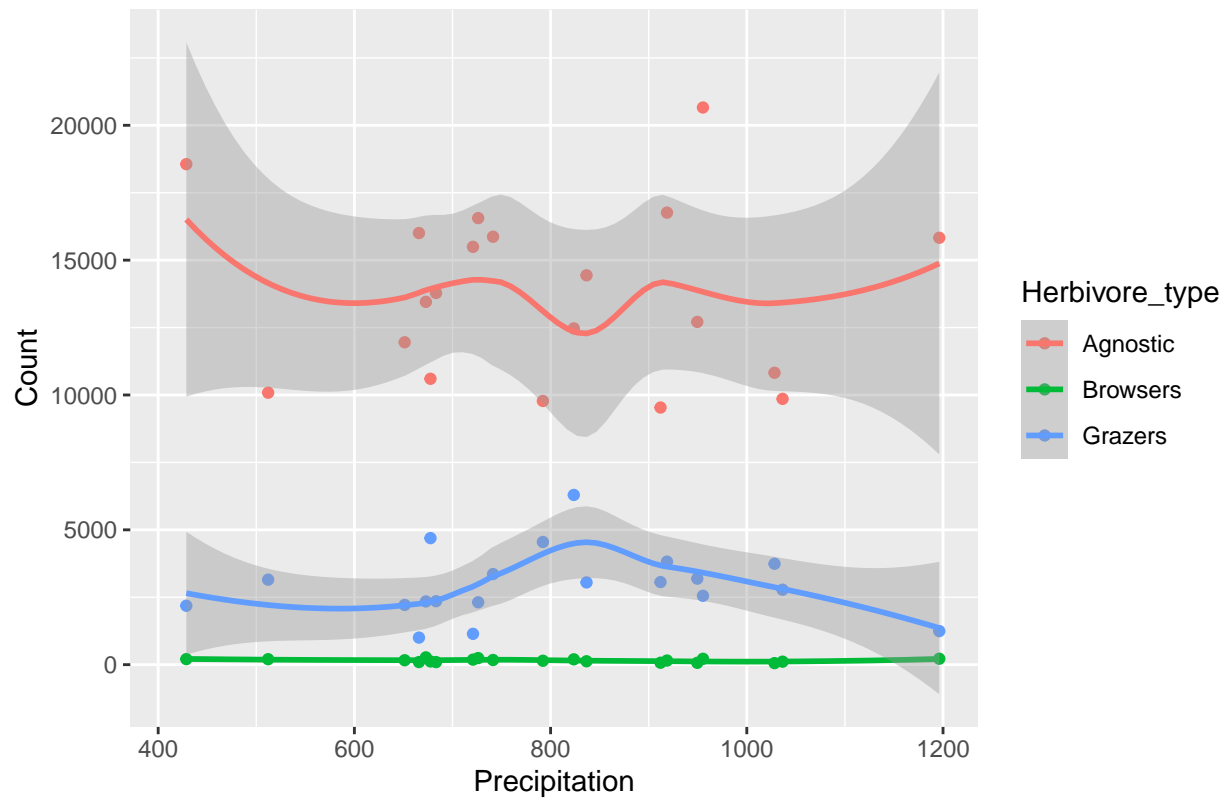
```
gc_precip_for_type_plot <- gc_precip %>% gather(c(11,12,13), key='Herbivore_type', value='Count')
ggplot(data=filter(gc_precip_for_type_plot, Area=="park")) + geom_point(mapping=aes(x=Precipitation, y=Count)) +
  geom_smooth(mapping=aes(x=Precipitation, y=Count, color=Herbivore_type), method='loess', se=TRUE)

## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'

## Warning: Removed 6 rows containing non-finite values (stat_smooth).

## Warning: Removed 6 rows containing missing values (geom_point).
```

Herbivore type counts vs park precipitation

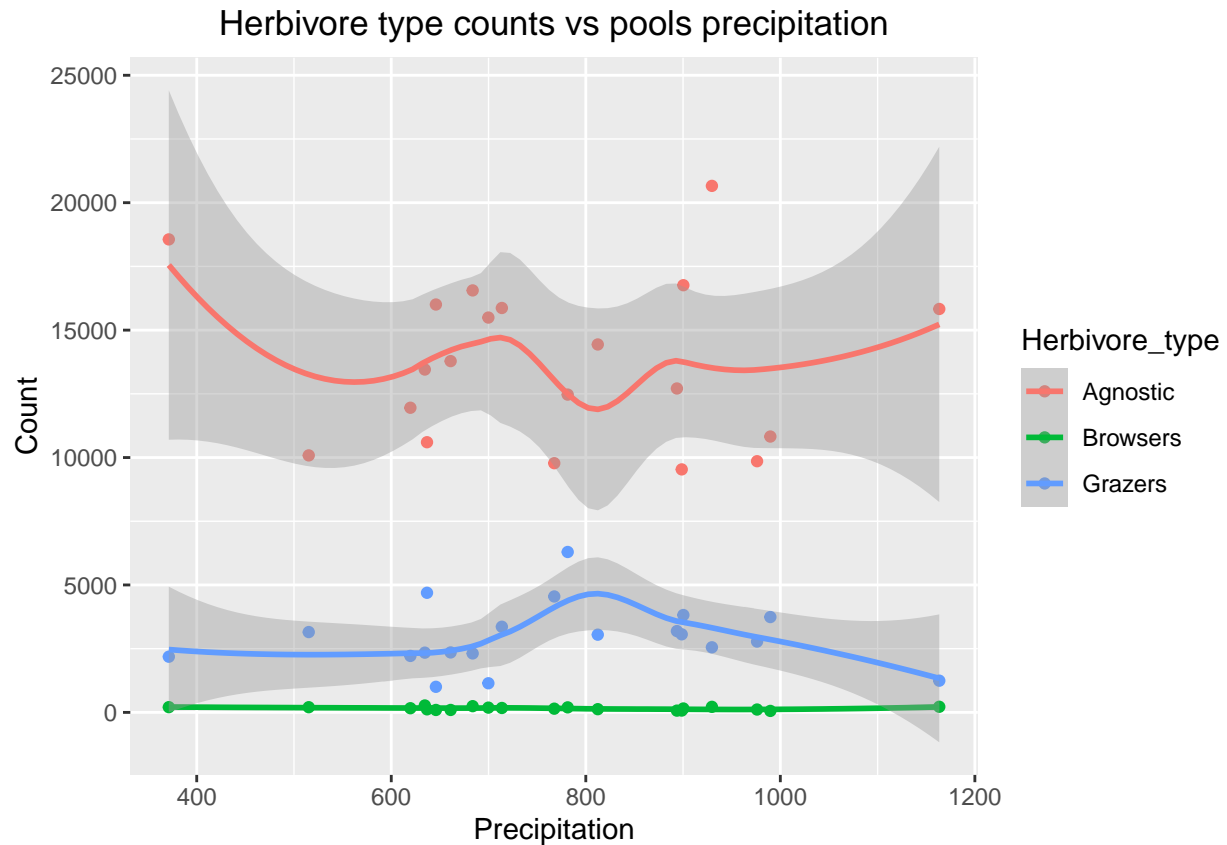


```
ggplot(data=filter(gc_precip_for_type_plot, Area=="pools")) + geom_point(mapping=aes(x=Precipitation, y=
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 6 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 6 rows containing missing values (geom_point).
```



I then graphed the park and pools precipitations against the normalized LSU values for each species, and park and pools precipitations against the normalized LSU values for each herbivore type.

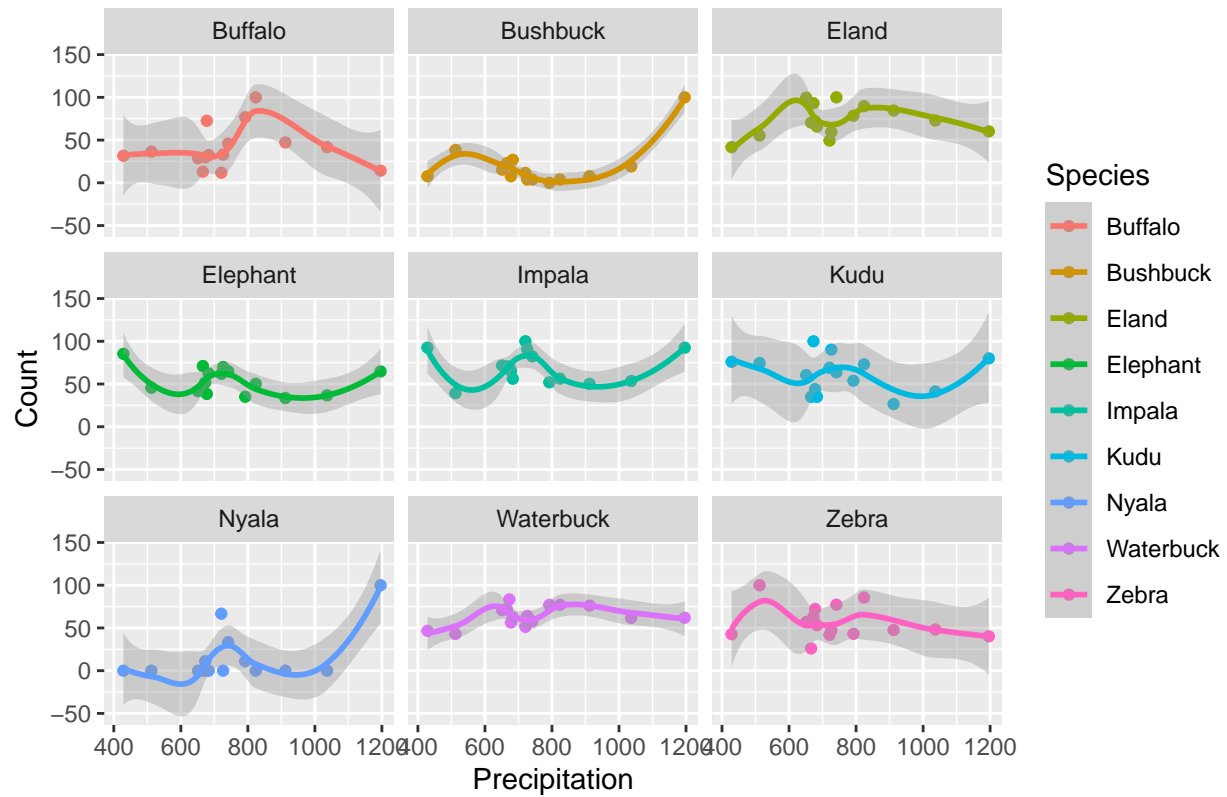
```
gc_precip_norm_spp_plot <- gc_norm_precip %>% gather(c(2,3,4,5,6,7,8,9,10), key='Species', value='Count')
ggplot(data=filter(gc_precip_norm_spp_plot, Area=="park")) + geom_point(mapping=aes(x=Precipitation, y=Count))

## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'

## Warning: Removed 63 rows containing non-finite values (stat_smooth).

## Warning: Removed 63 rows containing missing values (geom_point).
```

Normalized species counts vs park precipitation



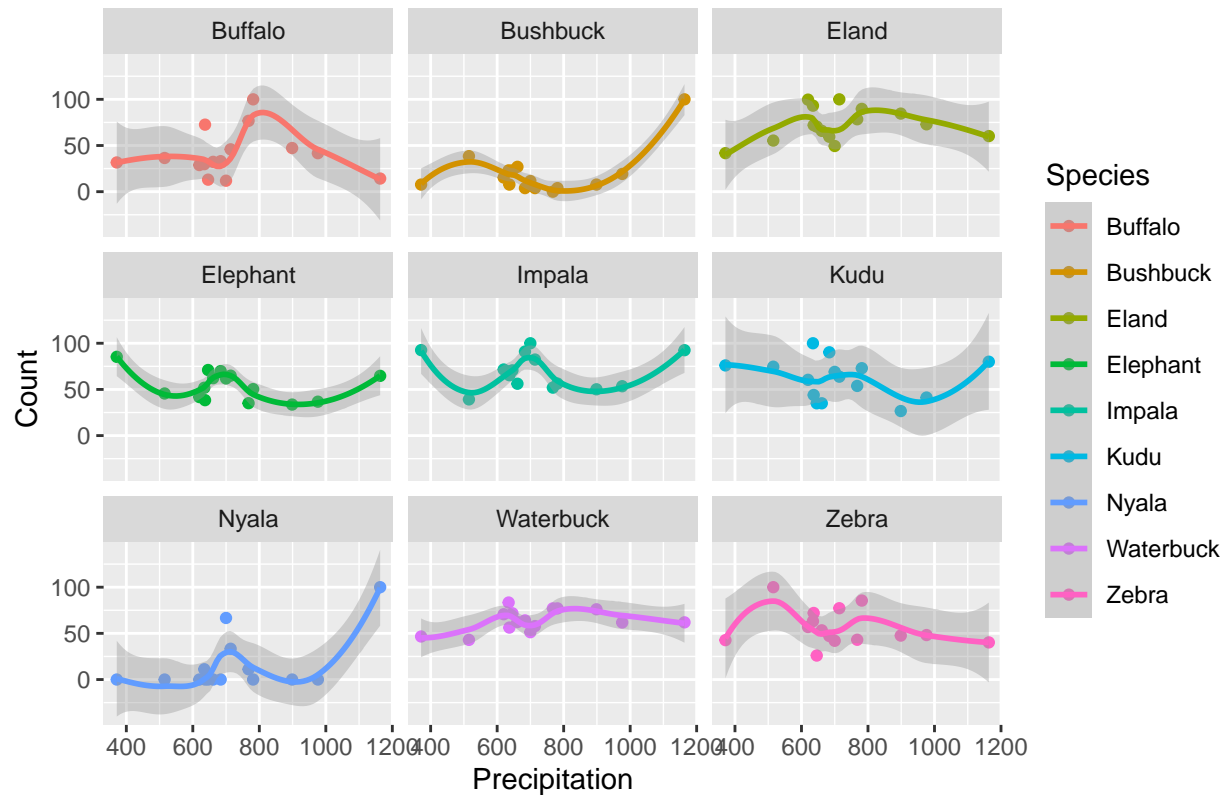
```
ggplot(data=filter(gc_precip_norm_spp_plot, Area=="pools")) + geom_point(mapping=aes(x=Precipitation, y=Count))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 63 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 63 rows containing missing values (geom_point).
```

