assignment_06_01_PothineniKalyan

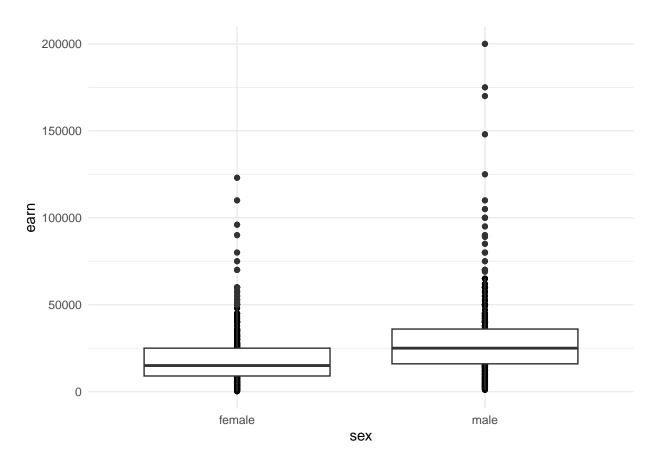
PothineniKalyan

2023-04-22

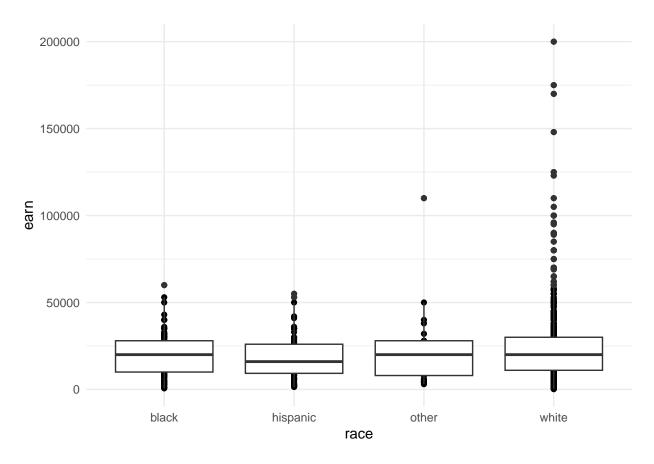
R Markdown

```
# Assignment: ASSIGNMENT 6.2
# Name: POthineni, Kalyan
# Date: 2023-04-20
## Load the ggplot2 package
library(ggplot2)
## Warning: package 'ggplot2' was built under R version 4.2.3
library(dplyr)
## Warning: package 'dplyr' was built under R version 4.2.3
##
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
##
      filter, lag
## The following objects are masked from 'package:base':
##
##
       intersect, setdiff, setequal, union
theme_set(theme_minimal())
## Load the `data/r4ds/heights.csv` to
heights_df <- read.csv("data/r4ds/heights.csv")
head(heights_df, n=5)
           height
                       sex ed age race
      earn
## 1 50000 74.42444
                      male 16 45 white
## 2 60000 65.53754 female 16 58 white
## 3 30000 63.62920 female 16 29 white
## 4 50000 63.10856 female 16 91 other
## 5 51000 63.40248 female 17 39 white
```

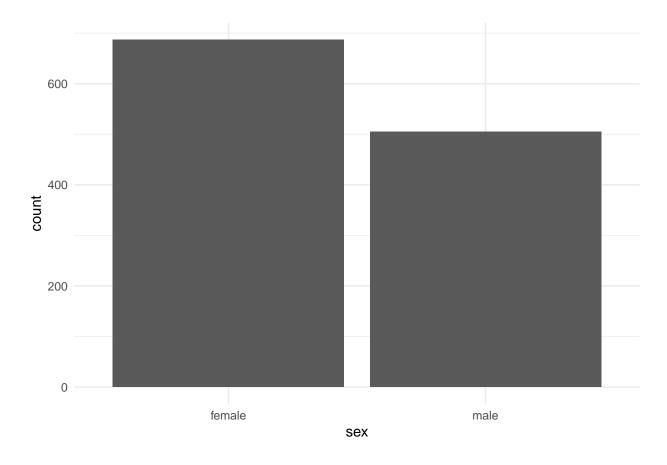
```
# https://ggplot2.tidyverse.org/reference/geom_boxplot.html
## Create boxplots of sex vs. earn and race vs. earn using `geom_point()` and `geom_boxplot()`
## sex vs. earn
ggplot(heights_df, aes(x=sex, y=earn)) + geom_point() + geom_boxplot()
```



