

R Analysis Example Replication C9

```
# Note: all data management and survey design setup code included in Chapter 5 document
# Chapter 9 ASDA2 Analysis Examples Replication

# Figure 9.2 Bar chart of work status NCS-R data
fig92 <- svymean( ~factor(WKSTAT3C), ncsrsvyp2, na.rm=T)
barplot(fig92, legend=c("Employed", "Unemployed", "NLF") , col=c("black", "grey60", "blue"))

# Tests for Potential Predictors of Work Status
svychisq(~WKSTAT3C+SEX, ncsrsvyp2, statistic="F")
svychisq(~WKSTAT3C+ald, ncsrsvyp2, statistic="F")
svychisq(~WKSTAT3C+mde, ncsrsvyp2, statistic="F")
svychisq(~WKSTAT3C+ED4CAT, ncsrsvyp2, statistic="F")
svychisq(~WKSTAT3C+ag4cat, ncsrsvyp2, statistic="F")
svychisq(~WKSTAT3C+MAR3CAT, ncsrsvyp2, statistic="F")

# Note: No Multinomial Logit Option in R Survey package

# Ordinal Regression with Russian Federation Data
# Read Data and set design variables
# Use C9 version of data for this example
rfdata_c9 <- read_sas("P:/ASDA 2/Data sets/ess6 russia/c9_russia_1jun2017.sas7bdat")
summary(rfdata_c9)

#create factor variables
rfdata_c9$marcatc <- factor(rfdata_c9$marcat, levels = 1:3, labels =c("Married", "Previous", "Never"))
rfsvy <- svydesign(strata=~stratify, id=~psu, weights=~PSPWGHT, data=rfdata_c9, nest=T)

ex936 <- svymean(~factor(stflife2), design=rfsvy, na.rm=T, se=T, deff=T, ci=T, keep.vars=T)
print(ex936)
barplot(ex936, legend=c("0-1", "2-4", "5", "6-8", "9-10") , col=c("black", "grey60", "blue", "red", "green"))

#ordinal logistic using satisfaction with life
summary(ex936_ordinal <- svyolr (factor(stflife2) ~ factor(agecat) + male + marcatc, design=rfsvy))
exp(ex936_ordinal$coef)

# HRS data for Poisson Regression using Number of Falls
#histogram of number of falls 24
svyhist(~numfalls24 , subset (hrssvyr, NAGE >=65), main="", col="grey80", xlab ="Histogram of Number of Falls
Past 24 Months")

# Poisson model
summary(hrs)

# Create variables for example
hrs$nage_c=(hrs$NAGE - 74.5)
hrs$bmi_c=(hrs$R11BMI - 27.7)
hrs$male=(hrs$GENDER ==1)

# Create Design Object and then subset for 65 Plus
hrssvyr <- svydesign(strata=~STRATUM, id=~SECU, weights=~NWGTR , data=hrs, nest=T)
```

```
summary(hrssvyr)
hrssvy65 <- subset(hrssvyr, age65p==1)

summary(ex947_poisson <- svyglm(numfalls24 ~ male + nage_c + arthritis + diabetes + bmi_c, design=hrssvy65,
family=quasipoisson(log)))
exp(ex947_poisson$coef)

# Negative binomial (not available with survey correction, dispersion is accounted for in svyglm, per lumley)
# Zero inflated negative binomial not available in R survey package
```

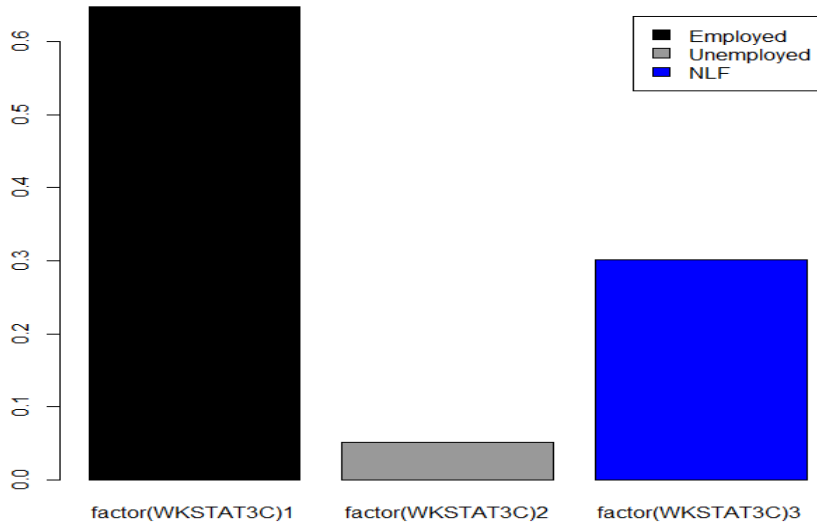
Output R Analysis Example Replication C9