Normalized species counts vs pools precipitation



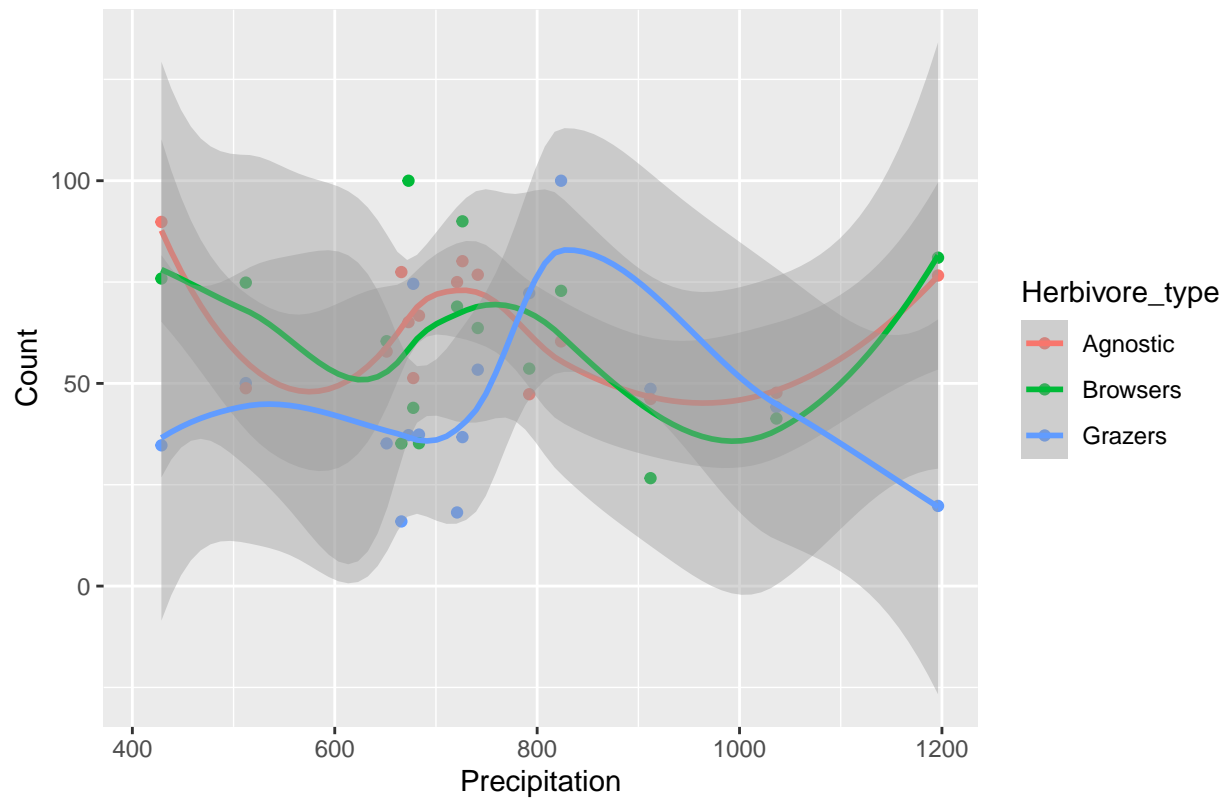
```
gc_precip_norm_type_plot <- gc_norm_precip %>% gather(c(11,12,13), key='Herbivore_type', value='Count')
ggplot(data=filter(gc_precip_norm_type_plot, Area=="park")) + geom_point(mapping=aes(x=Precipitation, y=
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 21 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 21 rows containing missing values (geom_point).
```


Normalized herbivore type counts vs park precipitation



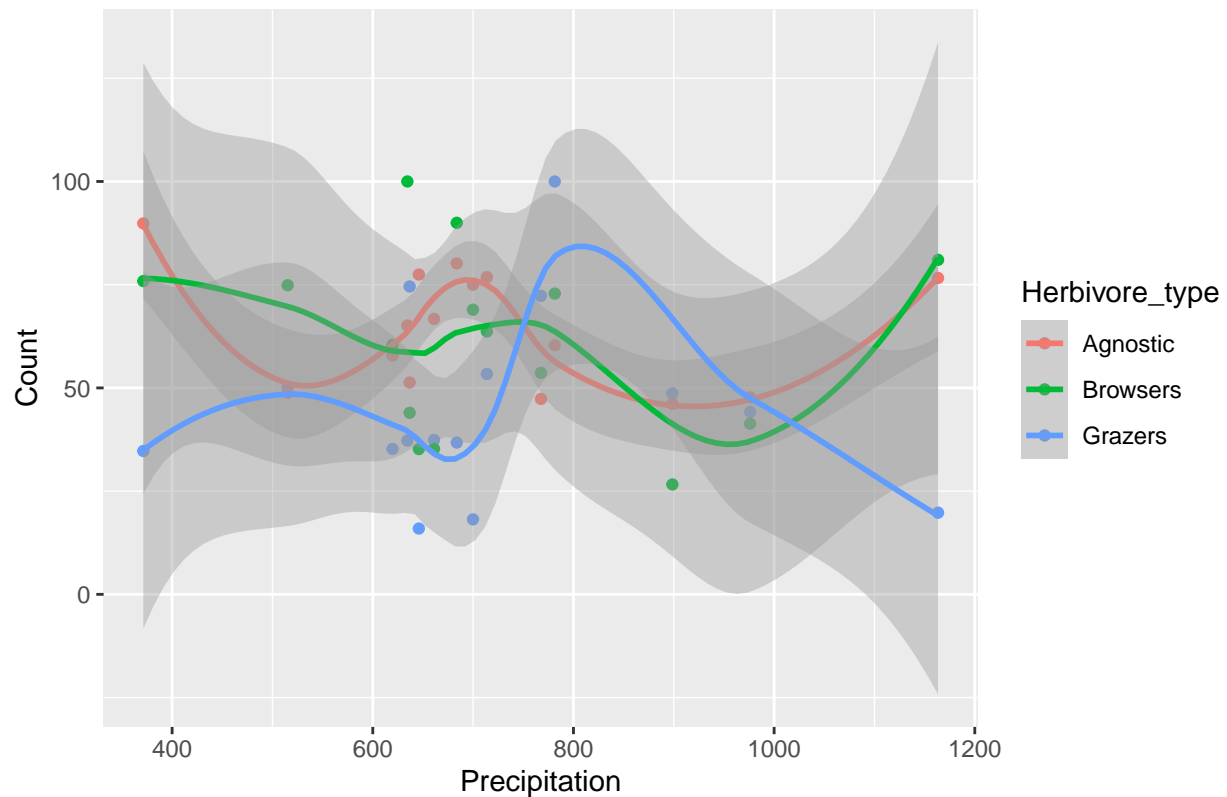
```
ggplot(data=filter(gc_precip_norm_type_plot, Area=="pools")) + geom_point(mapping=aes(x=Precipitation, y=Count, color=Herbivore_type))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 21 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 21 rows containing missing values (geom_point).
```

Normalized herbivore type counts vs pools precipitation



Correlation between soil moisture, drought index, vegetation, and water, and animal counts

I then graphed park and pools dry season climatic factors against LSU values for each species, and park and pools climatic factors against LSU values for each herbivore type.

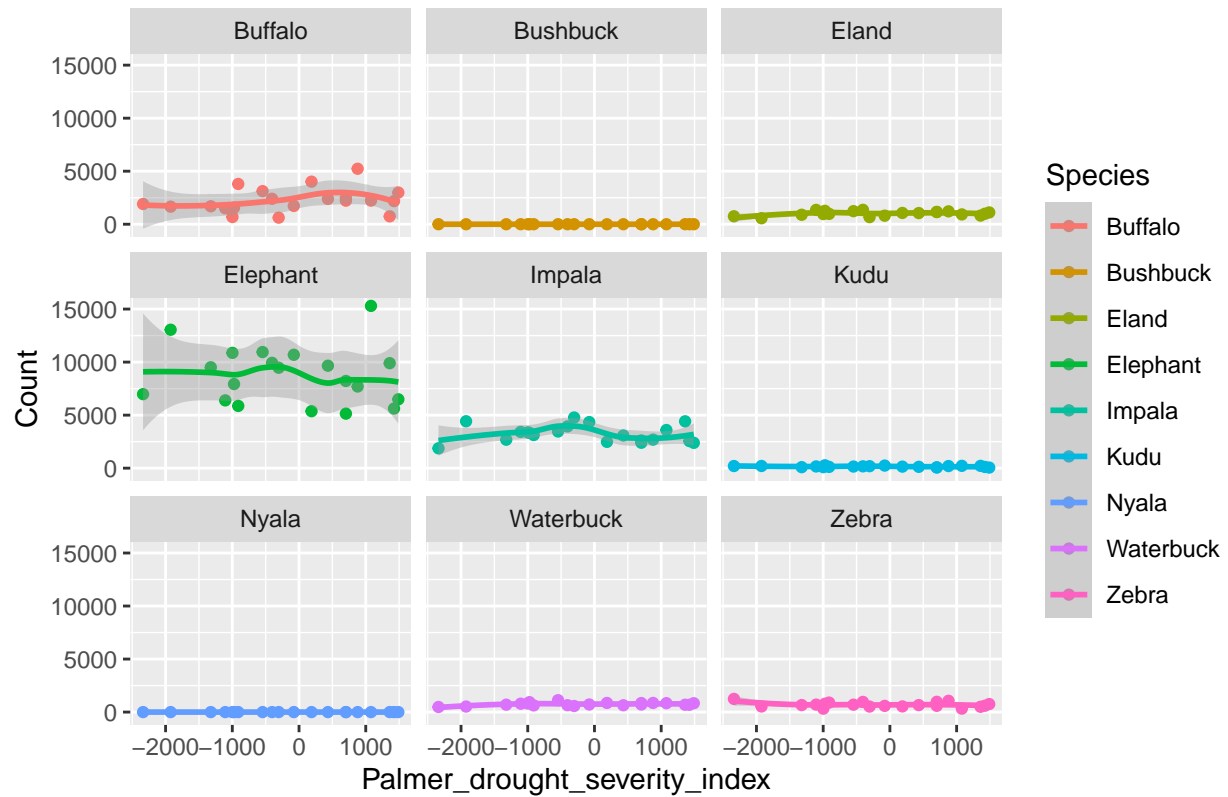
```
gc_clim_for_spp_plot <- gc_clim %>% gather(c(2,3,4,5,6,7,8,9,10), key='Species', value='Count')
ggplot(data=filter(gc_clim_for_spp_plot, Area=="park")) + geom_point(mapping=aes(x=Palmer_drought_sever

## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'

## Warning: Removed 18 rows containing non-finite values (stat_smooth).

## Warning: Removed 18 rows containing missing values (geom_point).
```

Species counts vs park drought index



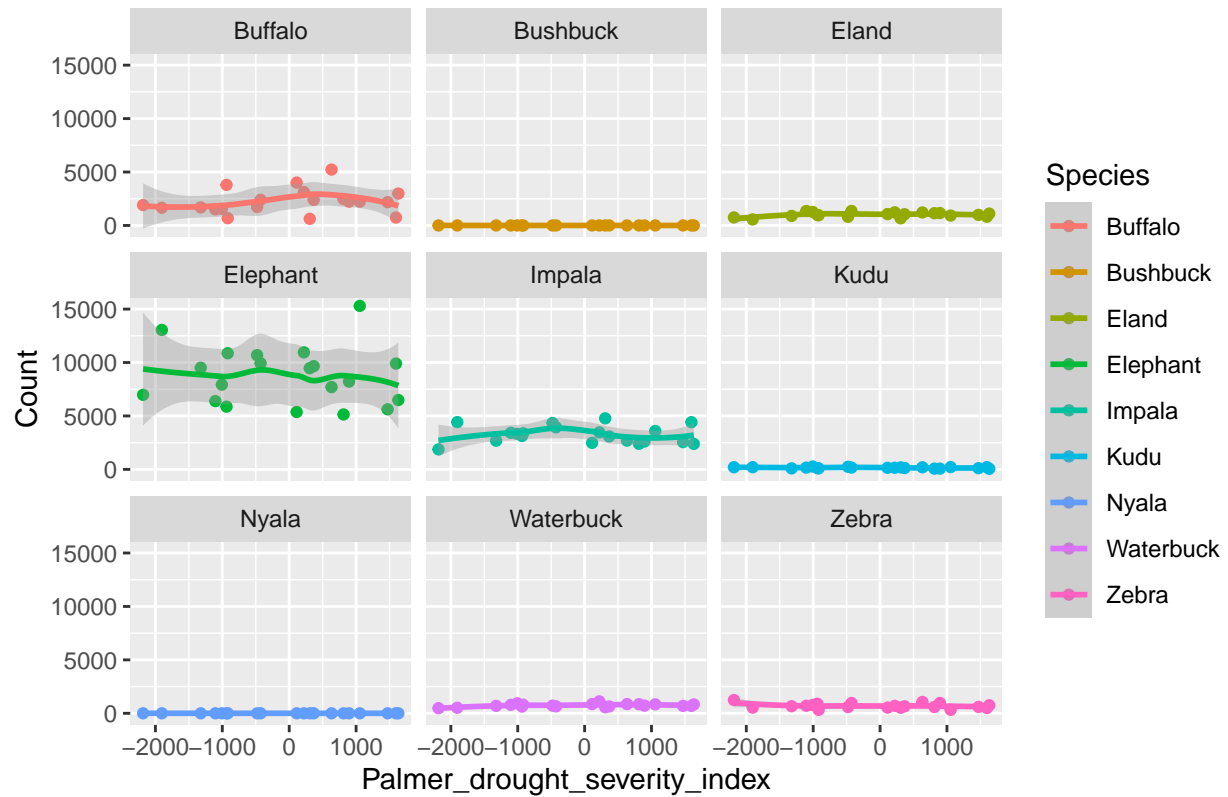
```
ggplot(data=filter(gc_clim_for_spp_plot, Area=="pools")) + geom_point(mapping=aes(x=Palmer_drought_seve
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 18 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 18 rows containing missing values (geom_point).
```

Species counts vs pools drought index



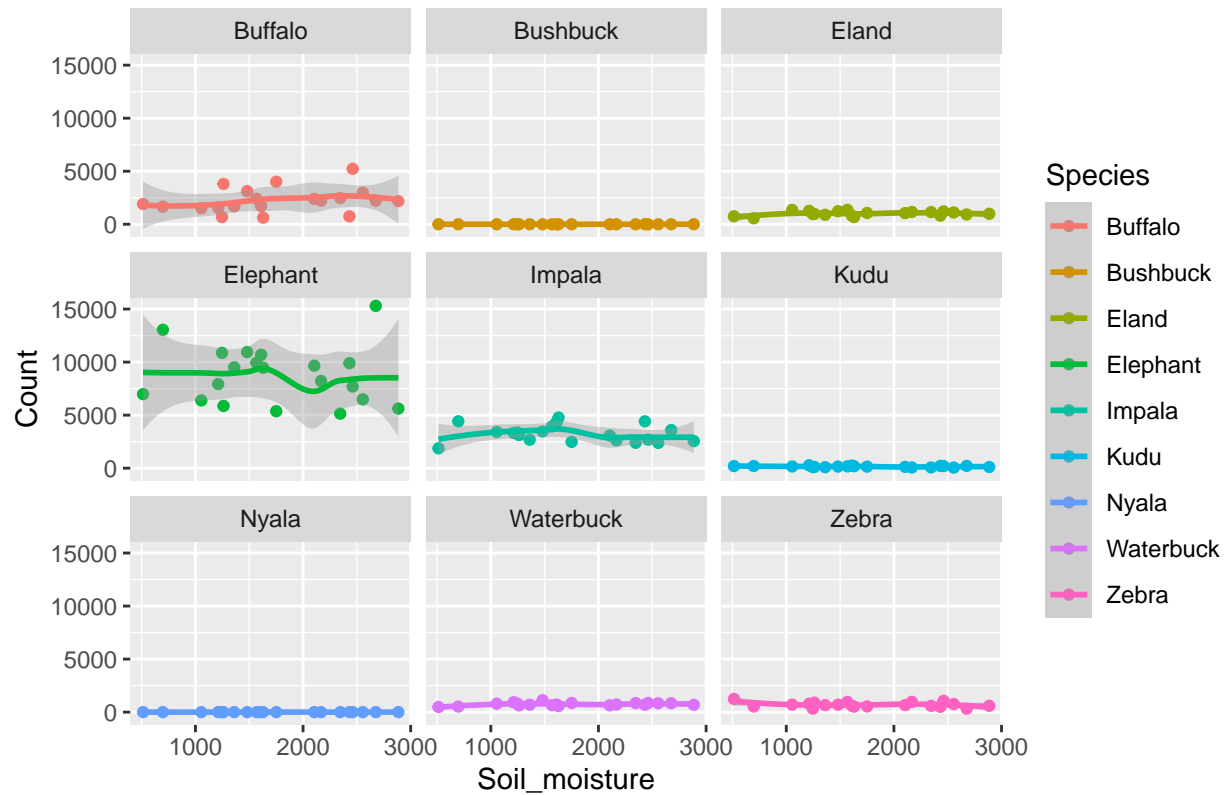
```
ggplot(data=filter(gc_clim_for_spp_plot, Area=="park")) + geom_point(mapping=aes(x=Soil_moisture, y=Count))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 18 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 18 rows containing missing values (geom_point).
```

