

Introduction to R

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Pocket calculator

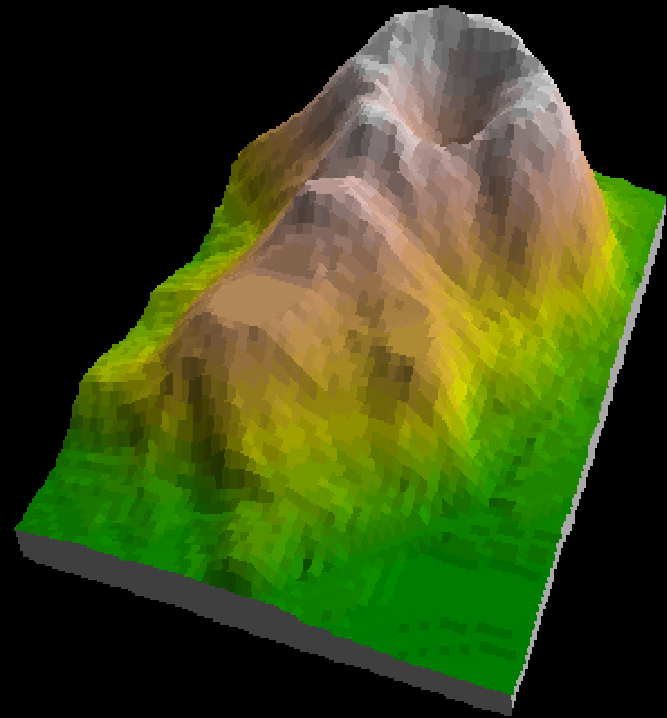
Data manipulation

Graphics

Pocket calculator

Data manipulation

Graphics



Pocket calculator

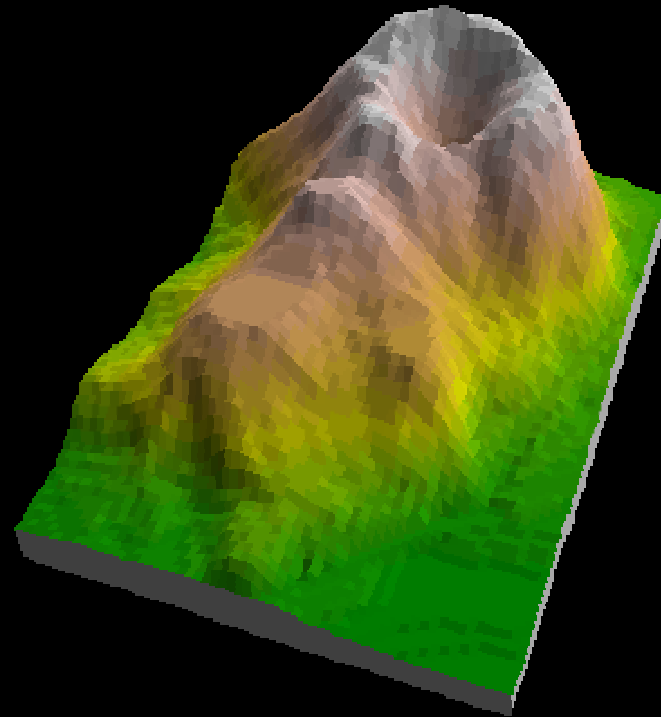
Data manipulation

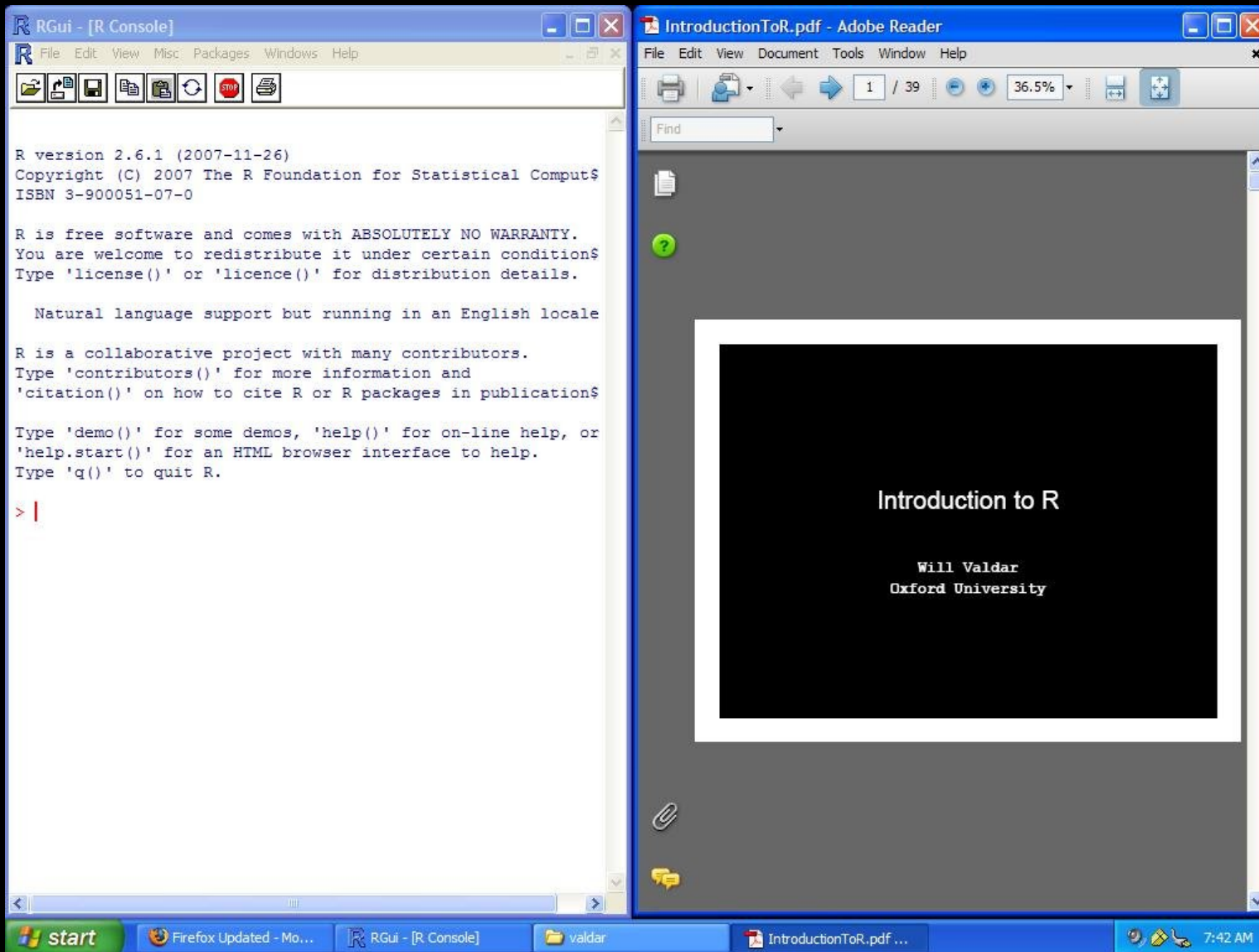
Graphics

Statistical modelling

Computationally
Intensive Statistics

Programming





R is a big calculator

Simple operations

`3*4`

`20 + 4`

`20/4`

`9^2`

`log(100, base=10)`

`2.5e2 / 2.5`

R is a big calculator

Simple operations

`3*4`

`20 + 4`

`20/4`

`9^2`

`log(100, base=10)`

`2.5e2 / 2.5`

Expressions

`(3*4)/(20+4)`

`3*(4/20)+4`

`(5/9)*(48-32)`

R is a big calculator

Variables

```
temp.f <- 48
```

```
temp.f
```

```
(5/9)*(temp.f-32)
```

```
temp.c <- (5/9)*(temp.f-32)
```

```
temp.c
```


R is a big calculator

Variables

```
temp.f <- 48
temp.f
(5/9)*(temp.f-32)
temp.c <- (5/9)*(temp.f-32)
temp.c
```

Write your own functions

```
will.fun <- function(x) { (5/9)*(x-32) }
will.fun(48)
will.fun(temp.f)
thing <- will.fun(26)
```

Data types

```
class(will.fun)
```

```
class(temp.c)
```

```
my.name <- "Will"
```

```
class(my.name)
```

Data types

```
class(will.fun)
```

```
class(temp.c)
```

```
my.name <- "Will"
```

```
class(my.name)
```

Vectors

```
a <- c("Will", "Ben")
```

```
x <- c(1,2,3)
```

```
sum(x)
```

```
mean(x)
```

```
range(x)
```

```
var(x)
```

```
1:3
```

Matrices

$$\mathbf{A} = \begin{bmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \\ 7 & 8 & 9 \end{bmatrix}, \quad \mathbf{B} = \begin{bmatrix} 5 & 1 & 1 \\ 5 & 2 & 14 \\ 5 & 3 & -1 \end{bmatrix}, \quad \mathbf{x} = \begin{bmatrix} 7 \\ 8 \\ 9 \end{bmatrix}$$

