

## R Analysis Example Replication C6

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# ASDA2 Chapter 6 analysis examples replication
# Example 6.1
(ex61 <- svymean(~factor(irregular), subhhanes, se=T, na.rm=T, deff=T, ci=T, keep.vars=T))
confint(ex61)
ex61p <- svyciprop(~factor(irregular), subhhanes, se=T, na.rm=T, deff=T, ci=T, keep.vars=T)
ex61p

# Example 6.2 NHANES ADULT DATA
ex62 <- svymean(~factor(racec), design=subhhanes, se=T, na.rm=T, deff=T, ci=T, keep.vars=T)
ex62
confint(ex62)

# Example 6.3 NHANES ADULT DATA
ex63 <- svymean(~factor(bp_catc), subhhanes, se=T, na.rm=T, deff=T, ci=T, keep.vars=T)
ex63
confint(ex63)

# EXAMPLE 6.4 ESS6 Russian Federation Data, Proportions of Russian 15+ by Marital Status
# GOF TOOL WITH PRE-SET PROPORTIONS (NOT AVAILABLE IN R)

rfdata <- read_sas("P:/ASDA 2/Data sets/ess6 russia/ess6_russia_20aug2016.sas7bdat")
summary(rfdata)

#create factor variables
rfdata$marcatc <- factor(rfdata$marcat, levels = 1:3, labels =c("Married", "Previous", "Never"))

rfsvy <- svydesign(strata=~stratify, id=~psu, weights=~PSPWGHT, data=rfdata, nest=T)

ex6_4 <- svymean(~factor(marcatc), design=rfsvy, na.rm=T, se=T, deff=T, ci=T, keep.vars=T)
print(ex6_4)

# Analysis Example 6.5 PIE AND BAR CHARTS

# Pie of Marital Status Russian Federation Data
ex6_5 <- svymean(~factor(marcatc), rfsvy, se=T, na.rm=T, deff=T, ci=T, keep.vars=T)
pie(ex6_5, col=c("black", "grey60", "blue", "red"), c("Married", "Previously Married", "Never Married"))

# Bar chart of marital status
barplot(ex6_5, legend=c("Married", "Previously Married", "Never Married"), col=c("black","blue", "red"))

# Analysis Example 6.6, NCS-R DATA
(ex6_6 <- svymean(~interaction (SEX, mde), ncsrsvyp1, se=T, na.rm=T, ci=T, keep.vars=T))
# obtain confidence intervals
confint(ex6_6)
# svyby analysis gives mean of mde by sex
(ex6_6a <- svyby(~mde, ~SEX, ncsrsvyp1, svymean, se=T, na.rm=T, ci=T, keep.vars=T))

#CODES FOR SEX 1=MALE 2=FEMALE
#svychisq provides a 2 by 2 chisq test (F)
svychisq(~SEX+mde, ncsrsvyp1, statistic="F")
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# Analysis Example 6.7 MEAN OF MDE OVER SEX AND LINEAR COMPARISON TEST
(ex6_7 <- svyby(~mde, ~sex, ncsrsvyp1, svymean, se=T, na.rm=T, ci=T, keep.vars=T))
svycontrast(ex6_7,list(avg=c(.5,.5), diff=c(1,-1)))

# Analysis Example 6.8 Independence of MDE and Gender
(ex6_8 <- svyby(~mde, ~SEX, ncsrsvyp1, svymean, se=T, na.rm=T, ci=T, keep.vars=T))
#CODES FOR SEX 1=MALE 2=FEMALE
#svychisq provides a 2 by 2 chisq test (F)
svychisq(~SEX+mde, ncsrsvyp1, statistic="F")

# Analysis Example 6.9 Independence of Education and Alcohol Dependence
ex6_9 <- svyby (~ald,-ed4catc, subset(ncsrsvyp2, AGE < 29 & !is.na(ED4CAT) & !is.na(ald)), svymean, na.rm=T,
ci=T)

# CODES FOR ED4CAT 1=0-11 2=12 3=13-15 4=16+ YEARS OF EDUCATION
print(ex6_9)
summary(ex6_9,statistic="ChiSq")
svychisq(~ald+ ed4catc, subset(ncsrsvyp2, AGE < 29 & !is.na(ED4CAT) & !is.na(ald)), na.rm=T, statistic = "F")

