

TidyTuesday World Spider Database

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```
spiders <- readr::read_csv('https://raw.githubusercontent.com/rfordatascience/tidyuesday/master/data/2021/2021-spiders.csv')
```

```
## Rows: 49785 Columns: 10
```

```
## -- Column specification -----
```

```
## Delimiter: ","
```

```
## chr (7): species_lsid, family, genus, species, subspecies, author, distribution
```

```
## dbl (3): speciesId, year, parentheses
```

```
##
```

```
## 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.
```

```
spiders <- spiders %>% janitor::clean_names()
```

```
spiders <- spiders %>% janitor::remove_empty(which = c("cols", "rows"))
```

Discoveries by Year: 2010 - 2021

```
spiders %>% count(year) %>%
```

```
  ggplot(aes(x=year, y=n)) + geom_line() +
```

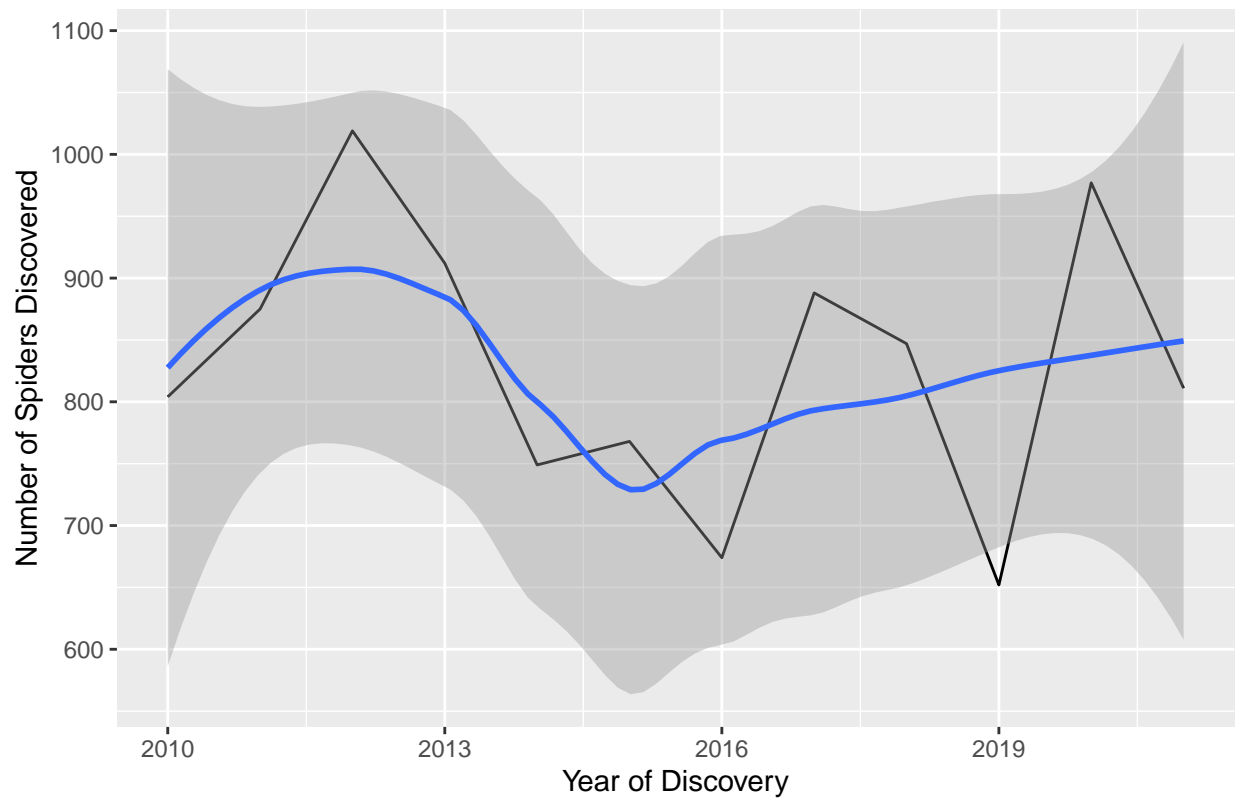
```
  labs(title = "Number of Spiders Discovered by Year",
```