```
A <- matrix(c(1:9), byrow=TRUE, ncol=3)
B <- cbind(c(5,5,5), 1:3, c(1,14,-1))
x <- c(7,8,9)
```

Matrices

$$\mathbf{A} = \begin{bmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \\ 7 & 8 & 9 \end{bmatrix}, \quad \mathbf{B} = \begin{bmatrix} 5 & 1 & 1 \\ 5 & 2 & 14 \\ 5 & 3 & -1 \end{bmatrix}, \quad \mathbf{x} = \begin{bmatrix} 7 \\ 8 \\ 9 \end{bmatrix}$$

```
A <- matrix(c(1:9), byrow=TRUE, ncol=3)
B <- cbind(c(5,5,5), 1:3, c(1,14,-1))
x <- c(7,8,9)
```

```
t(A)
```

```
diag(A)
```

```
A %*% B
```

```
A %*% t(B)
```

```
B.inv <- solve(B)
```

```
det(B)
```

```
median(x)
```

```
range(B)
```

```
max(x)
```

```
A + 1
```

```
chol(A+1)
```

Matrices

$$\mathbf{A} = \begin{bmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \\ 7 & 8 & 9 \end{bmatrix}, \quad \mathbf{B} = \begin{bmatrix} 5 & 1 & 1 \\ 5 & 2 & 14 \\ 5 & 3 & -1 \end{bmatrix}, \quad \mathbf{x} = \begin{bmatrix} 7 \\ 8 \\ 9 \end{bmatrix}$$

Subscripting is easy

`A[1,1]`

`A[3,2]`

`x[2]`

`A[3,]`

`A[,3]`

`A[3, c(1,2)]`

`A[, c(1,2)]`

`A[2,2] <- 1975`

Exercises 1

$$\mathbf{A} = \begin{bmatrix} 1 & 2 \\ 3 & 4 \end{bmatrix}, \quad \mathbf{B} = \begin{bmatrix} 11 & 12 \\ 21 & 22 \end{bmatrix}, \quad \mathbf{x} = \begin{bmatrix} 4 \\ 5.2 \end{bmatrix}$$

- 1) $(3.5 + 6) \times (7.9)^2$
- 2) $\mathbf{A} + \mathbf{B}$
- 3) \bar{x}
- 4) Convert \mathbf{B} from Farenheit to Centigrade
- * 5) $|\mathbf{AB}^T|$
- * 6) $\mathbf{A}^{-1}\mathbf{Bx}$
- * 7) $\mathbf{BA}(\mathbf{xx}^T)$

Answers 1

```
A <- matrix(1:4, byrow=TRUE, ncol=2)
```

```
B <- cbind(c(11,21), c(12,22))
```

```
x <- c(4, 5.2)
```