# Analysis Example 6.10 Logistic Regression MDE regressed on Gender
(ex6_10 <- svyglm (mde~sexm, design=ncsrsvyp1, family=quasibinomial))
summary(ex6_10)
#note can use exponent function with beta to obtain OR

# Figure 6.8 Bar Chart of Marital Status in Russian Federation data
fig6_8 <- svyby(~factor(marcac), ~GNDR, rfsvy, svymean, na.rm=T)

# Bar chart of marital status by gender
print(fig6_8)
barplot(fig6_8, legend=c("Married", "Previously Married", "Never Married"), col=c("black", "blue", "red"),
xlab=c("Male","Female") )

# Analysis Example 6.11 Independence of Gender and MDE while controlling for Age, not available in R Survey
Package

# Analysis Example 6.12 Loglinear Model Examining Relationship Between MDE and Male
#null model run first
null <-svyloglin(~mde+sexm,ncsrsvyp1)
summary(null)

# saturated model, update null model with interaction of mde and sexm
saturated <- update(null, ~.+mde:sexm)
summary(saturated)

# obtain F test
svychisq(~mde+sexm, ncsrsvyp1)
# obtain ChiSQ test
svychisq(~mde+sexm, ncsrsvyp1, statistic="Chisq")

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## Output R Analysis Example Replication C6

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> # Example 6.1
> (ex61 <- svymean(~factor(irregular), subnhanes, se=T, na.rm=T, deff=T, ci=T, keep.vars=T))
              mean      SE  DEff
factor(irregular)0 0.9835810 0.0016779 0.9367
factor(irregular)1 0.0164190 0.0016779 0.9367
> confint(ex61)
              2.5 %      97.5 %
factor(irregular)0 0.98029247 0.98686955
factor(irregular)1 0.01313045 0.01970753
> ex61p <- svyciprop(~factor(irregular), subnhanes, se=T, na.rm=T, deff=T, ci=T, keep.vars=T)
> ex61p
              2.5% 97.5%
factor(irregular) 0.0164 0.0132 0.02

> # Example 6.2 NHANES ADULT DATA
> ex62 <- svymean(~factor(racec), design=subnhanes, se=T, na.rm=T, deff=T, ci=T, keep.vars=T)
> ex62
              mean      SE  DEff
factor(racec)Mexican      0.079168 0.017251 22.9171
factor(racec)Other Hispanic 0.066224 0.015193 20.9551
factor(racec)White      0.659386 0.038892 37.8086
factor(racec)Black      0.117185 0.023370 29.6387
factor(racec)Other      0.078037 0.010917 9.3001
> confint(ex62)
              2.5 %      97.5 %
factor(racec)Mexican      0.04535738 0.11297837
factor(racec)Other Hispanic 0.03644730 0.09600144
factor(racec)White      0.58316044 0.73561243
factor(racec)Black      0.07138031 0.16298887
factor(racec)Other      0.05663959 0.09943387

