

GSS R: installation and first exploration

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```
if (!require(gssr)) {  
  if (!require(remotes)){  
    install.packages("remotes")  
  }  
  remotes::install_github("kjhealy/gssr")  
}
```

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! Objectives

Install and use package **gssr**

Get data for year 2018

The GSS is carried out every two years. It offers both *cross-sectional* data and *panel* data.

Package **gssr** offers a simple way to retrieve yearly data.

PhantomJS not found. You can install it with `webshot::install_phantomjs()`

gssr

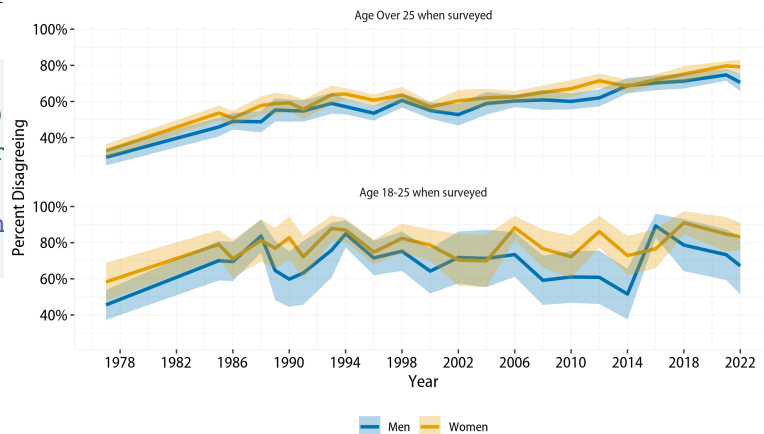


The General Social Survey Cumulative Data (1972-2022, release 2a) and Panel Data files packaged for easy use in R. The companion package to `gssr` (<https://github.com/kjhealy/gssr>) is `gssrdoc` (<https://kjhealy.github.io/gssrdoc>), which integrates the GSS codebook into R's help system. I recommend you install both packages.

We work again with General Social Survey (GSS) data. We take advantage of R package `gssr`

```
if (!require(gssr)) {  
  if (!require(remotes))  
    install.packages("remotes")  
  remotes::install_github("kjhealy/gssr")  
}
```

Disagreement with the statement, 'It is much better for everyone involved if the man is the achiever outside the home and the woman takes care of the home and family'



Kieran Healy <http://socviz.co>
Data source: General Social Survey

```
df_2018 <- gssr::gss_get_yr(2018)
```

Fetching: https://gss.norc.umd.edu/documents/stata/2018_stata.zip

Inspect the data

- How many observations?
- How many variables?
- Are the data tidy/messy?

```
dim(df_2018)
```

```
[1] 2348 1069
```

Numerical summaries for `age` and `agekdbrn`

The 2018 data provide (among too many other things) columns named `age` and `agekdbrn`. Get numerical summaries about these two columns.

```
df_2018 |>  
  dplyr::select(age, agekdbrn) |>  
  skimr::skim() |>  
  skimr::yank("numeric")
```

Variable type: numeric

skim_variable	n_missing	complete_rate	mean	sd	p0	p25	p50	p75	p100	hist
age	7	1.00	48.97	18.06	18	34	48	63	89	
agekdbrn	682	0.71	24.30	5.74	12	20	23	28	51	

Thanks to `gssr`, you can get meta-information about the columns

```
?aged
?agekdbnrn
?sex
```

How is `sex` encoded? Is it worth recoding it?

```
df_2018 |>
  mutate(sex=as_factor(sex)) |>
  skimr::skim(sex) |>
  skimr::yank("factor")
```

Variable type: factor

skim_variable	n_missing	complete_rate	ordered	n_unique	top_counts
sex	0	1	FALSE	2	fem: 1296, mal: 1052, don: 0, iap: 0

Histogram and density plots for age distribution/facet by sex