1) $(3.5+6)*((7.9)^2)$

2) $A + B$

3) $\text{mean}(x)$

4) $(5/9)*(B-32)$

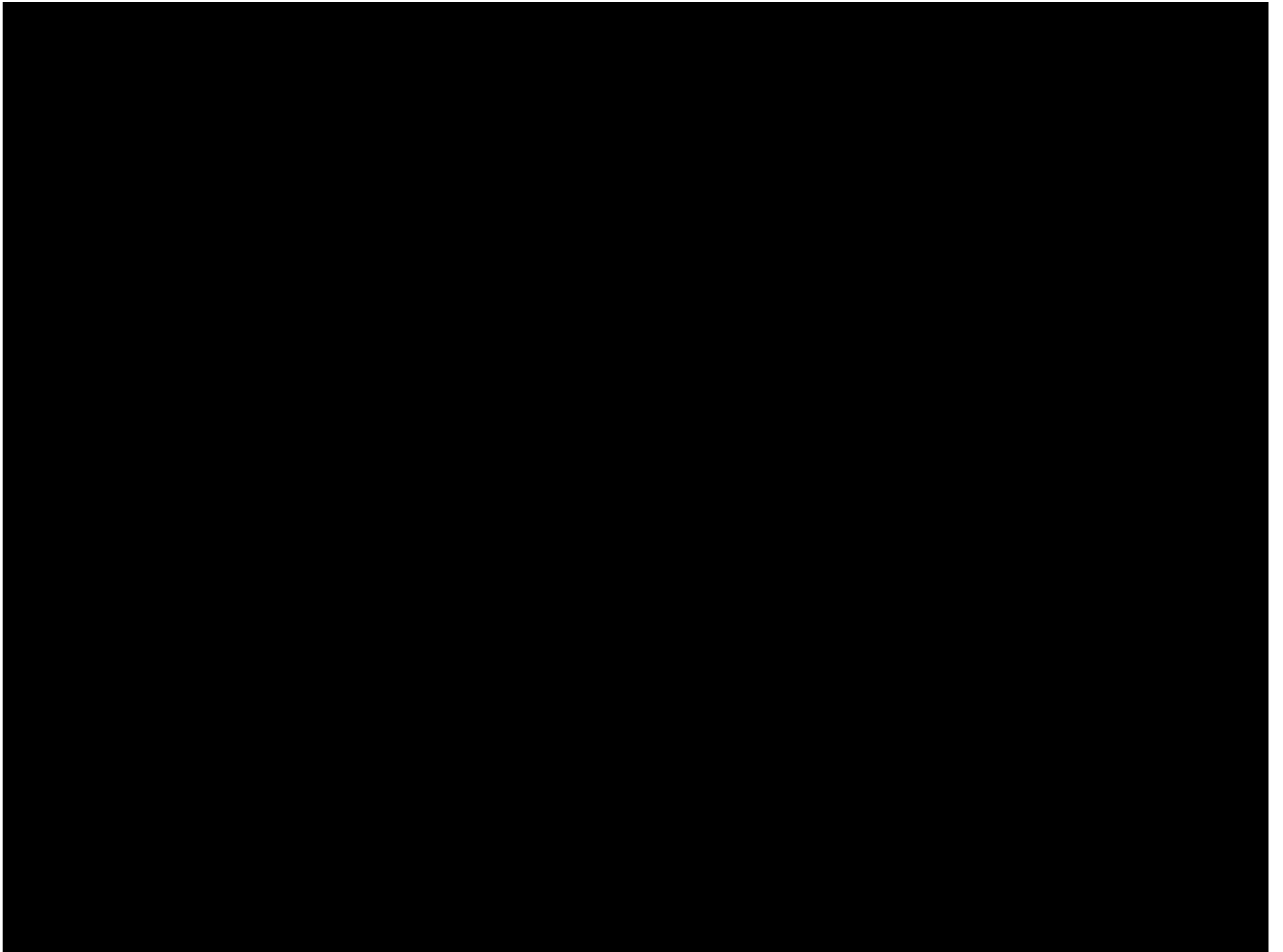
or

```
will.fun(B)
```

5) $\det(A \%*\% t(B))$

6) $\text{solve}(A) \%*\% B \%*\% x$

7) $B \%*\% A \%*\% (x \%*\% t(x))$



	A	B	C	D	E	F	G	H	I
1	sex	weight	country	age	snp.A	snp.B	family	phenotype	status
2	F	91.29	Europe	42	0	1	Family3	-2.6134576	FALSE
3	M	154.22	US	42	1	2	Family6	-7.7237116	FALSE
4	F	95.74	US	40	2	0	Family1	3.2361416	TRUE
5	F	110.25	Australia	30	0	1	Family10	-0.026201	TRUE
6	M	170.47	Australia	36	0	2	Family1	-11.132778	FALSE
7	M	127.57	US	45	1	1	Family6	-3.6016196	FALSE
8	M	125.56	Europe	44	1	1	Family9	-7.0376314	FALSE

data.frame

	A	B	C	D	E	F	G	H	I
1	sex	weight	country	age	snp.A	snp.B	family	phenotype	status
2	F	91.29	Europe	42	0	1	Family3	-2.6134576	FALSE
3	M	154.22	US	42	1	2	Family6	-7.7237116	FALSE
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8	M	125.56	Europe	44	1	1	Family9	-7.0376314	FALSE

```
data <- read.delim("F:valdar/disease.txt")  
class(data)  
head(data)
```

data.frame

	A	B	C	D	E	F	G	H	I
1	sex	weight	country	age	snp.A	snp.B	family	phenotype	status
2	F	91.29	Europe	42	0	1	Family3	-2.6134576	FALSE
3	M	154.22	US	42	1	2	Family6	-7.7237116	FALSE
4	F	95.74	US	40	2	0	Family1	3.2361416	TRUE
5	F	110.25	Australia	30	0	1	Family10	-0.026201	TRUE
6	M	170.47	Australia	36	0	2	Family1	-11.132778	FALSE
7	M	127.57	US	45	1	1	Family6	-3.6016196	FALSE
8	M	125.56	Europe	44	1	1	Family9	-7.0376314	FALSE

```
sex weight  country age snp.A snp.B  family  phenotype status
1   F   91.29   Europe  42     0     1  Family3  -2.61345764  FALSE
2   M  154.22     US    42     1     2  Family6  -7.72371165  FALSE
3   F   95.74     US    40     2     0  Family1   3.23614160   TRUE
4   F  110.25 Australia  30     0     1  Family10 -0.02620101  TRUE
5   M  170.47 Australia  36     0     2  Family1 -11.13277756  FALSE
6   M  127.57     US    45     1     1  Family6  -3.60161956  FALSE
```

a variable with other variables hanging off it

```
str(data)
```

```
'data.frame':  400 obs. of  9 variables:
 $ sex      : Factor w/ 2 levels "F","M": 1 2 1 1 2 ...
 $ weight   : num   91.3 154.2  95.7 110.2 170.5 ...
 $ country  : Factor w/ 3 levels "Australia","Europe",...: 2 3 ...
 $ age      : int   42 42 40 30 36 45 44 38 40 34 ...
 $ snp.A    : int   0 1 2 0 0 1 1 0 2 1 ...
 $ snp.B    : int   1 2 0 1 2 1 1 0 1 1 ...
 $ family   : Factor w/ 10 levels "Family1",...: 4 7 ...
 $ phenotype: num  -2.6135 -7.7237  3.2361 ...
 $ status   : logi  FALSE FALSE  TRUE  TRUE FALSE FALSE ...
```