> # Example 6.3 NHANES ADULT DATA
> ex63 <- svymean(~factor(bp_catc), subnhanes, se=T, na.rm=T, deff=T, ci=T, keep.vars=T)
> ex63
              mean      SE  DEff
factor(bp_catc)Normal      0.4722233 0.0155209 5.1762
factor(bp_catc)Pre-HBP      0.4279854 0.0120353 3.1685
factor(bp_catc)Stage 1 HBP 0.0797780 0.0058154 2.4669
factor(bp_catc)Stage 2 HBP 0.0200133 0.0043847 5.2494
> confint(ex63)
              2.5 %      97.5 %
factor(bp_catc)Normal      0.44180279 0.50264372
factor(bp_catc)Pre-HBP      0.40439669 0.45157418
factor(bp_catc)Stage 1 HBP 0.06837999 0.09117605
factor(bp_catc)Stage 2 HBP 0.01141946 0.02860712
```

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> # Example 6.4 ESS6 Russian Federation Data, Proportions of Russian 15+ by Marital Status
> # GOF TOOL WITH PRE-SET PROPORTIONS (NOT AVAILABLE IN R)

> rfdata <- read_sas("P://ASDA 2/Data sets/ess6 russia/ess6_russia_20aug2016.sas7bdat")
> #create factor variables
> rfdata$marcatc <- factor(rfdata$marcat, levels = 1:3, labels =c("Married", "Previous", "Never"))

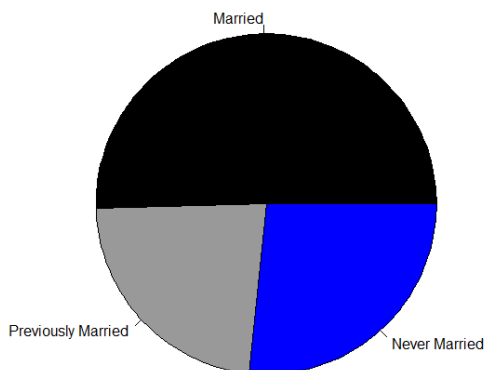
> rfsvy <- svydesign(strata=~stratify, id=~psu, weights=~PSPWGHT, data=rfdata, nest=T)

> ex6_4 <- svymean(~factor(marcatc), design=rfsvy, na.rm=T, se=T, deff=T, ci=T, keep.vars=T)
> print(ex6_4)

              mean      SE  DEff
factor(marcatc)Married  0.503860 0.012878 442.85
factor(marcatc)Previous 0.230066 0.011536 501.57
factor(marcatc)Never    0.266074 0.013401 613.97

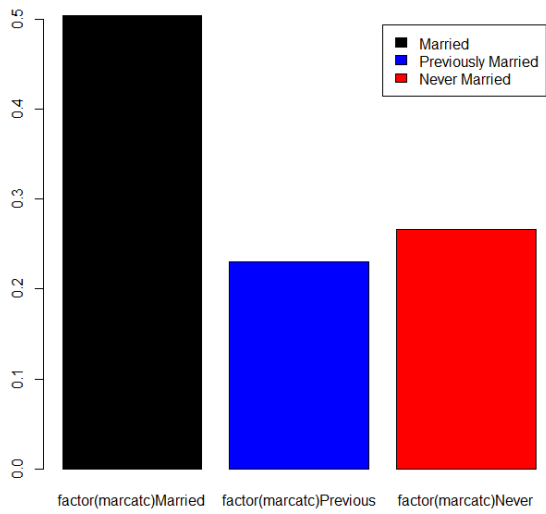
> # Pie of Marital Status Russian Federation Data
> ex6_5 <- svymean(~factor(marcatc), rfsvy, se=T, na.rm=T, deff=T, ci=T, keep.vars=T)
> pie(ex6_5, col=c("black", "grey60", "blue", "red"), c("Married", "Previously Married", "Never Married"))

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> # Bar chart of marital status
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> barplot(ex6_5, legend=c("Married", "Previously Married", "Never Married"), col=c("black", "blue", "red"))
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> # Analysis Example 6.6, NCS-R DATA
> (ex6_6 <- svymean(~interaction (SEX, mde), ncsrsvyp1, se=T, na.rm=T, ci=T, keep.vars=T))
              mean      SE
interaction(SEX, mde)1.0 0.406644 0.0070
interaction(SEX, mde)2.0 0.401644 0.0054
interaction(SEX, mde)1.1 0.072208 0.0034
interaction(SEX, mde)2.1 0.119504 0.0030
> # obtain confidence intervals
> confint(ex6_6)
              2.5 %      97.5 %
interaction(SEX, mde)1.0 0.39296383 0.42032513
interaction(SEX, mde)2.0 0.39113771 0.41215085
interaction(SEX, mde)1.1 0.06546993 0.07894551
interaction(SEX, mde)2.1 0.11356911 0.12543793

> # svyby analysis gives mean of mde by sex
> (ex6_6a <- svyby(~mde, ~SEX, ncsrsvyp1, svymean, se=T, na.rm=T, ci=T, keep.vars=T))
  SEX      mde      se
1   1 0.1507933 0.007747811
2   2 0.2293083 0.005647255
