

# Data

## Get Data

### ENTER RAW DATA:

1. Enter data in Excel (variables in columns, individuals in rows, first row has variable names, no spaces or special characters).
2. Save as "Comma Separated Values (\*.CSV)" file in your local directory/folder.

### DATA PROVIDED BY PROFESSOR:

1. Goto [Data Specific to MTH107 on Resources page](#).
2. Right-click on "data" link and save to your local directory/folder.

## Load CSV

1. Start script and save it in the same folder that contains the CSV file.
2. Select Session, Set Working Directory, To Source File Location menus.
3. Copy resulting `setwd()` code to script.
4. Use `read.csv()` to load data into `dfobj`.
5. Observe structure of data.frame.

```
dfobj <- read.csv("filename.csv")
```

```
str(dfobj)
```

## Filter Individuals

Individuals that meet a certain condition (or conditions) are filtered from the `dfobj` data.frame with `filterD()`.

```
newdf <- filterD(dfobj,cond)
```

where `cond` may be as follows with `var` replaced with a variable name and `value` replaced with a number or category level (if `value` is text then it must be in quotes):

<code>var == value</code>	# equal to
<code>var != value</code>	# not equal to
<code>var &gt; value</code>	# greater than
<code>var &gt;= value</code>	# greater than or equal
<code>var &lt; value</code>	# less than
<code>var &lt;= value</code>	# less than or equal
<code>var %in% c("val","val","val")</code>	# in the list
<code>cond   cond</code>	# either condition met
<code>cond, cond</code>	# both conditions met

Individual in row `rownum` is selected with:

```
dfobj[rownum,]
```

Individual in row `rownum` is excluded with:

```
dfobj[-rownum,]
```

# Exploratory Data Analysis

## Univariate

**QUANTITATIVE** – Summary statistics (mean, median, SD, IQR, etc.) and a histogram for the `qvar` variable.

```
hist(~qvar,data=dfobj,xlab="qvar label")
Summarize(~qvar,data=dfobj,digits=#)
```

**QUANTITATIVE BY GROUP** – Summary statistics and histograms for the `qvar` variable separated by groups in the `gvar` variable.

```
hist(qvar~gvar,data=dfobj,xlab="qvar label")
Summarize(qvar~gvar,data=dfobj,digits=#)
```

**CATEGORICAL** – Frequency and percentage tables and bar chart for the `cvar` variable.

```
(freq1 <- xtabs(~cvar,data=dfobj))
percTable(freq1,digits=#)
barplot(freq1,xlab="cvar label",
        ylab="Frequency")
```

## Bivariate

**QUANTITATIVE** – Correlation ( $r$ ) and scatterplot for the `qvarY` and `qvarX` variables.

```
plot(qvarY~qvarX,data=dfobj,
     ylab="yvar label",xlab="xvar label")
corr(~qvarY+qvarX,data=dfobj)
```

**CATEGORICAL** – Frequency and percentage tables for the `cvarRow` and `cvarCol` variables.

```
(freq2 <- xtabs(~cvarRow+cvarCol,
               data=dfobj))
percTable(freq2) # total/table %
percTable(freq2,margin=1) # row %
percTable(freq2,margin=2) # column %
```

# R CHEATSHEET • MTH107

## Class R FAQ

by Derek H. Ogle, revised Dec-17

## Models

### Normal Distributions

```
distrib(val,mean=meanval,sd=sdval,
        type="q",lower.tail=FALSE)
```

where

- `val` is a value of the quantitative variable or area (i.e., percentage as a proportion).
- `meanval` is population mean ( $\mu$ )
- `sdval` is standard deviation ( $\sigma$ ) or error (SE)
- `type="q"` is included for reverse calculations
- `lower.tail=FALSE` is included for "right-of" calculations

For SE use (where `nval`=sample size):

```
sd=sdval/sqrt(nval)
```

### Linear Regression

The best-fit line between the `rspvar` response and `expvar` explanatory variables.

```
(bfl <- lm(rspvar~expvar,data=dfobj))
```

A visual of the best-fit line.

```
fitPlot(bfl,ylab="rspvar lbl",xlab="expvar lbl")
```

The  $r^2$  value.

```
rSquared(bfl)
```

Predict a value of `rspvar` given a specific `expval` value of the `expvar`.

```
predict(bfl,data.frame(expvar=expval))
```