a variable with other variables hanging off it

```
str(data)
'data.frame':  400 obs. of  9 variables:
 $ sex      : Factor w/ 2 levels "F","M": 1 2 1 1 2  ...
 $ weight   : num   91.3 154.2  95.7 110.2 170.5  ...
 $ country  : Factor w/ 3 levels "Australia","Europe",...: 2 3  ...
 $ age      : int   42 42 40 30 36 45 44 38 40 34  ...
 $ snp.A    : int   0 1 2 0 0 1 1 0 2 1  ...
 $ snp.B    : int   1 2 0 1 2 1 1 0 1 1  ...
 $ family   : Factor w/ 10 levels "Family1",...: 4 7  ...
 $ phenotype: num   -2.6135  -7.7237   3.2361  ...
 $ status   : logi  FALSE FALSE  TRUE  TRUE FALSE FALSE ...
```

```
data$weight
```

```
data$weight[3:10]
```

```
mean(data$weight)
```

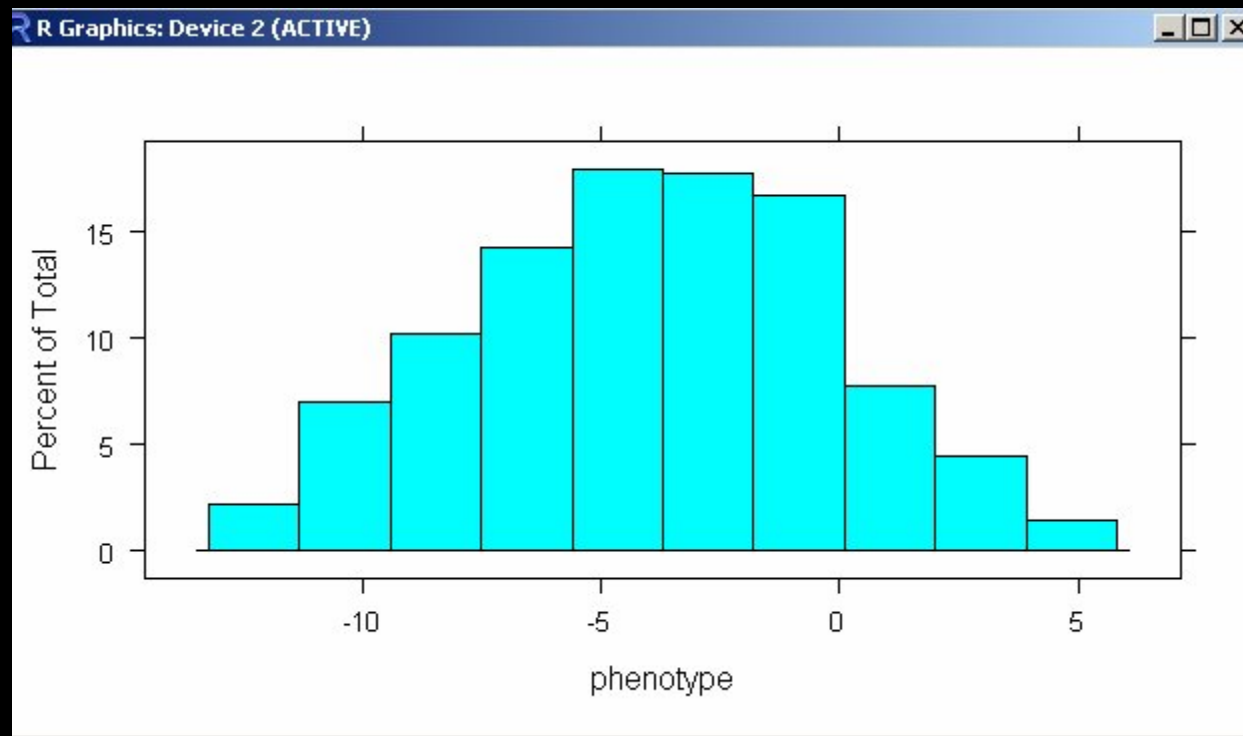
```
library(lattice)
```


histogram()

```
histogram( ~ data$phenotype )
```

or

```
histogram( ~ phenotype, data=data )
```

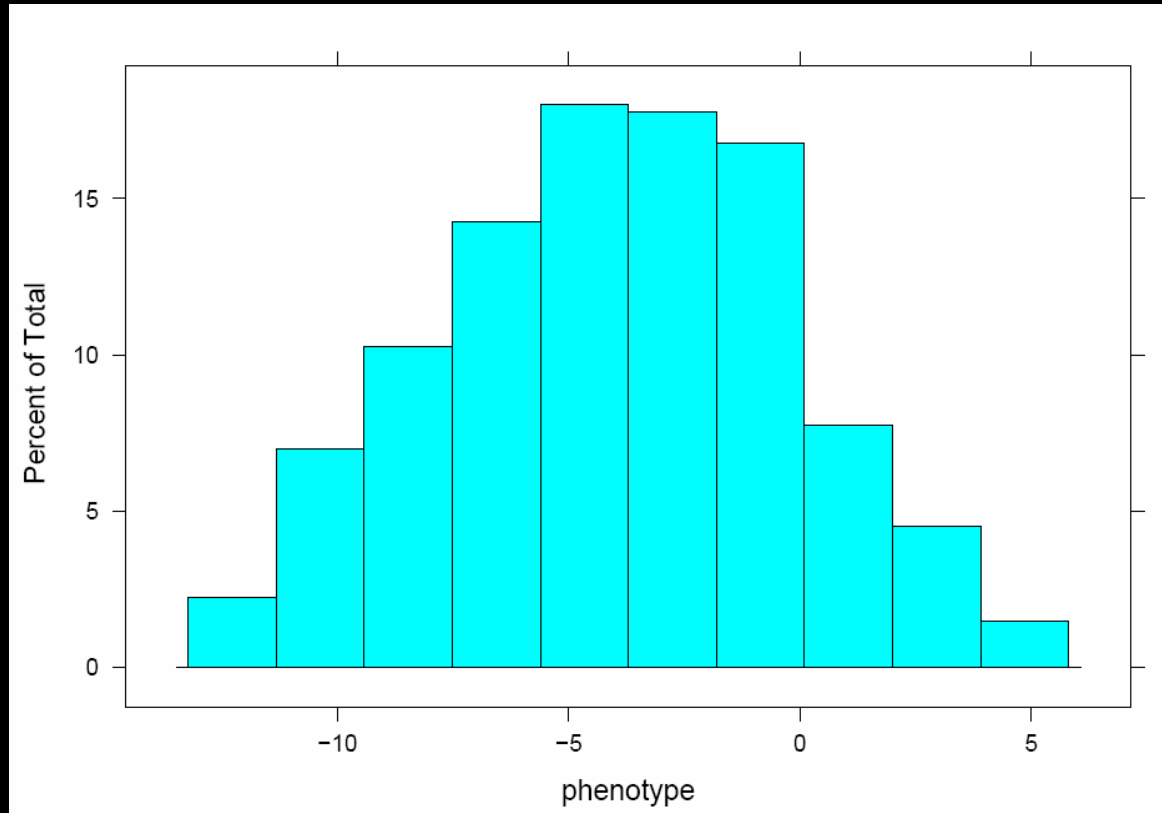


histogram()

```
pdf("c:myplot.pdf", width=7, height=5)
```

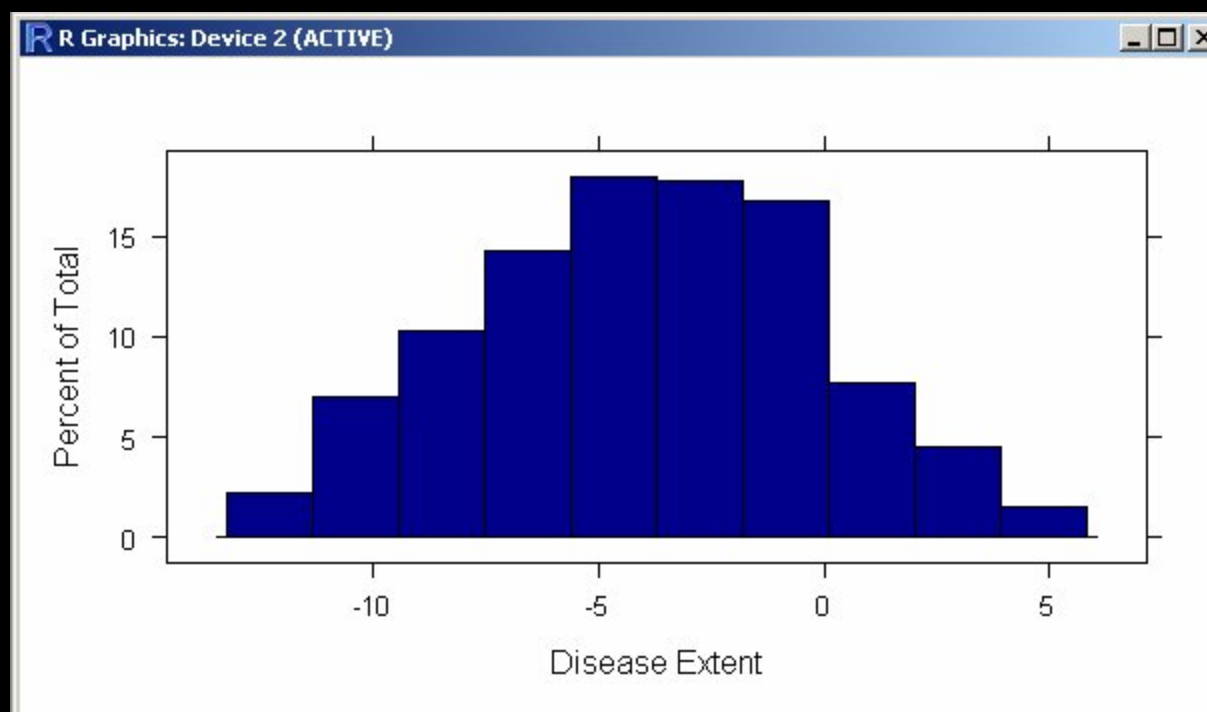
```
histogram( ~ phenotype, data=data)
```

```
dev.off()
```



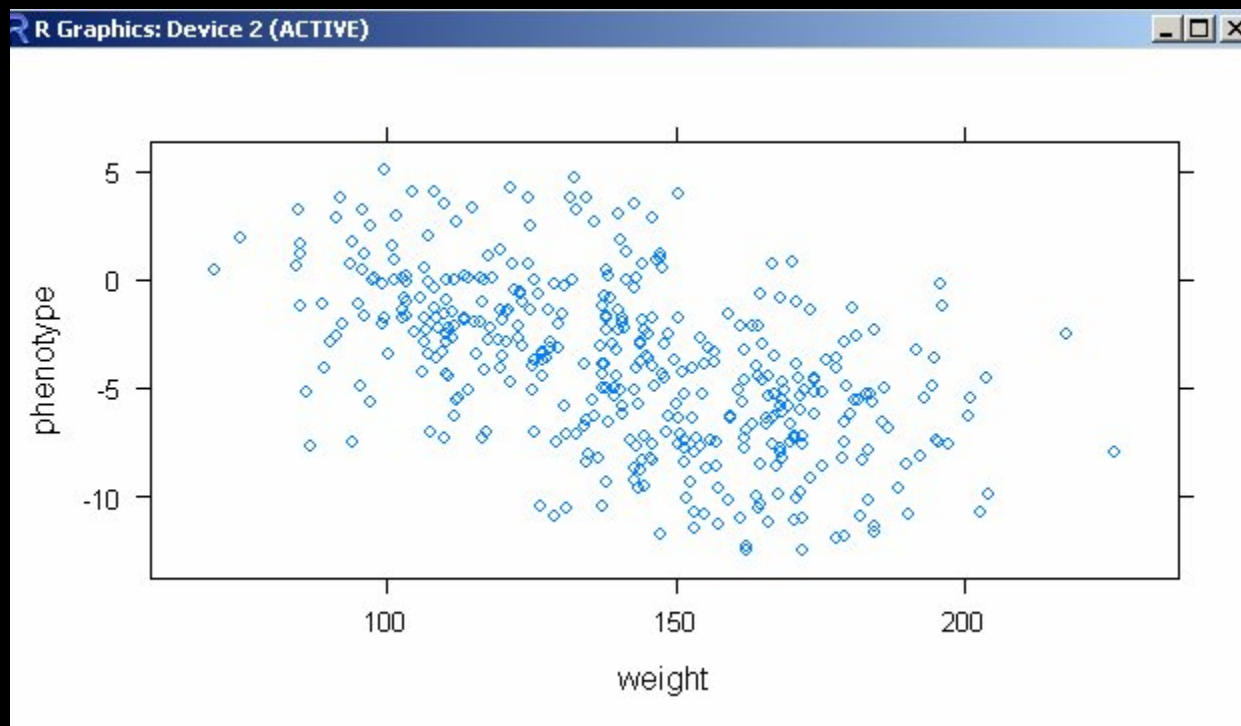
histogram()

```
histogram( ~ phenotype, data=data, col="darkblue",  
          xlab="Disease Extent")
```



