HW3 Problem 1

2025-06-26

R Markdown

This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word documents. For more details on using R Markdown see http://rmarkdown.rstudio.com.

When you click the **Knit** button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can embed an R code chunk like this:

```
library(gssr)
## Package loaded. To attach the GSS data, type data(gss_all) at the console.
## For the panel data and documentation, type e.g. data(gss_panel08_long) and data(gss_panel_doc).
## For help on a specific GSS variable, type ?varname at the console.
library(tidyverse)
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplvr
           1.1.4 v readr
                                   2.1.5
## v forcats 1.0.0
                       v stringr 1.5.1
                     v tibble
## v ggplot2 3.5.2
                                   3.2.1
## v lubridate 1.9.4
                       v tidyr
                                   1.3.1
## v 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 error
library(dplyr)
data(gss_all)
gss_2018 <- gss_get_yr(2018)
## Fetching: https://gss.norc.org/documents/stata/2018_stata.zip
dim(gss_2018)
## [1] 2348 1083
```

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##	[937]	"away11"	"where11"	"relate12"	"gender12"
##	[941]	"old12"	"mar12"	"relhhd1"	"relhhd2"
##	[945]	"relhhd3"	"relhhd4"	"relhhd5"	"relhhd6"
##	[949]	"relhhd7"	"relhhd8"	"relhhd9"	"relhhd10"
##	[953]	"relhhd11"	"relhhd12"	"hefinfo"	"hhrace"
##	[957]	"respnum"	"hhtype"	"hhtype1"	"famgen"
##	[961]	"rplace"	"rvisitor"	"visitors"	"relhh1"
##	[965]	"relhh2"	"relhh3"	"relhh4"	"relhh5"
##	[969]	"relhh6"	"relhh7"	"relhh8"	"relhh9"
##	[973]	"relhh10"	"relhh11"	"relhh12"	"relsp1"
##	[977]	"relsp2"	"relsp3"	"relsp4"	"relsp5"
##	[981]	"relsp6"	"relsp7"	"relsp8"	"relsp9"
##	[985]	"relsp10"	"relsp11"	"relsp12"	"dateintv"
##	[989]	"isco88"	"paisco88"	"maisco88"	"spisco88"
##	[993]	"isco08"	"paisco08"	"maisco08"	"spisco08"
##	[997]	"coisco08"	"sei10"	"sei10educ"	"sei10inc"
##	[1001]	"pasei10"	"pasei10educ"	"pasei10inc"	"masei10"
##	[1005]	"masei10educ"	"masei10inc"	"spsei10"	"spsei10educ"
##	[1009]	"spsei10inc"	"cosei10"	"cosei10educ"	"cosei10inc"
##	[1013]	"copres10"	"copres105plus"	"uswary"	"cohort"
##	[1017]	"zodiac"	"inthisp"	"intrace1"	"intrace2"
##	[1021]	"intrace3"	"whoelse1"	"whoelse2"	"whoelse3"
##	[1025]	"whoelse4"	"whoelse5"	"whoelse6"	"intid"
##	[1029]	"feeused"	"feelevel"	"lngthinv"	"intage"
##	[1033]	"intethn"	"mode"	"intsex"	"intyrs"
##	[1037]	"consent"	"adminconsent"	"ballot"	"version"
##	[1041]	"issp"	"formwt"	"sampcode"	"sample"
##	[1045]	"oversamp"	"phase"	"spanself"	"spanint"
##	[1049]	"spaneng"	"hlthstrt"	"huadd"	"huaddwhy"
##	[1053]	"dwellpre"	"kidsinhh"	"respond"	"incuspop"
##	[1057]	"neisafe"	"rlooks"	"rgroomed"	"rweight"
##	[1061]	"rhlthend"	"wtss"	"wtssnr"	"wtssall"
##	[1065]	"vstrat"	"vpsu"	"wtssps"	"wtssnrps"

```
## [1073] "cowrkslffam"
                            "cowksup"
                                             "subsamprate"
                                                              "hispanic_0022"
                                             "sibs 7222"
## [1077] "numemps 0422"
                            "uscitzn_0822"
                                                              "racecen1_0022"
## [1081] "racecen2_0022" "racecen3_0022" "earnrs_7222"
head(gss_2018)