```
p_age <- df_2018 |>
  mutate(sex=as_factor(sex)) |>
  ggplot() +
  aes(x=age) +
  facet_wrap(~ sex, )
```

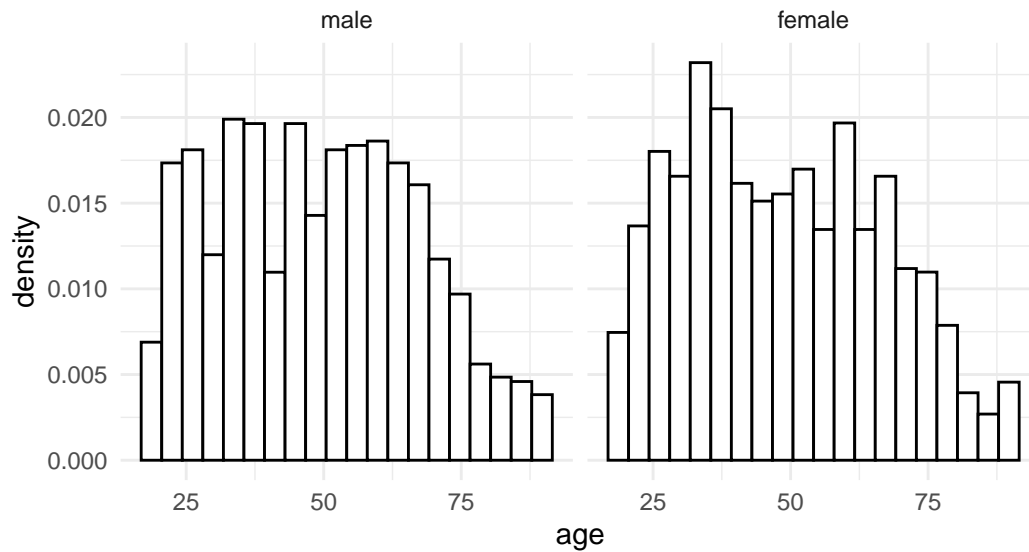
```
q_age <- df_2018 |>
  mutate(sex=as_factor(sex)) |>
  ggplot() +
  aes(x=age)
```

```
p_age +
  geom_histogram(aes(y=after_stat(density)),
                 fill="white",
                 color="black",
                 bins=20) +
  labs(
    title="GSS 2018",
    subtitle = "Age distribution of respondents"
  )
```

Warning: Removed 7 rows containing non-finite outside the scale range (``stat_bin()``).

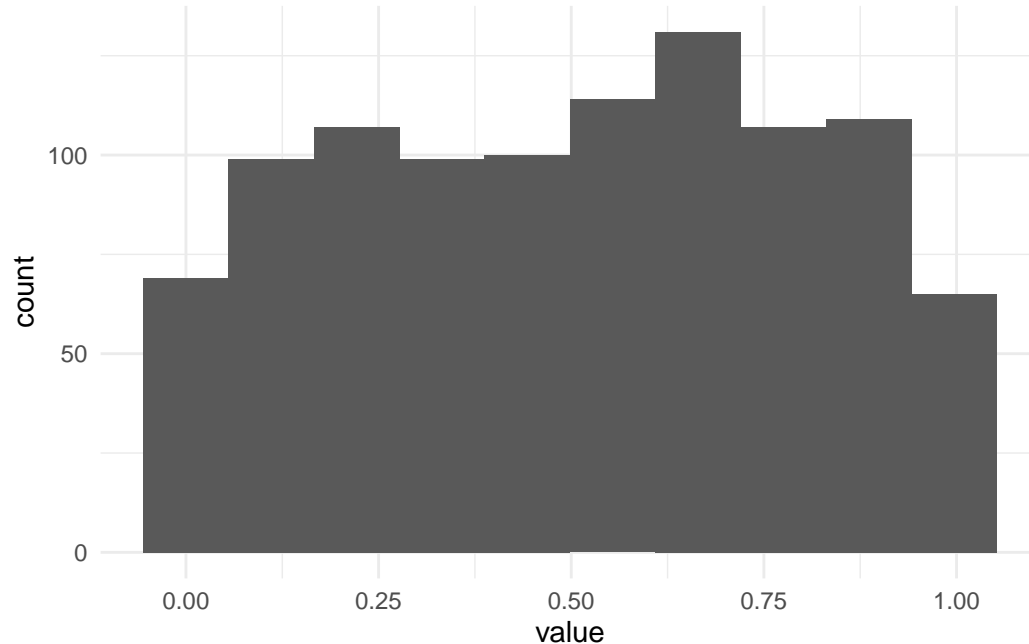
GSS 2018

Age distribution of respondents



- Play with number of bins
- Spot the irregular behavior of the histograms
- Something special at the right edge of both histograms