Species counts vs park soil moisture



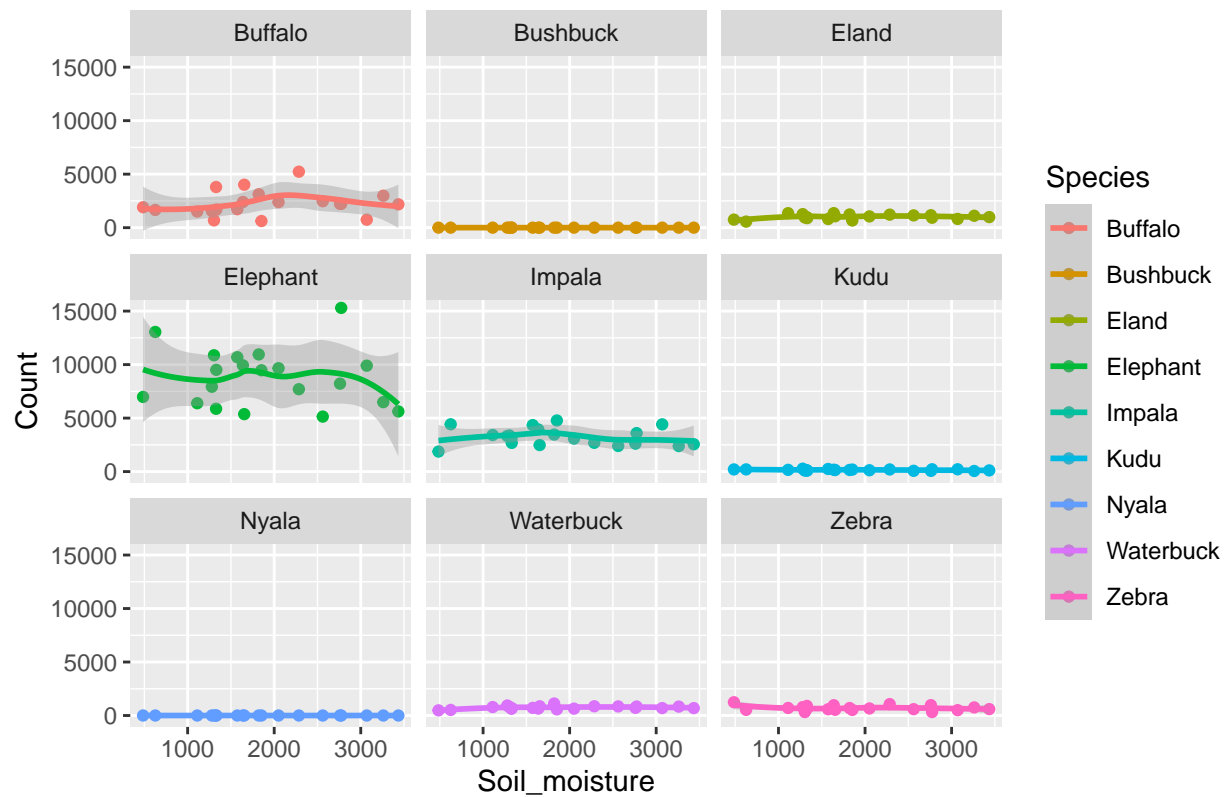
```
ggplot(data=filter(gc_clim_for_spp_plot, Area=="pools")) + geom_point(mapping=aes(x=Soil_moisture, y=Count, color=Species))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 18 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 18 rows containing missing values (geom_point).
```

Species counts vs pools soil moisture



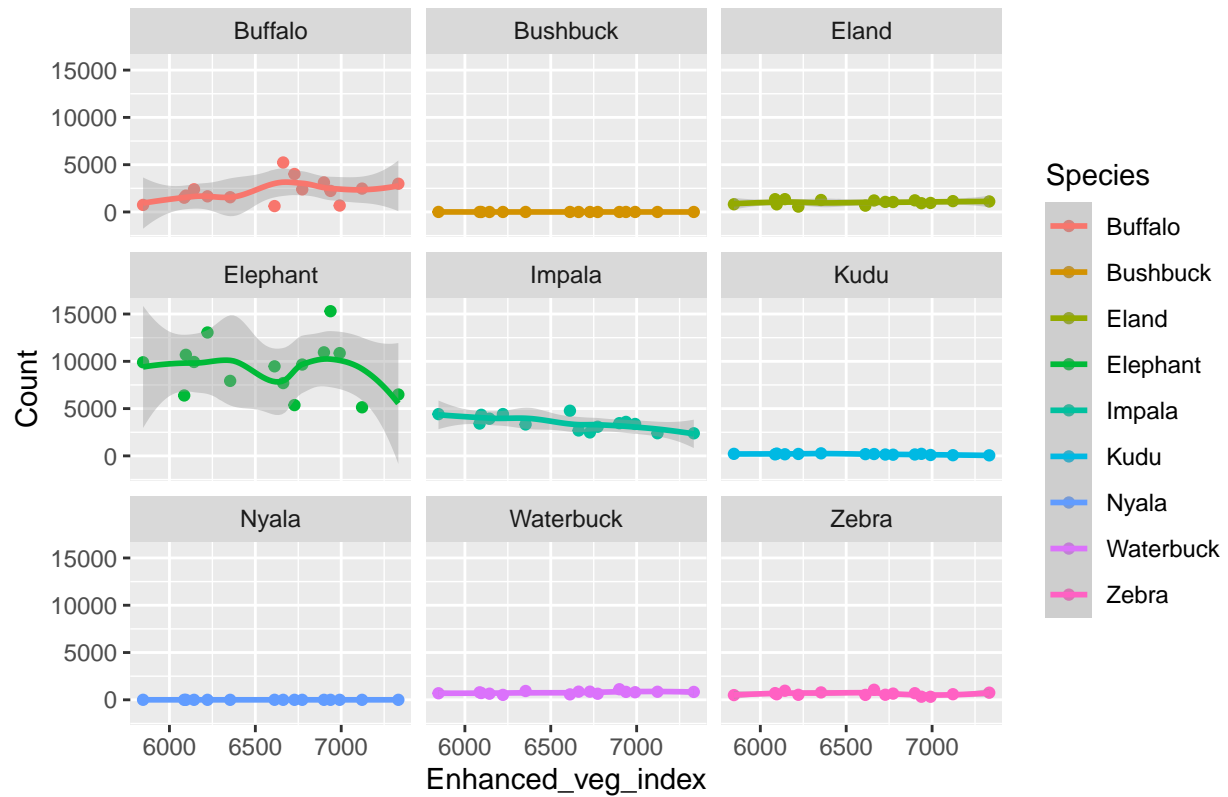
```
ggplot(data=filter(gc_clim_for_spp_plot, Area=="park" & Enhanced_veg_index > 0)) + geom_point(mapping=a
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 9 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 9 rows containing missing values (geom_point).
```

Species counts vs park vegetation index



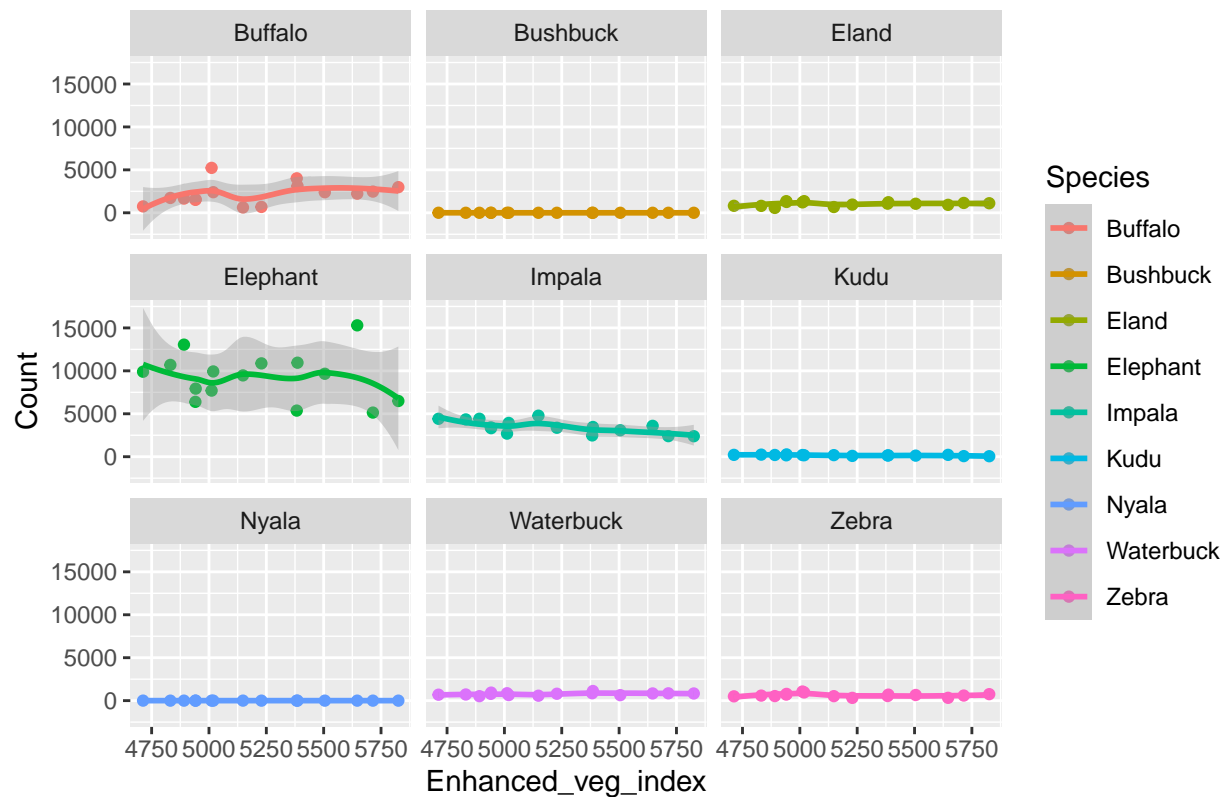
```
ggplot(data=filter(gc_clim_for_spp_plot, Area=="pools" & Enhanced_veg_index > 0)) + geom_point(mapping=
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 9 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 9 rows containing missing values (geom_point).
```

Species counts vs pools vegetation index



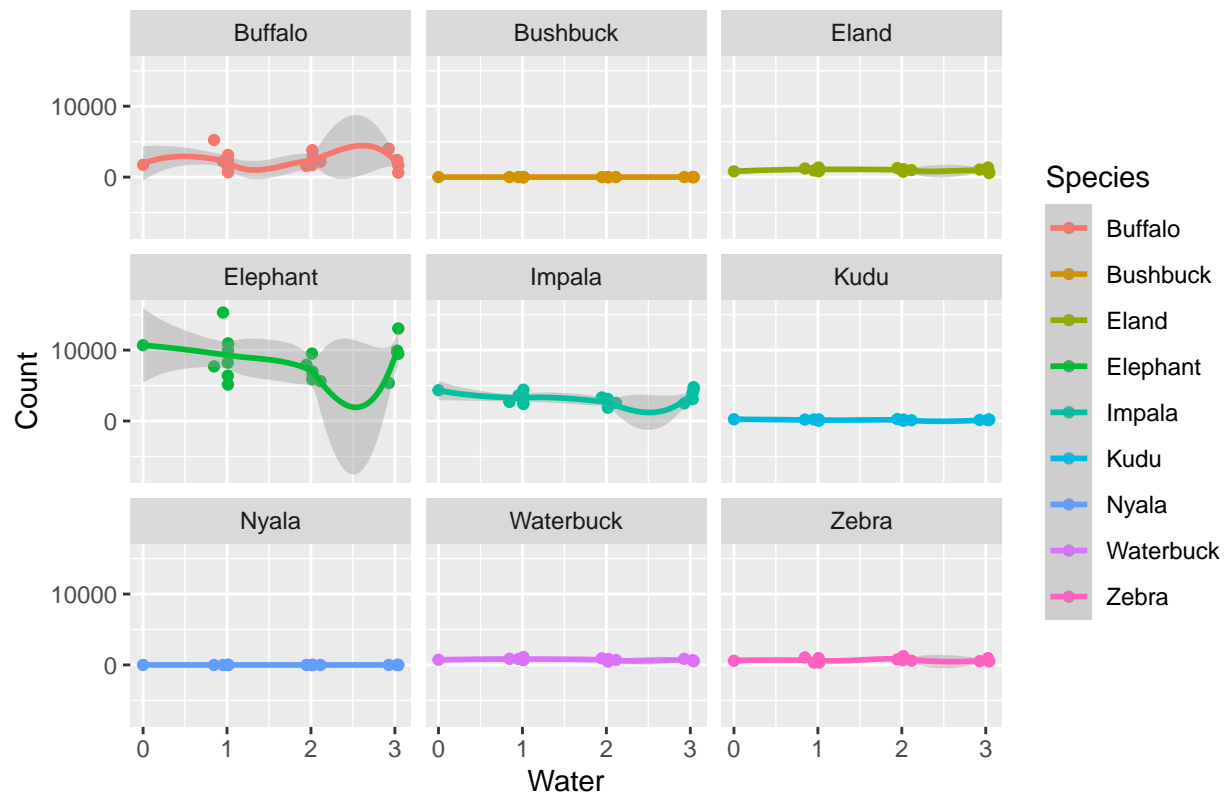
```
ggplot(data=filter(gc_clim_for_spp_plot, Area=="park")) + geom_point(mapping=aes(x=Water, y=Count, color=Species))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 18 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 18 rows containing missing values (geom_point).
```


Species counts vs park water



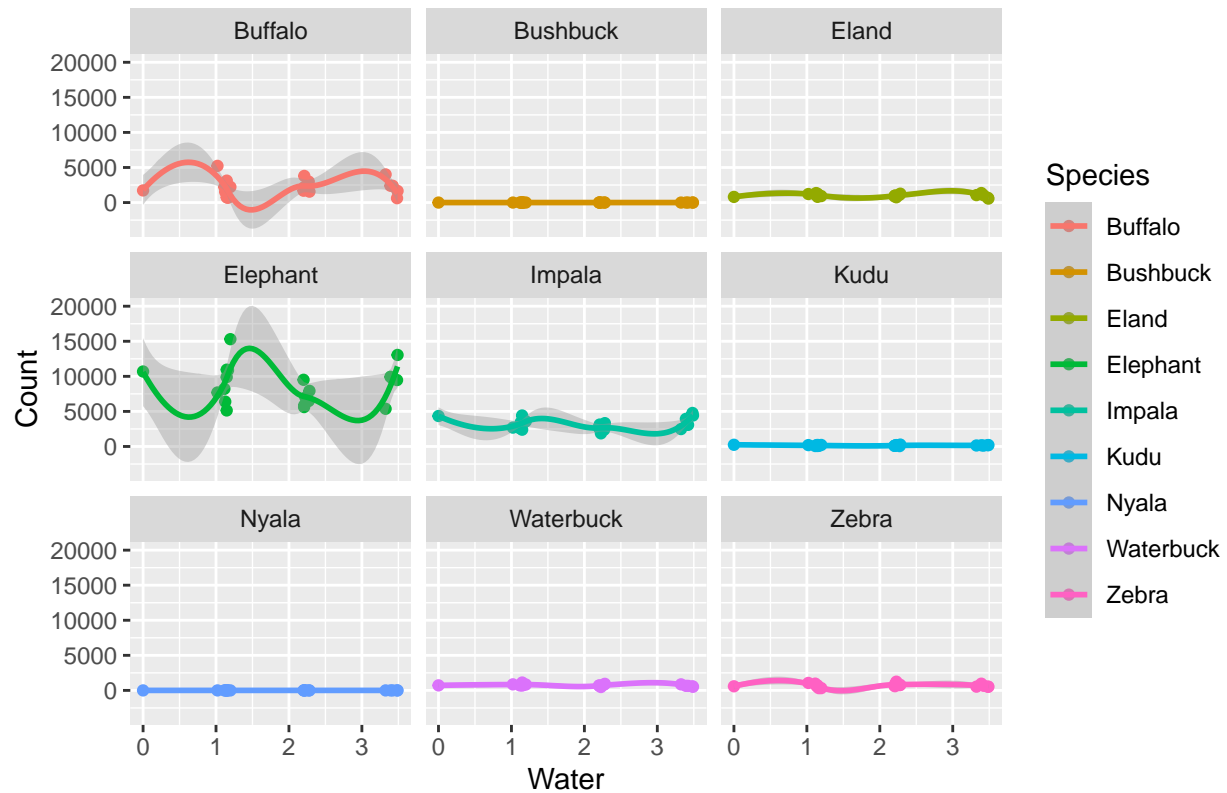
```
ggplot(data=filter(gc_clim_for_spp_plot, Area=="pools")) + geom_point(mapping=aes(x=Water, y=Count, col=Species))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 18 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 18 rows containing missing values (geom_point).
```

