

## PS792 Chapter 3 Basic Statistics

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In this R document, we will learn how to use R to run some basic statistics analysis. The data sets that will be used in this R document are mtcars, Arthritis, and UScrime. The first 6 rows of mtcars data are showed as follows.

```
data(mtcars)
head(mtcars)
```

##		mpg	cyl	disp	hp	drat	wt	qsec	vs	am	gear	carb
##	Mazda RX4	21.0	6	160	110	3.90	2.620	16.46	0	1	4	4
##	Mazda RX4 Wag	21.0	6	160	110	3.90	2.875	17.02	0	1	4	4
##	Datsun 710	22.8	4	108	93	3.85	2.320	18.61	1	1	4	1
##	Hornet 4 Drive	21.4	6	258	110	3.08	3.215	19.44	1	0	3	1
##	Hornet Sportabout	18.7	8	360	175	3.15	3.440	17.02	0	0	3	2
##	Valiant	18.1	6	225	105	2.76	3.460	20.22	1	0	3	1

### Descriptive statistics

Before running some descriptive statistics, we will substract three columns of data from the data frame mtcars. The three variables are mpg, hp, and wt as demonstrated as follows. The three columns of data are assigned to newdata, which is also a data frame.

```
myvars=c("mpg", "hp", "wt")
newdata=mtcars[myvars]
```

Using summary function you can have some descriptive statistics for each of three variables in the data frame newdata.

```
summary(newdata)
```

##	mpg	hp	wt
##	Min. :10.40	Min. : 52.0	Min. :1.513
##	1st Qu.:15.43	1st Qu.: 96.5	1st Qu.:2.581
##	Median :19.20	Median :123.0	Median :3.325
##	Mean :20.09	Mean :146.7	Mean :3.217
##	3rd Qu.:22.80	3rd Qu.:180.0	3rd Qu.:3.610
##	Max. :33.90	Max. :335.0	Max. :5.424

We may add more statistics for a variable such as standard deviation, skewness and kurtosis. The following R function mystats show the details that which parameters are added. Then you can use

```

mystats <- function(x, na.omit=FALSE){
  if (na.omit)
    x <- x[!is.na(x)]
  m <- mean(x)
  n <- length(x)
  s <- sd(x)
  skew <- sum((x-m)^3/s^3)/n
  kurt <- sum((x-m)^4/s^4)/n - 3
  return(c(n=n, mean=m, stdev=s, skew=skew, kurtosis=kurt))
}

#sapply(newdata,mystats)

```

With mystats function available, we can run one of variables mpg, hp, and wt in the data frame newdata as demonstrated as follows.

```

mpg.stat=with(newdata,mystats(mpg))

mpg.stat
##           n      mean      stdev      skew  kurtosis
## 32.000000 20.090625  6.026948  0.610655 -0.372766

hp.stat=with(newdata,mystats(hp))

hp.stat
##           n      mean      stdev      skew  kurtosis
## 32.0000000 146.6875000 68.5628685  0.7260237 -0.1355511

wt.stat=with(newdata,mystats(wt))

wt.stat
##           n      mean      stdev      skew  kurtosis
## 32.00000000  3.21725000 0.97845744  0.42314646 -0.02271075

```

We can combine the descriptive statistics for these three variables using dataframe function to make a data frame.

```

cars.stat=data.frame(mpg=mpg.stat, hp=hp.stat, wt=wt.stat)
cars.stat
##           mpg           hp           wt
## n      32.000000 32.0000000 32.0000000
## mean 20.090625 146.6875000  3.21725000
## stdev 6.026948 68.5628685  0.97845744
## skew  0.610655  0.7260237  0.42314646
## kurtosis -0.372766 -0.1355511 -0.02271075

```

If you have a large number of variables to be processed, then you may use loop to repeat the process.

```

wt.stat=with(newdata,mystats(wt))
nr=length(wt.stat)
nc=ncol(newdata)
STAT=matrix(0,nr,nc)
for(h in 1:nc){
  stat=mystats(newdata[,h])
  STAT[,h]=stat
}
colnames(STAT)=colnames(newdata)
rownames(STAT)=names(stat)

