

# Exploring bee-related spatial data

Vincent Cortes - Lauren Ponisio

## Conservation/ecology Topics

- Species distributions

## Computational Topics

- Convert a data frame to a spatial object.
  - Plot multiple spatial layers.
- 

## Lab part 1: Oregon bee atlas data exploration

a. Import the OBA data.

```
OBAdata <- read.csv("OBA_2018-2023.csv")
```

```
head(OBAdata)
```

```
##              Observation.No. Voucher.No. user_id    user_login
## 1 Andony_Melathopoulos:18.001.001          429964 amelathopoulos
## 2 Andony_Melathopoulos:18.002.001          429964 amelathopoulos
## 3 Andony_Melathopoulos:18.002.002          429964 amelathopoulos
## 4 Andony_Melathopoulos:18.002.003          429964 amelathopoulos
## 5 Andony_Melathopoulos:18.002.004          429964 amelathopoulos
## 6 Andony_Melathopoulos:18.002.005          429964 amelathopoulos
## Collector...First.Name Collector...First.Initial Collector...Last.Name
## 1              Andony              A.              Melathopoulos
## 2              Andony              A.              Melathopoulos
## 3              Andony              A.              Melathopoulos
## 4              Andony              A.              Melathopoulos
## 5              Andony              A.              Melathopoulos
## 6              Andony              A.              Melathopoulos
## Collectors taxon_kingdom_name Associated.plant...genus..species url
## 1 A.Melathopoulos
## 2 A.Melathopoulos
## 3 A.Melathopoulos
## 4 A.Melathopoulos
## 5 A.Melathopoulos
## 6 A.Melathopoulos
## Sample.ID Specimen.ID Collection.Day.1 Month.1 MonthJul MonthAb Year.1
## 1              NA              18      iii    March      3    2018
## 2              NA              20      iii    March      3    2018
## 3              NA              20      iii    March      3    2018
```

## 4	NA	20	iii	March	3	2018
## 5	NA	2	ix	September	9	2018
## 6	NA	2	ix	September	9	2018
##	Collection.Date	Time.1	Collection.Day.2	Month.2	Year.2	Collection.Day.2.Merge
## 1	3/18/2018					
## 2	3/20/2018					
## 3	3/20/2018					
## 4	3/20/2018					
## 5	9/2/2018					
## 6	9/2/2018					
##	Time.2	Collection.ID	Position.of.1st.digit	Collection.No.	Sample.No.	
## 1	A	Melathopoulos		1	1	
## 2	A	Melathopoulos		2	1	
## 3	A	Melathopoulos		2	2	
## 4	A	Melathopoulos		2	3	
## 5	A	Melathopoulos		2	4	
## 6	A	Melathopoulos		2	5	
##	Country	State	County		Location	
## 1	USA	Oregon	Benton		Corvallis, NW Orchard Ave	
## 2	USA	Oregon	Benton		Corvallis, NW Orchard Ave	
## 3	USA	Oregon	Benton		Corvallis, NW Orchard Ave	
## 4	USA	Oregon	Benton		Corvallis, NW Orchard Ave	
## 5	USA	Oregon	Clatsop	Clatskanie, Big Creek Mainline, Knob Point Road		
## 6	USA	Oregon	Clatsop	Clatskanie, Big Creek Mainline, Knob Point Road		
##	Abbreviated.Location	Collection.Site.Description		Team		
## 1	Astoria Maggie Johnson Rd			Melathopoulos		
## 2	Big Crk. Mainline Knob Pt Rd			Melathopoulos		
## 3	Big Crk. Mainline Knob Pt Rd			Melathopoulos		
## 4	Big Crk. Mainline Knob Pt Rd			Melathopoulos		
## 5	Big Crk. Mainline Knob Pt Rd			Melathopoulos		
## 6	Big Crk. Mainline Knob Pt Rd			Melathopoulos		
##	Habitat	Elevation..m.	Dec..Lat.	Dec..Long.	X	Collectionmethod
## 1			44.556	-123.285	NA	Net
## 2			44.567	-123.283	NA	Net
## 3			44.567	-123.283	NA	Net
## 4			44.567	-123.283	NA	Net
## 5			46.102	-123.506	NA	Net
## 6			46.102	-123.506	NA	Net
##	Collection.method.merge.field	Associated.plant...family				
## 1						
## 2						
## 3						
## 4						
## 5						
## 6						
##	Associated.plant...genus..species.1	Associated.plant...Inaturalist.URL				
## 1						
## 2						
## 3						
## 4						
## 5						
## 6						
##	Associated.plant	Assoc.plant.merge.field		Collectors.1		
## 1				Andony Melathopoulos		

```
## 2          Andony Melathopoulos
## 3          Andony Melathopoulos
## 4          Andony Melathopoulos
## 5          Andony Melathopoulos
## 6          Andony Melathopoulos
## Collector.1.abreviation Collector.2 Collector.3 Genus Species sex caste
## 1      A Melathopoulos      NA      NA
## 2      A Melathopoulos      NA      NA
## 3      A Melathopoulos      NA      NA
## 4      A Melathopoulos      NA      NA
## 5      A Melathopoulos      NA      NA
## 6      A Melathopoulos      NA      NA
## vol.det.Genus vol.det.Species vol.det.sex.caste Determined.By Date.Determined
## 1                                     NA
## 2                                     NA
## 3                                     NA
## 4                                     NA
## 5                                     NA
## 6                                     NA
## Verified.By Other.Determiner.s. Other.Dets.Sci..Name.s. Other.Dets..Date.s.
## 1      NA                                     NA      NA
## 2      NA                                     NA      NA
## 3      NA                                     NA      NA
## 4      NA                                     NA      NA
## 5      NA                                     NA      NA
## 6      NA                                     NA      NA
## Additional.Notes X.1
## 1      NA
## 2      NA
## 3      NA
## 4      NA
## 5      NA
## 6      NA
```