Species counts vs pools water



```
gc_clim_for_type_plot <- gc_clim %>% gather(c(11,12,13), key='Herbivore_type', value='Count')
ggplot(data=filter(gc_clim_for_type_plot, Area=="park")) + geom_point(mapping=aes(x=Palmer_drought_seve

## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'

## Warning: Removed 6 rows containing non-finite values (stat_smooth).

## Warning: Removed 6 rows containing missing values (geom_point).
```

Herbivore type counts vs park drought index

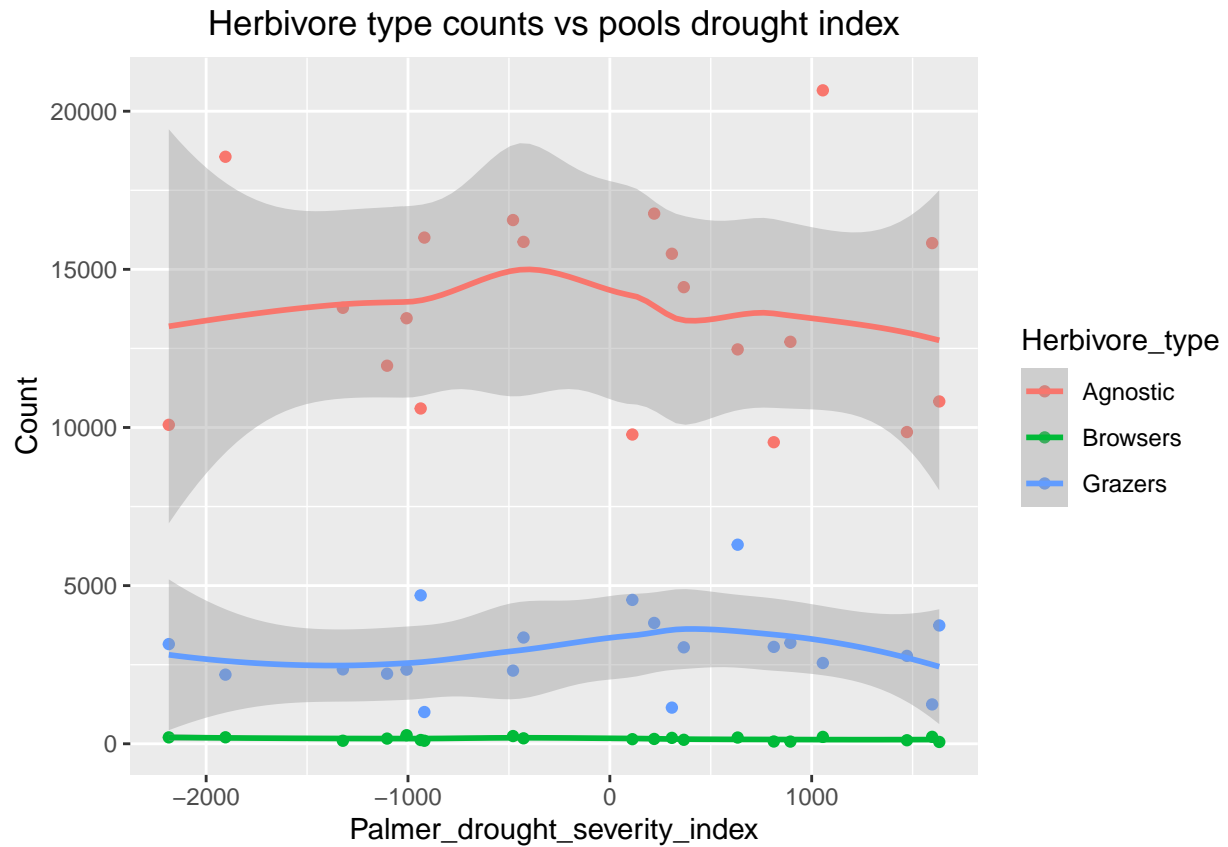


```
ggplot(data=filter(gc_clim_for_type_plot, Area=="pools")) + geom_point(mapping=aes(x=Palmer_drought_sev
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 6 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 6 rows containing missing values (geom_point).
```

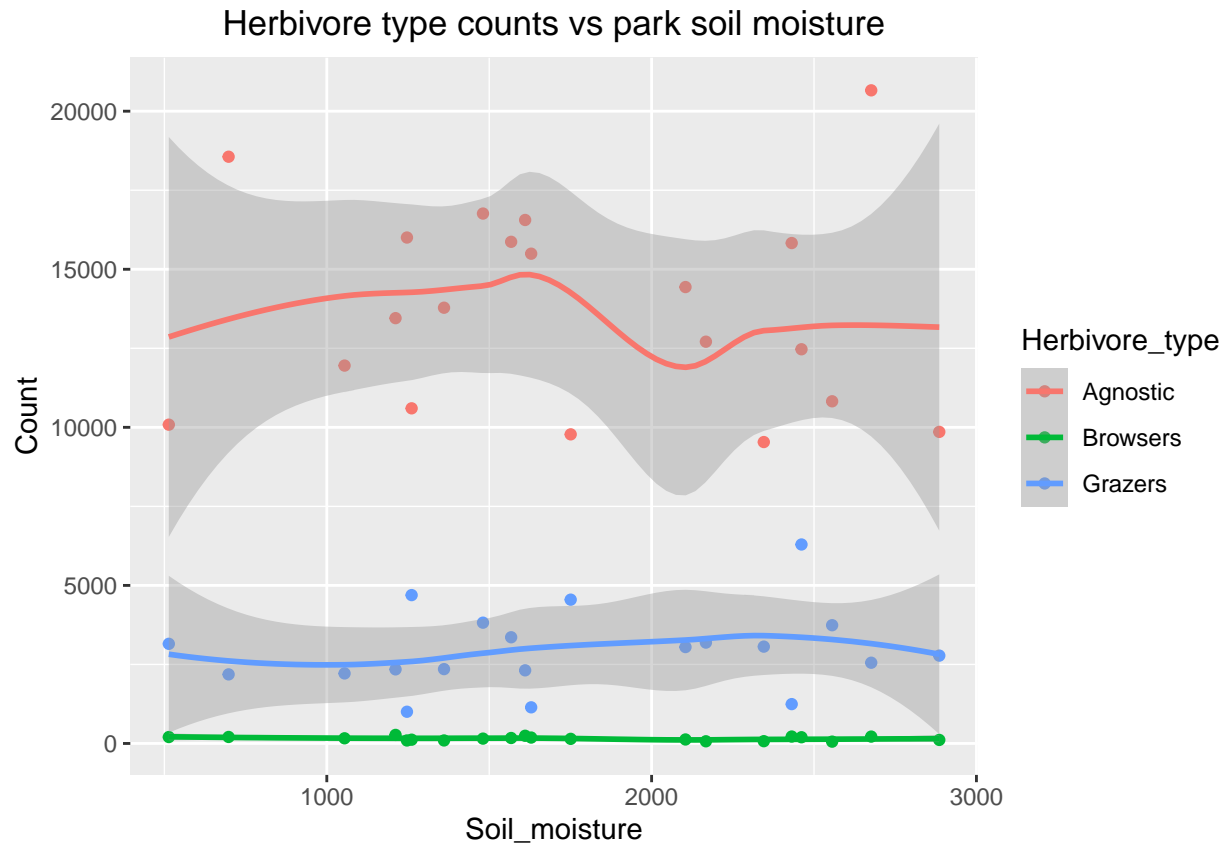


```
ggplot(data=filter(gc_clim_for_type_plot, Area=="park")) + geom_point(mapping=aes(x=Soil_moisture, y=Count))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 6 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 6 rows containing missing values (geom_point).
```

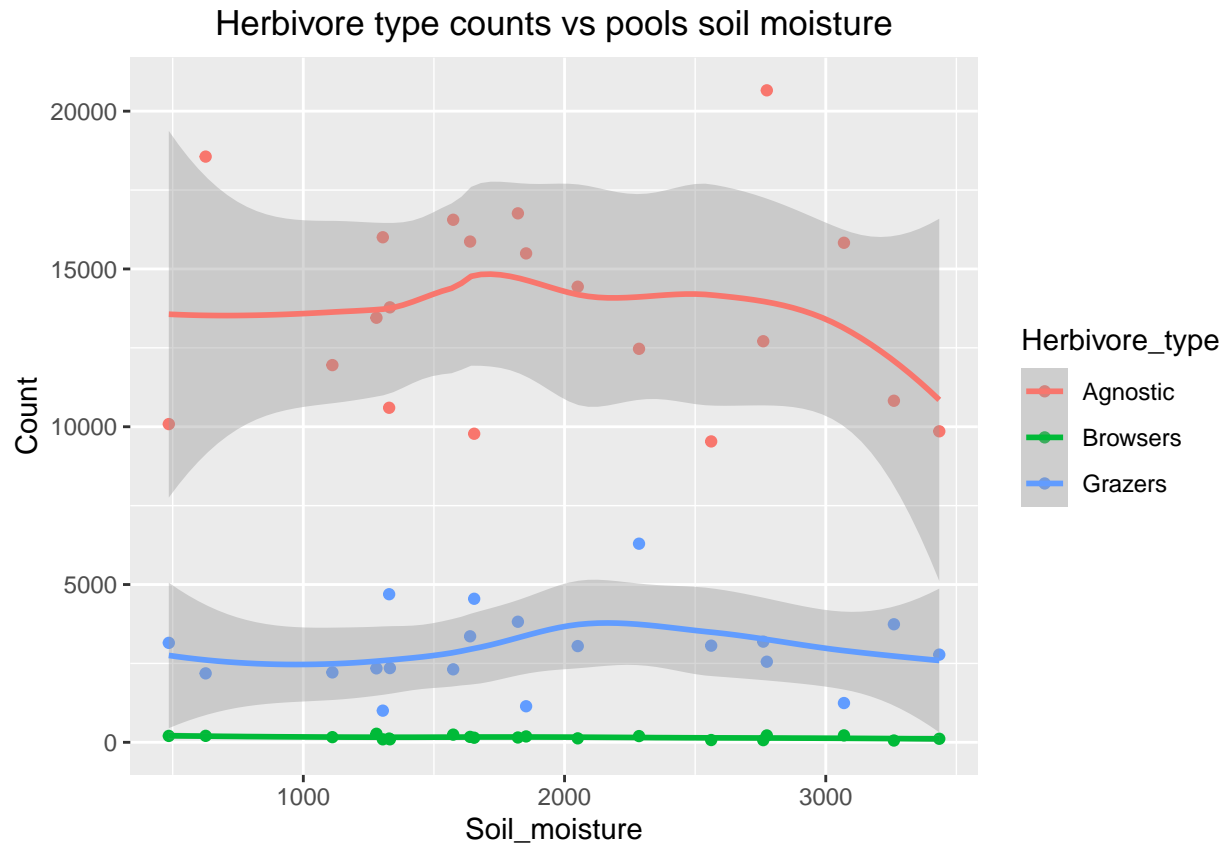


```
ggplot(data=filter(gc_clim_for_type_plot, Area=="pools")) + geom_point(mapping=aes(x=Soil_moisture, y=Count))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 6 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 6 rows containing missing values (geom_point).
```



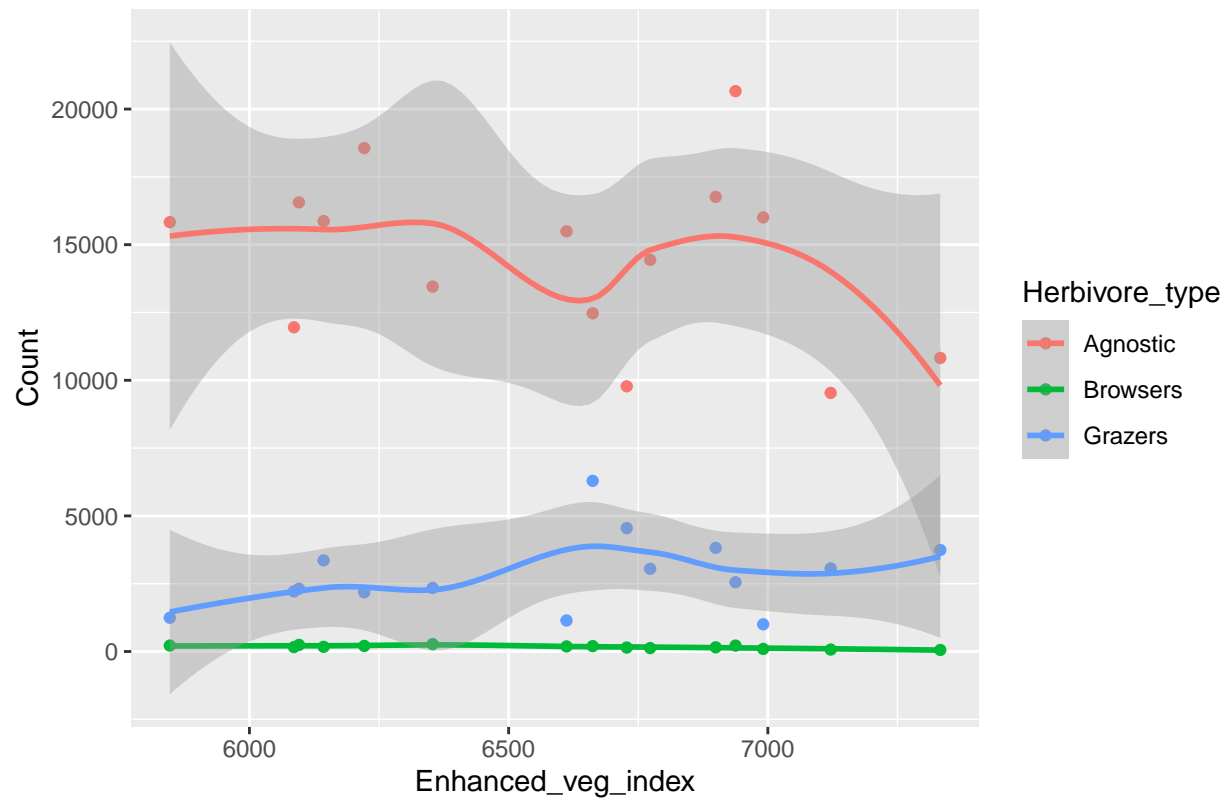
```
ggplot(data=filter(gc_clim_for_type_plot, Area=="park" & Enhanced_veg_index > 0)) + geom_point(mapping=
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 3 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 3 rows containing missing values (geom_point).
```