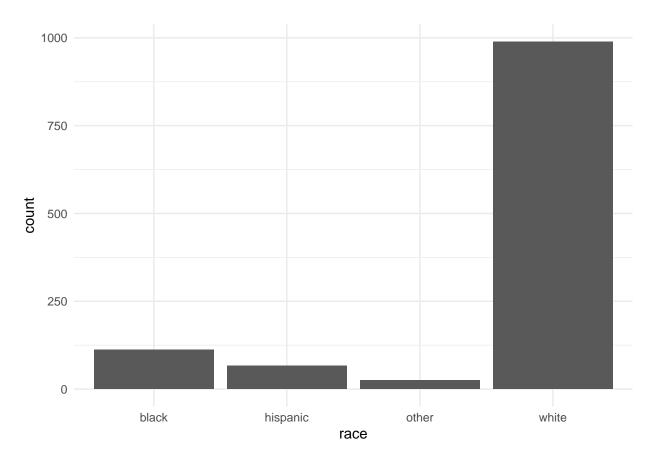
```
## race vs. earn
ggplot(heights_df, aes(x=race, y=earn)) + geom_point() + geom_boxplot()
```



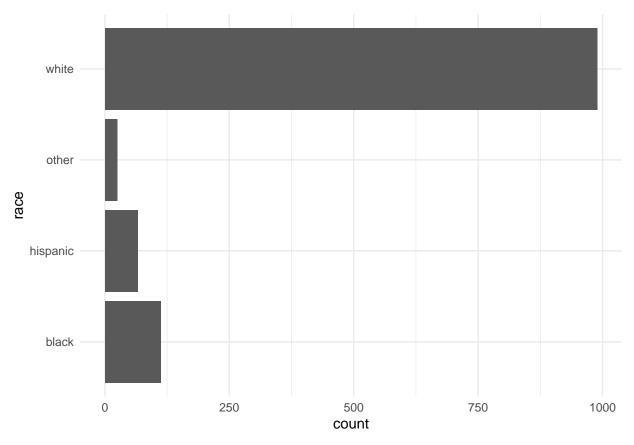
```
## https://ggplot2.tidyverse.org/reference/geom_bar.html
## Using `geom_bar()` plot a bar chart of the number of records for each `sex`
ggplot(heights_df, aes(sex)) + geom_bar()
```



Using `geom_bar()` plot a bar chart of the number of records for each race
ggplot(heights_df, aes(race)) + geom_bar()



Create a horizontal bar chart by adding `coord_flip()` to the previous plot
Fliiping horizontal to vertical and vise versa
ggplot(heights_df, aes(race)) + geom_bar() + coord_flip()



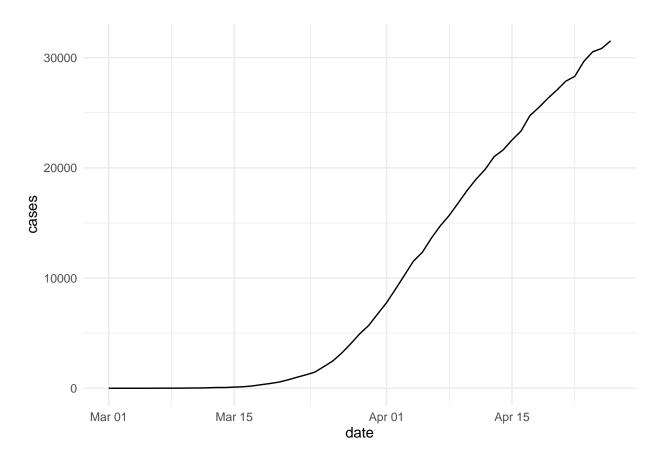
```
\# https://www.rdocumentation.org/packages/ggplot2/versions/3.3.0/topics/geom_path
## Load the file `"data/nytimes/covid-19-data/us-states.csv"` and
## assign it to the `covid_df` dataframe
covid_df <- read.csv("data/nytimes/covid-19-data/us-states.csv")</pre>
head(covid_df, n=5)
                      state fips cases deaths
##
           date
## 1 2020-01-21 Washington
                              53
                                      1
## 2 2020-01-22 Washington
                              53
                                      1
                                             0
## 3 2020-01-23 Washington
                              53
                                      1
                                             0
## 4 2020-01-24
                  Illinois
                              17
                                             0
## 5 2020-01-24 Washington
                              53
                                             0
                                     1
## Parse the date column using `as.Date()``
covid_df$date <- as.Date(covid_df$date)</pre>
head(covid_df$date, n=5)
## [1] "2020-01-21" "2020-01-22" "2020-01-23" "2020-01-24" "2020-01-24"
\hbox{\it \#\# Create three data} frames named `california\_df`, `ny\_df`, and `florida\_df`
## containing the data from California, New York, and Florida
distinct(covid_df, state)
```