```
        x="Year of Discovery", y="Number of Spiders Discovered") +
```

```
  geom_smooth(aes(x=year, y=n))
```

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

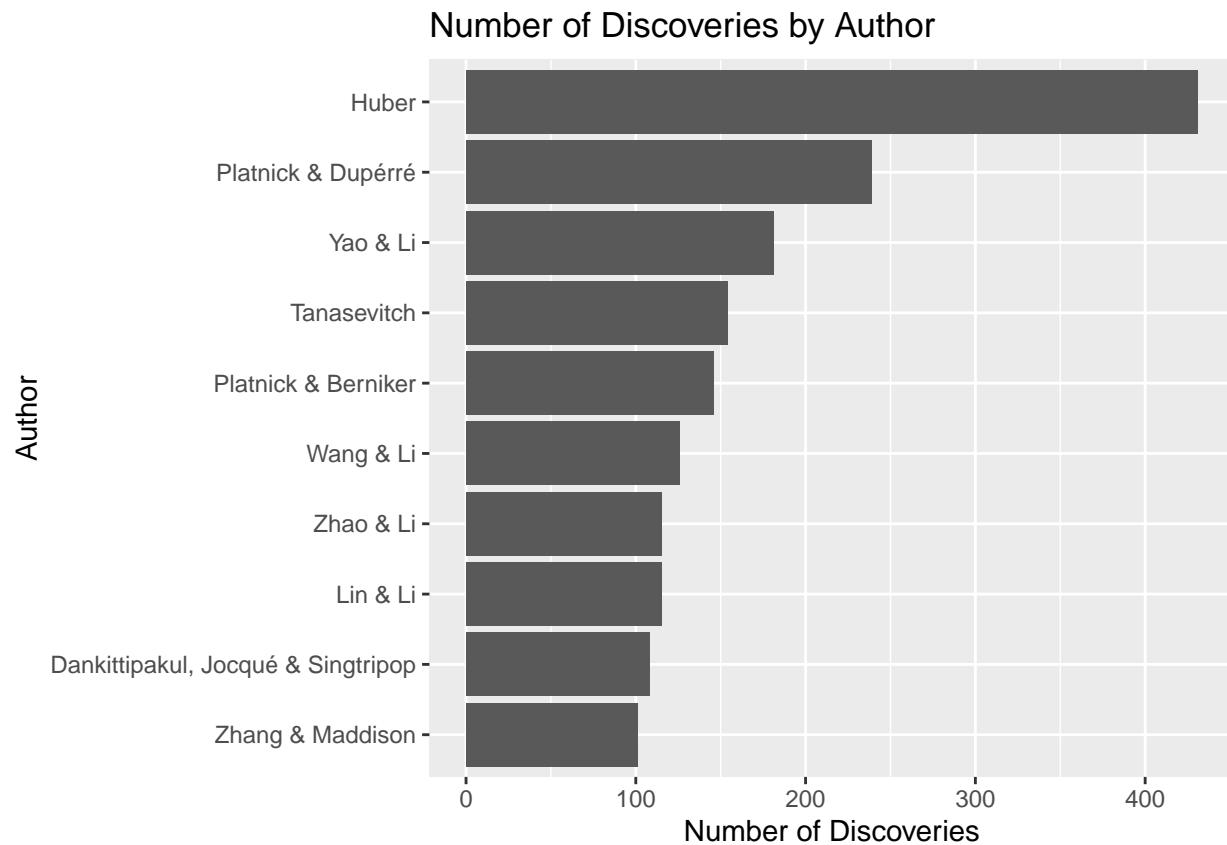
Number of Spiders Discovered by Year



Count of Spiders Discovered by Author