Herbivore type counts vs park vegetation index

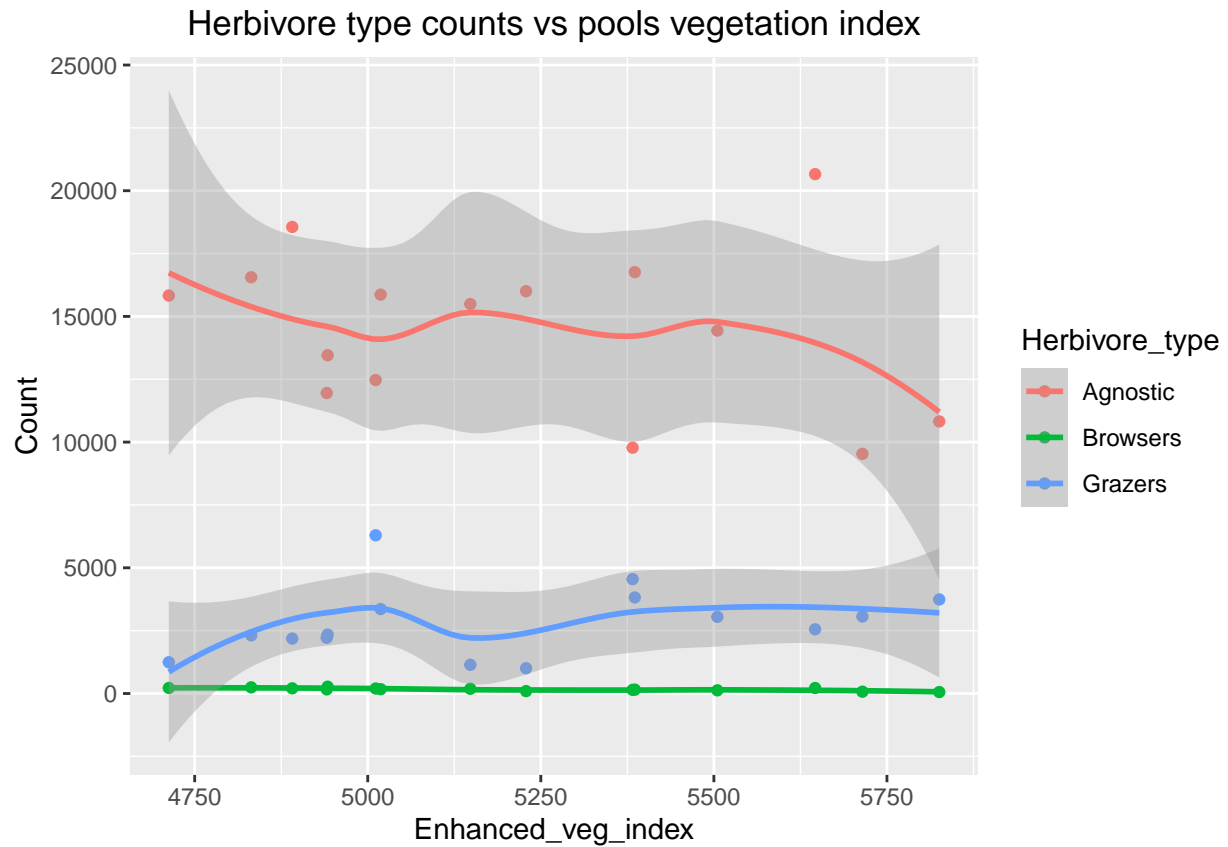


```
ggplot(data=filter(gc_clim_for_type_plot, Area=="pools" & Enhanced_veg_index > 0)) + geom_point(mapping=
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 3 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 3 rows containing missing values (geom_point).
```

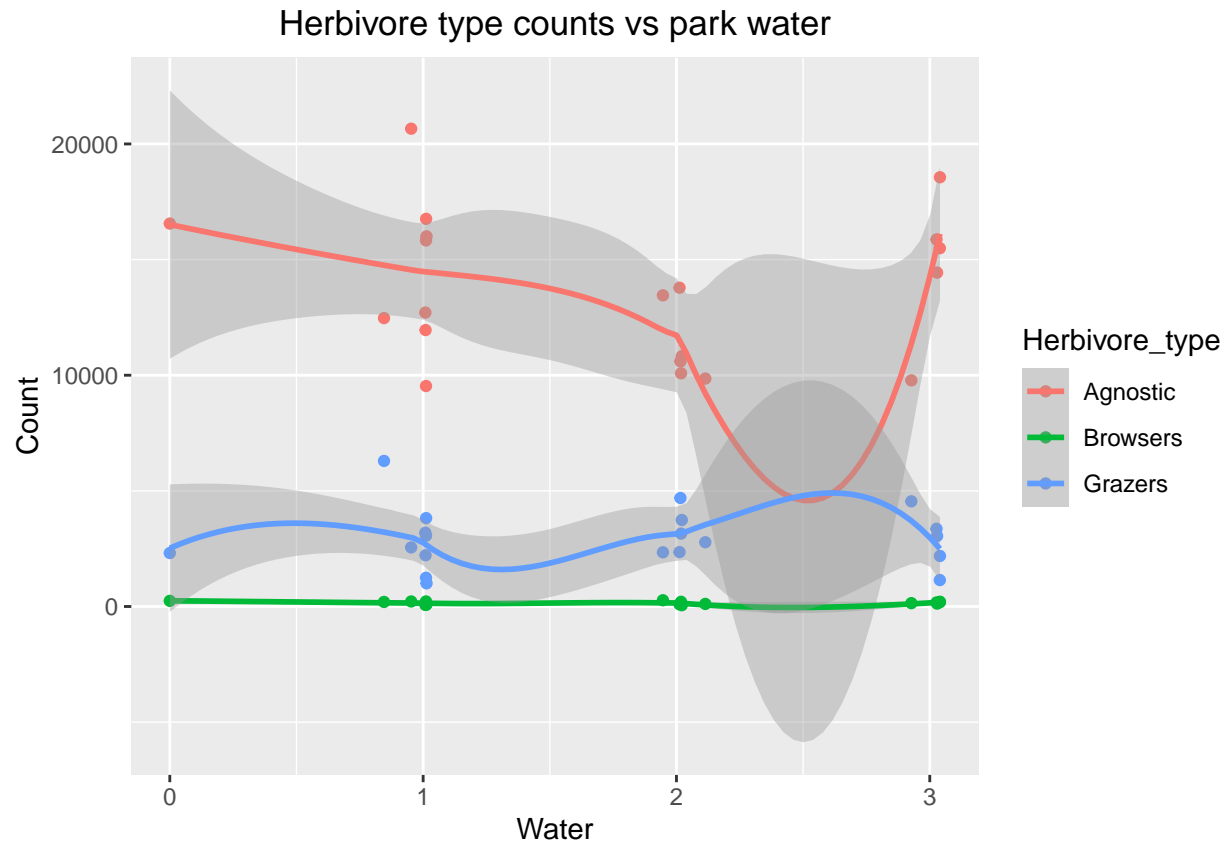


```
ggplot(data=filter(gc_clim_for_type_plot, Area=="park")) + geom_point(mapping=aes(x=Water, y=Count, col=Herbivore_type))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 6 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 6 rows containing missing values (geom_point).
```

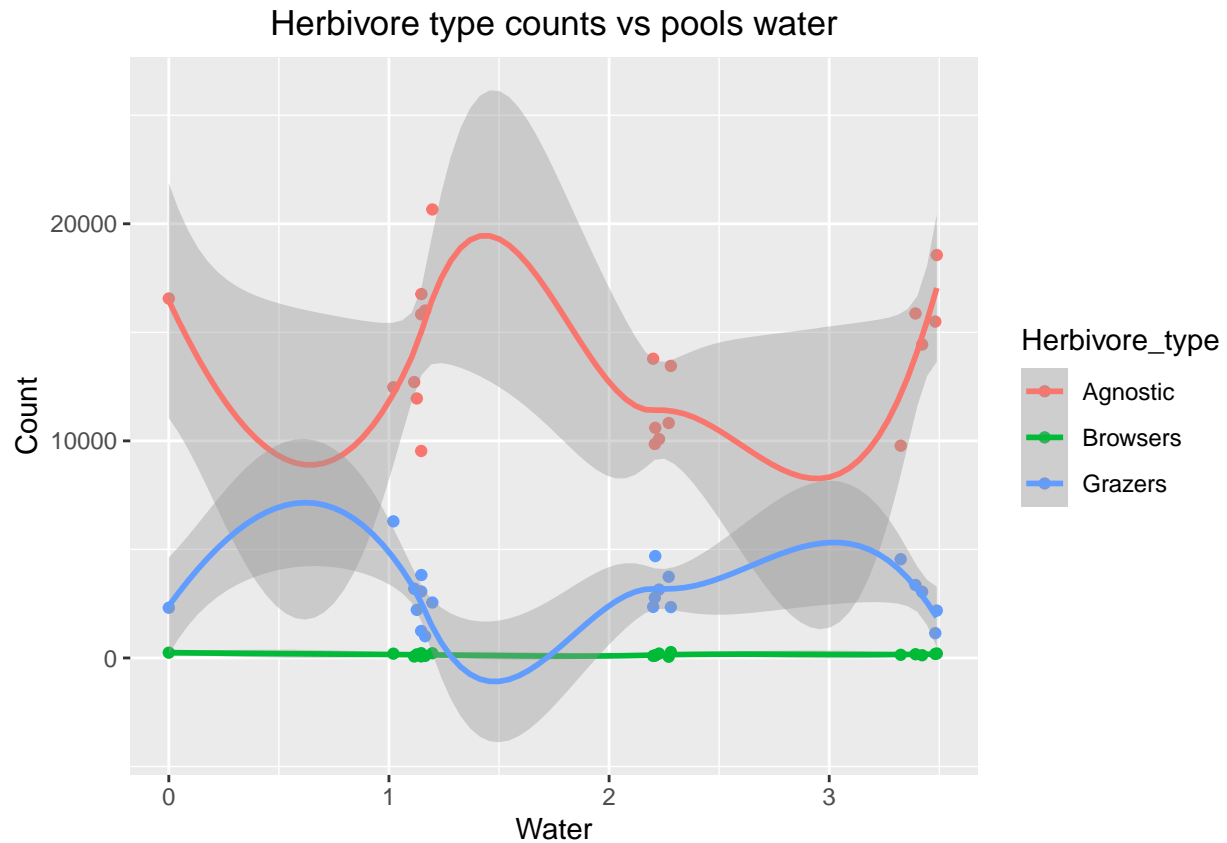



```
ggplot(data=filter(gc_clim_for_type_plot, Area=="pools")) + geom_point(mapping=aes(x=Water, y=Count, col=Herbivore_type))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 6 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 6 rows containing missing values (geom_point).
```



I then graphed park and pools dry season climatic factors against the normalized LSU values for each species, and park and pools climatic factors against the normalized LSU values for each herbivore type.

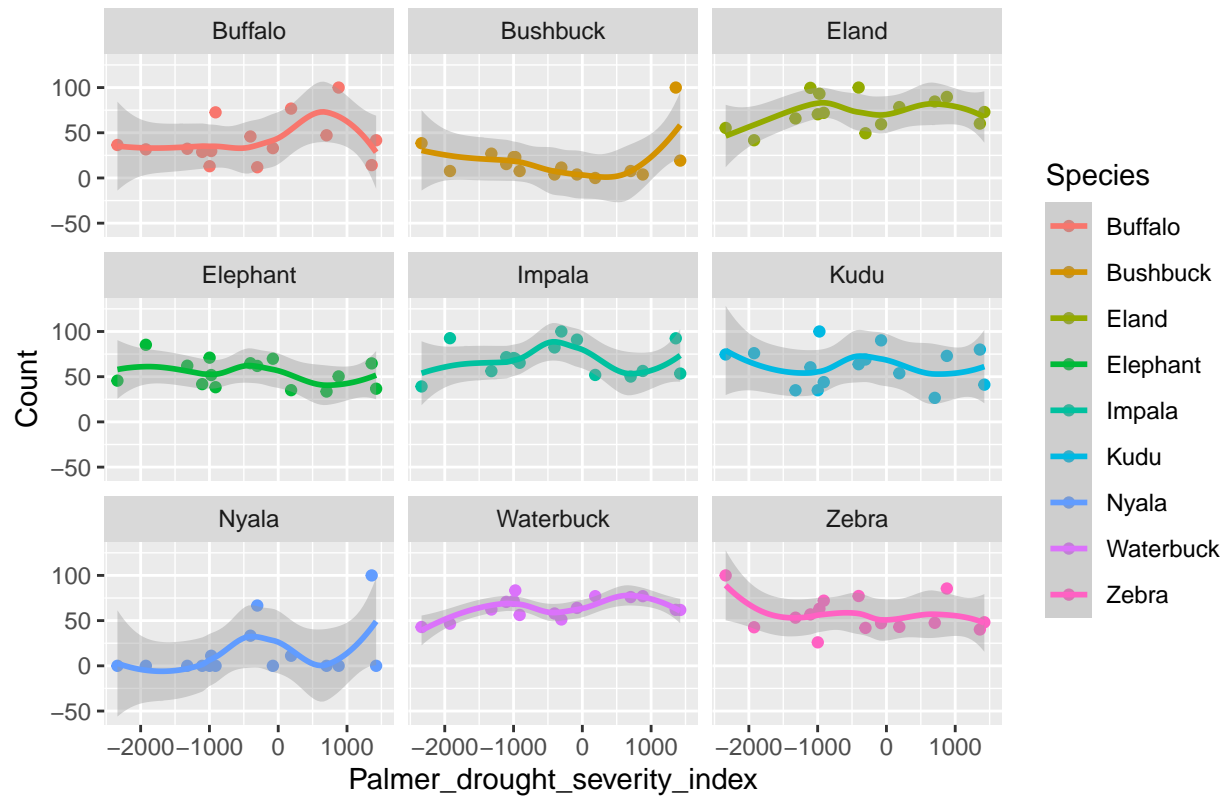
```
gc_clim_norm_spp_plot <- gc_norm_clim %>% gather(c(2,3,4,5,6,7,8,9,10), key='Species', value='Count')
ggplot(data=filter(gc_clim_norm_spp_plot, Area=="park")) + geom_point(mapping=aes(x=Palmer_drought_seve

## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'

## Warning: Removed 63 rows containing non-finite values (stat_smooth).

## Warning: Removed 63 rows containing missing values (geom_point).
```

Normalized species counts vs park drought index



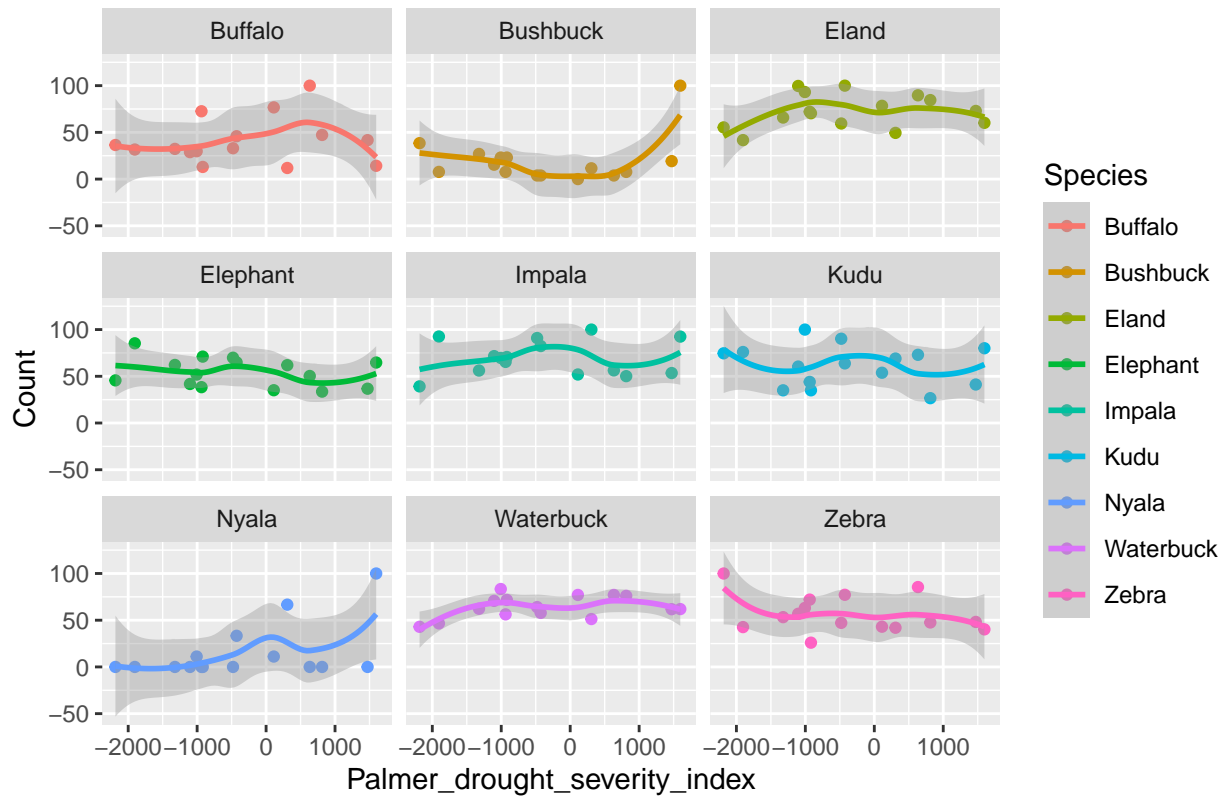
```
ggplot(data=filter(gc_clim_norm_spp_plot, Area=="pools")) + geom_point(mapping=aes(x=Palmer_drought_sev
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 63 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 63 rows containing missing values (geom_point).
```