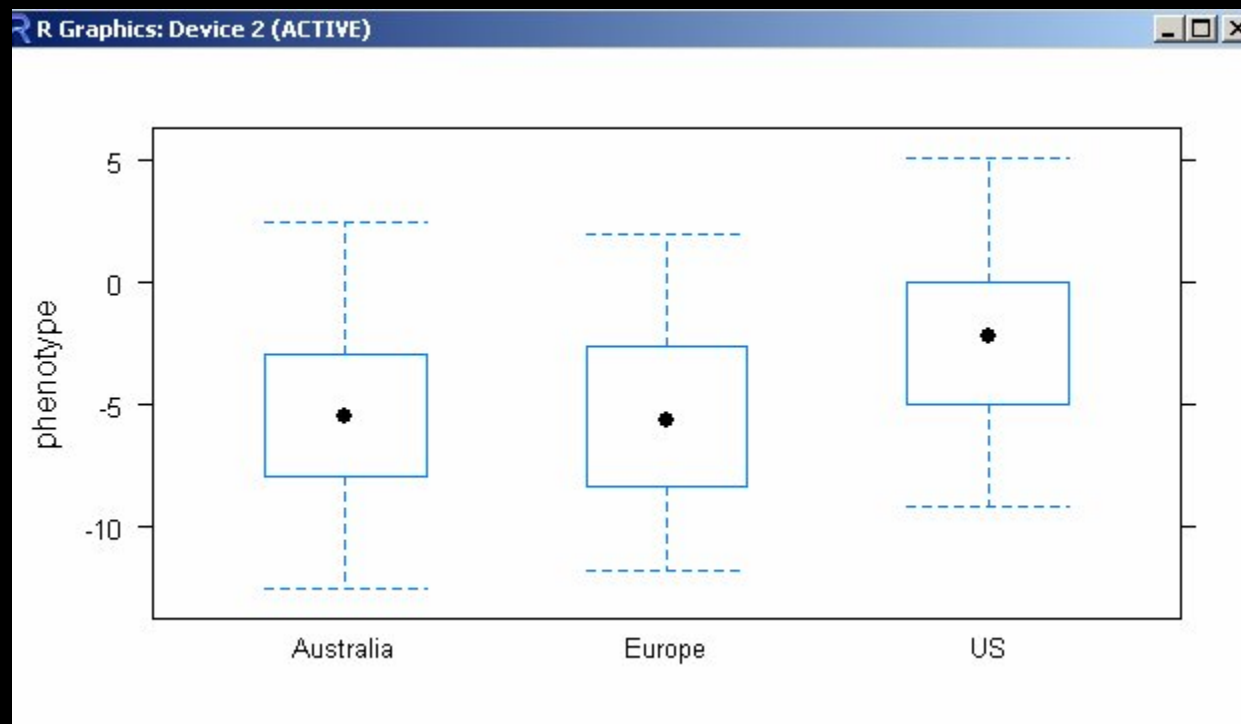
xyplot()

```
xyplot( phenotype ~ weight, data=data)
```

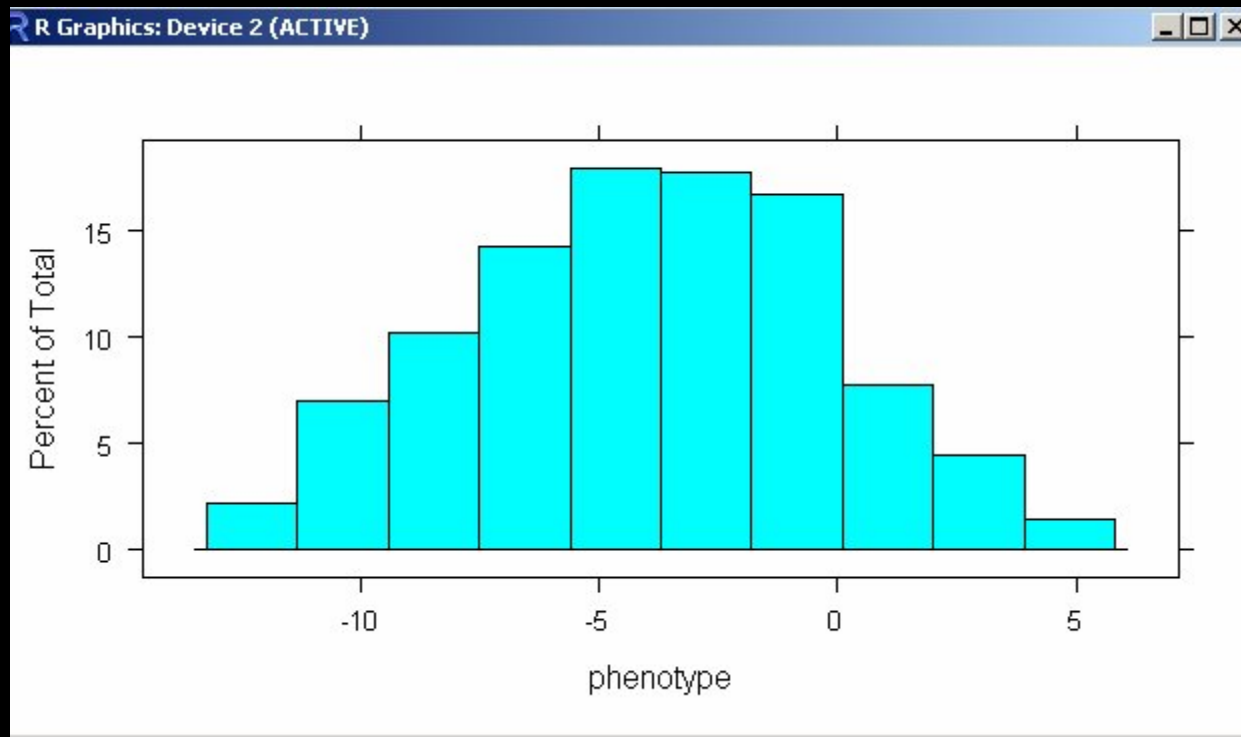


bwplot() , ie, box and whisker plot

```
bwplot( phenotype ~ country, data=data)
```