```
spiders %>% count(author, sort = T) %>% top_n(10) %>% ggplot(aes(x=reorder(author, n), y=n)) +
  geom_col() + coord_flip() +
  labs(title = "Number of Discoveries by Author",
        x = "Author", y = "Number of Discoveries")
```

Selecting by n

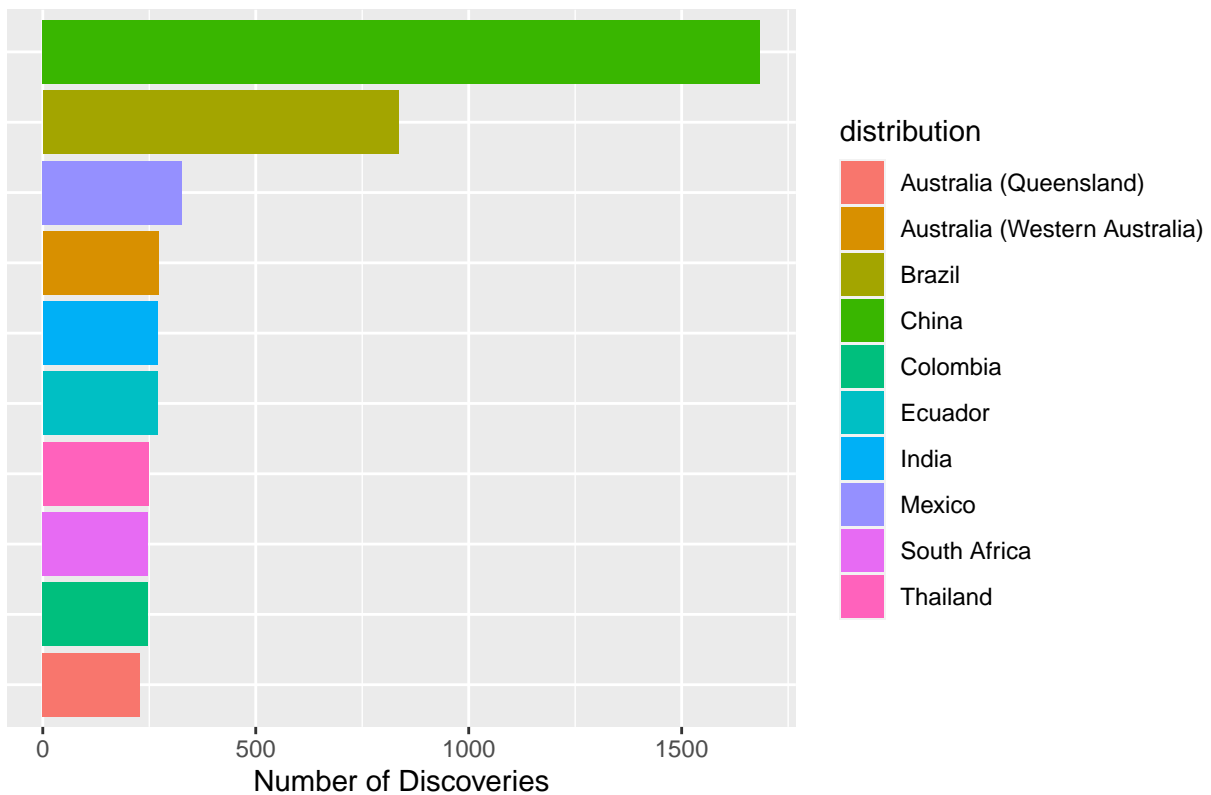


Distribution by Location: 2010-2021