```
enframe(x=runif(1000)) |>
  ggplot() +
  aes(x=value) +
  geom_histogram(bins=10)
```



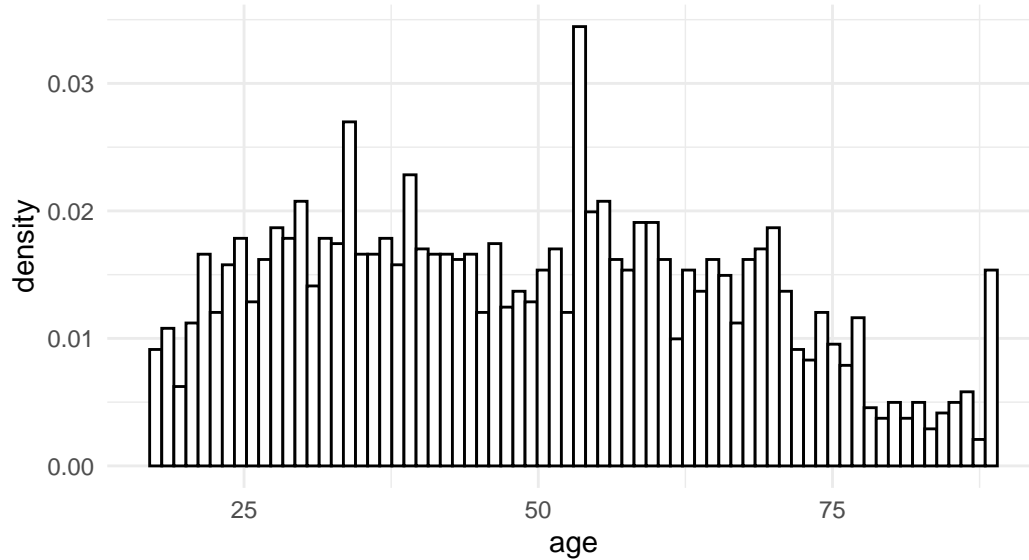
```
q_age +
  geom_histogram(aes(y=after_stat(density)),
    fill="white",
    color="black",
    bins=70) +
  labs(
    title="GSS 2018",
    subtitle = "Age distribution of respondents"
```

)

Warning: Removed 7 rows containing non-finite outside the scale range (`stat_bin()`).

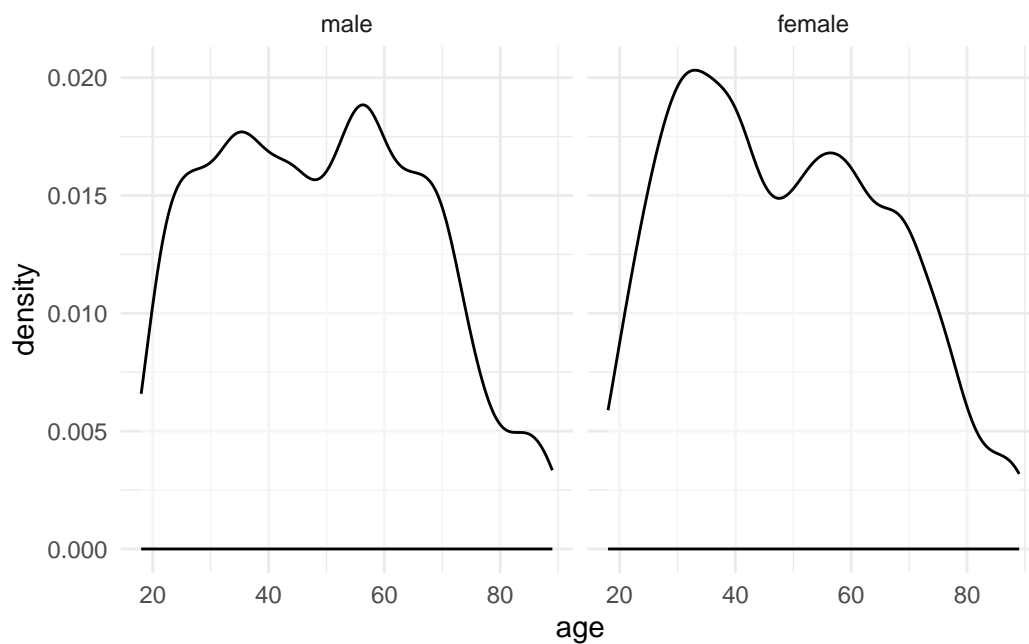
GSS 2018

Age distribution of respondents



