

Homework 3

Enter your name and EID here

Connor Hanna cdh3663

This homework is due on Feb. 7, 2022 at 11:00am. Please submit as a pdf file on Canvas.

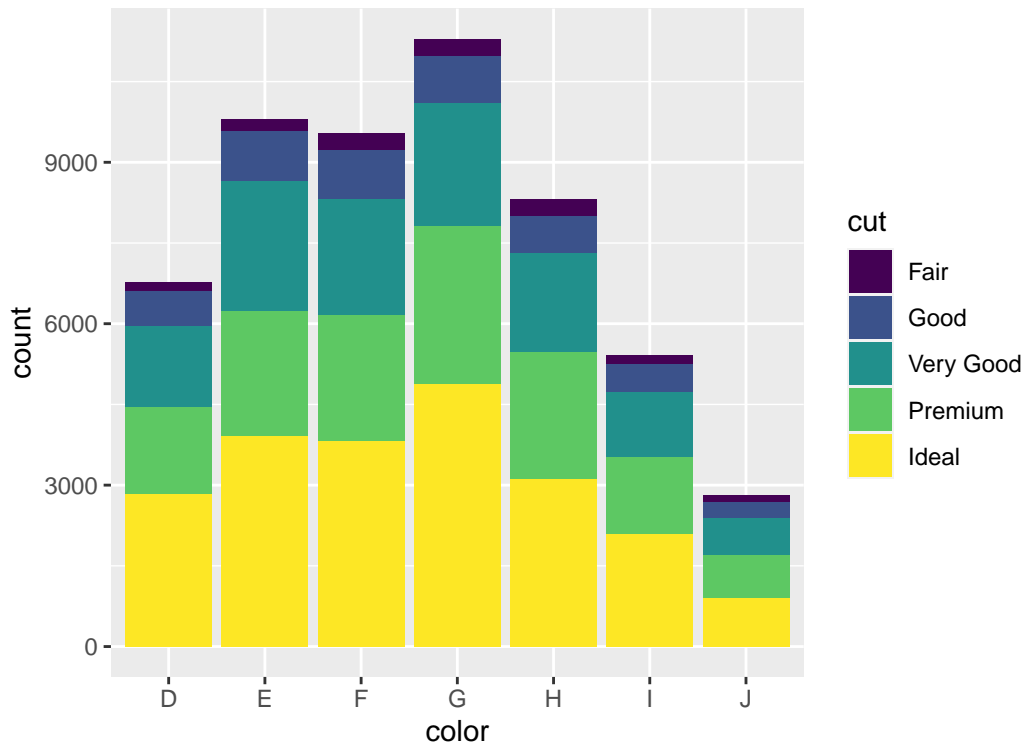
Problem 1: (2 pts) For problem 1, we will work with the `diamonds` dataset. See here for details: <https://ggplot2.tidyverse.org/reference/diamonds.html>.

```
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
```

- (a) 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.
- (b) In two sentences, explain when to use `geom_bar()` instead of `geom_col()`. Which of these functions requires only an `x` or `y` variable?

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



(b) `geom_col()` can be used to make bar graphs where observations contain values that need to be attributed to both an x and y axis, especially when each observation requires a separate column as in the `txhouse` data from HW 2. `geom_bar()` is used when counting observations containing certain values for a specified x or y variable.

