gss_cat.R

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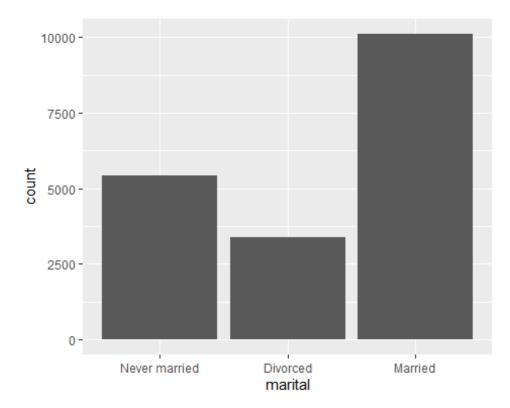
2025-04-07

```
library(tidyverse)
## Warning: package 'tidyverse' was built under R version 4.4.3
## Warning: package 'ggplot2' was built under R version 4.4.3
## Warning: package 'tidyr' was built under R version 4.4.3
## Warning: package 'readr' was built under R version 4.4.3
## Warning: package 'purrr' was built under R version 4.4.3
## Warning: package 'dplyr' was built under R version 4.4.3
## Warning: package 'forcats' was built under R version 4.4.3
## Warning: package 'lubridate' was built under R version 4.4.3
## — Attaching core tidyverse packages —
                                                                  - tidyverse
2.0.0 -
## √ dplyr
               1.1.4
                         √ readr
                                      2.1.5
## √ forcats
               1.0.0

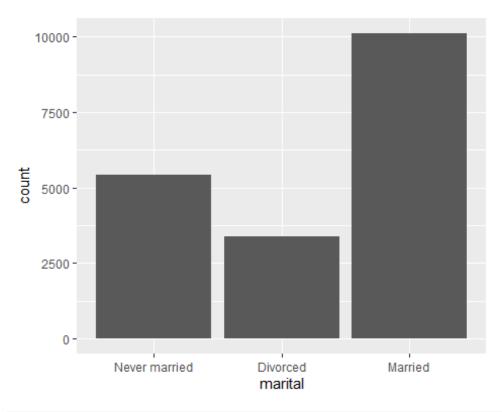
√ stringr

                                      1.5.1
## √ ggplot2 3.5.1
                         √ tibble
                                      3.2.1
## √ lubridate 1.9.4
                         √ tidyr
                                      1.3.1
## √ purrr
               1.0.4
## — Conflicts —
tidyverse_conflicts() —
## X dplyr::filter() masks stats::filter()
## X dplyr::lag() masks stats::lag()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all
conflicts to become errors
library(forcats)
library(ggplot2)
View(gss_cat)
#marital
my_data <- gss_cat %>%
  select(marital) %>%
  filter(marital %in% c("Married","Never married","Divorced")) %>%
  mutate(marital = fct drop(marital))
my table <- table(my data)</pre>
```

```
View(my_table)
#use chi -squared goodness of fit test
chisq.test(my_table)
##
##
   Chi-squared test for given probabilities
##
## data: my_table
## X-squared = 3784.1, df = 2, p-value < 2.2e-16
my_data%>%
  table()%>%
  chisq.test()
##
   Chi-squared test for given probabilities
##
##
## data: .
## X-squared = 3784.1, df = 2, p-value < 2.2e-16
#graphics
my_data%>%
 ggplot(aes(marital))+geom_bar()
```



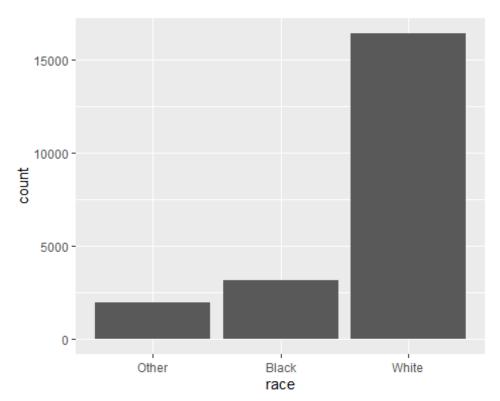
```
#use chi -squared goodness of Race
my_data <- gss_cat %>%
  select(marital) %>%
  filter(marital %in% c("Married","Never married","Divorced")) %>%
  mutate(marital = fct_drop(marital))
my_table <- table(my_data)</pre>
View(my_table)
#use chi -squared goodness of fit test
chisq.test(my_table)
##
## Chi-squared test for given probabilities
##
## data: my_table
## X-squared = 3784.1, df = 2, p-value < 2.2e-16
my_data%>%
 table()%>%
  chisq.test()
##
## Chi-squared test for given probabilities
##
## data:
## X-squared = 3784.1, df = 2, p-value < 2.2e-16
#graphics
my_data%>%
ggplot(aes(marital))+geom_bar()
```



```
#for race
my_data <- gss_cat %>%
  select(race) %>%
  filter(race %in% c("White", "Black", "Other")) %>%
  mutate(race = fct_drop(race))
my_table <- table(my_data)</pre>
View(my_table)
#use chi -squared goodness of fit test
chisq.test(my_table)
##
  Chi-squared test for given probabilities
##
##
## data: my_table
## X-squared = 17956, df = 2, p-value < 2.2e-16
my_data%>%
  table()%>%
  chisq.test()
##
## Chi-squared test for given probabilities
```

```
##
## data: .
## X-squared = 17956, df = 2, p-value < 2.2e-16
#graphics

my_data%>%
    ggplot(aes(race))+geom_bar()
```