```
p_age +
  stat_density(aes(y=after_stat(density)),
    fill="white",
    alpha=.5,
    color="black",
    bw = "sj",
    adjust = 1
  )
```

Warning: Removed 7 rows containing non-finite outside the scale range (`stat_density()`).

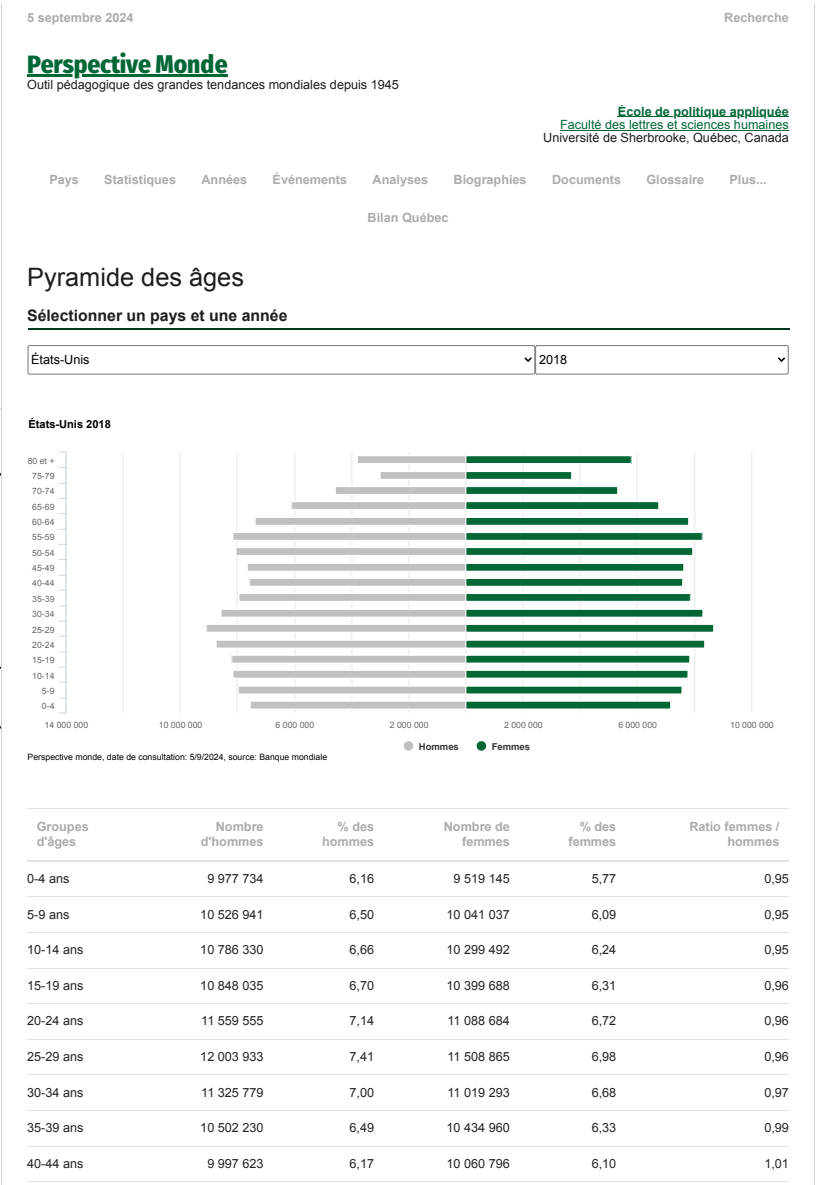


- Play with arguments `bw` and `adjust` of `stat_density`
- Same comments

Compare *sample* age distribution with *population* age distribution