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# Chapter 9 ASDA2 Analysis Examples Replication
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```
> # Figure 9.2 Bar chart of work status NCS-R data
```

```
> fig92 <- svymean( ~factor(WKSTAT3C), ncsrsvyp2, na.rm=T)
```

```
> barplot(fig92, legend=c("Employed", "Unemployed", "NLF"), col=c("black", "grey60", "blue"))
```



```

# Tests for Potential Predictors of Work Status
> svychisq(~WKSTAT3C+SEX, ncsrsvyp2, statistic="F")
    Pearson's X^2: Rao & Scott adjustment
data: svychisq(~WKSTAT3C + SEX, ncsrsvyp2, statistic = "F")
F = 27.329, ndf = 1.875, ddf = 78.748, p-value = 2.171e-09

> svychisq(~WKSTAT3C+ald, ncsrsvyp2, statistic="F")
    Pearson's X^2: Rao & Scott adjustment
data: svychisq(~WKSTAT3C + ald, ncsrsvyp2, statistic = "F")
F = 3.1249, ndf = 1.7248, ddf = 72.4410, p-value = 0.05716

> svychisq(~WKSTAT3C+mde, ncsrsvyp2, statistic="F")
    Pearson's X^2: Rao & Scott adjustment
data: svychisq(~WKSTAT3C + mde, ncsrsvyp2, statistic = "F")
F = 4.6693, ndf = 1.7348, ddf = 72.8610, p-value = 0.01605

> svychisq(~WKSTAT3C+ED4CAT, ncsrsvyp2, statistic="F")
    Pearson's X^2: Rao & Scott adjustment
data: svychisq(~WKSTAT3C + ED4CAT, ncsrsvyp2, statistic = "F")
F = 27.64, ndf = 5.1457, ddf = 216.1200, p-value < 2.2e-16

> svychisq(~WKSTAT3C+ag4cat, ncsrsvyp2, statistic="F")
    Pearson's X^2: Rao & Scott adjustment
data: svychisq(~WKSTAT3C + ag4cat, ncsrsvyp2, statistic = "F")
F = 113.49, ndf = 4.9646, ddf = 208.5100, p-value < 2.2e-16

> svychisq(~WKSTAT3C+MAR3CAT, ncsrsvyp2, statistic="F")
    Pearson's X^2: Rao & Scott adjustment
data: svychisq(~WKSTAT3C + MAR3CAT, ncsrsvyp2, statistic = "F")
F = 23.124, ndf = 3.1985, ddf = 134.3400, p-value = 1.229e-12

```

Note: No Multinomial Logit Option in R Survey package

```

# Ordinal Regression with Russian Federation Data
> # Read Data and set design variables
> # Use C9 version of data for this example

> rfdata_c9 <- read_sas("P:/ASDA 2/Data sets/ess6 russia/c9_russia_1jun2017.sas7bdat")

#create factor variables
> rfdata_c9$marcatc <- factor(rfdata_c9$marcat, levels = 1:3, labels =c("Married", "Previous", "Never"))
> rfsvy <- svydesign(strata=~stratify, id=~psu, weights=~PSPWGHT, data=rfdata_c9, nest=T)

> ex936 <- svymean(~factor(stflife2), design=rfsvy, na.rm=T, se=T, deff=T, ci=T, keep.vars=T)
Warning message:
In svymean.survey.design2(~factor(stflife2), design = rfsvy, na.rm = T, :
  Sample size greater than population size: are weights correctly scaled?
> print(ex936)