- b. Find the columns related to genus and species and paste them together (with a space between) using the function `paste()`. Name the new column `GenusSpecies`.

```
GenusSpecies <- paste(OBAdata$Genus, OBAdata$Species, sep=" ", recycle0 = FALSE)
head(GenusSpecies)
```

```
## [1] " " " " " " " " " " " "
```

- c. Use `sort()` and `unique()` to print the unique values of `GenusSpecies` in alphabetical order. How many species are there?

```
head(sort(unique(GenusSpecies)))
```

```
## [1] " " "Agapostemon "
## [3] "Agapostemon femoratus" "Agapostemon texanus"
## [5] "Agapostemon virescens " "Agapostemon femoratus"
```

Some specimens are not identified to species, only genus. How is this reflected in the data? In two weeks we will learn how to clean this up using regular expressions.

- d. So many bees, so little time. Count up the occurrences of each bee species, and subset the data to bees that have been seen at least two times. You can use the tidyverse or any other functions in R that you

like. How many “species” are there?

```
bee_counts <- OBAdata%>%
  mutate(GenusSpecies = paste(Genus, Species, sep=" ")) %>%
  group_by(GenusSpecies) %>%
  tally() %>%
  filter(n >= 2)
```

```
head(bee_counts)
```

```
## # A tibble: 6 x 2
##   GenusSpecies      n
##   <chr>          <int>
## 1 " "            92466
## 2 "Agapostemon "    261
## 3 "Agapostemon femoratus" 372
## 4 "Agapostemon texanus"  150
## 5 "Agapostemon virescens "  44
## 6 "Agapostemon femoratus" 165
```

```
num_species <- nrow(bee_counts)
num_species
```

```
## [1] 455
```

- e. Google a few bee names (that have been seen > 2 times) and find one with an a look that resonates with you.

What is the name of your bee?

Agapostemon texanus

Import the photos into Rmarkdown below (hint: googling bee name “discover life” or “inat” can often get you a photo. Many bees will no have any photos :(

## Lab part 2: Plotting the distrubution of your spirit bee.

How that have chosen your spirit bee, we would like to plot it’s distribution. What is the crs of the data? Annoyingly it is not described anywhere in the spreadsheet (always list your crs in your data) but it is the same as what inat uses because all bees have a georeferenced plant host. If the data is in lat long, it is “unprojected” so only a datum will be listed. DATUM: WGS84, unprojected lat long. EPSG code: 4326.

```
crs("EPSG:4326")
```

```
## [1] "GEOGCRS[\"WGS 84\", \n      ENSEMBLE[\"World Geodetic System 1984 ensemble\", \n      MEMBER[\"Wo
```