## Hypothesis Testing

### Quantitative

#### ONE SAMPLE:

```
z.test(dfobj$qvar,mu=mu0,alt=HA,
       conf.level=confval,sd=sdval)
t.test(dfobj$qvar,mu=mu0,alt=HA,
       conf.level=confval)
```

#### TWO SAMPLE:

```
levenesTest(qvar~gvar,data=dfobj)
t.test(qvar~gvar,data=dfobj,alt=HA,
       conf.level=confval,var.equal=TRUE)
```

- `qvar` is the quant. response variable in `dfobj`
- `mu0` is the population mean in  $H_0$
- `HA` is "two.sided" for a not equals, "less" for a less than, or "greater" for a greater than  $H_A$
- `confval` is the confidence level (e.g., 0.95)
- `sdval` is the popn. standard deviation ( $\sigma$ )
- `gvar` is a categorical variable in `dfobj` that identifies the groups
- `var.equal=TRUE` if the population variances are thought to be equal

### Categorical

#### ONE SAMPLE:

Goodness-of-fit test for observed frequencies in the `freq1` table and expected values (or proportions) in `exp.p`.

```
(gof <- chisq.test(freq1,p=exp.p,
                  rescale.p=TRUE,correct=FALSE))
```

#### TWO SAMPLE:

Chi-square for `freq2` two-way observed frequency table (with response variable in columns and groups in rows).

```
(chi <- chisq.test(freq2,correct=FALSE))
```

With the following follow-up analyses:

- Extract the expected values.

```
gof$expected or chi$expected
```

- Extract the residuals.

```
gof$residuals or chi$residuals
```

- Confidence intervals for goodness-of-fit.

```
gofCI(gof,digits=3)
```

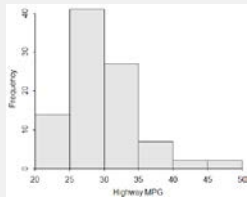
## Data

```
> library(NCStats)
> setwd("C:/aaaWork/Web/GitHub/NCMTH107")
> dfobj <- read.csv("93cars.csv")
> str(dfobj)
'data.frame':   93 obs. of  26 variables:
 $ Type      : Factor w/ 6 levels "Compact","Large",...: 4 ...
 $ HMPG      : int   31 25 26 26 30 31 28 25 27 25 ...
 $ Manual    : Factor w/ 2 levels "No","Yes": 2 2 2 2 1 ...
 $ Weight    : int  2705 3560 3375 3405 3640 2880 3470 ...
 $ Domestic  : Factor w/ 2 levels "No","Yes": 1 1 1 1 1 2 ...

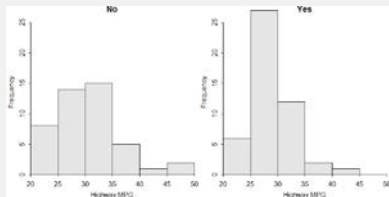
> justSporty <- filterD(dfobj,Type=="Sporty")
> justHMPGgt30 <- filterD(dfobj,HMPG>30)
> noDomestics <- filterD(dfobj,Domestic!="Yes")
> Sprty_Small <- filterD(dfobj,Type %in% c("Sporty","Small"))
```

## Univariate EDA

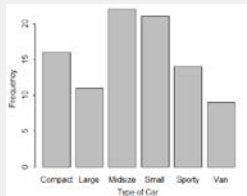
```
> Summarize(~HMPG,data=dfobj,digits=1)
      n mean sd min Q1 median Q3 max
93.0  29.1  5.3 20.0 26.0 28.0 31.0 50.0
> hist(~HMPG,data=dfobj,xlab="Highway MPG")
```



```
> Summarize(HMPG~Domestic,data=dfobj,digits=1)
Domestic n mean sd min Q1 median Q3 max
1 No 45 30.1 6.2 21 25 30 33 50
2 Yes 48 28.1 4.2 20 26 28 30 41
> hist(HMPG~Domestic,data=dfobj,xlab="Highway MPG")
```

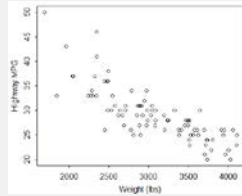


```
> ( freq1 <- xtabs(~Type,data=dfobj) )
Type
Compact Large Midsize Small Sporty Van
16      11      22      21      14      9
> percTable(freq1,digits=1)
Type
Compact Large Midsize Small Sporty Van Sum
17.2    11.8    23.7    22.6    15.1    9.7 100.1
> barplot(freq1,xlab="Type of Car",ylab="Frequency")
```