## # A tibble: 6 x 1,083
                                               hrs2
                                                                        wrkslf wrkgovt
     year
                   id wrkstat
                                  hrs1
                                                            evwork
                                                                        <dbl+l> <dbl+l>
     <dbl+1bl> <dbl> <dbl+1bl> <dbl+1bl>
                                               <dbl+lbl>
                                                            <dbl+lbl>
                    1 3 [with a~ NA(i) [iap]
                                                            NA(i) [iap] 2 [som~ 2 [pri~
## 1 2018
                                                  41
                                                                1 [yes] 2 [som~ 2 [pri~
## 2 2018
                    2 5 [retire~ NA(i) [iap] NA(i) [iap]
## 3 2018
                    3 1 [workin~
                                     40
                                              NA(i) [iap] NA(i) [iap] 2 [som~ 2 [pri~
                                               NA(i) [iap] NA(i) [iap] 2 [som~ 2 [pri~
## 4 2018
                    4 1 [workin~
                                     40
## 5 2018
                    5 5 [retire~ NA(i) [iap] NA(i) [iap]
                                                                1 [yes] 2 [som~ 2 [pri~
                    6 5 [retire~ NA(i) [iap] NA(i) [iap]
                                                                1 [yes] 2 [som~ 2 [pri~
## 6 2018
## # i 1,075 more variables: occ10 <dbl+lbl>, prestg10 <dbl+lbl>,
       prestg105plus <dbl+lbl>, indus10 <dbl+lbl>, marital <dbl+lbl>,
       martype <dbl+lbl>, divorce <dbl+lbl>, widowed <dbl+lbl>,
       spwrksta <dbl+lbl>, sphrs1 <dbl+lbl>, sphrs2 <dbl+lbl>, spevwork <dbl+lbl>,
       cowrksta <dbl+lbl>, cowrkslf <dbl+lbl>, coevwork <dbl+lbl>,
## #
       cohrs1 <dbl+lbl>, cohrs2 <dbl+lbl>, spwrkslf <dbl+lbl>, spocc10 <dbl+lbl>,
## #
       sppres10 <dbl+lbl>, sppres105plus <dbl+lbl>, spind10 <dbl+lbl>, ...
view(gss_2018)
gss_subset <- gss_2018 %>%
  select(bible, trustsci) %>%
  na.omit()
gss_subset <- gss_2018 %>%
  select(bible, trustsci) %>%
  filter(!is.na(bible), !is.na(trustsci)) %>%
  filter(trustsci %in% c(1, 2, 3)) %>%
  mutate(
    bible_binary = ifelse(bible == 1, 1, 0)
str(gss_subset)
## tibble [767 x 3] (S3: tbl_df/tbl/data.frame)
                   : \  \, \mathtt{dbl+lbl} \  \, [1:767] \  \, 1, \  \, 2, \  \, 1, \  \, 1, \  \, 1, \  \, 1, \  \, 1, \  \, 1, \  \, 2, \  \, 1, \  \, 2, \  \, 2, \  \, 2, \  \, 1, \  \, 2, \  \, 1, \  \, 2, \  \, \ldots
##
    $ bible
##
                       : chr "Feelings about the bible"
      ..@ format.stata: chr "%29.0g"
##
##
                      : Named num [1:16] 1 2 3 4 NA NA NA NA NA NA ...
      ... - attr(*, "names")= chr [1:16] "word of god" "inspired word" "ancient book" "other (vol)" ...
##
                   : dbl+lbl [1:767] 1, 3, 3, 3, 1, 2, 2, 3, 2, 2, 3, 2, 3, 2, 2, 3, 3, 2, ...
    $ trustsci
                      : chr "We trust too much in science"
##
      ..@ label
##
      ..@ format.stata: chr "%29.0g"
##
                     : Named num [1:17] 1 2 3 4 5 NA NA NA NA NA ...
      ... - attr(*, "names")= chr [1:17] "strongly agree" "agree" "neither agree nor disagree" "disagr
## $ bible_binary: num [1:767] 1 0 1 1 1 1 1 0 1 ...
```

