in_class_week2_sols.R

moffer

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```
library(tidyverse)
## -- Attaching packages -----
                                              ----- tidyverse 1.3.1 --
## v ggplot2 3.3.6
                     v purrr
                               0.3.4
## v tibble 3.1.7
                     v dplyr
                               1.0.9
## v tidyr
            1.2.0
                     v stringr 1.4.0
## v readr
            2.1.2
                     v forcats 0.5.1
## -- Conflicts -----
                                            ## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                   masks stats::lag()
# download the data from this website: https://data.cdc.gov/NCHS/Provisional-COVID-19-Deaths-by-Race-an
# and read it in locally, using `read_csv()`
dat <- read_csv('.../data/Provisional_COVID-19_Deaths_by_Race_and_Hispanic_Origin__and_Age.csv')</pre>
## Rows: 6489 Columns: 13
## -- Column specification -------
## Delimiter: ","
## chr (7): Data as of, Start Date, End Date, State, Age group, Race and Hispan...
## dbl (6): COVID-19 Deaths, Total Deaths, Pneumonia Deaths, Pneumonia and COVI...
## 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.
# check out the data using `dat`, `summary(dat)`, and `View(dat)`.
dat
## # A tibble: 6,489 x 13
##
     `Data as of` `Start Date` `End Date` State
                                                    'Age group' 'Race and Hisp~'
##
                                         <chr>>
     <chr>
                  <chr>
                              <chr>
                                                    <chr>
                                                                <chr>
                 01/01/2020 10/01/2022 United Sta~ All Ages
## 1 10/05/2022
                                                                Total Deaths
## 2 10/05/2022
                 01/01/2020 10/01/2022 United Sta~ All Ages
                                                                Non-Hispanic Wh~
## 3 10/05/2022
                 01/01/2020 10/01/2022 United Sta~ Under 1 ye~ Non-Hispanic Wh~
## 4 10/05/2022
                 01/01/2020
                              10/01/2022 United Sta~ 0-17 years
                                                                Non-Hispanic Wh~
                              10/01/2022 United Sta~ 1-4 years
## 5 10/05/2022
                 01/01/2020
                                                                Non-Hispanic Wh~
## 6 10/05/2022
                 01/01/2020
                              10/01/2022 United Sta~ 5-14 years Non-Hispanic Wh~
## 7 10/05/2022
                 01/01/2020
                              10/01/2022 United Sta~ 15-24 years Non-Hispanic Wh~
## 8 10/05/2022
                 01/01/2020
                              10/01/2022 United Sta~ 18-29 years Non-Hispanic Wh~
## 9 10/05/2022
                 01/01/2020
                              10/01/2022 United Sta~ 25-34 years Non-Hispanic Wh~
## 10 10/05/2022
                  01/01/2020
                              10/01/2022 United Sta~ 30-49 years Non-Hispanic Wh~
## # ... with 6,479 more rows, and 7 more variables: `COVID-19 Deaths` <dbl>,
     `Total Deaths` <dbl>, `Pneumonia Deaths` <dbl>,