- a. Extract the X and Y locations for your species only from the data and create a spatial object. Don’t forget to set the CRS! Hint 1: consider what other data you would like to keep as attributes, for example what flower they were foraging on. Hint 2: Remember the lat is y and long is x. Hint 3: You may want to rename the column names you can use, colnames() and reassign the names, since the ones in the oba data spreadsheet are really ugly.

```
# Filter the data for Agapostemon texanus
spirit_bee_data <- OBAdata %>%
  filter(paste(Genus, Species, sep=" ") == "Agapostemon texanus") %>%
  select(Longitude = Dec..Long., Latitude = Dec..Lat., Flower = Associated.plant)
```

```
# Renames columns
colnames(spirit_bee_data) <- c("Longitude", "Latitude", "Flower")
```



Figure 1: Photo of *Agapostemon texanus*

```
# Convert the filtered data to a spatial object
spirit_bee_sf <- st_as_sf(spirit_bee_data, coords = c("Longitude", "Latitude"), crs = 4326)

# Display the spatial object
print(spirit_bee_sf)
```

```
## Simple feature collection with 710 features and 1 field
## Geometry type: POINT
## Dimension: XY
## Bounding box: xmin: -124.408 ymin: 42.004 xmax: -116.935 ymax: 45.875
## Geodetic CRS: WGS 84
## First 10 features:
```

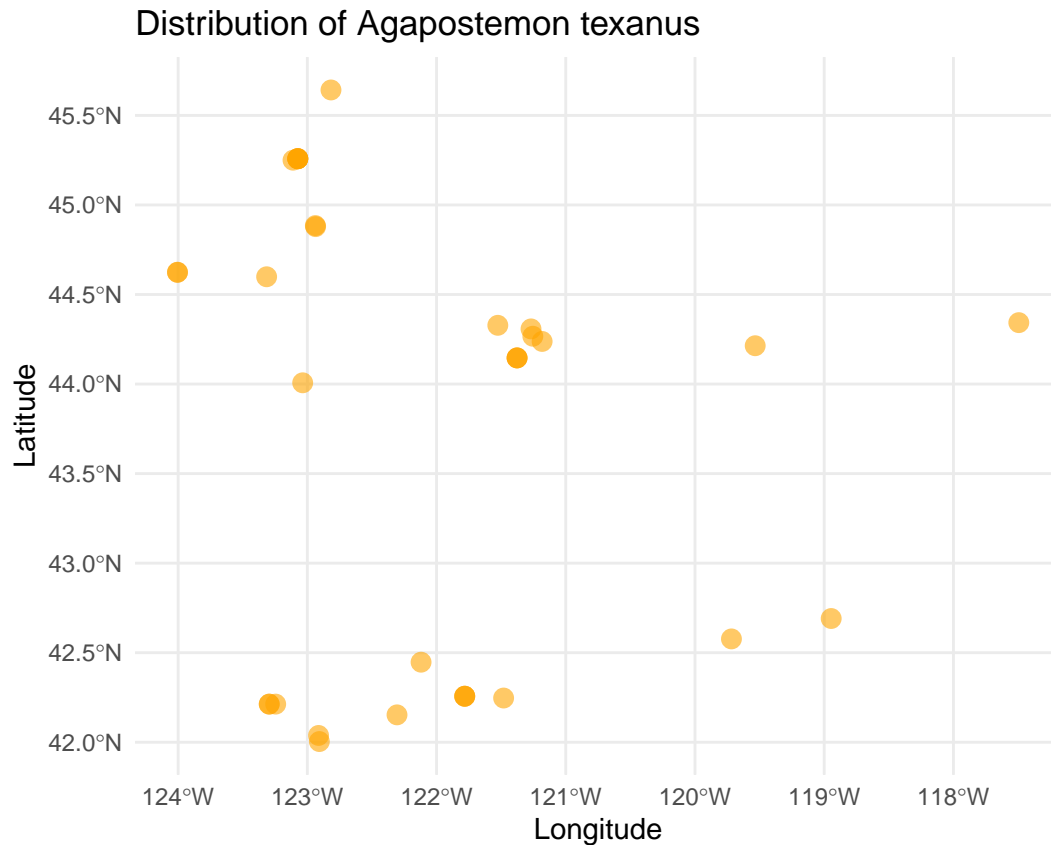
```
##           Flower          geometry
## 1              POINT (-122.814 45.643)
## 2              POINT (-122.814 45.643)
## 3              POINT (-122.816 45.644)
## 4 Symphoricarpos albus POINT (-122.817 45.642)
## 5              POINT (-123.074 45.258)
## 6              POINT (-123.074 45.258)
## 7      Sidalcea sp. POINT (-123.111 45.25)
## 8   Heracleum maximum POINT (-123.074 45.258)
## 9              POINT (-123.135 45.202)
## 10             POINT (-123.06 45.257)
```

```
spirit_bee_sf <- spirit_bee_sf[spirit_bee_sf$Flower != "",]
spirit_bee_sf <- spirit_bee_sf[spirit_bee_sf$Flower != "Salix sp., Achillea millefolium (Yarrow) and Co",]
sort(unique(spirit_bee_sf$Flower))
```

```
## [1] "Alcea rosea" "Berberis aquifolium"
## [3] "Calochortus macrocarpus" "Chrysothamnus nauseosus"
## [5] "Cucurbita sp." "Ericameria nauseosa"
## [7] "Eriogonum sp." "Eriophyllum lanatum"
## [9] "Eriophyllum lanatum (Linear daisy)" "Eschscholzia californica"
## [11] "Grindelia stricta stricta" "Heracleum maximum"
## [13] "Ilex sp." "Jaumea carnosa"
## [15] "Leucanthemum vulgare" "Lonicera tatarica"
## [17] "Potentilla gracilis" "Rubus parviflorus"
## [19] "Rudbeckia hirta" "Senecio hydrophilus"
## [21] "Sidalcea campestris" "Sidalcea sp."
## [23] "Symphoricarpos albus" "Symphyotrichum oolentangiense"
## [25] "Trifolium repens" "Wyethia sp."
```

