Homework 2

Lupe Antonio

10/2/2023

Problem 1

```
#loading pols-month data
pols_month <- read_csv("data_hw2/fivethirtyeight_datasets/pols-month.csv")</pre>
```

```
#cleaning the data
pols_month <- janitor::clean_names(pols_month) %>%
  #breaking up variable mon into integer variables: year, month, & day
  separate(mon, into = c("year", "month", "day"), convert = TRUE) %>%
  mutate(
    #replacing month num with month name
   month = case_match(
     month,
     1 ~ "january",
      2 ~ "february",
      3 ~ "march",
      4 ~ "april",
      5 ~ "may",
      6 ~ "june",
      7 ~ "july",
     8 ~ "august",
     9 ~ "september",
     10 ~ "october",
      11 ~ "november",
      12 ~ "december"),
    #creating president variable
   president = case_when(
      prez_dem == 1 & prez_gop == 0 ~ "democratic",
      prez_dem == 0 & prez_gop == 1 ~ "republican")) %>%
  #removing prez_dem, prez_gop, and day variables
  select(-prez_dem, -prez_gop, -day)
```

```
#loading snp.csv data
snp <- read_csv("data_hw2/fivethirtyeight_datasets/snp.csv") %>%
  #cleaning data
  janitor::clean_names() %>%
  #fixing the dates
  mutate(date = lubridate::mdy(date)) %>%
  #breaking up date variable into integer variables: year, month, day
  separate(date, into = c("year", "month", "day"), convert = TRUE) %>%
```

```
mutate(
  #fixing the year problem
 year = ifelse(year > 2023, year - 100, year),
  #replacing month num with month name
 month = case_match(
   month,
   1 ~ "january",
   2 ~ "february",
   3 ~ "march",
   4 ~ "april",
   5 ~ "may",
   6 ~ "june",
   7 ~ "july",
   8 ~ "august",
   9 ~ "september",
   10 ~ "october",
   11 ~ "november",
    12 ~ "december")) %>%
#removing day variable
select(-day)
```

```
#loading unemployment.csv data
unemployment <- read_csv("data_hw2/fivethirtyeight_datasets/unemployment.csv") %>%
  #cleaning the data
  janitor::clean_names() %>%
  #going from "wide" to "long" format
  pivot_longer(jan:dec,
              names_to = "month",
               values_to = "unemp_perctg") %>%
  #fixing month names
  mutate(
   month = case_match(
      month,
      "jan" ~ "january",
     "feb" ~ "february",
      "mar" ~ "march",
      "apr" ~ "april",
      "may" ~ "may",
      "jun" ~ "june",
     "jul" ~ "july",
     "aug" ~ "august",
      "sep" ~ "september",
      "oct" ~ "october",
      "nov" ~ "november",
      "dec" ~ "december"
```

```
#merging snp into pols
final_data <- left_join(pols_month, snp)</pre>
```

Joining with 'by = join_by(year, month)'

```
final_data <- left_join(final_data, unemployment)
## Joining with 'by = join_by(year, month)'</pre>
```

The pols_month data has a total of 822 observations and 9 variables. This dataset contains information regarding the party affiliations of governors and senators, and the current president (at that time) between 1947 to 2015. The snp data has a total of 787 observations and 3 variables. It contains information from the years 1950 to 2015. Additionally, the unemployment dataset has a total of 816 observations and 3 variables. It contains information from the years 1948 to 2015.

Problem 2

```
#loading Mr. Trash Wheel data
mr_trash_wheel <- read_excel("data_hw2/Trash_Wheel_Collection_Data.xlsx", sheet = "Mr. Trash Wheel") %>
  #cleaning data
  janitor::clean_names() %>%
  #removing x15 & x16 columns
  select(-x15, -x16) %>%
  #creating new homes_powered variable
  mutate(
   homes_powered = (weight_tons * 500)/30) %>%
  #adding trash wheel variable
  mutate(trash_wheel = "Mr. Trash Wheel") %>%
  #filter NA's in dumpster
  filter(!is.na(dumpster)) %>%
  #making year numeric
  mutate(year = as.numeric(year))
#loading Professor Trash Wheel data
professor_trash_wheel <- read_excel("data_hw2/Trash_Wheel_Collection_Data.xlsx", sheet = "Professor Tra
  #cleaning data
  janitor::clean_names() %>%
  #adding trash wheel variable
  mutate(trash_wheel = "Professor Trash Wheel") %>%
  #filter NA's in dumpster
  filter(!is.na(dumpster)) %>%
  #making year numeric
  mutate(year = as.numeric(year))
#loading Gwynnda data
gwynnda_trash_wheel <- read_excel("data_hw2/Trash_Wheel_Collection_Data.xlsx", sheet = "Gwynnda Trash W.
  #cleaning data
  janitor::clean_names() %>%
  #adding trash wheel variable
  mutate(trash_wheel = "Gwynnda Trash Wheel") %>%
  #filter NA's in dumpster
  filter(!is.na(dumpster)) %>%
  #making year numeric
```

mutate(year = as.numeric(year))

```
#combining datasets
trash_wheels_tidy <- bind_rows(mr_trash_wheel, professor_trash_wheel, gwynnda_trash_wheel)</pre>
```

The final tidy dataset contains information from mr_trash_wheel, professor_trash_wheel, and gwynnda_trash_wheel. It has a total of 845 observations and 15 variables. This dataset contains information regarding the amount of trash per different trash type between 2014 and 2023 for each type of trash wheel.