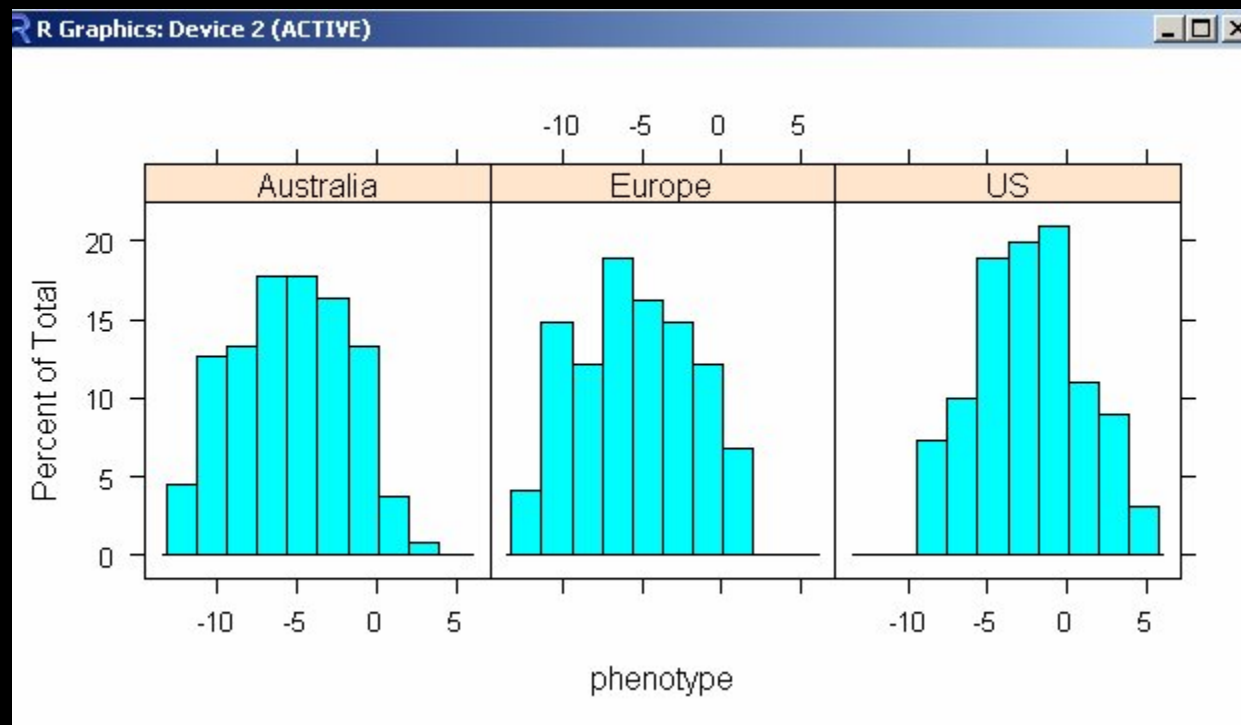


```
histogram( ~ phenotype, data=data)
```



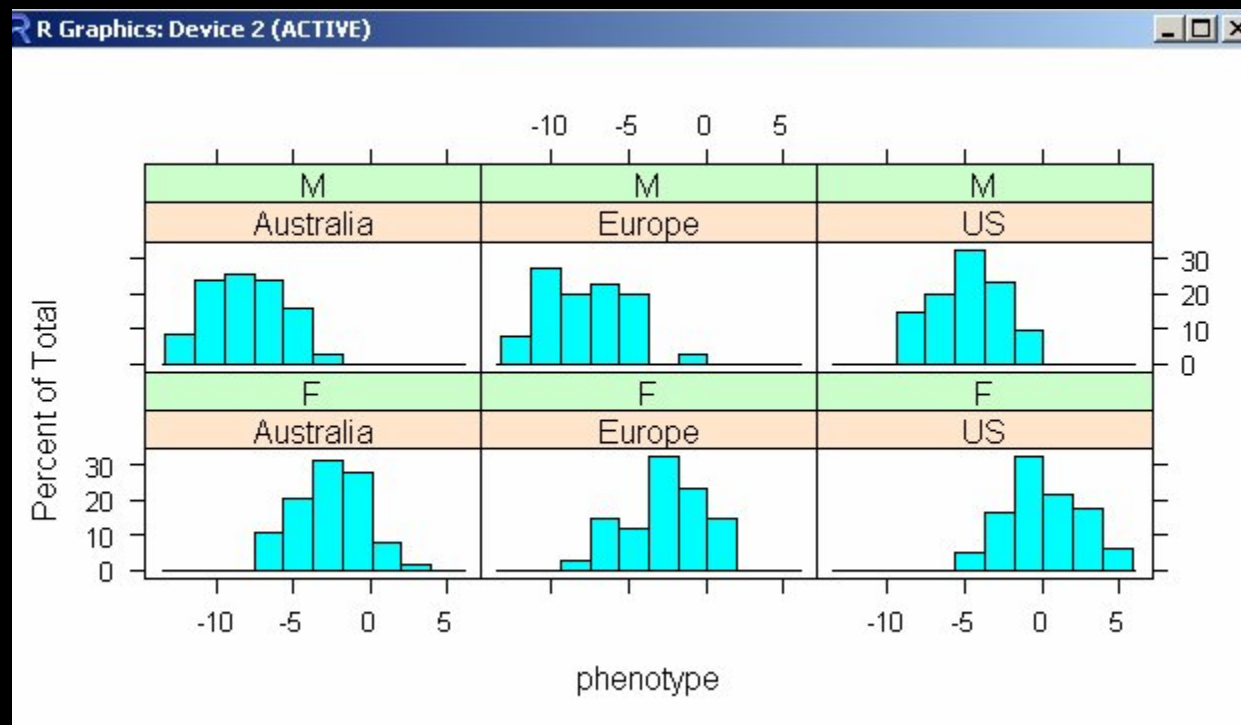
Conditioning

```
histogram( ~ phenotype | country, data=data)
```