```
spiders %>% count(distribution, sort = T) %>% top_n(10) %>%
  ggplot(aes(x=reorder(distribution, n), y=n)) + geom_col(aes(fill=distribution)) +
  theme(axis.title.y=element_blank(),
        axis.text.y=element_blank(),
        axis.ticks.y=element_blank()) +
  labs(title = "Top 10 Countries with Discoveries: 2010 - 2021",
        y="Number of Discoveries") + coord_flip()
```

Selecting by n

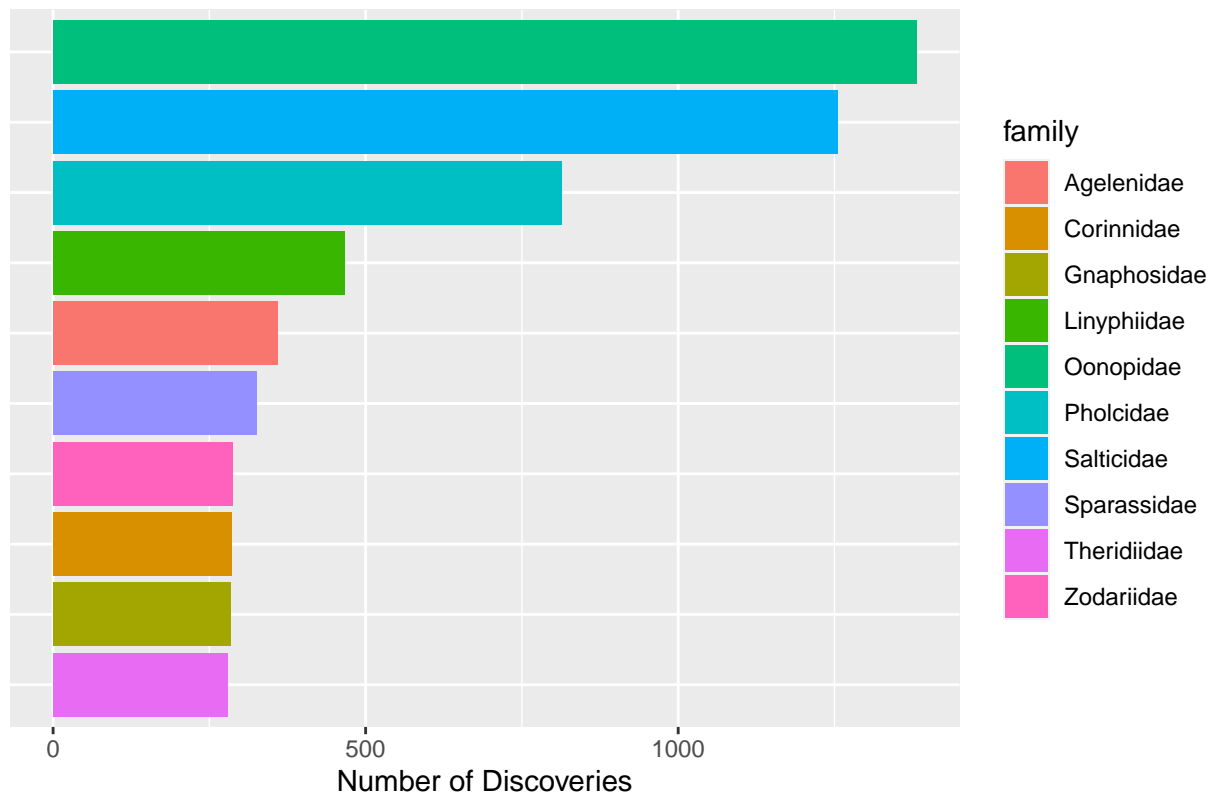
Top 10 Countries with Discoveries: 2010 – 2021