b. Plot your exciting bee data!

```
ggplot(data = spirit_bee_sf) +
  geom_sf(color = "orange", size = 3, alpha = 0.6) +
  labs(title = "Distribution of Agapostemon texanus",
       x = "Longitude",
       y = "Latitude") +
  theme_minimal()
```



Not so exciting without some kind of background...

Luckily we can download basemaps into R using the `map_data` function in `ggplot` (among many others). There is an example for retrieving the Oregon county polygons.

```
or <- map_data("county", "oregon") %>%
  select(long = long, lat, group, id = subregion)
```

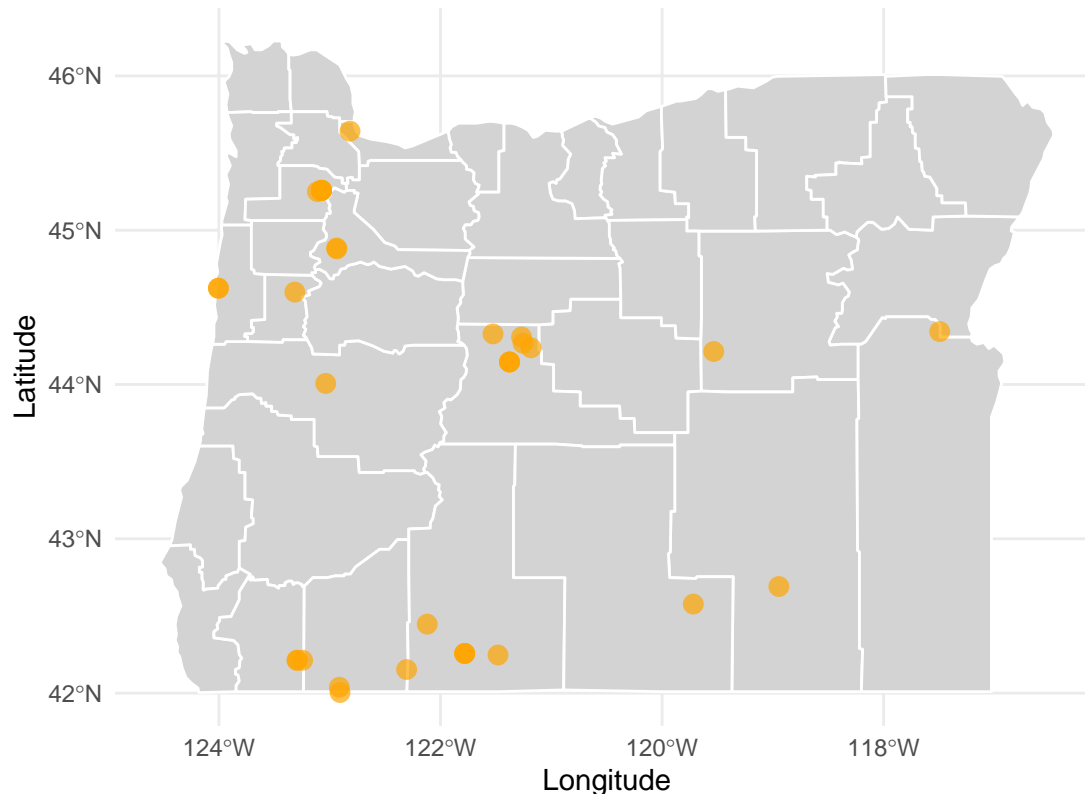
c. Add your species's points to your choice or an Oregon basemap.

```
library(ggspatial)

ggplot() +
  geom_polygon(data = or, aes(x = long, y = lat, group = group),
    fill = "lightgrey", color = "white") +

  geom_sf(data = spirit_bee_sf, color = "orange", size = 3, alpha = 0.7) +
  labs(title = "Distribution of Agapostemon texanus in Oregon",
    x = "Longitude",
    y = "Latitude") +
  theme_minimal() +
  coord_sf()
```

## Distribution of *Agapostemon texanus* in Oregon



# Lab part 3:

### Cartography

- a. Here is your moment to explore your cartographic skills.
  1. Add another spatial layer relevant to your final project and tweak the Oregon map in anyway that is useful/visually appealing. You may need to crop that layer to the extent of your species's distribution.
  2. Color your points according to some data attribute and add a legend (month collected, county, collector, associated plant, whatever you think is interesting). You may need to circle back to 2.1 to save additional attributes when you converted the dataframe to a spatial object.
  3. Fine-tune your map: add a title, make sure the legend label makes sense, add a scale bar (google “add scale bar map ggplot” and choose your favorite package). All maps must always have a scale bar. You can add a N arrow as well, though some cartographers argue that is only necessary if N isn't at the top of the map.