Conditioning

```
histogram( ~ phenotype | sex * country, data=data)
```



Conditioning

Examples:

```
xypplot( phenotype ~ weight | sex, data=data)
```

```
bwplot( phenotype ~ country | snp.A, data=data)
```

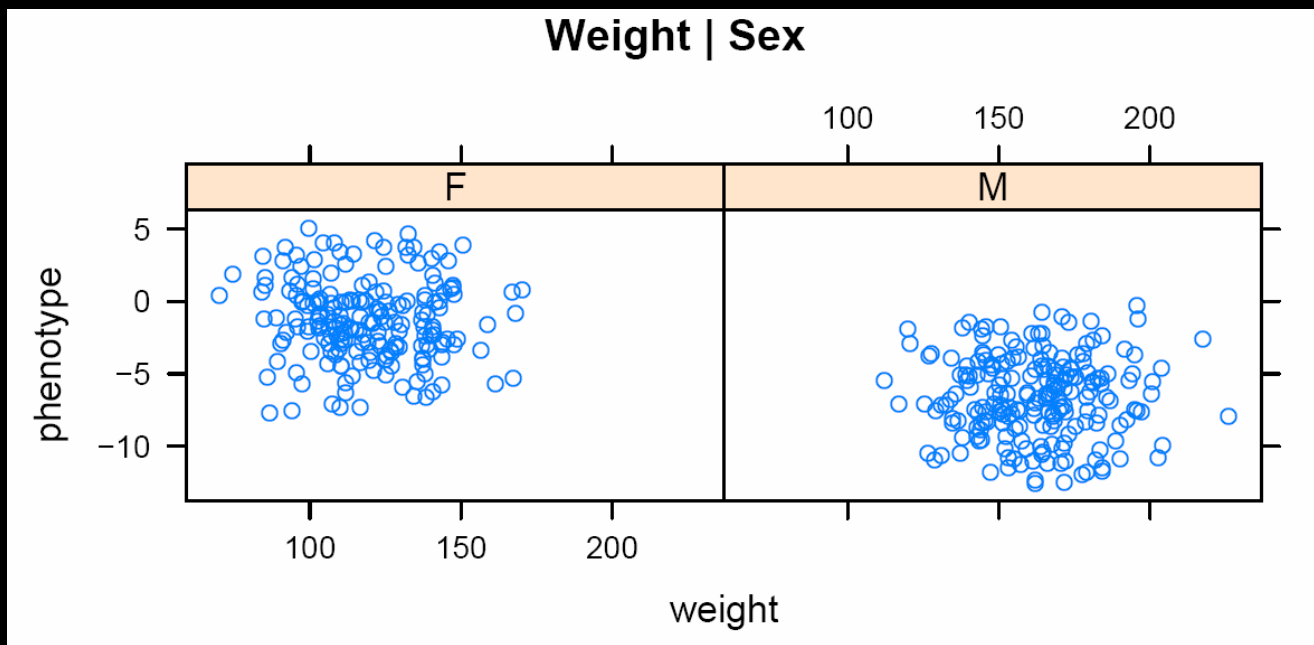
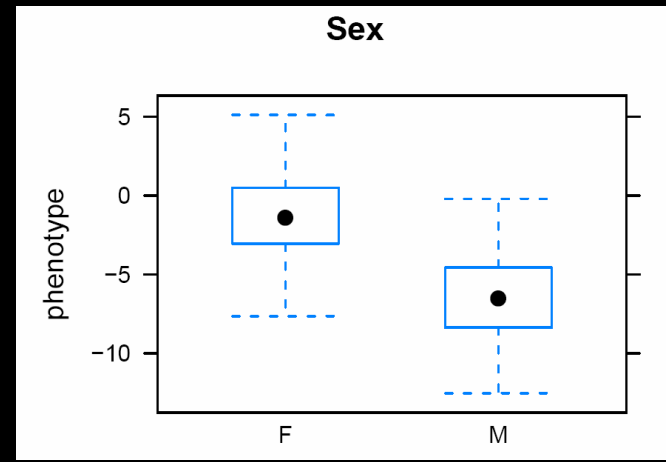
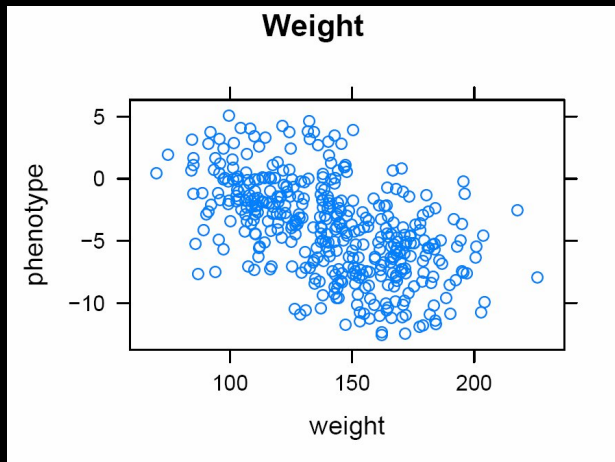
Exercises 2

- 1) What do you think the status variable means and how is it defined? (*conditional histogram*)
- 2) How does `snp.B` affect the phenotype? (*cond. hist*)
- 3) Is there an effect of age on the phenotype? Are you sure? (*conditional xyplot*)
- 4) Does weight affect the phenotype? Does sex? Does sex affect weight? Does weight affect sex? What's going on here? (*conditional xyplot*)

Answers 2

- 1) `histogram(~ phenotype | status, data=data)`
- 2) `xyplot(phenotype ~ snp.B, data=data)`
`histogram(~ phenotype | snp.B, data=data)`
- 3) `xyplot(phenotype ~ age, data=data)`
`xyplot(phenotype ~ age | sex, data=data)`
- 4) `bwplot(phenotype ~ sex, data=data)`
`xyplot(phenotype ~ weight, data=data)`
`xyplot(phenotype ~ weight | sex, data=data)`

Answers 2: question 4



Further reading

Books:

Introductory Statistics with R

by Peter Dalgaard

Modern Applied Statistics with S

by Venables and Ripley

Web sites:

<http://www.r-project.org/>

<http://wiki.r-project.org/rwiki/doku.php>

<http://news.gmane.org/gmane.comp.lang.r.general>

Graph gallery:

<http://addictedtor.free.fr/graphiques/>