

Homework 3

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This homework is due on Jan. 31, 2023 at 11:00pm. Please submit as a pdf file on Canvas.

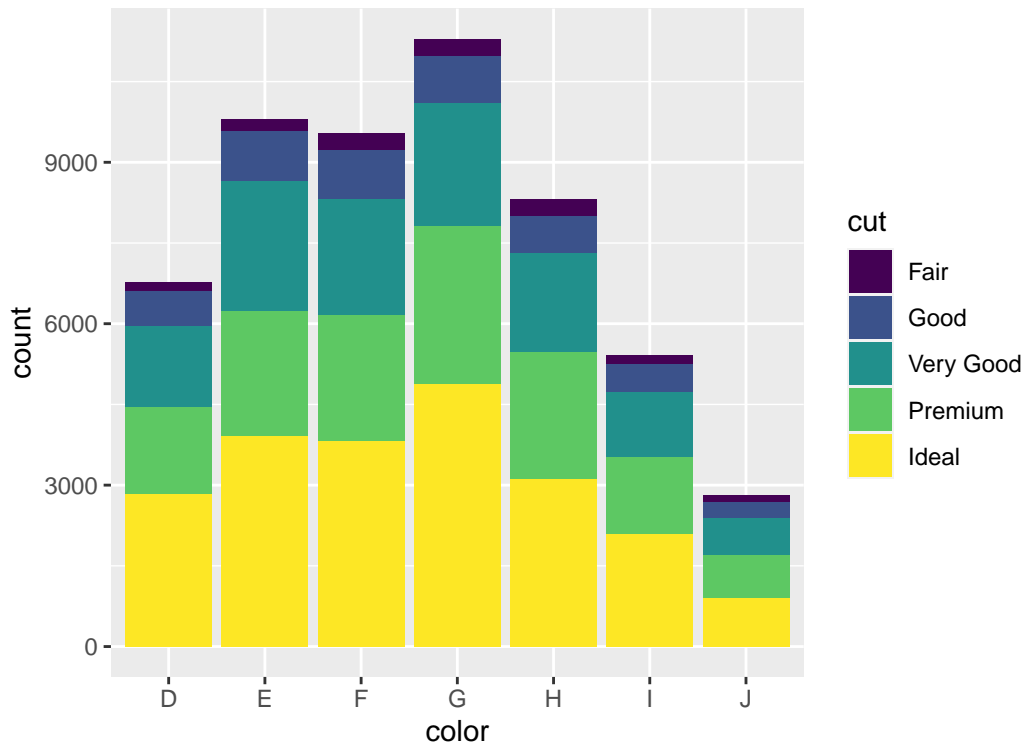
Problem 1: (4 pts) For problem 1, we will work with the `diamonds` dataset. See here for details: <https://ggplot2.tidyverse.org/reference/diamonds.html>.

`diamonds`

```
## # A tibble: 53,940 x 10
##   carat cut      color clarity depth table price      x      y      z
##   <dbl> <ord>    <ord> <ord>    <dbl> <dbl> <int> <dbl> <dbl> <dbl>
## 1  0.23 Ideal    E      SI2     61.5    55   326   3.95   3.98   2.43
## 2  0.21 Premium E      SI1     59.8    61   326   3.89   3.84   2.31
## 3  0.23 Good     E      VS1     56.9    65   327   4.05   4.07   2.31
## 4  0.29 Premium I      VS2     62.4    58   334   4.2    4.23   2.63
## 5  0.31 Good     J      SI2     63.3    58   335   4.34   4.35   2.75
## 6  0.24 Very Good J      VVS2     62.8    57   336   3.94   3.96   2.48
## 7  0.24 Very Good I      VVS1     62.3    57   336   3.95   3.98   2.47
## 8  0.26 Very Good H      SI1     61.9    55   337   4.07   4.11   2.53
## 9  0.22 Fair     E      VS2     65.1    61   337   3.87   3.78   2.49
## 10 0.23 Very Good H      VS1     59.4    61   338   4      4.05   2.39
## # ... with 53,930 more rows
```

- Use `ggplot` to make a bar plot of the total diamond count per `color` and show the proportion of each `cut` within each `color` category.
- In two sentences, explain when to use `geom_bar()` instead of `geom_col()`. Which of these functions requires only an `x` or `y` variable?

```
ggplot(diamonds, aes(color, fill = cut)) + geom_bar()
```



One would use `geom_bar()` instead of `geom_col()` when graphing bar plots that rely on count with bar heights proportional to the number of cases in each group. The `geom_bar()` function requires only an `x` or `y` variable as it only needs one, while `geom_col()` needs both.