```
knitr::include_url("https://perspective.usherbrooke.ca/bilan/servlet/BMPagePyramide/USA/2018")
```

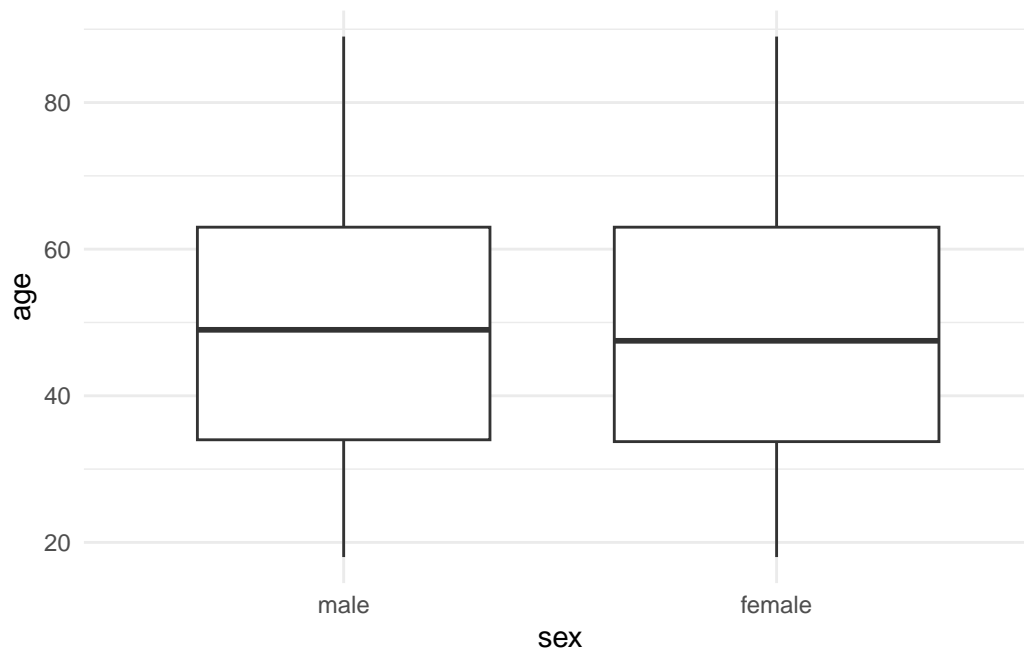
Sherbrooke University offers visual information about the age structure of population of a wide range of countries. Following demographic usage, the age structure is presented through an age pyramid. Note that an age pyramid is a special kind of histogram



Parallel boxplots of age with respect to sex

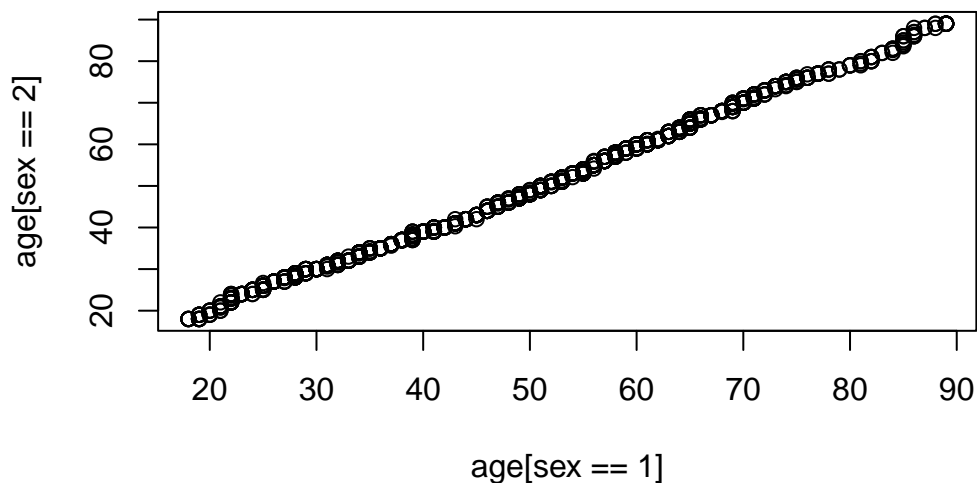
```
df_2018 |>
  mutate(sex=as_factor(sex)) |>
  ggplot() +
  aes(y=age, x=sex) +
  geom_boxplot(varwidth = T) +
  xlab("sex")
```

Warning: Removed 7 rows containing non-finite outside the scale range (``stat_boxplot()``).



QQplot comparing sample male and female age distributions