```
spiders %>% count(family, sort = T) %>% top_n(10) %>%
  ggplot(aes(x=reorder(family, n), y=n)) + geom_col(aes(fill=family)) +
  theme(axis.title.y=element_blank(),
        axis.text.y=element_blank(),
        axis.ticks.y=element_blank()) +
  labs(title = "Top 10 Families: 2010 - 2021",
        y="Number of Discoveries") + coord_flip()
```

Selecting by n

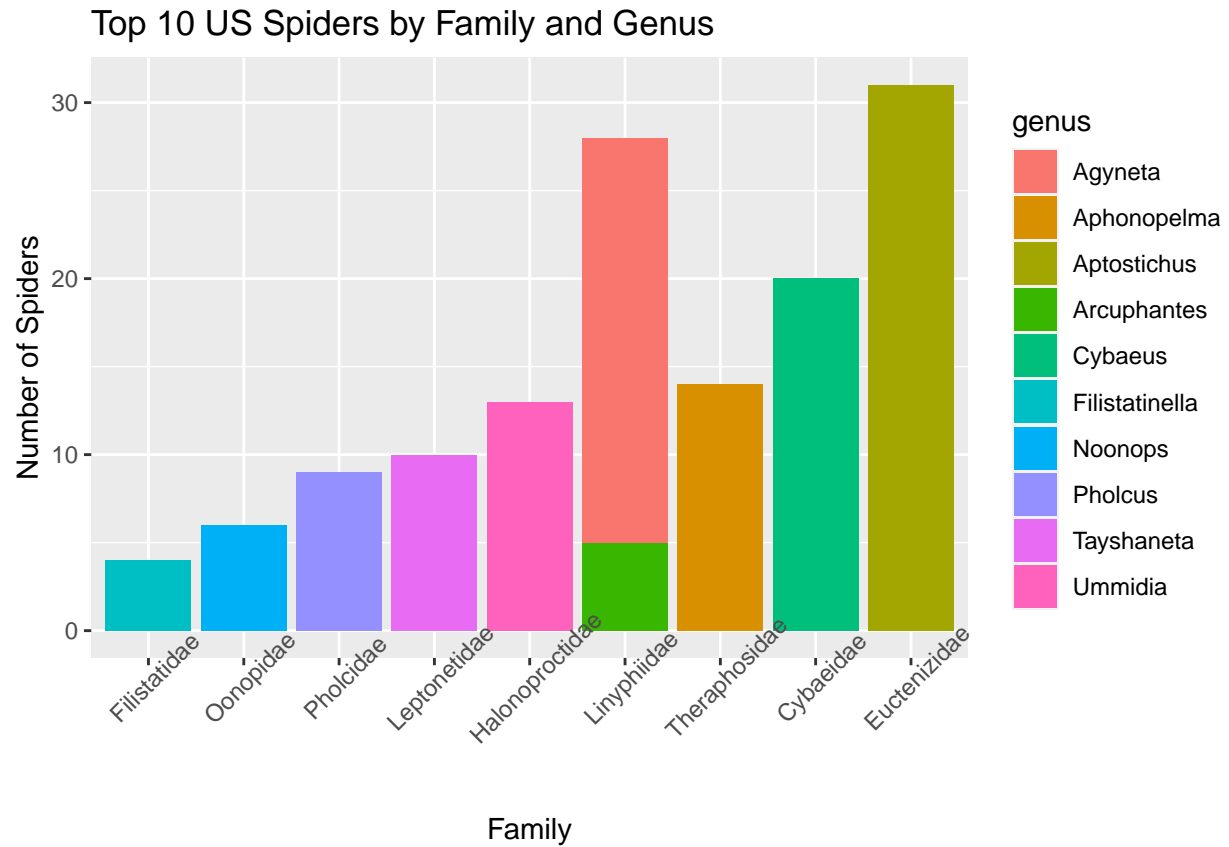
Top 10 Families: 2010 – 2021



