

In *R* numbers can be used in vectors and in matrices. We begin by creating a vector and then doing some elementary operations on it.

```
> w<-c(1,2,4)
> w[3]<-3
> w

[1] 1 2 3

> w1<-c(w,4)+4
> w1

[1] 5 6 7 8

> w2<-1:4
> w1*w2

[1] 5 12 21 32
```

Next we generate a random sample of size 50 from a normal distribution with mean 100 and standard deviation 7 and apply some of *R*'s built in functions to the result.

```
> x<-rnorm(50,100,7)
> x[c(1,3)]

[1] 94.62999 101.39920

> mean(x)

[1] 99.25023

> var(x)

[1] 62.40661

> min(x)

[1] 82.3573

> median(x)

[1] 98.77586

> quantile(x)

      0%      25%      50%      75%     100%
82.35730 94.87472 98.77586 105.88346 111.89912

> as.numeric(quantile(x,c(0.25,0.6)))
```

```
[1] 94.87472 99.44699
```

In the above `hist(x)` would make a histogram of the values in x .

In the next bit of code we generate observations from 5 independent binomial distributions and play around with matrix notation.

```
> N<-c(10,20,30,40,50)
> p<-seq(0.1,0.9,length=5)
> y<-rbinom(5,N,p)
> M1<-rbind(N,y)
> dim(M1)
```

```
[1] 2 5
```

```
> M1[2,]
```

```
[1] 1 6 13 30 46
```

```
> M1[2,5]
```

```
  y
46
```

```
> apply(M1,1,mean)
```

```
  N  y
30.0 19.2
```

Suppose 5 subjects in an experiment received the treatment and 4 belonged to the control group. In the following x identifies which group the subject belonged to and y is the measurement of interest.

```
> x<-c(1,1,2,1,2,2,1,2,2)
> y<-c(11,9,15,14,11,14,12,15,16)
> y[x==1]
```

```
[1] 11 9 14 12
```

```
> dum<-split(y,x) #this creates a list
> length(dum)
```

```
[1] 2
```

```
> dum[[1]]
```

```
[1] 11 9 14 12
```

```
> dum[[1]][2]
```

```
[1] 9
```

```
> sapply(dum,mean)
```

```
 1  2
11.5 14.2
```