  4. Write a figure caption for your map explaining any interesting trends you see.
  5. Export you cropped layer to a .shp so you can use it again for your final project.
  6. Push this lab to your github repo (just the .Rmd, don't push the data!)

```
fire_data <- terra::rast("HolidayFarm_SBS_final.tif")
fire_data <- terra::project(fire_data, st_crs(spirit_bee_sf)$wkt)
fire_data_cropped <- terra::crop(fire_data, terra::ext(spirit_bee_sf))
fire_df <- as.data.frame(fire_data_cropped, xy = TRUE, na.rm = TRUE)

or_sf <- st_as_sf(map_data("county", "oregon"), coords = c("long", "lat"), crs = 4326)
fire_data_oregon <- terra::crop(fire_data, terra::ext(or_sf))
fire_data_cropped <- terra::crop(fire_data_oregon, terra::ext(spirit_bee_sf))

ggplot() +
  # Oregon counties map
  geom_polygon(data = or, aes(x = long, y = lat, group = group),
```



```

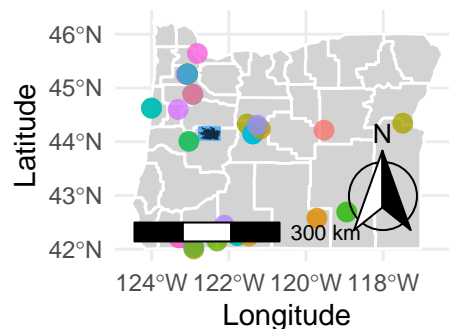
    fill = "lightgrey", color = "white") +
geom_raster(data = fire_df, aes(x = x, y = y, fill = HolidayFarm_SBS_final)) +
# Spirit bee data points
geom_sf(data = spirit_bee_sf, aes(color = Flower), size = 3, alpha = 0.8) +
# Add a scale bar
annotation_scale(location = "bl", width_hint = 0.5) +
# Add a north arrow
annotation_north_arrow(location = "br", which_north = "true",
                        style = north_arrow_fancy_orienteering) +
labs(title = "Distribution of Agapostemon texanus and Wildfire Data",
     subtitle = "Colored by Associated Plant",
     color = "Associated Plant",
     fill = "Fire Intensity",
     x = "Longitude",
     y = "Latitude") +
theme_minimal() +

theme(
  legend.position = "bottom", # Position legend at the top
  plot.title = element_text(hjust = 0.5, face = "bold", size = 14, margin = margin(b = 10)),
  plot.subtitle = element_text(hjust = 0.5, size = 12)
) +
coord_sf()