```
with(df_2018,
      qqplot(x=age[sex==1], y=age[sex==2]))
```



Make your own qqplot

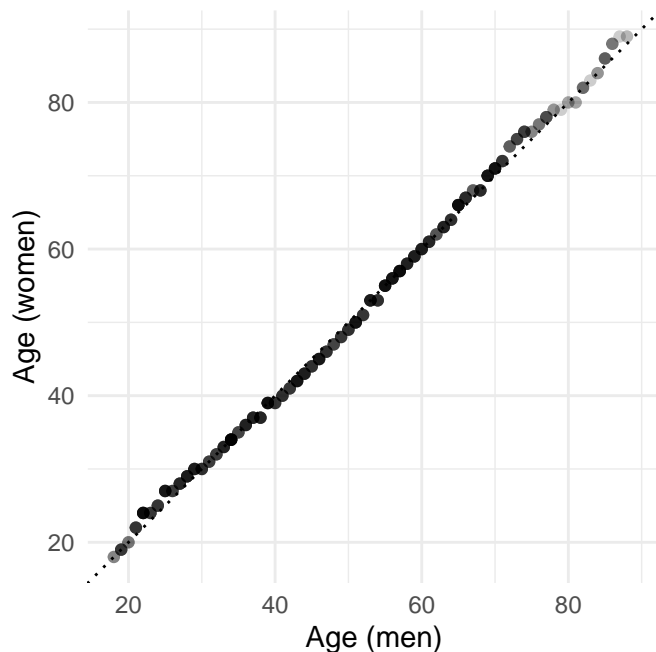
```
cdf_age_2018_1 <- ecdf(df_2018$age[df_2018$sex==1])

tb <- df_2018 |>
  dplyr::filter(sex==2) |>
  dplyr::select(age) |>
  mutate(Fn=rank(age, ties.method = "max")/n()) |>
  distinct() |>
  arrange(age)

eqf_age_2018_2 <- with(tb,
  stepfun(x=Fn, y=c(age, max(age)), right = T, f = 1))
```

```
filter(df_2018, sex==1) |>
  ggplot() +
  aes(x=age, y=eqf_age_2018_2(cdf_age_2018_1(age))) +
  geom_point(alpha=.1, fill="white") +
  geom_abline(intercept = 0, slope=1, linetype="dotted") +
  coord_fixed() +
  xlab("Age (men)") +
  ylab("Age (women)")
```

Warning: Removed 15 rows containing missing values or values outside the scale range (`geom_point()`).



```
# data(gss_all)
```

```
data(gss_dict)
```

```
gss_dict |>
  filter(variable=="age")
```

```
# A tibble: 1 x 13
  pos variable label      missing var_doc_label value_labels var_text years
<int> <chr>    <chr>        <int> <chr>          <chr>      <chr>  <list>
1    90 age      age of re~    769 age of respo~ [89] 89 or ~ 13. Res~ <tibble>
# i 5 more variables: var_yrtab <list>, var_ballots <list>, col_type <chr>,
#   var_type <chr>, var_na_codes <chr>
```

```
# gss_which_years(gss_all, c("age", "agekdbrn"))
```

Scatterplot for age and agekdbrn, facet by sex ‘

Working with gss_sub