Normalized species counts vs pools drought index



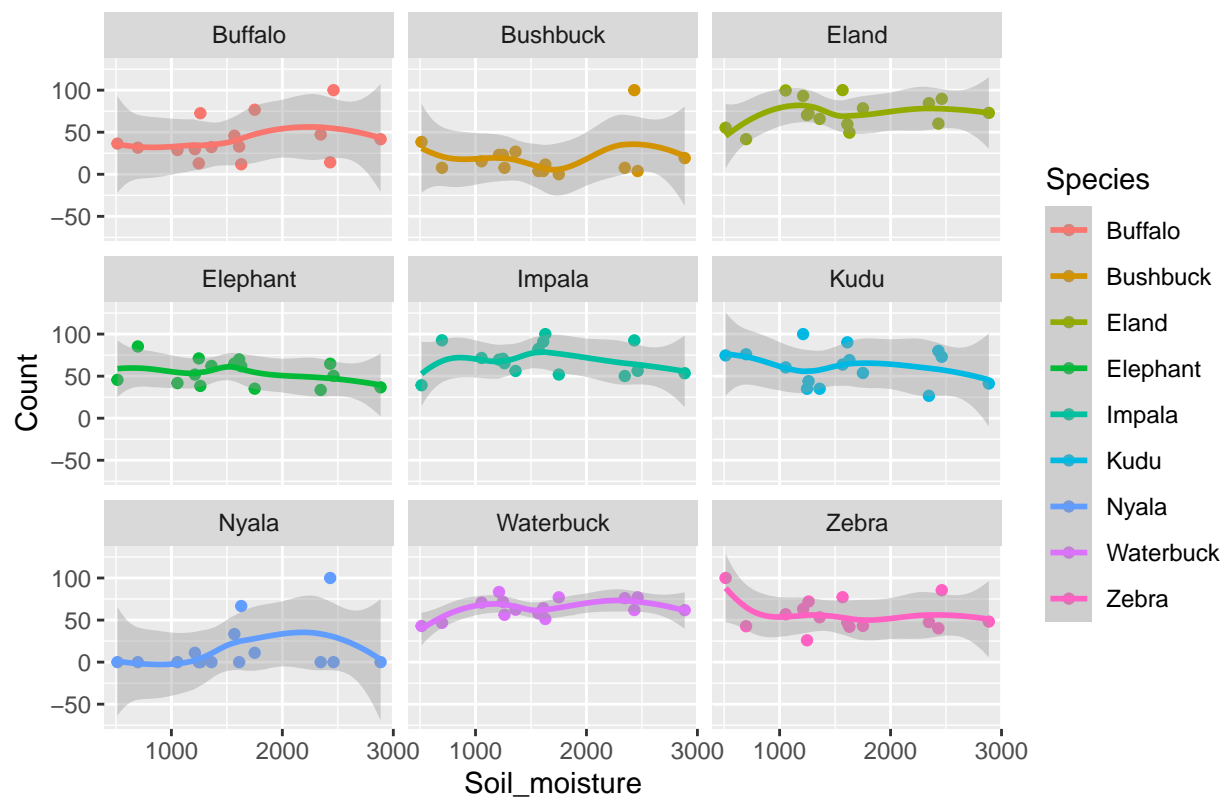
```
ggplot(data=filter(gc_clim_norm_spp_plot, Area=="park")) + geom_point(mapping=aes(x=Soil_moisture, y=Count))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 63 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 63 rows containing missing values (geom_point).
```

Normalized species counts vs park soil moisture



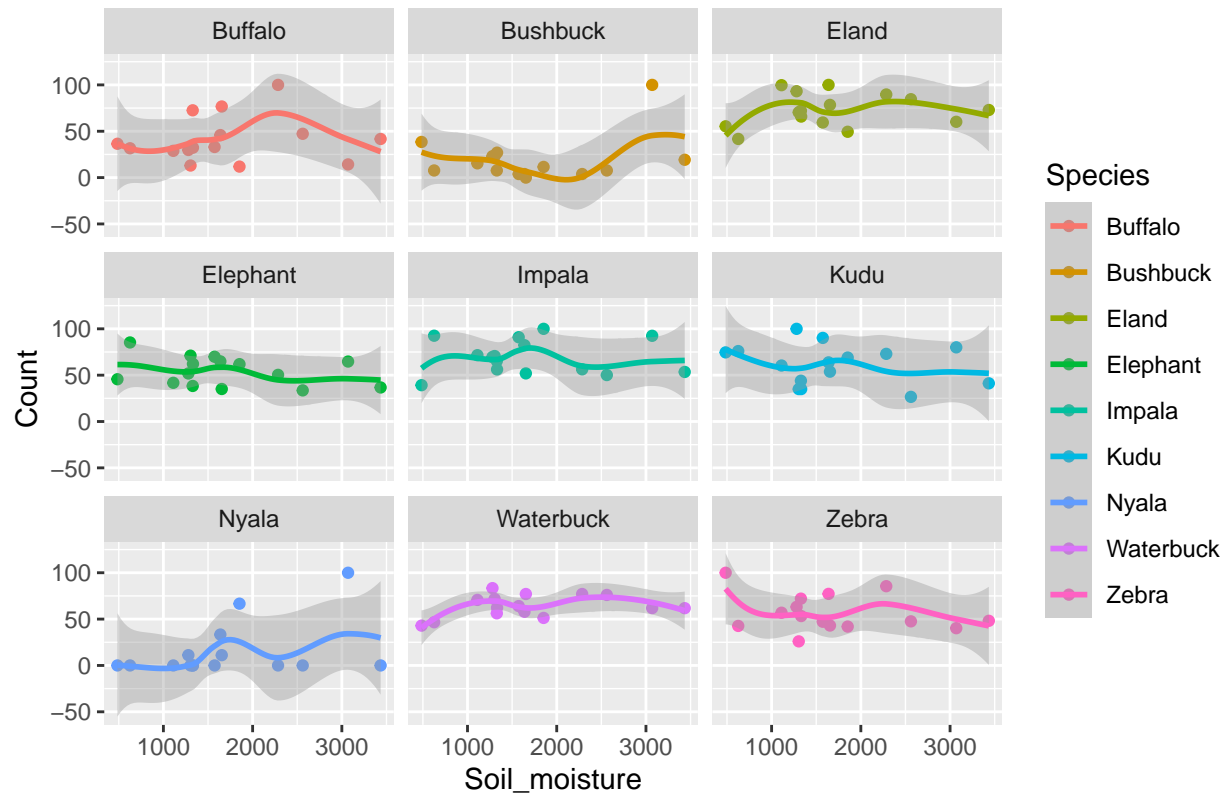
```
ggplot(data=filter(gc_clim_norm_spp_plot, Area=="pools")) + geom_point(mapping=aes(x=Soil_moisture, y=Count))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 63 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 63 rows containing missing values (geom_point).
```

Normalized species counts vs pools soil moisture



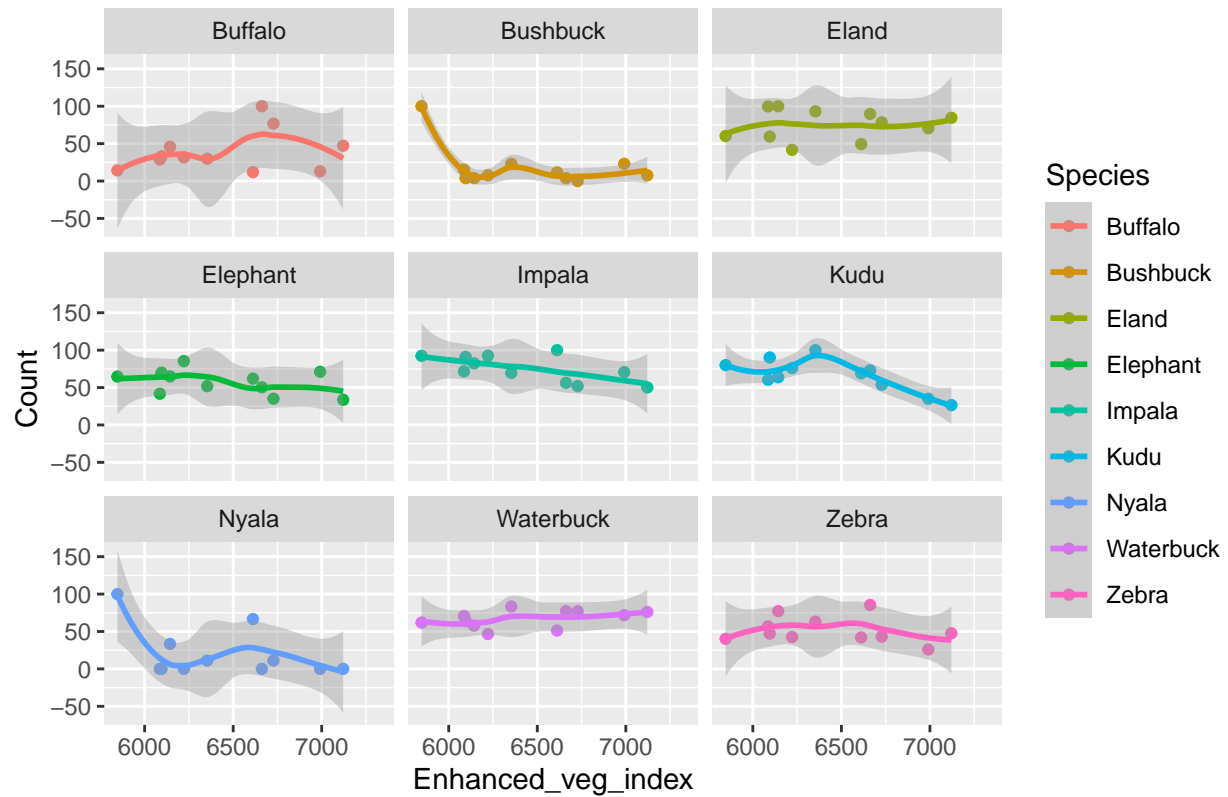
```
ggplot(data=filter(gc_clim_norm_spp_plot, Area=="park" & Enhanced_veg_index > 0)) + geom_point(mapping=
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 45 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 45 rows containing missing values (geom_point).
```

Normalized species counts vs park vegetation index



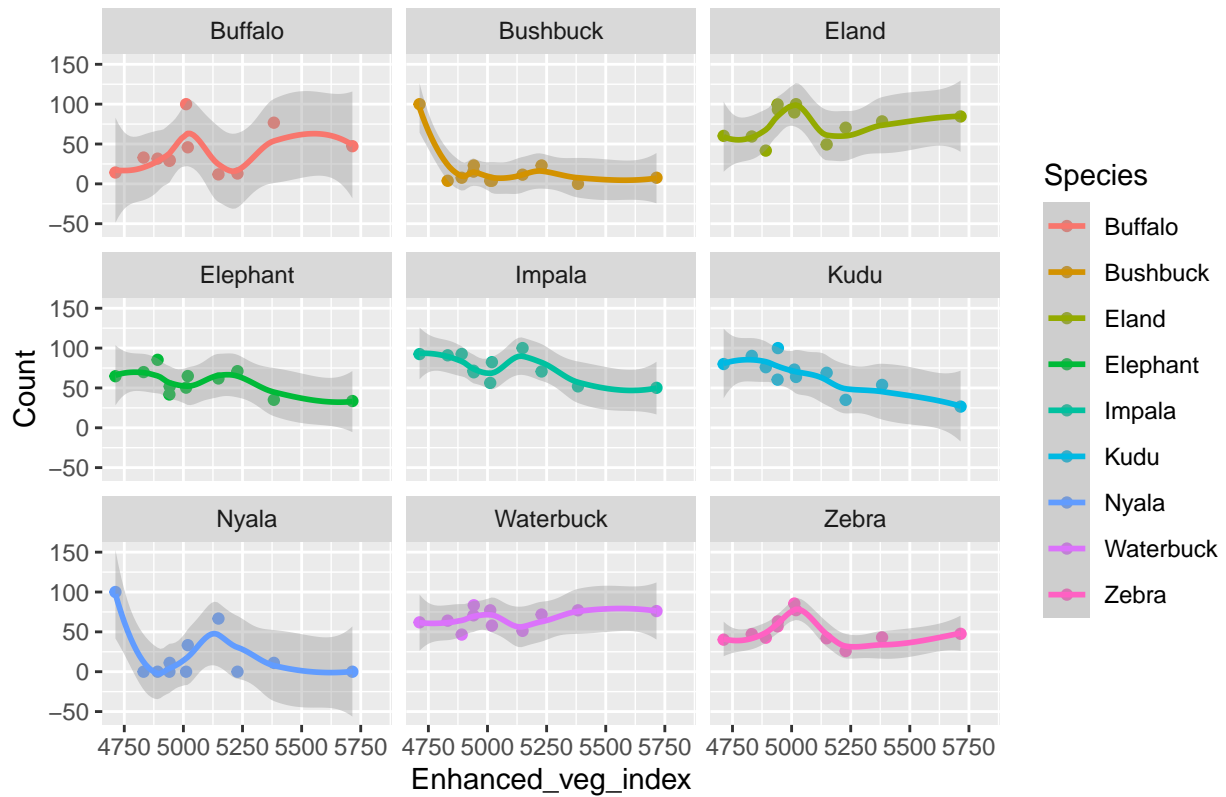
```
ggplot(data=filter(gc_clim_norm_spp_plot, Area=="pools" & Enhanced_veg_index > 0)) + geom_point(mapping=
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 45 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 45 rows containing missing values (geom_point).
```