Problem 2: (4 pts) For problem 2 and 3, we will work with the dataset `OH_pop` that contains Ohio state demographics and has been derived from the `midwest` dataset provided by `ggplot2`. See here for details of the original dataset: <https://ggplot2.tidyverse.org/reference/midwest.html>. `OH_pop` contains two columns: `county` and `poptotal` (the county's total population), and it only contains counties with at least 100,000 inhabitants.

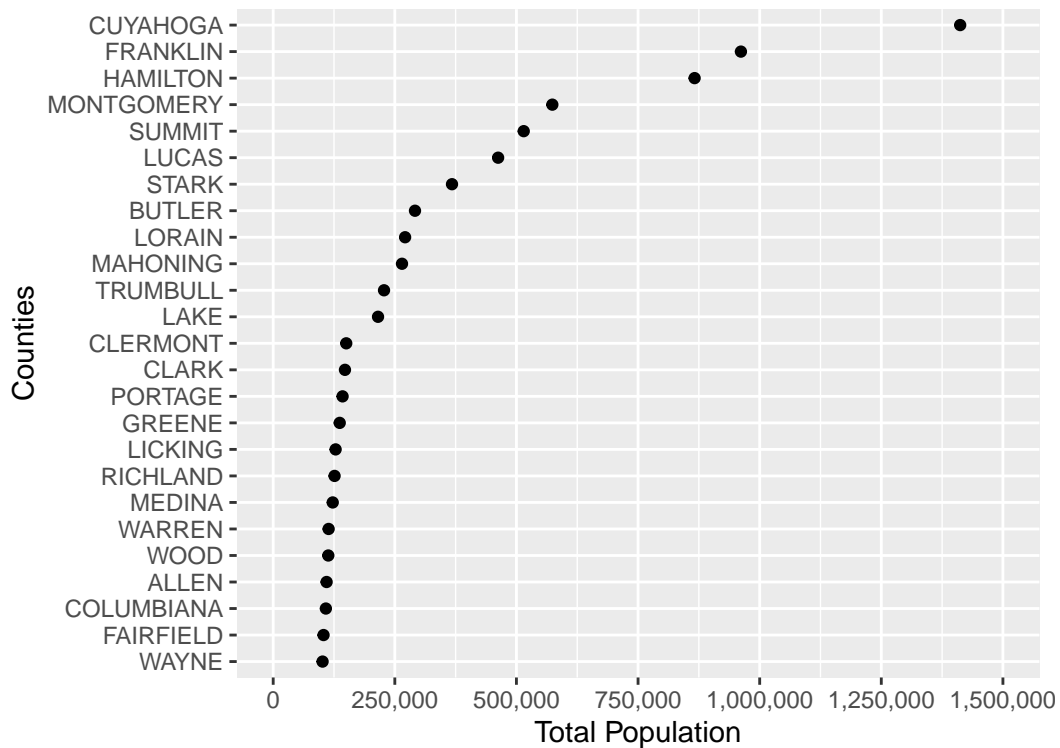
`OH_pop`

```
## # A tibble: 25 x 2
##   county    poptotal
##   <chr>      <int>
## 1 CUYAHOGA  1412140
## 2 FRANKLIN   961437
## 3 HAMILTON   866228
## 4 MONTGOMERY 573809
## 5 SUMMIT     514990
## 6 LUCAS      462361
## 7 STARK      367585
## 8 BUTLER     291479
## 9 LORAIN     271126
## 10 MAHONING  264806
## # ... with 15 more rows
```

- Use `ggplot` to make a scatter plot of `county` vs total population (column `poptotal`) and order the counties by increasing population.
- Rename the axes and set appropriate limits, breaks and labels. Note: Do not use `xlab()` or `ylab()` to

label the axes.

```
ggplot(OH_pop, aes(poptotal, fct_reorder(county, poptotal))) +
  geom_point() +
  scale_x_continuous(name = "Total Population",
    limits = c(0, 1500000),
    breaks = seq(0, 1500000, 250000),
    labels = c("0", "250,000", "500,000", "750,000", "1,000,000", "1,250,000",
      "1,500,000")) +
  scale_y_discrete(name = "Counties")
```



Problem 3: (2 pts)

(a) Modify the plot from Problem 2 by changing the scale for `poptotal` to logarithmic.

(b) Adjust the limits, breaks and labels for the logarithmic scale.

```
ggplot(OH_pop, aes(poptotal, fct_reorder(county, poptotal))) +
  geom_point() +
  scale_x_log10(name = "Total Population",
    limits = c(100000, 1500000),
    breaks = c(100000, 250000, 500000, 1000000, 1500000),
    labels = c("100,000", "250,000", "500,000", "1,000,000", "1,500,000")) +
  scale_y_discrete(name = "Counties")
```