From the available data, the total weight (in tons) of trash collected by Professor Trash Wheel was 216.26. The total weight (in tons) of trash collected by Mr. Trash Wheel was 1875.1. Similarly, the total weight (in tons) of trash collected by Gwynnda Trash Wheel was 451.65. Additionally, the total number of cigarette butts collected by Gwynnda in July of 2021 was 1.63×10^4 .

Problem 3

```
#loading the baseline dataset
baseline_demos <- read_csv("data_hw2/data_mci/MCI_baseline.csv", skip = 1) %>%
    #cleaning dataset
    janitor::clean_names() %>%
    #renaming the column names
    rename(
        'baseline_age' = 'current_age',
        'education_years' = 'education',
        'apoe4_carrier' = 'apoe4',
        'onset_age' = 'age_at_onset')
```

```
#continuation of tidying data
baseline_demos <- baseline_demos%>%
  mutate(
    #changing sex values
    sex = case when(
      sex == 0 ~ "female",
      sex == 1 ~ "male"),
    #changing APOE4 carrier values
    apoe4_carrier = case_when(
      apoe4_carrier == 0 ~ "non-carrier",
      apoe4_carrier == 1 ~ "carrier"),
    #adding NA's in blank spaces
    onset_age = case_when(
      onset_age == "." ~ NA,
      TRUE ~ as.numeric(onset_age)
   )) %>%
  #filtering no MCI
  filter(onset_age > baseline_age | is.na(onset_age))
```

```
## Warning: There was 1 warning in 'mutate()'.
## i In argument: 'onset_age = case_when(onset_age == "." ~ NA, TRUE ~
## as.numeric(onset_age))'.
## Caused by warning:
## ! NAs introduced by coercion
```

The baseline_demos dataset was imported and tidyied up by renaming column names when necessary, mutating the sex variable values into female or male, and similarly the apoe4_carrier into carrier or non-carrier. Additionally, NA's were needed to onset_age variable, and we filtered for participants without MCI at baseline. The resulting dataset contains a total of 479 observations and 6 variables. There was a total of 483 participants recruited and of those 479 developed MCI. The average baseline age was 65.0286013. The proportion of women in the study that were APOE4 carriers were $1.3883 \times 10^4 / 479$.

```
## Rows: 487 Columns: 6
## -- Column specification ------
## Delimiter: ","
## chr (5): Baseline, Time 2, Time 4, Time 6, Time 8
## dbl (1): Study ID
##
## 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.
```

The amyloid_dataset dataset was imported and tidyled up by renaming column names when necessary, and transforming the dataset from 'wide' format to 'long' format. The resulting dataset contains a total of 2435 observations and 3 variables.

```
#observations in the baseline_demos dataset, but not in amyloid_dataset
anti_join(baseline_demos, amyloid_dataset, by = 'id')
```

```
## # A tibble: 8 x 6
##
        id baseline_age sex
                                 education_years apoe4_carrier onset_age
##
                   <dbl> <chr>
                                           <dbl> <chr>
                                                                     <dbl>
     <dbl>
## 1
                    58.4 female
                                               20 non-carrier
                                                                      66.2
        14
                    64.7 male
                                               16 non-carrier
                                                                      68.4
## 2
        49
## 3
        92
                    68.6 female
                                               20 non-carrier
                                                                      NA
## 4
       179
                    68.1 male
                                               16 non-carrier
                                                                      NA
## 5
       268
                    61.4 female
                                                                      67.5
                                               18 carrier
                    63.8 female
## 6
       304
                                               16 non-carrier
                                                                      NA
                    59.3 female
                                               16 non-carrier
## 7
       389
                                                                      NA
## 8
       412
                    67
                         male
                                               16 carrier
                                                                      NA
```

Of the participants that were in the baseline_demos dataset, most were non-carriers of APOE4, and most of their onset age is missing.

```
#observations in amyloid_dataset, but not in baseline_demos dataset
anti_join(amyloid_dataset, baseline_demos, by = 'id')
```

```
## # A tibble: 80 x 3
##
          id time_elapsed biomarkers_ratio
       <dbl> <chr>
##
                           <chr>
##
          72 baseline
                           0.106965463
    1
##
    2
          72 time 2
                           <NA>
                           0.107266218
##
    3
          72 \text{ time } 4
##
          72 time 6
                           0.106665207
         72 time 8
                           <NA>
##
    5
##
    6
        234 baseline
                           0.110521689
##
    7
        234 time_2
                           0.110988335
        234 \ time_4
                           0.110318671
        234 \text{ time}_6
                           0.107334344
##
    9
        234 \ time_8
                           0.108868811
## 10
## # i 70 more rows
```

Based on these findings, we can see that the participants that were not in the baseline_demoshas simialr biomarker ratios.

```
#dataset with common observations
common_amyloid_dataset <- inner_join(baseline_demos, amyloid_dataset, by = "id")
#export csv
write_csv(common_amyloid_dataset, "data_hw2/common_amyloid_dataset.csv")</pre>
```

After combining both datasets, the common_amyloid_dataset is produced. This dataset contains at total of 2355 observations and 8 variables. This dataset contains information of participants' biomarkers ratio obtained at different times/stages in the study, and shows whether the participant is a APOE4 carrier or not, the participants' sex, ages at baseline and onset, and their years of education.