```
data("gss_sub")
```

```
gss_sub |>
  glimpse()
```


Rows: 72,390

Columns: 20

```

$ year      <dbl+lbl> 1972, 1972, 1972, 1972, 1972, 1972, 1972, 1972, 1972, 197~
$ id        <dbl> 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18~
$ ballot    <dbl+lbl> NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), N~
$ age       <dbl+lbl> 23, 70, 48, 27, 61, 26, 28, 27, 21, 30, 30, 56, 54, 49, 4~
$ race      <dbl+lbl> 1, 1, 1, 1, 1, 1, 1, 1, 1, 2, 2, 2, 2, 2, 2, 2, 1, 1, 1, ~
$ sex       <dbl+lbl> 2, 1, 2, 2, 2, 1, 1, 1, 2, 2, 2, 1, 1, 2, 1, 1, 1, 2, 2, ~
$ degree    <dbl+lbl> 3, 0, 1, 3, 1, 1, 1, 3, 1, 1, 1, 0, 0, 0, 0, 1, 1, 0, 3, ~
$ padeg     <dbl+lbl> 0, 0, 0, 3, 0, 3, 3, 3, ~
$ madeg     <dbl+lbl> NA(i), 0, 0, 1, 0, 4, 1, 1, ~
$ relig     <dbl+lbl> 3, 2, 1, 5, 1, 1, 2, 3, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, ~
$ polviews  <dbl+lbl> NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), N~
$ fefam     <dbl+lbl> NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), N~
$ vpsu      <dbl+lbl> NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), N~
$ vstrat    <dbl+lbl> NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), N~
$ oversamp  <dbl+lbl> 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, ~
$ formwt    <dbl+lbl> NA(y), NA(y), NA(y), NA(y), NA(y), NA(y), NA(y), NA(y), N~
$ wtssall   <dbl+lbl> 0.4446, 0.8893, 0.8893, 0.8893, 0.8893, 0.4446, 0.4446, 0~
$ wtssps    <dbl+lbl> 0.6631963, 0.9173700, 0.8974125, 1.0663408, 0.9443237, 0~
$ sampcode  <dbl+lbl> NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), NA(i), N~
$ sample    <dbl+lbl> 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, ~

```

```

gss_sub |>
  head()

```

A tibble: 6 x 20

```

  year      id ballot      age  race  sex  degree padeg  madeg
  <dbl+lbl> <dbl> <dbl+lbl> <dbl+> <dbl+lbl> <dbl+lbl> <dbl+lbl> <dbl+lbl> <dbl+lbl>
1 1972      1 NA(i) [iap] 23    1 [whi~ 2 [fem~ 3 [bac~ 0 [les~ NA(i) [iap]
2 1972      2 NA(i) [iap] 70    1 [whi~ 1 [mal~ 0 [les~ 0 [les~ 0 [les~
3 1972      3 NA(i) [iap] 48    1 [whi~ 2 [fem~ 1 [hig~ 0 [les~ 0 [les~
4 1972      4 NA(i) [iap] 27    1 [whi~ 2 [fem~ 3 [bac~ 3 [bac~ 1 [hig~
5 1972      5 NA(i) [iap] 61    1 [whi~ 2 [fem~ 1 [hig~ 0 [les~ 0 [les~
6 1972      6 NA(i) [iap] 26    1 [whi~ 1 [mal~ 1 [hig~ 3 [bac~ 4 [gra~
# i 11 more variables: relig <dbl+lbl>, polviews <dbl+lbl>, fefam <dbl+lbl>,
#   vpsu <dbl+lbl>, vstrat <dbl+lbl>, oversamp <dbl+lbl>, formwt <dbl+lbl>,
#   wtssall <dbl+lbl>, wtssps <dbl+lbl>, sampcode <dbl+lbl>, sample <dbl+lbl>

```

```

gss_sub |>
  dplyr::select(-id, -year) |>
  summarise(across(everything(), n_distinct)) |>
  pivot_longer(cols = everything(), names_to="name_col", values_to = "n_distct") |>
  arrange(n_distct) |>
  filter(n_distct < 15) |>
  left_join(gss_dict, by=c("name_col"="variable"))

```

A tibble: 11 x 14