```

```
data.frame(STAT)
```

```

##           mpg           hp           wt
## n          32.000000  32.0000000 32.0000000
## mean       20.090625 146.6875000  3.2172500
## stdev       6.026948  68.5628685  0.9784574
## skew        0.610655   0.7260237  0.4231464
## kurtosis   -0.372766  -0.1355511 -0.0227107

```

You may also convert the above code as another R function MYSTAT.

```

MYSTAT=function(data){
  cn=ncol(data)
  stat=mystats(data[,1])
  rn=length(stat)
  STAT=matrix(0,rn,cn)
  for(h in 1:cn){
    stat=mystats(data[,h])
    STAT[,h]=stat
  }
  colnames(STAT)=colnames(data)
  rownames(STAT)=names(stat)
  STAT=data.frame(STAT)

  return(STAT)
}

```

```

stat=MYSTAT(newdata)
stat

```

```

##           mpg           hp           wt
## n          32.000000  32.0000000 32.0000000
## mean       20.090625 146.6875000  3.2172500
## stdev       6.026948  68.5628685  0.9784574
## skew        0.610655   0.7260237  0.4231464
## kurtosis   -0.372766  -0.1355511 -0.0227107

```

## More methods

There are a couple of other functions packed in different R packages. For example, the `describe` function in the R package `Hmisc`, `stat.desc` in package `pastecs`, `describe` in package `psych`. It is very important to install these packages before using it. For detailed information, please refer to these functions in corresponding R packages by yourselves.

```
require(Hmisc)
```

```
## Loading required package: Hmisc
```

```
## Loading required package: lattice
```

```
## Loading required package: survival
```

```
## Loading required package: Formula
```

```
## Loading required package: ggplot2
```

```
##
```

```
## Attaching package: 'Hmisc'
```

```
## The following objects are masked from 'package:base':
```

```
##
```

```
##      format.pval, units
```

```
describe(newdata)
```

```
## newdata
```

```
##
```

```
## 3 Variables      32 Observations
```

```
## -----
```

```
-
```

```
## mpg
```

```
##      n missing distinct      Info      Mean      Gmd      .05      .10
```

```
##      32      0       25    0.999    20.09    6.796    12.00    14.34
```

```
##      .25      .50      .75      .90      .95
```

```
##    15.43    19.20    22.80    30.09    31.30
```

```
##
```

```
## lowest : 10.4 13.3 14.3 14.7 15.0, highest: 26.0 27.3 30.4 32.4 33.9
```

```
## -----
```

```
-
```

```
## hp
```

```
##      n missing distinct      Info      Mean      Gmd      .05      .10
```

```
##      32      0       22    0.997    146.7    77.04    63.65    66.00
```

```
##      .25      .50      .75      .90      .95
```

```
##    96.50   123.00   180.00   243.50   253.55
```

```
##
```

```
## lowest : 52 62 65 66 91, highest: 215 230 245 264 335
```

```
## -----
```

```
-
```

```
## wt
```

```
##          n missing distinct      Info      Mean      Gmd      .05      .10
##          32         0        29      0.999      3.217      1.089      1.736      1.956
##          .25        .50        .75        .90        .95
##          2.581      3.325      3.610      4.048      5.293
##
## lowest : 1.513 1.615 1.835 1.935 2.140, highest: 3.845 4.070 5.250 5.345
## 5.424
```

```
## -----
-
```

```
library(pastecs)
```

```
## Loading required package: boot
```

```
##
```

```
## Attaching package: 'boot'
```

```
## The following object is masked from 'package:survival':
```

```
##
```

```
##      aml
```

```
## The following object is masked from 'package:lattice':
```

```
##
```

```
##      melanoma
```

```
stat.desc(newdata)
```

```
##          mpg          hp          wt
## nbr.val      32.0000000  32.0000000  32.0000000
## nbr.null      0.0000000  0.0000000  0.0000000
## nbr.na        0.0000000  0.0000000  0.0000000
## min          10.4000000  52.0000000  1.5130000
## max          33.9000000  335.0000000  5.4240000
## range        23.5000000  283.0000000  3.9110000
## sum          642.9000000  4694.0000000 102.9520000
## median        19.2000000  123.0000000  3.3250000
## mean         20.0906250  146.6875000  3.2172500
## SE.mean        1.0654240   12.1203173  0.1729685
## CI.mean.0.95   2.1729465   24.7195501  0.3527715
## var           36.3241028  4700.8669355  0.9573790
## std.dev        6.0269481   68.5628685  0.9784574
## coef.var        0.2999881    0.4674077  0.3041285
```

```
library(psych)
```

```
##
```

```
## Attaching package: 'psych'
```

```
## The following object is masked from 'package:boot':
```

```
##
```

```
##      logit
```

```
## The following object is masked from 'package:Hmisc':
##
## describe

## The following objects are masked from 'package:ggplot2':
##
## %+%, alpha
```

```
describe(newdata)
```

```
##      vars  n   mean    sd median trimmed   mad   min    max   range skew
## mpg     1 32  20.09  6.03  19.20   19.70   5.41 10.40   33.90  23.50  0.61
## hp      2 32 146.69 68.56 123.00  141.19  77.10 52.00  335.00 283.00  0.73
## wt      3 32   3.22  0.98   3.33   3.15   0.77  1.51   5.42   3.91  0.42
##      kurtosis    se
## mpg      -0.37  1.07
## hp      -0.14 12.12
## wt      -0.02  0.17
```