US Discoveries by Family and Genus

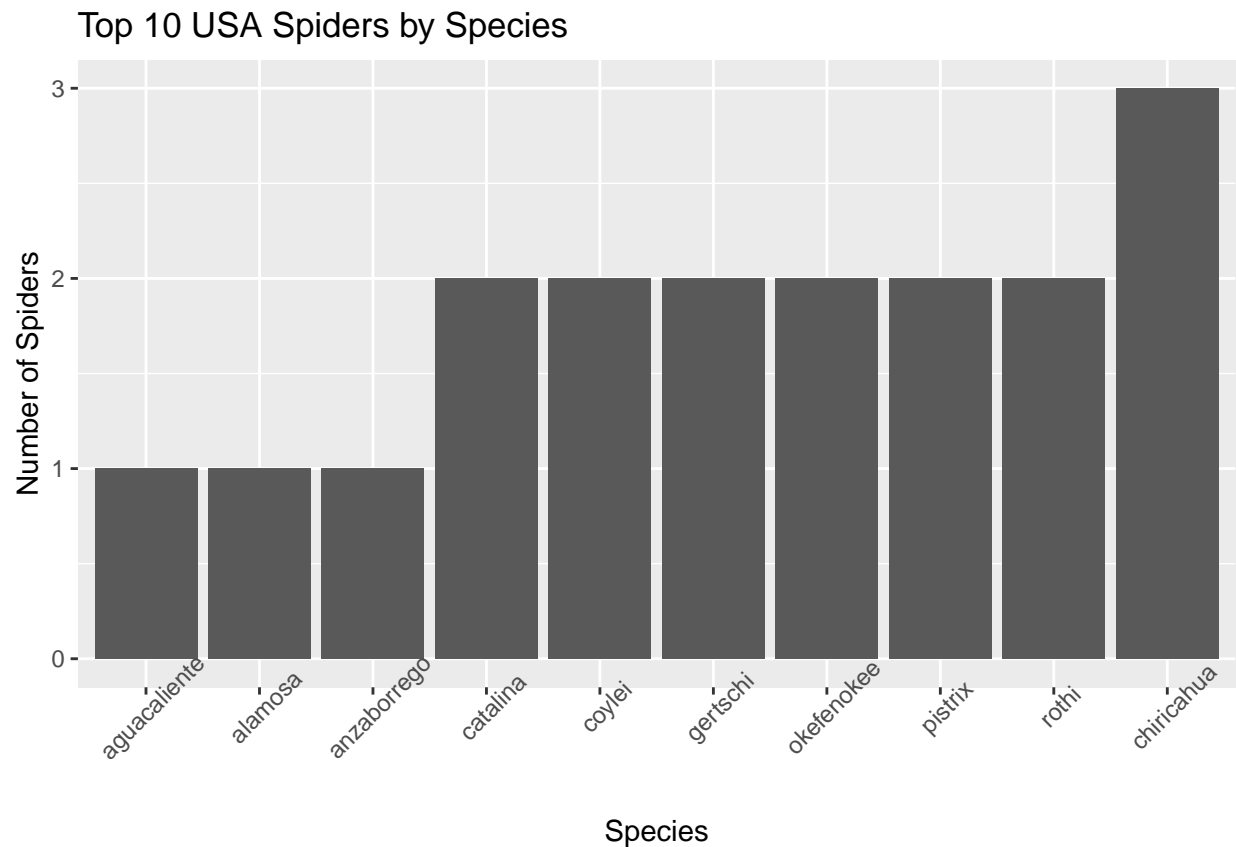
```
spiders %>% filter(distribution == "USA") %>% count(family, genus, sort = T) %>%
  top_n(10) %>%
  ggplot() + geom_col(aes(x=reorder(family, n), y=n, fill=genus)) +
  theme(axis.text.x = element_text(angle = 45)) +
  labs(title = "Top 10 US Spiders by Family and Genus", x="Family",
        y="Number of Spiders")
```

Selecting by n



US Discoveries by Species