"spwrkslffam"

"cowrkslf2"

[1069] "fileversion"

"spwrkslf2"

```
head(gss_subset)
## # A tibble: 6 x 3
##
   bible
                       trustsci
                                                       bible_binary
     <dbl+lbl>
                      <dbl+1b1>
                                                              <dbl>
## 1 1 [word of god] 1 [strongly agree]
                                                                  1
## 2 2 [inspired word] 3 [neither agree nor disagree]
                                                                  0
                       3 [neither agree nor disagree]
## 3 1 [word of god]
                                                                  1
## 4 1 [word of god]
                       3 [neither agree nor disagree]
## 5 1 [word of god] 1 [strongly agree]
                                                                  1
## 6 1 [word of god] 2 [agree]
                                                                  1
summary(gss_subset)
##
        bible
                       trustsci
                                     bible_binary
          :1.000 Min.
## Min.
                           :1.000
                                           :0.0000
                                    Min.
## 1st Qu.:1.000 1st Qu.:2.000
                                    1st Qu.:0.0000
## Median :2.000 Median :2.000
                                    Median :0.0000
## Mean :1.692 Mean :2.296
                                    Mean :0.4237
## 3rd Qu.:2.000
                    3rd Qu.:3.000
                                    3rd Qu.:1.0000
## Max. :4.000 Max. :3.000
                                    Max. :1.0000
mean_trust <- gss_subset %>%
  group_by(bible_binary) %>%
  summarise(mean_trustsci = mean(trustsci, na.rm = TRUE))
print(mean_trust)
## # A tibble: 2 x 2
    bible_binary mean_trustsci
                     <dbl>
##
            <dbl>
## 1
                0
                          2.50
## 2
                           2.01
                1
group1 <- gss_subset %>% filter(bible_binary == 1)
group0 <- gss_subset %>% filter(bible_binary == 0)
n1 <- nrow(group1)</pre>
n0 <- nrow(group0)</pre>
mean1 <- mean(group1$trustsci)</pre>
mean0 <- mean(group0$trustsci)</pre>
sd1 <- sd(group1$trustsci)</pre>
sd0 <- sd(group0$trustsci)</pre>
diff_in_means <- mean0 - mean1</pre>
se_diff \leftarrow sqrt((sd1^2 / n1) + (sd0^2 / n0))
z_score <- diff_in_means / se_diff</pre>
# Safe printing to LaTeX/PDF
cat("Difference in means:", round(diff_in_means, 3), "\n")
```

```
## Difference in means: 0.492
cat("Standard error:", round(se_diff, 3), "\n")
## Standard error: 0.051
cat("Z-score:", round(z_score, 3), "\n")
## Z-score: 9.602
model <- lm(trustsci ~ bible_binary, data = gss_subset)</pre>
summary(model)
##
## Call:
## lm(formula = trustsci ~ bible_binary, data = gss_subset)
## Residuals:
                 1Q Median
       Min
                                  3Q
                                          Max
## -1.50452 -0.50452 -0.01231 0.49548 0.98769
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.50452 0.03274 76.498 <2e-16 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.6883 on 765 degrees of freedom
## Multiple R-squared: 0.1113, Adjusted R-squared: 0.1101
## F-statistic: 95.77 on 1 and 765 DF, p-value: < 2.2e-16
library(lmtest)
## Loading required package: zoo
## Attaching package: 'zoo'
## The following objects are masked from 'package:base':
##
##
      as.Date, as.Date.numeric
library(sandwich)
coeftest(model, vcov = vcovHC(model, type = "HC1"))
##
## t test of coefficients:
                Estimate Std. Error t value Pr(>|t|)
##
```