## Descriptive statistics by group

```
#aggregate(newdata, by=List(am=mtcars$am), mean)
```

```
dstat=function(x)apply(x,mystats)
by(newdata,mtcars$am,dstat)
```

```
## mtcars$am: 0
##              mpg              hp              wt
## n          19.00000000 19.00000000 19.00000000
## mean        17.14736842 160.26315789  3.7688947
## stdev         3.83396639  53.90819573  0.7774001
## skew          0.01395038 -0.01422519  0.9759294
## kurtosis     -0.80317826 -1.20969733  0.1415676
## -----
## mtcars$am: 1
##              mpg              hp              wt
## n          13.00000000 13.00000000 13.00000000
## mean        24.39230769 126.8461538  2.4110000
## stdev         6.16650381  84.0623243  0.6169816
## skew          0.05256118  1.3598859  0.2103128
## kurtosis     -1.45535200  0.5634635 -1.1737358
```

## Additional methods by group

```
require(doBy)
```

```
## Loading required package: doBy
```

```
summaryBy(mpg+hp+wt~am,data=mtcars,FUN=mystats)
```

```
##   am mpg.n mpg.mean mpg.stdev  mpg.skew mpg.kurtosis hp.n  hp.mean
## 1  0   19 17.14737  3.833966 0.01395038 -0.8031783  19 160.2632
## 2  1   13 24.39231  6.166504 0.05256118 -1.4553520  13 126.8462
```

```
##   hp.stdev      hp.skew hp.kurtosis wt.n  wt.mean  wt.stdev  wt.skew
## 1 53.90820 -0.01422519 -1.2096973  19 3.768895 0.7774001 0.9759294
## 2 84.06232 1.35988586  0.5634635  13 2.411000 0.6169816 0.2103128
##   wt.kurtosis
## 1    0.1415676
## 2   -1.1737358
```

```
require(psych)
describeBy(newdata, list(am=mtcars$am))
```

```
##
## Descriptive statistics by group
## am: 0
##      vars  n   mean    sd median trimmed   mad   min    max   range  skew
## mpg     1 19  17.15  3.83  17.30   17.12  3.11 10.40   24.40  14.00  0.01
## hp      2 19 160.26 53.91 175.00  161.06 77.10 62.00 245.00 183.00 -0.01
## wt      3 19   3.77  0.78   3.52   3.75  0.45  2.46   5.42   2.96  0.98
##      kurtosis    se
## mpg    -0.80  0.88
## hp     -1.21 12.37
## wt      0.14  0.18
## -----
## am: 1
##      vars  n   mean    sd median trimmed   mad   min    max   range  skew
## mpg     1 13  24.39  6.17  22.80   24.38  6.67 15.00   33.90  18.90  0.05
## hp      2 13 126.85 84.06 109.00  114.73 63.75 52.00 335.00 283.00 1.36
## wt      3 13   2.41  0.62   2.32   2.39  0.68  1.51   3.57   2.06  0.21
##      kurtosis    se
## mpg    -1.46  1.71
## hp      0.56 23.31
## wt     -1.17  0.17
```