```
spiders %>% filter(distribution == "USA") %>% count(species, sort = T) %>%
  head(10) %>%
  ggplot() + geom_col(aes(x=reorder(species, n), y=n)) +
  theme(axis.text.x = element_text(angle = 45)) +
  labs(title = "Top 10 USA Spiders by Species", x="Species",
        y="Number of Spiders")
```

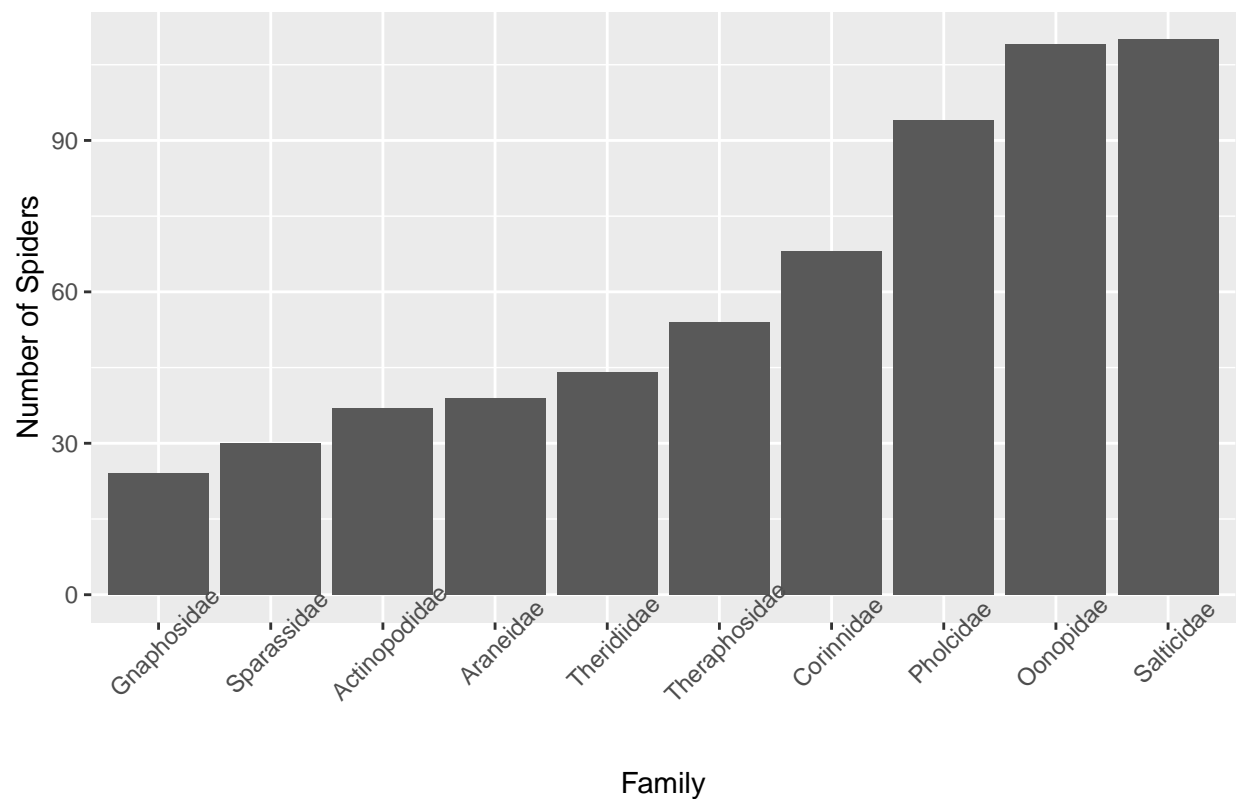


Brazil