Normalized species counts vs pools vegetation index



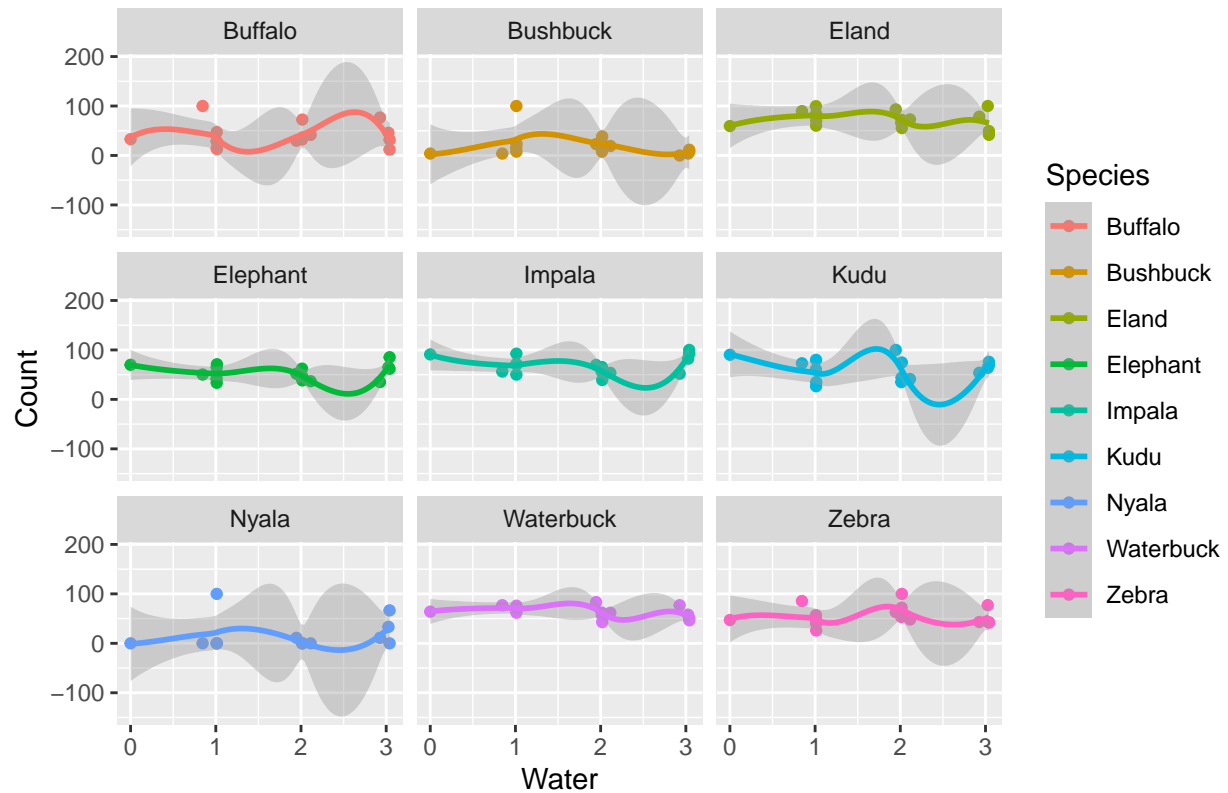
```
ggplot(data=filter(gc_clim_norm_spp_plot, Area=="park")) + geom_point(mapping=aes(x=Water, y=Count, col=Species))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 63 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 63 rows containing missing values (geom_point).
```


Normalized species counts vs park water



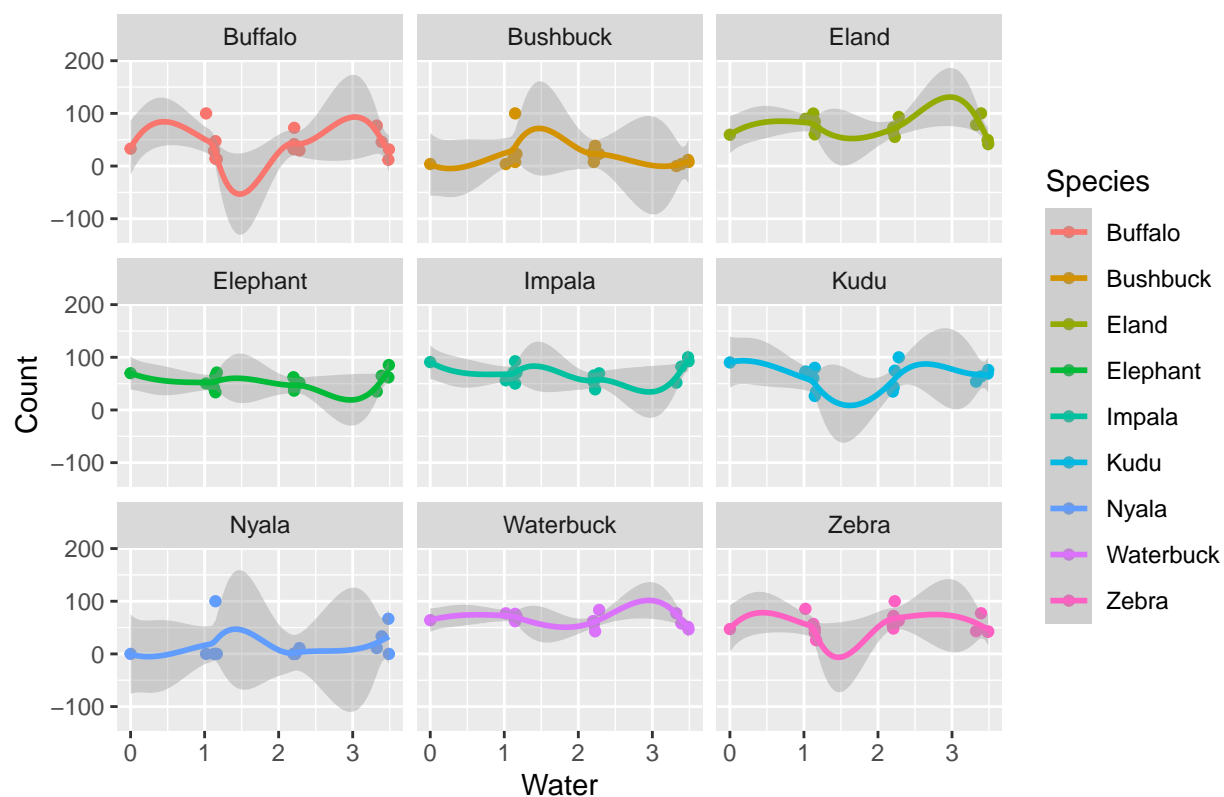
```
ggplot(data=filter(gc_clim_norm_spp_plot, Area=="pools")) + geom_point(mapping=aes(x=Water, y=Count, col=Species))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 63 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 63 rows containing missing values (geom_point).
```

Normalized species counts vs pools water



```
gc_clim_norm_type_plot <- gc_norm_clim %>% gather(c(11,12,13), key='Herbivore_type', value='Count')
ggplot(data=filter(gc_clim_norm_type_plot, Area=="park")) + geom_point(mapping=aes(x=Palmer_drought_sev

## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'

## Warning: Removed 21 rows containing non-finite values (stat_smooth).

## Warning: Removed 21 rows containing missing values (geom_point).
```

Normalized herbivore type counts vs park drought index



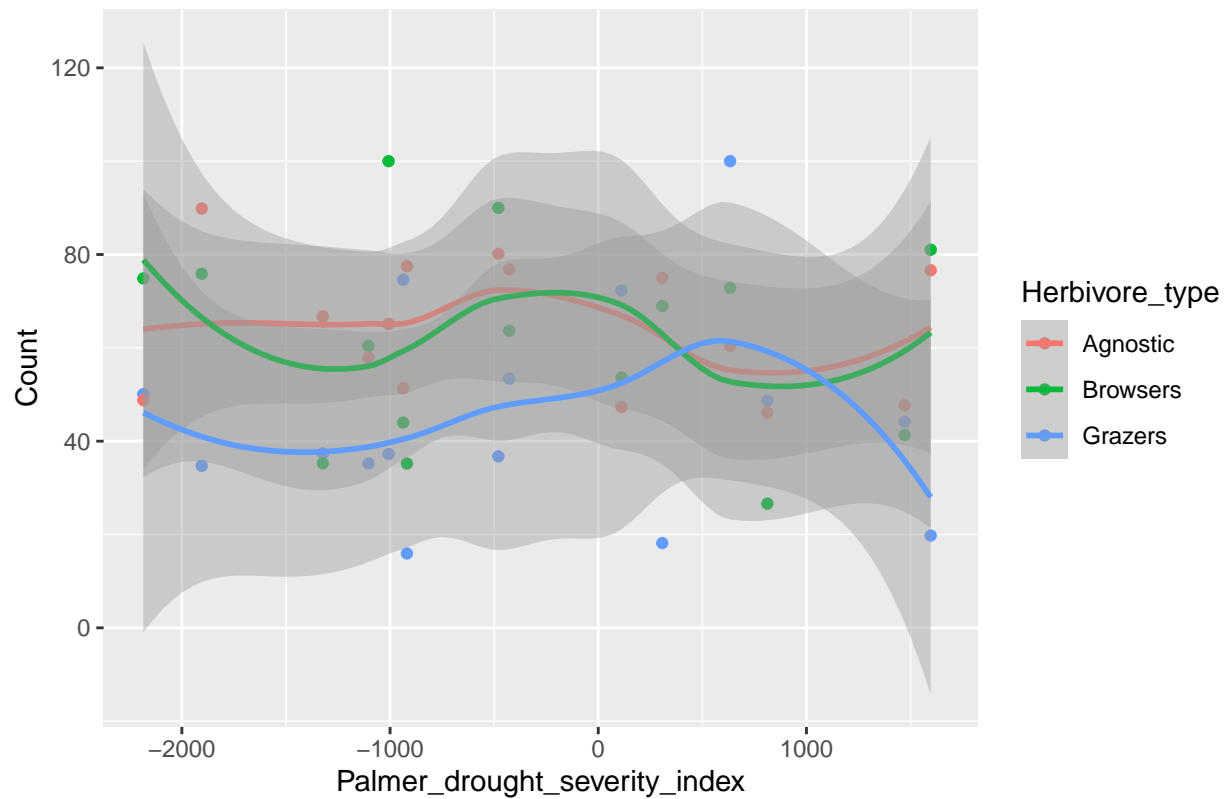
```
ggplot(data=filter(gc_clim_norm_type_plot, Area=="pools")) + geom_point(mapping=aes(x=Palmer_drought_severity_index, y=Count, color=Herbivore_type))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 21 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 21 rows containing missing values (geom_point).
```

Normalized herbivore type counts vs pools drought index



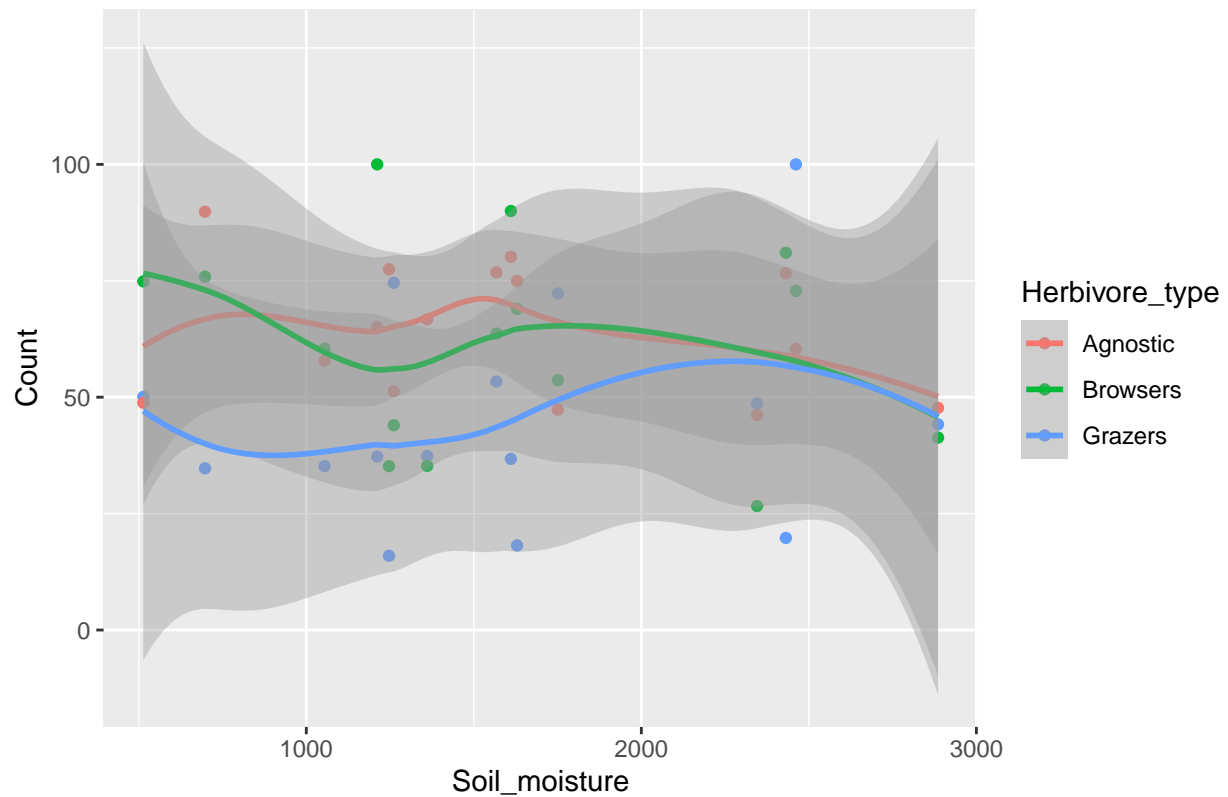
```
ggplot(data=filter(gc_clim_norm_type_plot, Area=="park")) + geom_point(mapping=aes(x=Soil_moisture, y=C
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 21 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 21 rows containing missing values (geom_point).
```

Normalized herbivore type counts vs park soil moisture



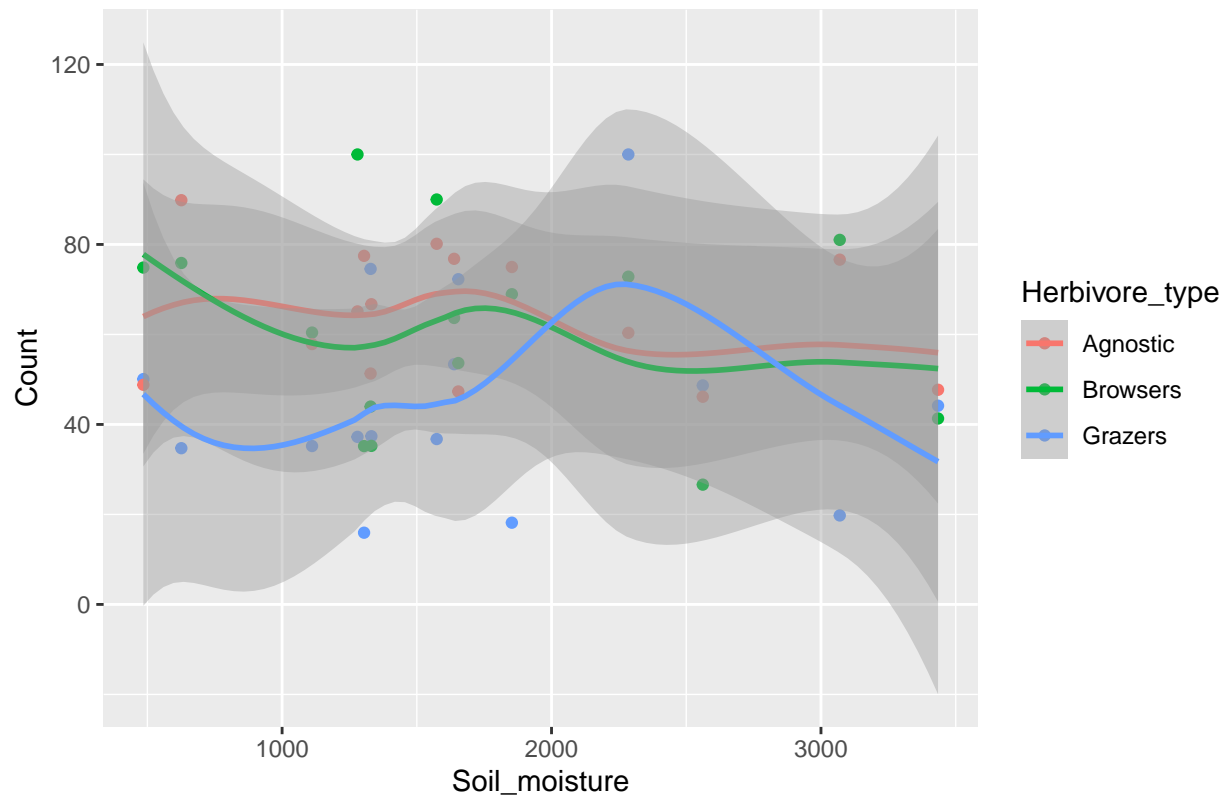
```
ggplot(data=filter(gc_clim_norm_type_plot, Area=="pools")) + geom_point(mapping=aes(x=Soil_moisture, y=
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 21 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 21 rows containing missing values (geom_point).
```