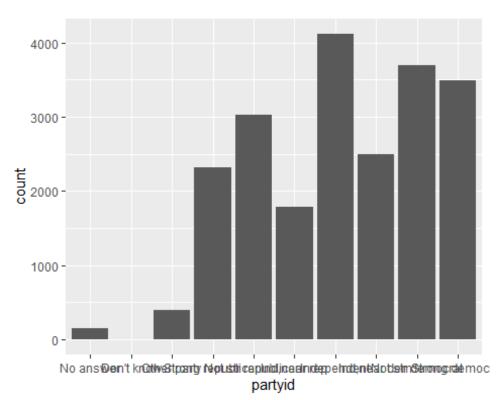
```
#use chi -squared goodness of Race
my_data <- gss_cat %>%
    select(race) %>%
    filter(race %in% unique(race)) %>%
    mutate(race = fct_drop(race))

my_table <- table(my_data)
View(my_table)

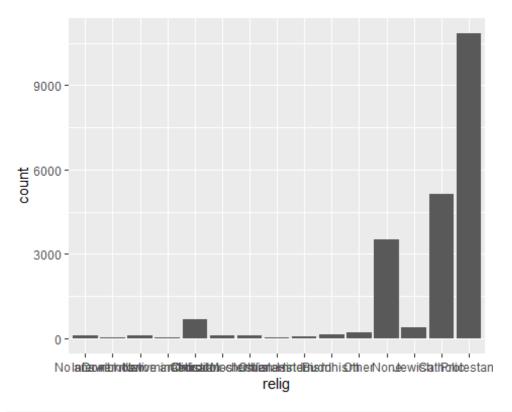
#for partyid

my_data <- gss_cat %>%
    select(partyid) %>%
    filter(partyid %in% unique(partyid)) %>%
    mutate(partyid = fct_drop(partyid))
my_table <- table(my_data)
```

```
View(my_table)
#use chi -squared goodness of fit test
chisq.test(my_table)
##
##
   Chi-squared test for given probabilities
##
## data: my_table
## X-squared = 9676.9, df = 9, p-value < 2.2e-16
my_data%>%
  table()%>%
  chisq.test()
##
   Chi-squared test for given probabilities
##
##
## data:
## X-squared = 9676.9, df = 9, p-value < 2.2e-16
#graphics
my_data%>%
 ggplot(aes(partyid))+geom_bar()
```

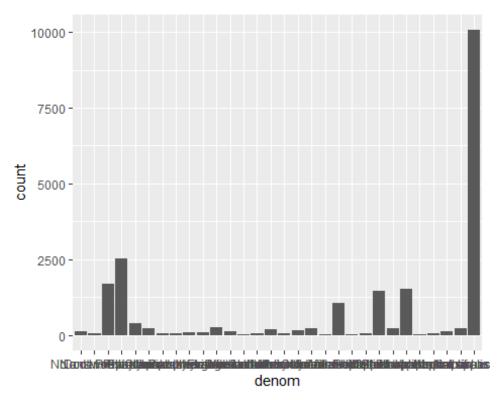


```
#use chi -squared goodness of partyid
my_data <- gss_cat %>%
  select(partyid) %>%
  filter(partyid %in% unique(partyid)) %>%
  mutate(partyid = fct_drop(partyid))
my_table <- table(my_data)</pre>
View(my_table)
#Relig
my_data <- gss_cat %>%
  select(relig) %>%
  filter(relig %in% unique(relig)) %>%
  mutate(relig = fct_drop(relig))
my_table <- table(my_data)</pre>
View(my_table)
#use chi -squared goodness of fit test
chisq.test(my_table)
##
## Chi-squared test for given probabilities
##
## data: my table
## X-squared = 88171, df = 14, p-value < 2.2e-16
my_data%>%
 table()%>%
  chisq.test()
##
## Chi-squared test for given probabilities
##
## data:
## X-squared = 88171, df = 14, p-value < 2.2e-16
#graphics
my_data%>%
ggplot(aes(relig))+geom_bar()
```



```
#use chi -squared goodness of Religion
my_data <- gss_cat %>%
  select(relig) %>%
  filter(relig %in% unique(relig)) %>%
  mutate(relig = fct_drop(relig))
my_table <- table(my_data)</pre>
View(my_table)
#denom
my_data <- gss_cat %>%
  select(denom) %>%
  filter(denom %in% unique(denom)) %>%
  mutate(denom = fct_drop(denom))
my_table <- table(my_data)</pre>
View(my_table)
#use chi -squared goodness of fit test
chisq.test(my_table)
```

```
##
    Chi-squared test for given probabilities
##
##
## data: my_table
## X-squared = 141814, df = 29, p-value < 2.2e-16
my_data%>%
  table()%>%
  chisq.test()
##
##
   Chi-squared test for given probabilities
##
## data:
## X-squared = 141814, df = 29, p-value < 2.2e-16
#graphics
my_data%>%
 ggplot(aes(denom))+geom_bar()
```



```
#use chi -squared goodness of denom
my_data <- gss_cat %>%
   select(denom) %>%
   filter(denom %in% unique(denom)) %>%
   mutate(denom = fct_drop(denom))
```

```
my_table <- table(my_data)
View(my_table)</pre>
```