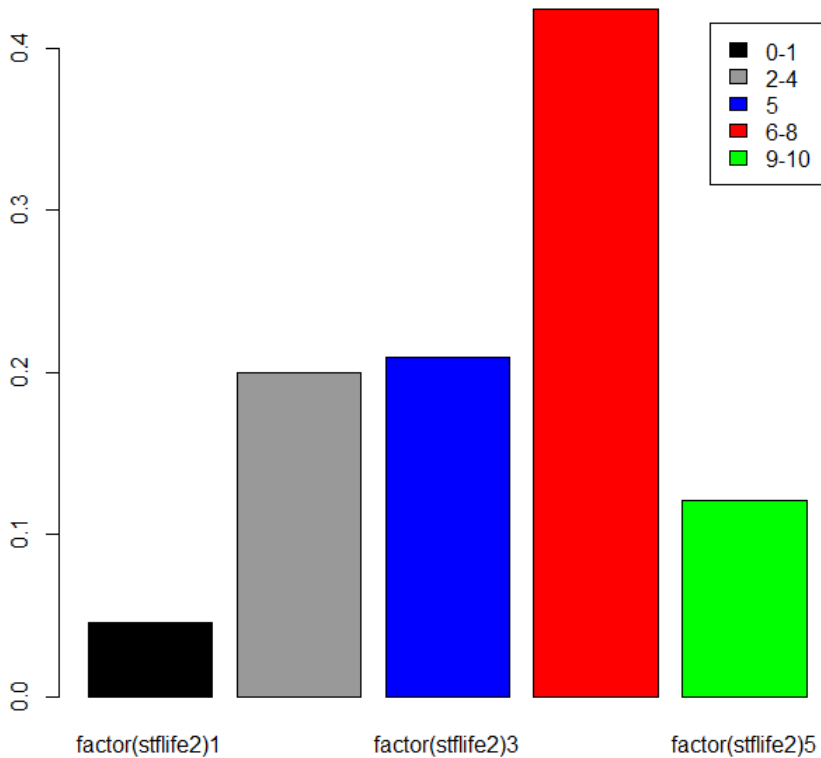
```

	mean	SE	DEff
factor(stflife2)1	0.0452316	0.0065542	NA
factor(stflife2)2	0.2001318	0.0135063	NA
factor(stflife2)3	0.2092193	0.0122451	NA
factor(stflife2)4	0.4240569	0.0174742	NA
factor(stflife2)5	0.1213603	0.0092441	NA

```

> barplot(ex936, legend=c("0-1", "2-4", "5", "6-8", "9-10") , col=c("black", "grey60", "blue", "red", "green"))

```

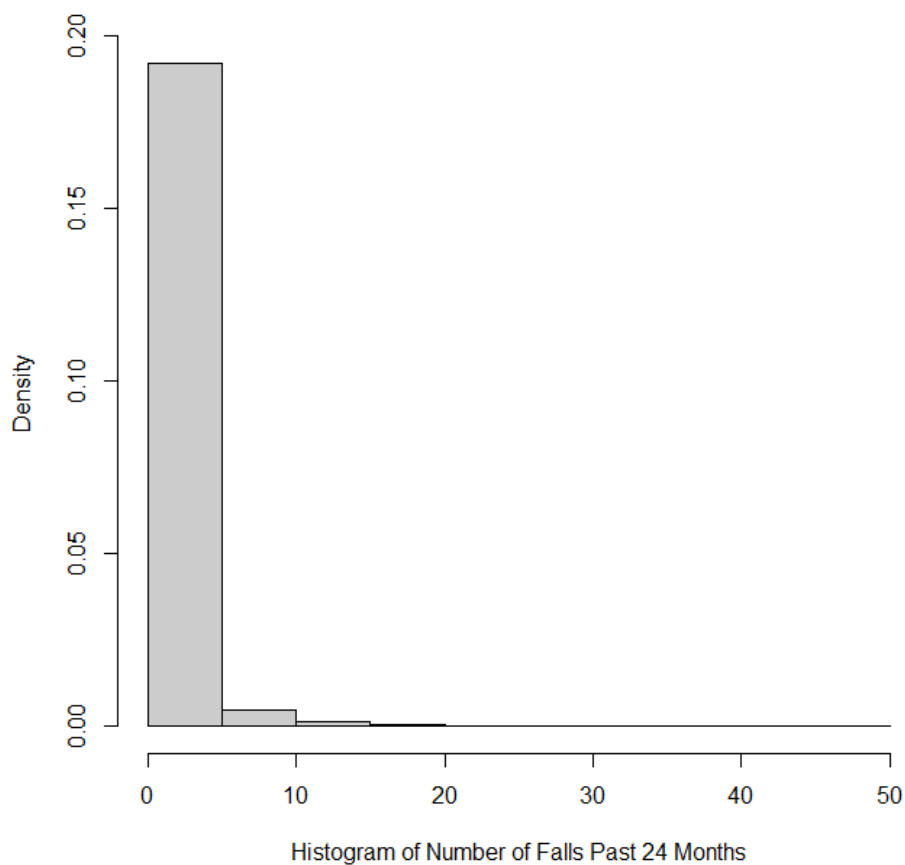