Normalized herbivore type counts vs pools soil moisture



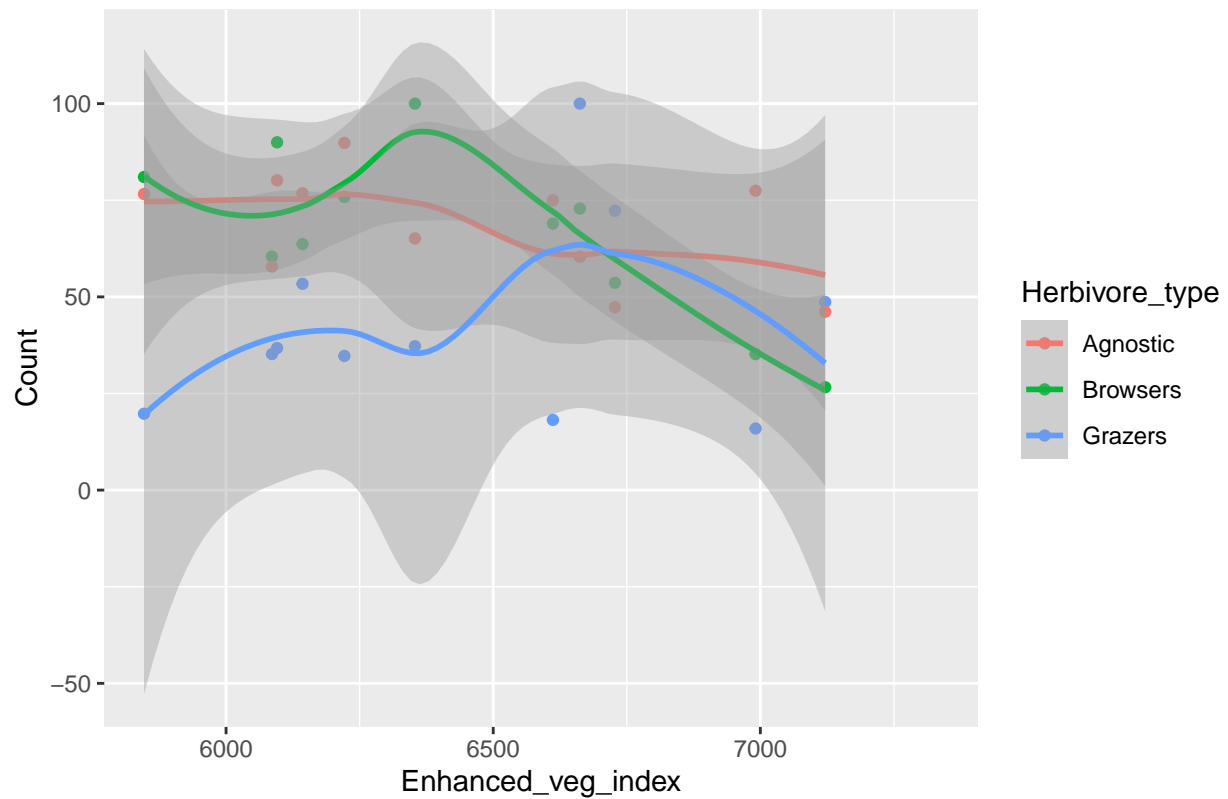
```
ggplot(data=filter(gc_clim_norm_type_plot, Area=="park" & Enhanced_veg_index > 0)) + geom_point(mapping=
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 15 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 15 rows containing missing values (geom_point).
```

Normalized herbivore type counts vs park vegetation index



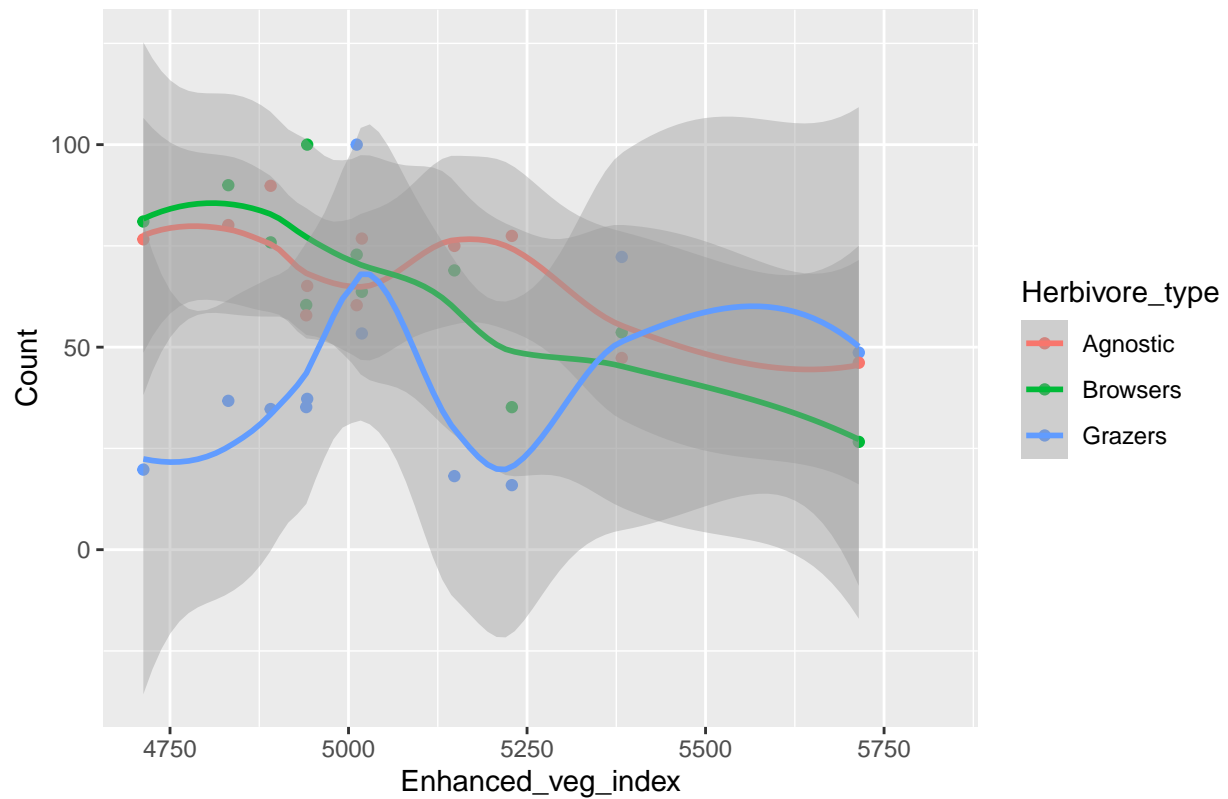
```
ggplot(data=filter(gc_clim_norm_type_plot, Area=="pools" & Enhanced_veg_index > 0)) + geom_point(mapping=
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 15 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 15 rows containing missing values (geom_point).
```

Normalized herbivore type counts vs pools vegetation index



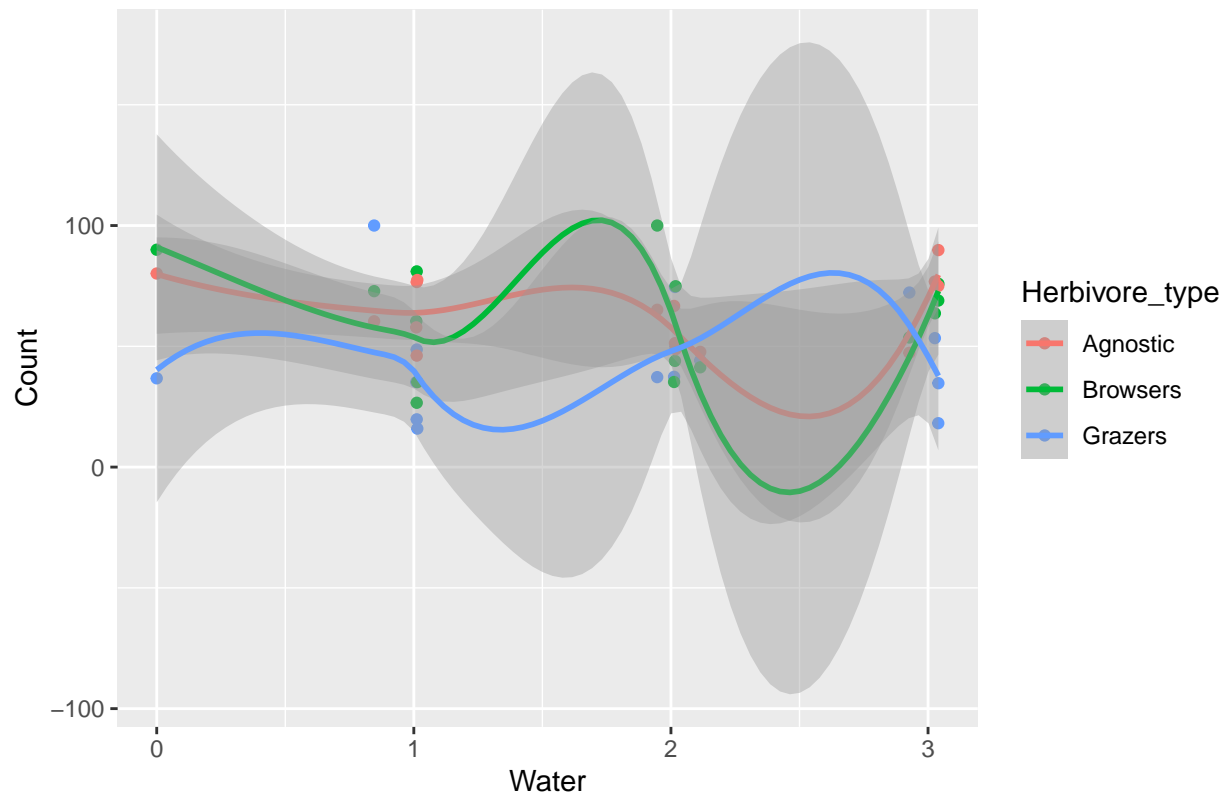
```
ggplot(data=filter(gc_clim_norm_type_plot, Area=="park")) + geom_point(mapping=aes(x=Water, y=Count, col=Herbivore_type))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 21 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 21 rows containing missing values (geom_point).
```


Normalized herbivore type counts vs park water

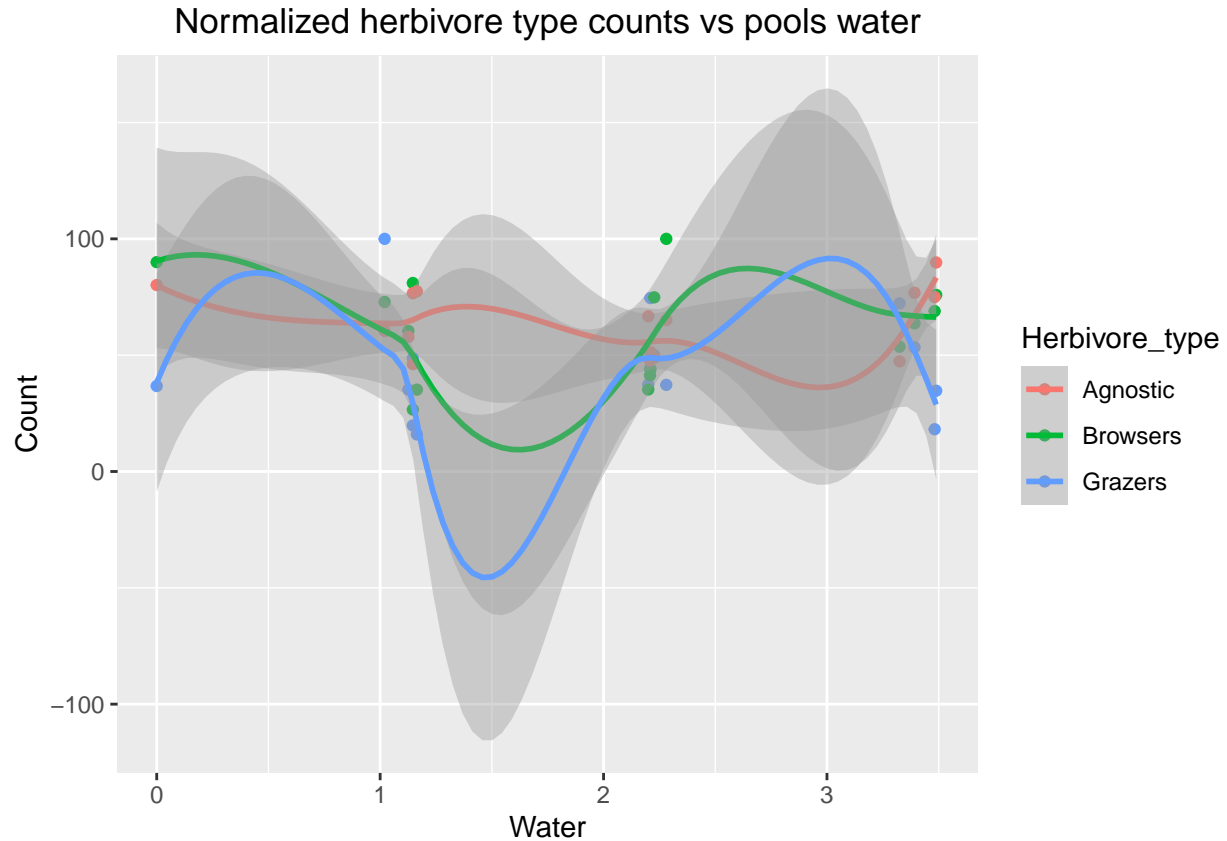


```
ggplot(data=filter(gc_clim_norm_type_plot, Area=="pools")) + geom_point(mapping=aes(x=Water, y=Count, color=Herbivore_type))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 21 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 21 rows containing missing values (geom_point).
```



Discussion

Previous wet season precipitation

The amount of precipitation during the previous wet season did not appear to be strongly correlated with game counts (individual species or herbivore type). This may be due to the large time period between the previous wet season and the game count, variation in climatic factors (such as temperature, wind and solar radiation) during this interim period may have confounded any relationship.

Soil moisture, drought index, vegetation and water

There did not appear to be any easily discernible correlation between drought index and water, and game counts. Soil moisture may have and a slight negative correlation with agnostic herbivores and browsers, but not for grazers. This correlation was present for both park and pools soil moisture. There was a strongly negative correlation between the enhanced vegetation index (pool and park) and browsers, and a less strong negative correlation between enhanced vegetation index (pool and park) and agnostic herbivores. This appears to indicate that extent of local migrations is more influenced by presence/absence of vegetation than by the presence/absence of water. Please note that although the extent of browsers local migration is more varied, this does not mean that browsers are more likely to migrate depending on water or vegetation. It may be the case that grazers always migrate depending on the season, despite what the climatic or vegetation conditions are, thus resulting in more uniform game counts from year to year. This study only looks at the variation of migratory patterns, not whether species/herbivore groups are more likely to exhibit season local migrations.