## Frequency and contingency tables

```
require(vcd)
```

```
## Loading required package: vcd
```

```
## Loading required package: grid
```

```
head(Arthritis)
```

```
##   ID Treatment  Sex Age Improved
## 1 57   Treated Male  27     Some
## 2 46   Treated Male  29     None
## 3 77   Treated Male  30     None
## 4 17   Treated Male  32   Marked
## 5 36   Treated Male  46   Marked
## 6 23   Treated Male  58   Marked
```

```
names(Arthritis)
```

```
## [1] "ID"          "Treatment" "Sex"        "Age"        "Improved"

dim(Arthritis)

## [1] 84  5

str(Arthritis)

## 'data.frame':  84 obs. of  5 variables:
## $ ID      : int  57 46 77 17 36 23 75 39 33 55 ...
## $ Treatment: Factor w/ 2 levels "Placebo","Treated": 2 2 2 2 2 2 2 2 2 2 ...
## $ Sex      : Factor w/ 2 levels "Female","Male": 2 2 2 2 2 2 2 2 2 2 ...
## $ Age      : int  27 29 30 32 46 58 59 59 63 63 ...
## $ Improved : Ord.factor w/ 3 levels "None"<"Some"<...: 2 1 1 3 3 3 1 3 1 1 ...
```

## One-way tables

Use table function can make an one-way table easily.

```
mytable = with(Arthritis,table(Improved))
mytable

## Improved
##   None   Some Marked
##    42    14    28

mytable=table(Arthritis$Improved)
mytable

##
##   None   Some Marked
##    42    14    28
```

We can use prop.table to convert the count table generated from table function.

```
pstable=prop.table(mytable)*100
pstable

##
##   None   Some Marked
## 50.00000 16.66667 33.33333
```

## Two-way tables

Table function can be used to generate a two-way table and to generate a corresponding two-way proportional table. For example

```
mytable=with(Arthritis,table(Treatment,Improved))
mytable
```



```
##           Improved
## Treatment None Some Marked
## Placebo   29   7   7
## Treated   13   7  21

prop.table(mytable)

##           Improved
## Treatment      None      Some      Marked
## Placebo 0.34523810 0.08333333 0.08333333
## Treated 0.15476190 0.08333333 0.25000000
```

We can use `margin.table` function to generate a margin table for each row or column.

```
margin.table(mytable,1)

## Treatment
## Placebo Treated
##      43      41

margin.table(mytable,2)

## Improved
## None Some Marked
##  42  14  28

prop.table(mytable,1)

##           Improved
## Treatment      None      Some      Marked
## Placebo 0.6744186 0.1627907 0.1627907
## Treated 0.3170732 0.1707317 0.5121951

prop.table(mytable,2)

##           Improved
## Treatment      None      Some      Marked
## Placebo 0.6904762 0.5000000 0.2500000
## Treated 0.3095238 0.5000000 0.7500000

addmargins(prop.table(mytable,2),1)

##           Improved
## Treatment      None      Some      Marked
## Placebo 0.6904762 0.5000000 0.2500000
## Treated 0.3095238 0.5000000 0.7500000
## Sum      1.0000000 1.0000000 1.0000000
```

Another method for creating two-way tables is the `CrossTable()` function on the `gmodels` package. The `CrossTable()` function produces two-ways tables modeled after PROC FREQ in SAS or CROSSTABS in SPSS. The following listing showed an example.

```
require(gmodels)
```

```
## Loading required package: gmodels
```

```
mytable=with(Arthritis, CrossTable(Treatment,Improved))
```

```
##
```

```
##
```

```
## Cell Contents
```

```
## |-----|
## |                      N
## | Chi-square contribution
## |      N / Row Total
## |      N / Col Total
## |      N / Table Total
## |-----|
```

```
##
```

```
##
```

```
## Total Observations in Table:  84
```

```
##
```

```
##
```