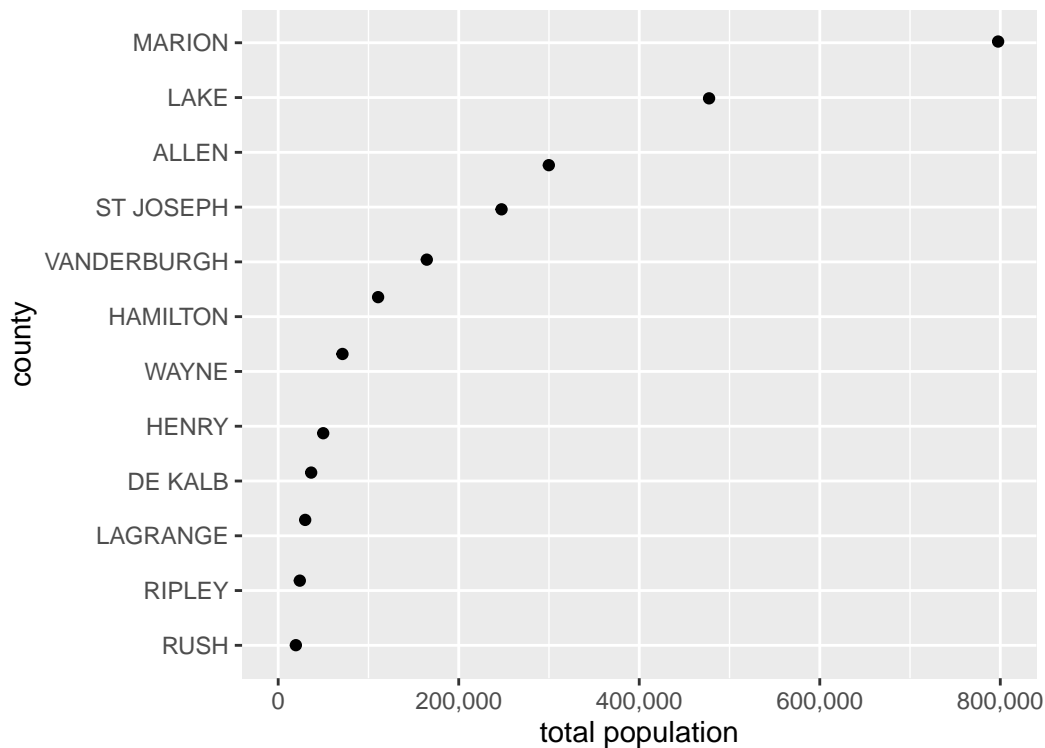
Problem 2: (4 pts) For problems 2 and 3, we will work with the dataset `IL_pop` that contains Illinois state demographics and has been derived from the `midwest` dataset provided by `ggplot2`. See here for details of the original dataset: <https://ggplot2.tidyverse.org/reference/midwest.html>. `IL_pop` contains two columns: `county` and `poptotal` (the county's total population).

`IL_pop`

```
## # A tibble: 12 x 2
##   county      poptotal
##   <chr>      <int>
## 1 MARION      797159
## 2 LAKE        475594
## 3 ALLEN       300836
## 4 ST JOSEPH   247052
## 5 VANDERBURGH 165058
## 6 HAMILTON    108936
## 7 WAYNE       71951
## 8 HENRY       48139
## 9 DE KALB     35324
## 10 LAGRANGE    29477
## 11 RIPLEY      24616
## 12 RUSH        18129
```

- 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(IL_pop, aes(poptotal, fct_reorder(county, poptotal))) +
  geom_jitter() +
  scale_x_continuous(
    name = "total population",
    limits = c(0, 800000),
    breaks = c(0, 200000, 400000, 600000, 800000),
    labels = c("0", "200,000", "400,000", "600,000", "800,000")
  ) +
  scale_y_discrete(
    name = "county"
  )
)
```



Problem 3: (4 pts)

- Modify the plot from Problem 2 by changing the scale for `poptotal` to logarithmic.
- Adjust the limits, breaks and labels for the logarithmic scale.

```
ggplot(IL_pop, aes(poptotal, fct_reorder(county, poptotal))) +
  geom_jitter() +
  scale_x_log10(
    name = "total population",
    limits = c(1e4, 1e6),

```

```

breaks = c(1e4, 3.16e4, 1e5, 3.16e5, 1e6),
labels = c("10,000", "31,600", "100,000", "316,000", "1,000,000")
) +
scale_y_discrete(
  name = "county"
)

```