## Bivariate EDA

```
> plot(HMPG~Weight,data=dfobj,ylab="Highway MPG")
      xlab="Weight (lbs)")
```

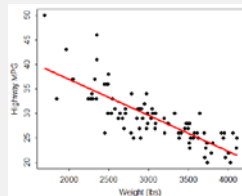


```
> corr(HMPG~Weight,data=dfobj)
[1] -0.8106581

> ( freq2 <- xtabs(~Domestic+Manual,data=dfobj) )
Manual
Domestic No Yes
No        6 39
Yes       26 22
> percTable(freq2,digits=1)
Manual
Domestic No Yes Sum
No        6.5 41.9 48.4
Yes       28.0 23.7 51.7
Sum       34.5 65.6 100.1
> percTable(freq2,margin=1,digits=1)
Manual
Domestic No Yes Sum
No       13.3 86.7 100.0
Yes      54.2 45.8 100.0
> percTable(freq2,margin=2,digits=1)
Manual
Domestic No Yes
No       18.8 63.9
Yes      81.2 36.1
Sum     100.0 100.0
```

## Linear Regression

```
> ( bfl <- lm(HMPG~Weight,data=dfobj) )
Coefficients:
(Intercept)      Weight
51.601365      -0.007327
> fitPlot(bfl,xlab="Weight (lbs)",ylab="Highway MPG")
```



```
> rSquared(bfl)
[1] 0.6571665
> predict(bfl,data.frame(Weight=3000))
29.62019
```

## Hypothesis Tests

```
> z.test(dfobj$HMPG,mu=26,alt="greater",conf.level=0.95,sd=6)
z = 4.9601, n = 93, Std. Dev = 6.000, Std. Dev of the sample
mean = 0.622, p-value = 3.523e-07
alternative hypothesis: true mean is greater than 26
95 percent confidence interval:
 28.06264      Inf
sample estimates:
mean of dfobj$HMPG
29.08602

> t.test(dfobj$HMPG,mu=26,alt="two.sided",conf.level=0.99)
t = 5.5818, df = 92, p-value = 2.387e-07
alternative hypothesis: true mean is not equal to 26
99 percent confidence interval:
 27.63178 30.54026
sample estimates:
mean of x
29.08602

> levenesTest(HMPG~Domestic,data=dfobj)
      Df F value Pr(>F)
group 1  5.3595 0.02286 *
      91

> t.test(HMPG~Manual,data=dfobj,alt="less",conf.level=0.99,
var.equal=TRUE)
t = -4.2183, df = 91, p-value = 2.904e-05
alt. hypothesis: true difference in means is less than 0
99 percent confidence interval:
 -Inf -1.980103
sample estimates:
mean in group No mean in group Yes
26.12500      30.63934

> exp <- c(1,1,1,1,1,1)/6
> (gof<-chisq.test(freq1,p=exp,rescale.p=TRUE,correct=FALSE))
X-squared = 8.871, df = 5, p-value = 0.1143

> gof$expected
Compact Large Midsize Small Sporty Van
15.5    15.5    15.5    15.5    15.5    15.5

> gof$residuals
Compact Large Midsize Small Sporty Van
0.12700 -1.14300  1.65100  1.39700 -0.38100 -1.65100

> gofCI(gof,digits=3)
p.obs p.LCI p.UCI p.exp
Compact 0.172 0.109 0.261 0.167
Large   0.118 0.067 0.199 0.167
Midsize 0.237 0.162 0.332 0.167
Small   0.226 0.153 0.321 0.167
Sporty  0.151 0.092 0.237 0.167
Van     0.097 0.052 0.174 0.167

> ( chi <- chisq.test(freq2,correct=FALSE) )
Pearson's Chi-squared test with freq2
X-squared = 17.1588, df = 1, p-value = 3.438e-05

> chi$expected
Manual
Domestic No Yes
No 15.48387 29.51613
Yes 16.51613 31.48387

> chi$residuals
Manual
Domestic No Yes
No -2.410160 1.745645
Yes 2.333627 -1.690214
```