Treatment	Improved			Row Total
	None	Some	Marked	
Placebo	29	7	7	43
	2.616	0.004	3.752	
	0.674	0.163	0.163	0.512
	0.690	0.500	0.250	
	0.345	0.083	0.083	
Treated	13	7	21	41
	2.744	0.004	3.935	
	0.317	0.171	0.512	0.488
	0.310	0.500	0.750	
	0.155	0.083	0.250	
Column Total	42	14	28	84
	0.500	0.167	0.333	

```
##
```

```
##
```

```
mytable
```

```
## $t
```

```
##      y
```

```
## x      None Some Marked
```

```
## Placebo  29   7    7
```

```
## Treated  13   7   21
```

```
##
```

```
## $prop.row
```

```
##      y
```

```
## x      None      Some      Marked
```

```
## Placebo 0.6744186 0.1627907 0.1627907
## Treated 0.3170732 0.1707317 0.5121951
##
## $prop.col
##      y
## x      None      Some      Marked
## Placebo 0.6904762 0.5000000 0.2500000
## Treated 0.3095238 0.5000000 0.7500000
##
## $prop.tbl
##      y
## x      None      Some      Marked
## Placebo 0.34523810 0.08333333 0.08333333
## Treated 0.15476190 0.08333333 0.25000000
```

## Tests of Independence

R provides several methods of testing the independence of categorical variables. The three tests described in this section are the chi-square test of independence, the Fisher exact test, and the Cochran-Mantel-Haenszel test.

### Chi-square test

```
mytable=with(Arthritis, table(Treatment,Improved))
chisq.test(mytable)
```

```
##
## Pearson's Chi-squared test
##
## data: mytable
## X-squared = 13.055, df = 2, p-value = 0.001463
```

```
mytable=with(Arthritis, table(Treatment,Sex))
chisq.test(mytable)
```

```
##
## Pearson's Chi-squared test with Yates' continuity correction
##
## data: mytable
## X-squared = 0.38378, df = 1, p-value = 0.5356
```

### Fisher's exact test

```
mytable=with(Arthritis, table(Treatment,Improved))
fisher.test(mytable)
```

```
##
## Fisher's Exact Test for Count Data
##
## data: mytable
## p-value = 0.001393
## alternative hypothesis: two.sided
```

```
mytable=with(Arthritis, table(Treatment,Sex))
fisher.test(mytable)

##
## Fisher's Exact Test for Count Data
##
## data: mytable
## p-value = 0.4763
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
## 0.5320442 4.3286798
## sample estimates:
## odds ratio
## 1.500984
```

### COCHRAN-MANTEL-HAENSZEL test

```
mytable=with(Arthritis, table(Treatment,Improved,Sex))
mantelhaen.test(mytable)

##
## Cochran-Mantel-Haenszel test
##
## data: mytable
## Cochran-Mantel-Haenszel M^2 = 14.632, df = 2, p-value = 0.0006647
```

### Measures of association

```
require(vcd)
mytable=with(Arthritis, table(Treatment,Improved))
assocstats(mytable)

##              X^2 df  P(> X^2)
## Likelihood Ratio 13.530  2 0.0011536
## Pearson          13.055  2 0.0014626
##
## Phi-Coefficient   : NA
## Contingency Coeff.: 0.367
## Cramer's V        : 0.394
```