```

  name_col n_distct  pos label      missing var_doc_label value_labels var_text
  <chr>      <int> <int> <chr>      <int> <chr>      <chr>      <chr>
1 sex          3    125 responde~    112 respondents ~ [1] male; [~ 23. Cod~
2 race          4    126 race of ~    107 race of resp~ [1] white; ~ 24. Wha~
3 ballot        5   6072 ballot u~   21875 ballot used ~ [1] ballot ~ 1659. B~
4 fefam          5    784 better f~   37259 better for m~ [1] strongl~ 252. No~

```

```
5 oversamp      5  6078 weights ~      0 weights for ~ [1] not 198~ None
6 degree        6    98 r's high~    196 r's highest ~ [0] less th~ 19. If ~
7 padeg         6    99 father's~ 17881 father's hig~ [0] less th~ 20. If ~
8 madeg         6   100 mothers ~   8971 mothers high~ [0] less th~ 21. If ~
9 polviews      8   227 think of~   9672 think of sel~ [1] extreme~ 67a. We~
10 sample       11  6077 sampling~  4032 sampling fra~ [1] 1960 sa~ 1664. T~
11 relig        14   336 r's reli~   437 r's religiou~ [1] protest~ 104. Wh~
# i 6 more variables: years <list>, var_yrtab <list>, var_ballots <list>,
#   col_type <chr>, var_type <chr>, var_na_codes <chr>
```

Education through generations

What kind of information do we get through variables `degree` and `padeg`?

```
?degree
?padeg
```

Compute contingency table for `degree` and `padeg`

```
tab_degree_padeg <- gss_sub |>
  dplyr::select(degree, padeg) |>
  mutate(across(everything(), as_factor)) |>
  table()
```

```
tab_degree_padeg |>
  chisq.test()
```

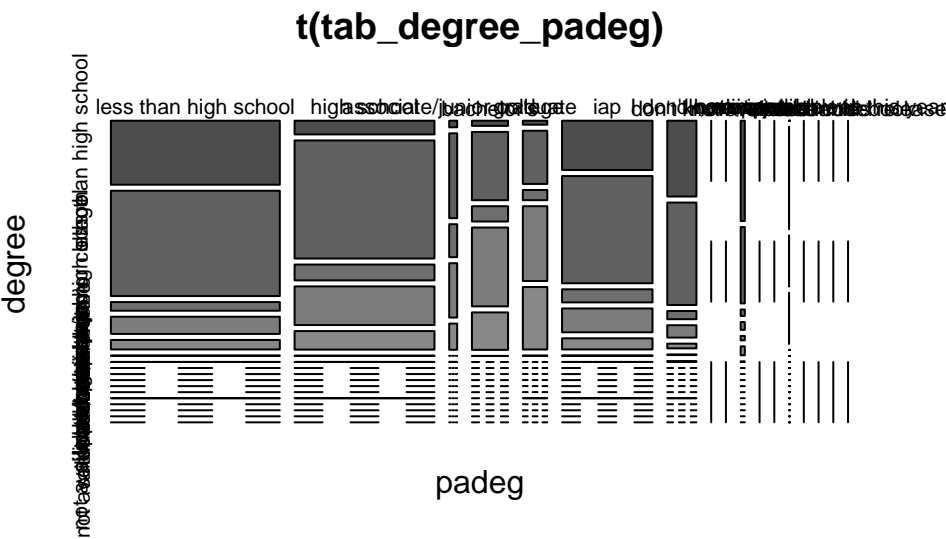
Warning in `chisq.test(tab_degree_padeg)`: Chi-squared approximation may be incorrect

Pearson's Chi-squared test

```
data:  tab_degree_padeg
X-squared = NaN, df = 256, p-value = NA
```

Visualize contingency table for `degree` and `padeg`

```
tab_degree_padeg |>
  t() |>
  mosaicplot(color = T)
```



Rearrange the levels of degree and pdeg