```

## Distribution of Agapostemon texanus and Wildfire Data

Colored by Associated Plant



● Alcea rosea	● Eriogonum sp.	● Ilex sp.
● Berberis aquifolium	● Eriophyllum lanatum	● Jaumea carnosa
● Calochortus macrocarpus	● Eriophyllum lanatum (Linear daisy)	● Leucanthemum vulgare
● Chrysothamnus nauseosus	● Eschscholzia californica	● Lonicera tatarica
● Cucurbita sp.	● Grindelia stricta stricta	● Potentilla gracilis
● Ericameria nauseosa	● Heracleum maximum	● Rubus parviflorus

We are looking forward to seeing the maps you create!

## Lab part 4: Spatial summary statistics

For your final projects, you will likely need to come up with summary statistics that describes the areas around where bees are captured. a. Using the distribution of your chosen bee and the spatial layer you imported in 2.6, extract a meaningful summary statistics from your spatial layer within a buffer of 500, 750 1000 km. b. Create a plot that illustrates this summary data (box plot, barplot, scatter plot, histogram). c. Create a map of your cropped spatial data.

*#alittle confused how to work with this fire dataset, I think this this incorrect, maybe if I used a be*

*#Found this package online, that loads data faster???*

```
library(exactextractr)
```

*# Create buffers*

```
buffer_500 <- st_buffer(spirit_bee_sf, dist = 50000) # 50 km
```

```
buffer_750 <- st_buffer(spirit_bee_sf, dist = 80000) # 80 km
```

```
buffer_1000 <- st_buffer(spirit_bee_sf, dist = 100000) # 100 km
```

*# Extracts the mean fire intensity for each buffer distance*

```
fire_mean_500 <- exact_extract(fire_data_cropped, buffer_500, fun = "mean")
```

```
## |
```

```
fire_mean_750 <- exact_extract(fire_data_cropped, buffer_750, fun = "mean")
```

```
## |
```

```
fire_mean_1000 <- exact_extract(fire_data_cropped, buffer_1000, fun = "mean")
```

```
## |
```

*# This will combine results into a DF*

```
fire_summary <- data.frame(
  Distance = rep(c("50 km", "80 km", "100 km"),
    times = c(length(fire_mean_500), length(fire_mean_750), length(fire_mean_1000))),
  FireIntensity = c(fire_mean_500, fire_mean_750, fire_mean_1000)
)
```

```
ggplot() +
```

```
  geom_polygon(data = or, aes(x = long, y = lat, group = group),
```

```
    fill = "lightgrey", color = "white") +
```

```
  geom_raster(data = fire_df, aes(x = x, y = y, fill = HolidayFarm_SBS_final)) +
```

```
  geom_sf(data = buffer_500, fill = "blue", alpha = 0.2) +
```

```
  geom_sf(data = buffer_750, fill = "green", alpha = 0.2) +
```

```
  geom_sf(data = buffer_1000, fill = "red", alpha = 0.2) +
```

```
  geom_sf(data = spirit_bee_sf, color = "orange", size = 3) +
```

```
  annotation_scale(location = "bl", width_hint = 0.5) +
```

```
  annotation_north_arrow(location = "br", which_north = "true",
```

```
    style = north_arrow_fancy_orienteering) +
```

```
  labs(title = "Fire Intensity around Agapostemon texanus Observations",
```

```
    fill = "Fire Intensity",
```