##	state
## 1	Washington
## 2	Illinois
## 3	California
## 4	Arizona
## 5	Massachusetts
## 6	Wisconsin
## 7	Texas
## 8	Nebraska
## 9	Utah
## 10	Oregon
## 11	Florida
## 12	New York
## 13	Rhode Island
## 14	Georgia
## 15	New Hampshire
## 16	North Carolina
## 17	New Jersey
## 18	Colorado
## 19	Maryland
## 20	Nevada
## 21	Tennessee
## 22	Hawaii
## 23	Indiana
## 24	Kentucky
## 25	Minnesota
## 26	Oklahoma
## 27	Pennsylvania
## 28	South Carolina
## 29	District of Columbia
## 30	Kansas
## 31	Missouri
## 32	Vermont
## 33	Virginia
## 34	Connecticut
## 35	Iowa
## 36	Louisiana
## 37	Ohio
## 38	Michigan
## 39	South Dakota
## 40	Arkansas
## 41	Delaware
## 42	Mississippi
## 43	New Mexico
## 44	North Dakota
## 45	Wyoming
## 46	Alaska
## 47	Maine
## 48	Alabama
## 49	Idaho
## 50	Montana
## 50	Puerto Rico
## 51	Virgin Islands
	Virgin Islands Guam
## 53	Guam

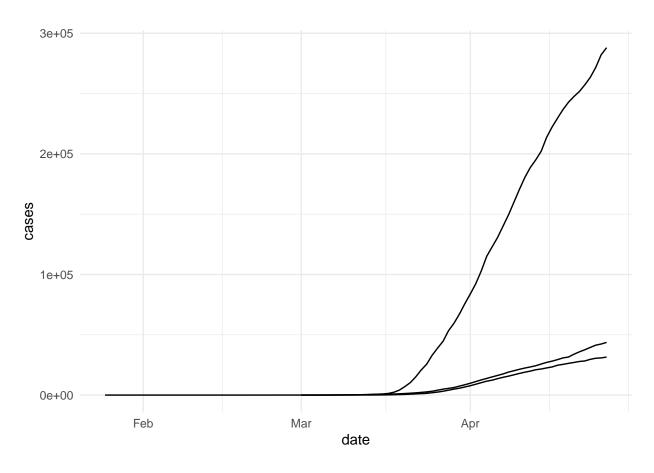
```
## 54 West Virginia
## 55 Northern Mariana Islands
```

```
california_df <- covid_df[ which( covid_df$state == "California"), ]
ny_df <- covid_df[ which( covid_df$state == "New York"), ]
florida_df <- covid_df[ which( covid_df$state == "Florida"), ]

## Plot the number of cases in Florida using `geom_line()`
##group=1 to plot all the data points as a single group
ggplot(data=florida_df, aes(x=date, y=cases, group=1)) + geom_line()</pre>
```



```
## Add lines for New York and California to the plot
ggplot(data=florida_df, aes(x=date, group=1)) +
  geom_line(aes(y = cases)) +
  geom_line(data=ny_df, aes(y = cases)) +
  geom_line(data=california_df, aes(y = cases))
```



```
## Use the colors "darkred", "darkgreen", and "steelblue" for Florida, New York, and California
ggplot(data=florida_df, aes(x=date, group=1)) +
  geom_line(aes(y = cases), color = "darkred") +
  geom_line(data=ny_df, aes(y = cases), color="darkgreen") +
  geom_line(data=california_df, aes(y = cases), color= "steelblue")
```