     `Pneumonia and COVID-19 Deaths` <dbl>, `Influenza Deaths` <dbl>,
```

`Pneumonia, Influenza, or COVID-19 Deaths` <dbl>, Footnote <chr> summary(dat) End Date ## Data as of Start Date State ## Length:6489 Length:6489 Length:6489 Length:6489 ## Class : character Class : character Class : character Class : character Mode : character Mode : character Mode :character Mode : character ## ## ## ## ## Age group Race and Hispanic Origin Group COVID-19 Deaths Length:6489 ## Length:6489 Min. Class : character Class : character 1st Qu.: 0 ## Mode :character Mode :character 12 Median: ## Mean : 1058 ## 3rd Qu.: 97 ## Max. :1055967 ## :1995 NA's Total Deaths Pneumonia Deaths Pneumonia and COVID-19 Deaths ## ## Min. : 0 Min. : 0 Min. : 0 ## 1st Qu.: 24 1st Qu.: 0 1st Qu.: ## Median : 106 Median : Median : 11 ## Mean 8034 Mean : 985 Mean 517 ## 3rd Qu.: 770 42 3rd Qu.: 88 3rd Qu.: ## Max. :9225883 Max. :969256 Max. :536751 ## NA's :1269 NA's :2068 NA's :1783 ## Influenza Deaths Pneumonia, Influenza, or COVID-19 Deaths Footnote ## Min. : 0.00 Min. : 0.0 Length:6489 0.00 1st Qu.: 0.0 Class :character ## 1st Qu.: 0.00 ## Median: Median : 18.0 Mode :character 10.79 1518.4 ## Mean Mean ## 3rd Qu.: 0.00 3rd Qu.: 137.8 ## Max. :12381.00 Max. :1499035.0 ## NA's :1419 NA's :2059 # View(dat) # filter the data so that you're just using data for the United states as a whole, # and drop observations where the column `Race and Hispanic Origin Group` is 'Total Deaths' dat <- dat %>% filter(State == 'United States', `Race and Hispanic Origin Group` != 'Total Deaths') # Consider the age groups in this data. Do you want to keep all of them? # Filter out any age groups you don't want to keep. table(dat\$`Age group`) ## ## 0-17 years 1-4 years 15-24 years 18-29 years ## 8 ## 30-49 years 25-34 years 35-44 years 45-54 years ##

55-64 years

65-74 years

8

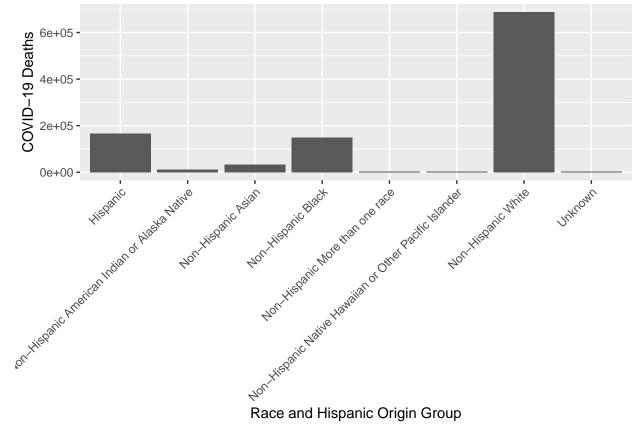
50-64 years

8

5-14 years

##

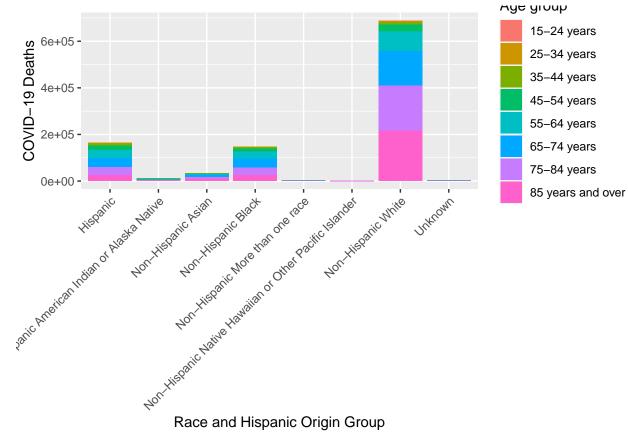
```
##
##
         75-84 years 85 years and over
                                                All Ages
                                                               Under 1 year
##
dat <- dat %>%
  filter(`Age group` %in% c('15-24 years', '25-34 years', '35-44 years',
                            '45-54 years', '55-64 years', '65-74 years',
                            '75-84 years', '85 years and over', 'All Ages'))
# create a data set, `dat0`, that is just the data where the column `Age group`
# is 'All Ages'
dat0 <- dat %>%
  filter(`Age group` == 'All Ages')
# create another data set, `dat1`, that is the date where the column `Age group`
# is anything but 'All Ages'
dat1 <- dat %>%
  filter(`Age group` != 'All Ages')
# in dat0, plot COVID-19 deaths by `Race and Hispanic Origin Group` using `geom_col`.
# put the x axis labels at 45 degrees (you can google how to do this)
ggplot(dat0, aes(y = `COVID-19 Deaths`,
                x = `Race and Hispanic Origin Group`)) +
  geom_col() +
  theme(axis.text.x = element_text(angle = 45, hjust=1))
```



Race and Hispanic Origin Group

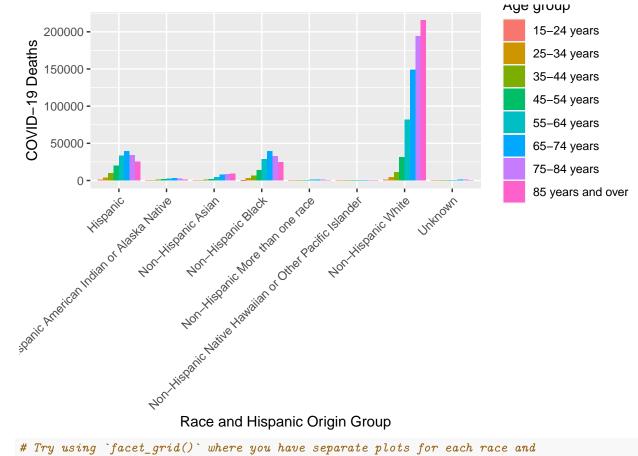
in dat1, plot COVID-19 deaths by `Race and Hispanic Origin Group` using `geom_col`, # AND set the fill by `Age group`