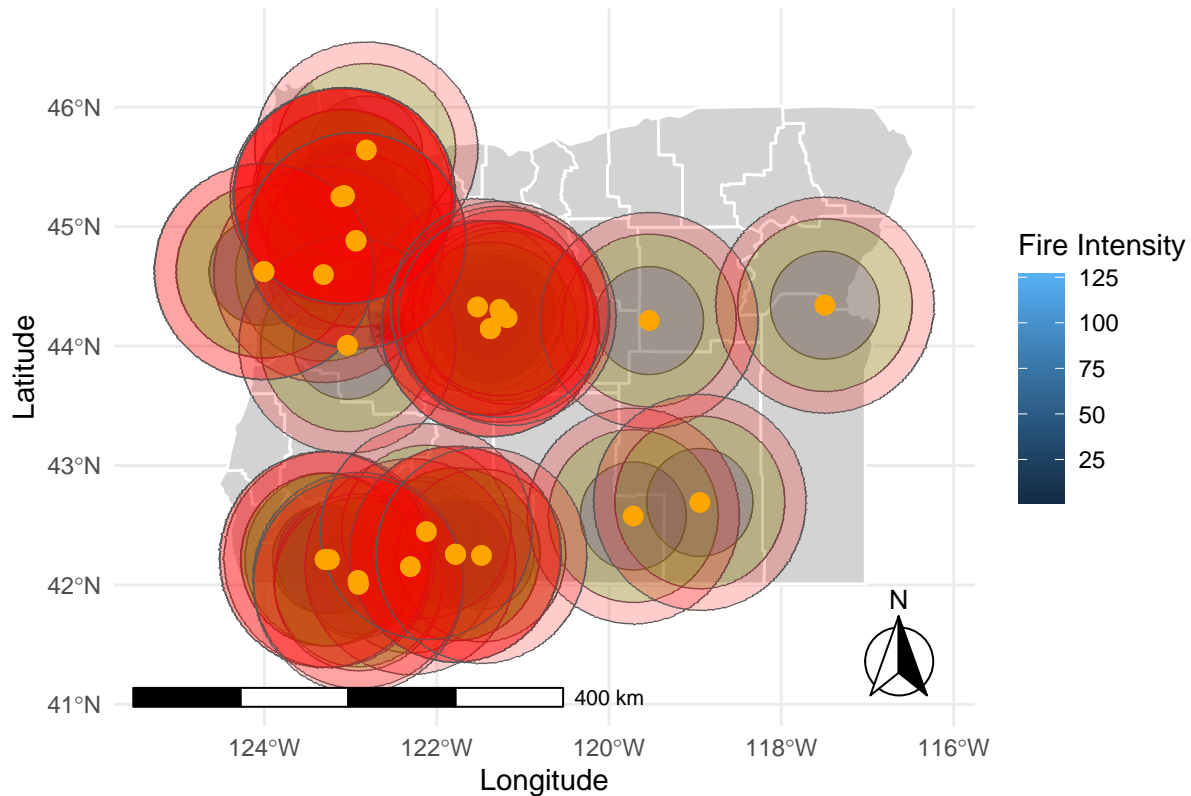
```
    color = "Buffer Distance",
```

```
x = "Longitude",
y = "Latitude") +

theme_minimal() +
coord_sf()
```

## Scale on map varies by more than 10%, scale bar may be inaccurate

### Fire Intensity around *Agapostemon texanus* Observations



```
ggplot(fire_summary, aes(x = reorder(Distance, -as.numeric(Distance)), y = FireIntensity, fill = Distance)) +
  geom_boxplot() +
  labs(title = "Mean Fire Intensity within Buffers of Agapostemon texanus",
       x = "Buffer Distance",
       y = "Mean Fire Intensity") +
  theme_minimal() +
  theme(legend.position = "none")
```

## Warning in tapply(X = X, INDEX = x, FUN = FUN, ...): NAs introduced by coercion

## Warning: Removed 83 rows containing non-finite outside the scale range

## (`stat\_boxplot()`).