>
> #CODES FOR SEX 1=MALE 2=FEMALE
> #svychisq provides a 2 by 2 chisq test (F)
> svychisq(~SEX+mde, ncsrsvyp1, statistic="F")
      Pearson's X^2: Rao & Scott adjustment
data: svychisq(~SEX + mde, ncsrsvyp1, statistic = "F")
F = 57.978, ndf = 1, ddf = 42, p-value = 1.947e-09

> # Analysis Example 6.7 MEAN OF MDE OVER SEX AND LINEAR COMPARISON TEST
> (ex6_7 <- svyby(~mde, ~sex, ncsrsvyp1, svymean, se=T, na.rm=T, ci=T, keep.vars=T))
      sex      mde      se
Male   Male 0.1507933 0.007747811
Female Female 0.2293083 0.005647255
> svycontrast(ex6_7,list(avg=c(.5,.5), diff=c(1,-1)))
      contrast      SE
avg    0.190051 0.0048
diff  -0.078515 0.0096
Warning message:
In vcov.svyby(stat) : Only diagonal elements of vcov() available

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> # Analysis Example 6.8 Independence of MDE and Gender
> (ex6_8 <- svyby(~mde, ~SEX, ncsrsvyp1, svymean, se=T, na.rm=T, ci=T, keep.vars=T))
  SEX      mde      se
1   1 0.1507933 0.007747811
2   2 0.2293083 0.005647255
> #CODES FOR SEX 1=MALE 2=FEMALE
> #svychisq provides a 2 by 2 chisq test (F)
> svychisq(~SEX+mde, ncsrsvyp1, statistic="F")
  Pearson's X^2: Rao & Scott adjustment
data: svychisq(~SEX + mde, ncsrsvyp1, statistic = "F")
F = 57.978, ndf = 1, ddf = 42, p-value = 1.947e-09

> # Analysis Example 6.9 Independence of Education and Alcohol Dependence
> ex6_9 <- svyby (~ald, ~ed4catc, subset(ncsrsvyp2, AGE < 29 & !is.na(ED4CAT) & !is.na(ald)), svymean, na.rm=T,
ci=T)
>
> # CODES FOR ED4CAT 1=0-11 2=12 3=13-15 4=16+ YEARS OF EDUCATION
> print(ex6_9)
  ed4catc      ald      se
0-11     0-11 0.09128575 0.02937999
12         12 0.04855850 0.01345971
13-15    13-15 0.04895775 0.01004206
16+       16+ 0.06903765 0.01364029
> summary(ex6_9,statistic="ChiSq")
  ed4catc      ald      se
0-11 :1  Min.   :0.04856  Min.   :0.01004
12   :1  1st Qu.:0.04886  1st Qu.:0.01261
13-15:1  Median :0.05900  Median :0.01355
16+  :1  Mean    :0.06446  Mean    :0.01663
      3rd Qu.:0.07460  3rd Qu.:0.01758
      Max.   :0.09129  Max.   :0.02938
> svychisq(~ald+ ed4catc, subset(ncsrsvyp2, AGE < 29 & !is.na(ED4CAT) & !is.na(ald)), na.rm=T, statistic = "F")
  Pearson's X^2: Rao & Scott adjustment
data: svychisq(~ald + ed4catc, subset(ncsrsvyp2, AGE < 29 & !is.na(ED4CAT) & !is.na(ald)), na.rm = T,
statistic = "F")
F = 1.6498, ndf = 2.7506, ddf = 112.7800, p-value = 0.1858

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> # Analysis Example 6.10 Logistic Regression MDE regressed on Gender
> (ex6_10 <- svyglm (mde~sexm, design=ncsrsvyp1, family=quasibinomial))
Stratified 1 - level Cluster Sampling design (with replacement)
With (84) clusters.
svydesign(strata = ~SESTRAT, id = ~SECLUSTER, weights = ~NCSRWTSH,
  data = ncsr, nest = T)
Call: svyglm(formula = mde ~ sexm, design = ncsrsvyp1, family = quasibinomial)
Coefficients:
(Intercept)          sexm
    -1.2122         -0.5162
Degrees of Freedom: 9281 Total (i.e. Null);  41 Residual
Null Deviance:      9072
Residual Deviance: 8979      AIC: NA
> summary(ex6_10)
Call:
svyglm(formula = mde ~ sexm, design = ncsrsvyp1, family = quasibinomial)
Survey design:
svydesign(strata = ~SESTRAT, id = ~SECLUSTER, weights = ~NCSRWTSH,
  data = ncsr, nest = T)
Coefficients:
              Estimate Std. Error t value Pr(>|t|)
(Intercept) -1.21222    0.03195 -37.935 < 2e-16 ***
sexm         -0.51617    0.06820  -7.568 2.63e-09 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for quasibinomial family taken to be 1.000108)
Number of Fisher Scoring iterations: 4
> #note can use exponent function with beta to obtain OR

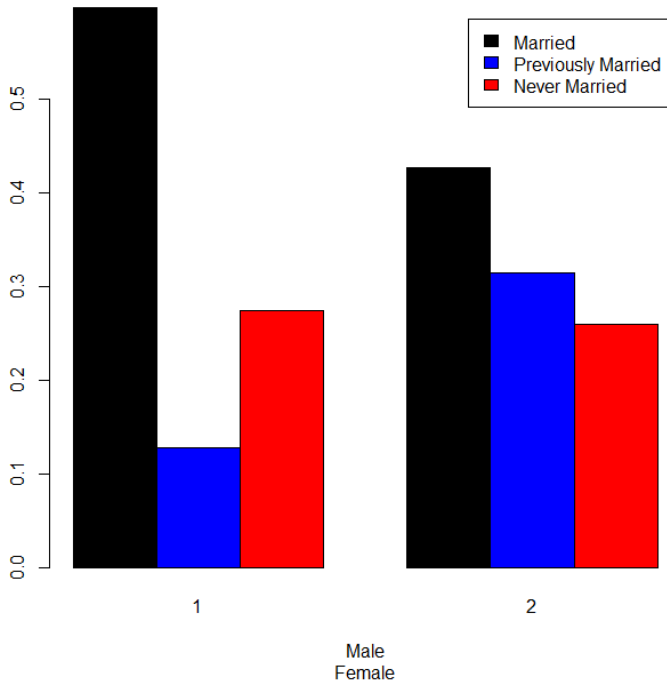
```

```

> # Figure 6.8 Bar Chart of Marital Status in Russian Federation data
> fig6_8 <- svyby(~factor(marcatc), ~GNDR, rfsvy, svymean, na.rm=T)

> # Bar chart of marital status by gender
> print(fig6_8)
  GNDR factor(marcatc)Married factor(marcatc)Previous factor(marcatc)Never se.factor(marcatc)Married
se.factor(marcatc)Previous se.factor(marcatc)Never
1  1      0.5972364      0.1281528      0.2746108      0.01707498
0.01197434      0.01631450
2  2      0.4261892      0.3148384      0.2589724      0.01632922
0.01610291      0.01722477
> barplot(fig6_8, legend=c("Married", "Previously Married", "Never Married"), col=c("black", "blue", "red"),
xlab=c("Male","Female") )

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> # Analysis Example 6.11 Independence of Gender and MDE while controlling for Age, not available in R Survey
Package

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> # Analysis Example 6.12 Loglinear Model Examining Relationship Between MDE and Male
> #null model run first
> null <-svyloglin(~mde+sexm,ncsrsvyp1)
> summary(null)
Loglinear model: svyloglin(~mde + sexm, ncsrsvyp1)
           coef      se      p
mde1  0.71946455 0.01573591 0.000000e+00
sexm1  0.04232084 0.01065001 7.073982e-05
>
> # saturated model, update null model with interaction of mde and sexm
> saturated <- update(null, ~.+mde:sexm)
> summary(saturated)
Loglinear model: update(null, ~. + mde:sexm)
           coef      se      p
mde1      0.7351533 0.01716099 0.000000e+00
sexm1      0.1228566 0.01151009 1.349733e-26
mde1:sexm1 -0.1290428 0.01705080 3.786148e-14
>
> # obtain F test
> svychisq(~mde+sexm, ncsrsvyp1)
      Pearson's X^2: Rao & Scott adjustment
data:  svychisq(~mde + sexm, ncsrsvyp1)
F = 57.978, ndf = 1, ddf = 42, p-value = 1.947e-09
> # obtain ChiSQ test
> svychisq(~mde+sexm, ncsrsvyp1, statistic="Chisq")
      Pearson's X^2: Rao & Scott adjustment
data:  svychisq(~mde + sexm, ncsrsvyp1, statistic = "Chisq")
X-squared = 92.15, df = 1, p-value = 2.65e-14

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