### Covariance and Correlation

```
states=state.x77[,1:6]

cov(states)

##              Population      Income  Illiteracy  Life Exp  Murder
## Population 19931683.7588 571229.7796 292.8679592 -407.8424612 5663.523714
## Income      571229.7796 377573.3061 -163.7020408 280.6631837 -521.894286
## Illiteracy   292.8680 -163.7020 0.3715306 -0.4815122 1.581776
## Life Exp    -407.8425 280.6632 -0.4815122 1.8020204 -3.869480
## Murder      5663.5237 -521.8943 1.5817755 -3.8694804 13.627465
## HS Grad     -3551.5096 3076.7690 -3.2354694 6.3126849 -14.549616
```

```
##              HS Grad
## Population -3551.509551
## Income      3076.768980
## Illiteracy  -3.235469
## Life Exp     6.312685
## Murder      -14.549616
## HS Grad      65.237894
```

```
cor(states)
```

```
##              Population      Income Illiteracy      Life Exp      Murder
## Population  1.00000000  0.2082276  0.1076224 -0.06805195  0.3436428
## Income      0.20822756  1.0000000 -0.4370752  0.34025534 -0.2300776
## Illiteracy  0.10762237 -0.4370752  1.0000000 -0.58847793  0.7029752
## Life Exp    -0.06805195  0.3402553 -0.5884779  1.00000000 -0.7808458
## Murder      0.34364275 -0.2300776  0.7029752 -0.78084575  1.0000000
## HS Grad     -0.09848975  0.6199323 -0.6571886  0.58221620 -0.4879710
##              HS Grad
## Population -0.09848975
## Income      0.61993232
## Illiteracy -0.65718861
## Life Exp    0.58221620
## Murder     -0.48797102
## HS Grad     1.00000000
```

```
cor(states, method="spearman")
```

```
##              Population      Income Illiteracy      Life Exp      Murder
## Population  1.00000000  0.1246098  0.3130496 -0.1040171  0.3457401
## Income      0.1246098  1.0000000 -0.3145948  0.3241050 -0.2174623
## Illiteracy  0.3130496 -0.3145948  1.0000000 -0.5553735  0.6723592
## Life Exp    -0.1040171  0.3241050 -0.5553735  1.0000000 -0.7802406
## Murder      0.3457401 -0.2174623  0.6723592 -0.7802406  1.0000000
## HS Grad     -0.3833649  0.5104809 -0.6545396  0.5239410 -0.4367330
##              HS Grad
## Population -0.3833649
## Income      0.5104809
## Illiteracy -0.6545396
## Life Exp    0.5239410
## Murder     -0.4367330
## HS Grad     1.0000000
```

```
x = states[,c("Population", "Income", "Illiteracy", "HS Grad")]
y = states[,c("Life Exp", "Murder")]
```

```
cor(x,y)
```

```
##              Life Exp      Murder
## Population -0.06805195  0.3436428
## Income      0.34025534 -0.2300776
```

```
## Illiteracy -0.58847793  0.7029752
## HS Grad    0.58221620 -0.4879710

cor.test(states[,1],states[,3])

##
## Pearson's product-moment correlation
##
## data:  states[, 1] and states[, 3]
## t = 0.74999, df = 48, p-value = 0.4569
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## -0.1759976  0.3747441
## sample estimates:
##      cor
## 0.1076224

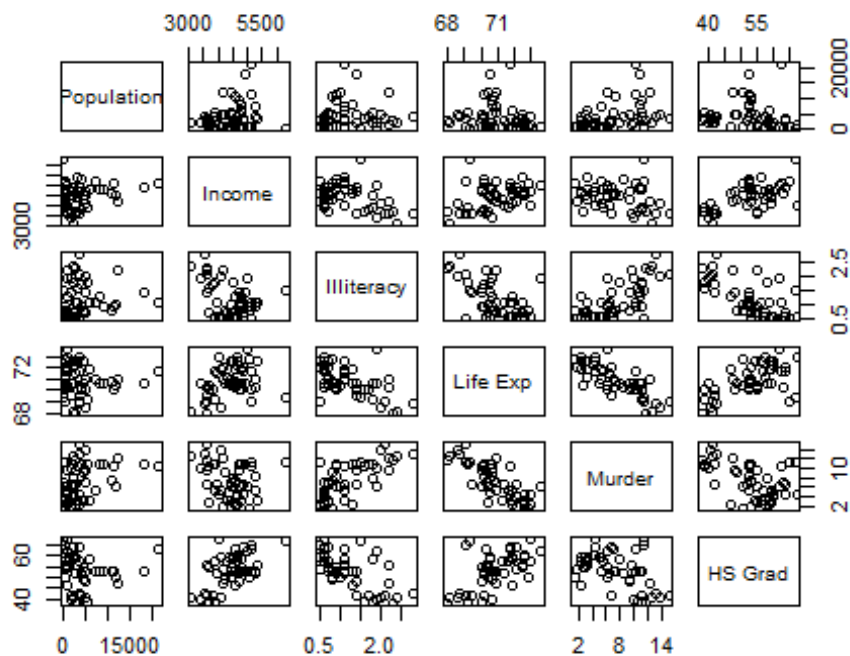
require(psych)
corr.test(states,use="complete")

## Call:corr.test(x = states, use = "complete")
## Correlation matrix
##      Population Income Illiteracy Life Exp Murder HS Grad
## Population      1.00  0.21      0.11  -0.07  0.34  -0.10
## Income          0.21  1.00     -0.44   0.34 -0.23  0.62
## Illiteracy      0.11 -0.44      1.00  -0.59  0.70  -0.66
## Life Exp       -0.07  0.34     -0.59   1.00 -0.78  0.58
## Murder         0.34 -0.23      0.70  -0.78  1.00  -0.49
## HS Grad       -0.10  0.62     -0.66   0.58 -0.49  1.00
## Sample Size
## [1] 50
## Probability values (Entries above the diagonal are adjusted for multiple
## tests.)
##      Population Income Illiteracy Life Exp Murder HS Grad
## Population      0.00  0.59      1.00   1.0  0.10      1
## Income          0.15  0.00      0.01   0.1  0.54      0
## Illiteracy      0.46  0.00      0.00   0.0  0.00      0
## Life Exp       0.64  0.02      0.00   0.0  0.00      0
## Murder         0.01  0.11      0.00   0.0  0.00      0
## HS Grad        0.50  0.00      0.00   0.0  0.00      0
##
## To see confidence intervals of the correlations, print with the
short=FALSE option
```