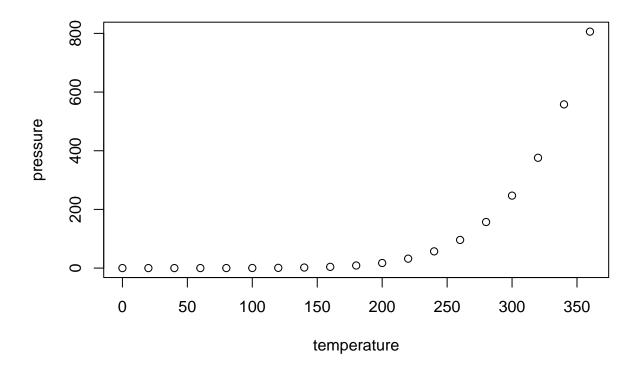
```
## (Intercept) 2.504525  0.030937  80.957 < 2.2e-16 ***
## bible_binary -0.492217  0.051257 -9.603 < 2.2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1

# Coefficient matches the difference in means
summary(cars)</pre>
```

```
##
        speed
                          dist
##
    Min.
            : 4.0
                    Min.
                               2.00
##
    1st Qu.:12.0
                    1st Qu.: 26.00
    Median:15.0
                    Median : 36.00
##
##
    Mean
            :15.4
                    Mean
                            : 42.98
    3rd Qu.:19.0
                    3rd Qu.: 56.00
##
##
    Max.
            :25.0
                    Max.
                            :120.00
```

Including Plots

You can also embed plots, for example:



Note that the echo = FALSE parameter was added to the code chunk to prevent printing of the R code that generated the plot.

```
\label{eq:continuous} \begin{split} & library(gssr) \ library(tidyverse) \ library(dplyr) \\ & data(gss\_all) \end{split}
```

```
gss_2018 < -gss_get_yr(2018)
dim(gss_2018) colnames(gss_2018) head(gss_2018) view(gss_2018)
gss_subset <- gss_2018 %>% select(bible, trustsci) %>% na.omit()
gss_subset <- gss_2018 %>% select(bible, trustsci) %>% filter(!is.na(bible), !is.na(trustsci)) %>% filter(!is.na(bible), !is.na(bible), !is.na(trustsci)) %>% filter(!is.na(bible), !is.na(bible), !is.na(bibl
ter(trustsci \%in\% c(1, 2, 3)) \%>\% mutate(bible binary = ifelse(bible == 1, 1, 0))
str(gss_subset) head(gss_subset) summary(gss_subset)
mean_trust <- gss_subset %>% group_by(bible_binary) %>% summarise(mean_trustsci = mean(trustsci,
na.rm = TRUE)) print(mean_trust)
group1 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <- gss\_subset \%>\% \ filter(bible\_binary == 1) \ group0 <-
n1 < -nrow(group1) n0 < -nrow(group0)
mean1 < -mean(group1trustsci)mean0 < -mean(group0trustsci)
sd1 < -sd(group1trustsci)sd0 < -sd(group0trustsci)
# Calculate values
diff in means <- mean0 - mean1
se_diff \leftarrow sqrt((sd1^2 / n1) + (sd0^2 / n0))
z_score <- diff_in_means / se_diff</pre>
# Safe printing to LaTeX/PDF
cat("Difference in means:", round(diff_in_means, 3), "\n")
## Difference in means: 0.492
cat("Standard error:", round(se_diff, 3), "\n")
## Standard error: 0.051
cat("Z-score:", round(z_score, 3), "\n")
## Z-score: 9.602
model <- lm(trustsci ~ bible_binary, data = gss_subset)</pre>
summary(model)
##
## Call:
## lm(formula = trustsci ~ bible_binary, data = gss_subset)
## Residuals:
                                                                                                                                          3Q
##
                              Min
                                                                     1Q
                                                                                   Median
                                                                                                                                                                         Max
## -1.50452 -0.50452 -0.01231 0.49548 0.98769
##
## Coefficients:
##
                                                             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.50452 0.03274 76.498
                                                                                                                                                                                 <2e-16 ***
## bible_binary -0.49222
                                                                                                          0.05030 - 9.786
                                                                                                                                                                                 <2e-16 ***
```

```
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Residual standard error: 0.6883 on 765 degrees of freedom
## Multiple R-squared: 0.1113, Adjusted R-squared: 0.1101
## F-statistic: 95.77 on 1 and 765 DF, p-value: < 2.2e-16
library(lmtest)
library(sandwich)
coeftest(model, vcov = vcovHC(model, type = "HC1"))
##
## t test of coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.504525 0.030937 80.957 < 2.2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
# Coefficient matches the difference in means
```