```
spiders %>% filter(distribution == "Brazil") %>% count(family, sort = T) %>%
  top_n(10) %>%
  ggplot() + geom_col(aes(x=reorder(family, n), y=n)) +
  theme(axis.text.x = element_text(angle = 45)) +
  labs(title = "Top 10 Brazil Spiders by Family", x="Family",
        y="Number of Spiders")
```

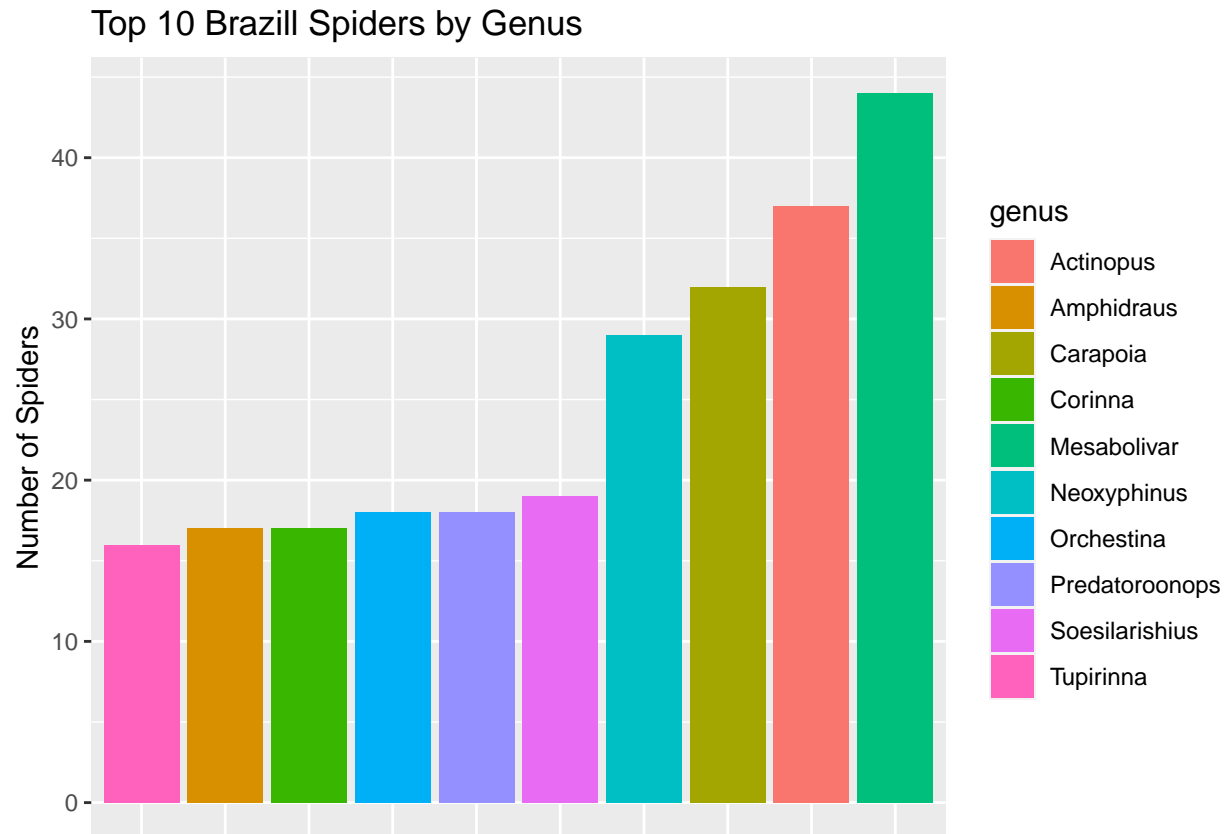
Selecting by n

Top 10 Brizal Spiders by Family



```
spiders %>% filter(distribution == "Brazil") %>% count(genus, sort = T) %>%
  top_n(10) %>%
  ggplot() + geom_col(aes(x=reorder(genus, n), y=n, fill=genus)) +
  theme(axis.text.x = element_text(angle = 45)) +
  labs(title = "Top 10 Brazill Spiders by Genus", x="Genus",
        y="Number of Spiders") +
  theme(axis.title.x=element_blank(),
        axis.text.x=element_blank(),
        axis.ticks.x=element_blank())
```

Selecting by n



Brazil's Top 10 Spiders by Species

```
spiders %>% filter(distribution == "Brazil") %>% count(species, sort = T) %>%
  top_n(10) %>%
  ggplot() + geom_col(aes(x=reorder(species, n), y=n, fill=species)) +
  theme(axis.text.x = element_text(angle = 45)) +
  labs(title = "Top 10 Brazill Spiders by Species", x="Species",
       y="Number of Spiders") +
  theme(axis.title.x=element_blank(),
        axis.text.x=element_blank(),
        axis.ticks.x=element_blank())
```

Selecting by n

Top 10 Brazil Spiders by Species