## Visualizing correlations

We may use the pairs function to visualize the correlations among different examples

```
pairs(states)
```



## t-tests

The most common activity in research is the comparison of two groups. There are two types of t-tests. The 1st one is independent t-test and the second one is paired t-test.

```
require(MASS)

## Loading required package: MASS

t.test(Prob~So, data=UScrime)

##
## Welch Two Sample t-test
##
## data: Prob by So
## t = -3.8954, df = 24.925, p-value = 0.0006506
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.03852569 -0.01187439
## sample estimates:
## mean in group 0 mean in group 1
## 0.03851265 0.06371269

Input = ("
Time Student Score
Before a 65
Before b 75
```

```

Before c      86
Before d      69
Before e      60
Before f      81
Before g      88
Before h      53
Before i      75
Before j      73
After  a      77
After  b      98
After  c      92
After  d      77
After  e      65
After  f      77
After  g     100
After  h      73
After  i      93
After  j      75
")

```

```

dat1 = read.table(textConnection(Input),header=TRUE)
dat1=with(dat1,dat1[order(Time,Student),])

```

```

library(psych)

```

```

headTail(dat1)

```

```

##      Time Student Score
## 11  After      a      77
## 12  After      b      98
## 13  After      c      92
## 14  After      d      77
## ... <NA>    <NA>    ...
## 7   Before      g      88
## 8   Before      h      53
## 9   Before      i      75
## 10  Before      j      73

```

```

str(dat1)

```

```

## 'data.frame':  20 obs. of  3 variables:
## $ Time   : Factor w/ 2 levels "After","Before": 1 1 1 1 1 1 1 1 1 1 ...
## $ Student: Factor w/ 10 levels "a","b","c","d",...: 1 2 3 4 5 6 7 8 9 10
## ...
## $ Score  : int  77 98 92 77 65 77 100 73 93 75 ...

```

```

summary(dat1)

```

```

##      Time      Student      Score
## After :10  a          :2  Min.    : 53.0
## Before:10  b          :2  1st Qu.: 72.0

```



```

##           c      :2   Median : 76.0
##           d      :2   Mean    : 77.6
##           e      :2   3rd Qu.: 86.5
##           f      :2   Max.    :100.0
##          (Other):8

t.test(Score ~ Time,
       data=dat1,
       paired = TRUE,
       conf.level = 0.95)

##
## Paired t-test
##
## data:  Score by Time
## t = 3.8084, df = 9, p-value = 0.004163
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
##   4.141247 16.258753
## sample estimates:
## mean of the differences
##                   10.2

```