```

> summary(ex936_ordinal <- svyolr (factor(stflife2) ~ factor(agecat) + male + marcatc, design=rfsvy))
Call:
svyolr(factor(stflife2) ~ factor(agecat) + male + marcatc, design = rfsvy)
Coefficients:
                Value Std. Error  t value
factor(agecat)2 -0.5293373 0.13614998 -3.887899
factor(agecat)3 -0.7455312 0.14332027 -5.201855
factor(agecat)4 -0.8080856 0.16561383 -4.879337
male              -0.1096234 0.09523049 -1.151138
marcatcPrevious  -0.2088768 0.10542193 -1.981341
marcatcNever     -0.1371932 0.13208274 -1.038692
Intercepts:
      Value   Std. Error t value
1|2  -3.7111   0.2144  -17.3066
2|3  -1.7928   0.1668  -10.7469
3|4  -0.8348   0.1592   -5.2433
4|5   1.3841   0.1536    9.0095
(69 observations deleted due to missingness)
> exp(ex936_ordinal$coef)
factor(agecat)2 factor(agecat)3 factor(agecat)4      male marcatcPrevious
      0.5889952      0.4744822      0.4457105      0.8961715      0.8114952
marcatcNever

```

```
> # HRS data for Poisson Regression using Number of Falls
> #histogram of number of falls 24
> svyhist(~numfalls24 , subset (hrssvyr, NAGE >=65), main="", col="grey80", xlab ="Histogram of Number of Falls
Past 24 Months")
```



```
> # Poisson model

> # Create variables for example
> hrs$nage_c=(hrs$NAGE - 74.5)
> hrs$bmi_c=(hrs$R11BMI - 27.7)
> hrs$male=(hrs$GENDER ==1)

> # Create Design Object and then subset for 65 Plus
> hrssvyr <- svydesign(strata=~STRATUM, id=~SECU, weights=~NWGTR , data=hrs, nest=T)
> hrssvy65 <- subset(hrssvyr, age65p==1)
```

```
> summary(ex947_poisson <- svyglm(numfalls24 ~ male + nage_c + arthritis + diabetes + bmi_c, design=hrssvy65, family=quasipoisson(log)))
```

Call:

```
svyglm(formula = numfalls24 ~ male + nage_c + arthritis + diabetes +  
       bmi_c, design = hrssvy65, family = quasipoisson(log))
```

Survey design:

```
subset(hrssvyr, age65p == 1)
```

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	-0.635588	0.074371	-8.546	3.33e-11	***
maleTRUE	0.257180	0.080487	3.195	0.002470	**
nage_c	0.014666	0.004425	3.314	0.001753	**
arthritis	0.736177	0.077384	9.513	1.26e-12	***
diabetes	0.247533	0.070193	3.526	0.000938	***
bmi_c	0.004126	0.008402	0.491	0.625615	

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for quasipoisson family taken to be 8.755473)

Number of Fisher Scoring iterations: 6

Warning messages:

1: In summary.glm(g) :

observations with zero weight not used for calculating dispersion

2: In summary.glm(glm.object) :

observations with zero weight not used for calculating dispersion

> exp(ex947_poisson\$coef)

(Intercept)	maleTRUE	nage_c	arthritis	diabetes	bmi_c
0.5296241	1.2932785	1.0147744	2.0879372	1.2808623	1.0041347

> # NEGATIVE BINOMIAL (NOT AVAILABLE WITH SURVEY CORRECTION, DISPERSION IS ACCOUNTED FOR IN SVYGLM, PER LUMLEY)

> # ZERO INFLATED NEGATIVE BINOMIAL NOT AVAILABLE IN R SURVEY PACKAGE