```
ggplot(dat1, aes(y = `COVID-19 Deaths`,
                x = `Race and Hispanic Origin Group`,
               fill = `Age group`)) +
  geom_col() +
 theme(axis.text.x = element_text(angle = 45, hjust=1))
```

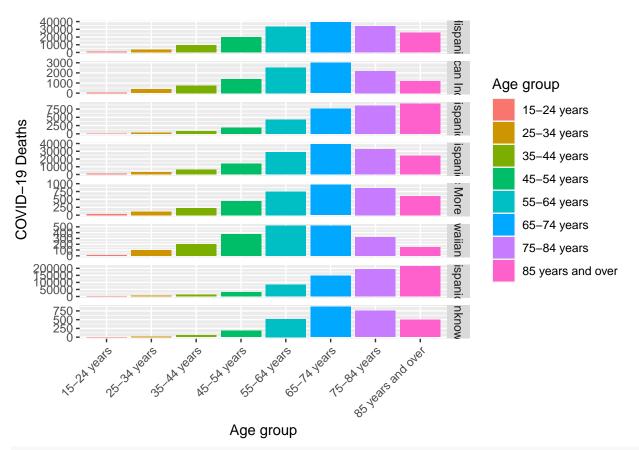


Race and Hispanic Origin Group

```
# Do the last plot again, but try using the argument `position = 'dodge'` in
# `geom_col()`. Which do you like better? Why?
ggplot(dat1, aes(y = `COVID-19 Deaths`,
                 x = `Race and Hispanic Origin Group`,
                fill = `Age group`)) +
  geom_col(position = 'dodge') +
  theme(axis.text.x = element_text(angle = 45, hjust=1))
```



```
# Try using `facet_grid()` where you have separate plots for each race and
# hispanic origin group, with COVID deaths on the Y axis, and age on the X axis
ggplot(dat1, aes(y = `COVID-19 Deaths`,
                 x = `Age group`,
                 fill = `Age group`)) +
  facet_grid(vars(`Race and Hispanic Origin Group`), scales = 'free') +
  geom_col(position = 'identity') +
  theme(axis.text.x = element_text(angle = 45, hjust=1))
```



- $\hbox{\it\#Which plot do you think is the best way to summarize this data? Why?}$
- # What other data would you need to say something about the relative risk for # by age and race/hispanic origin?
- # Use ggsave() to save your best plot, and email it to mollyow@uchicago.edu # with this script, and the names of students in your group.