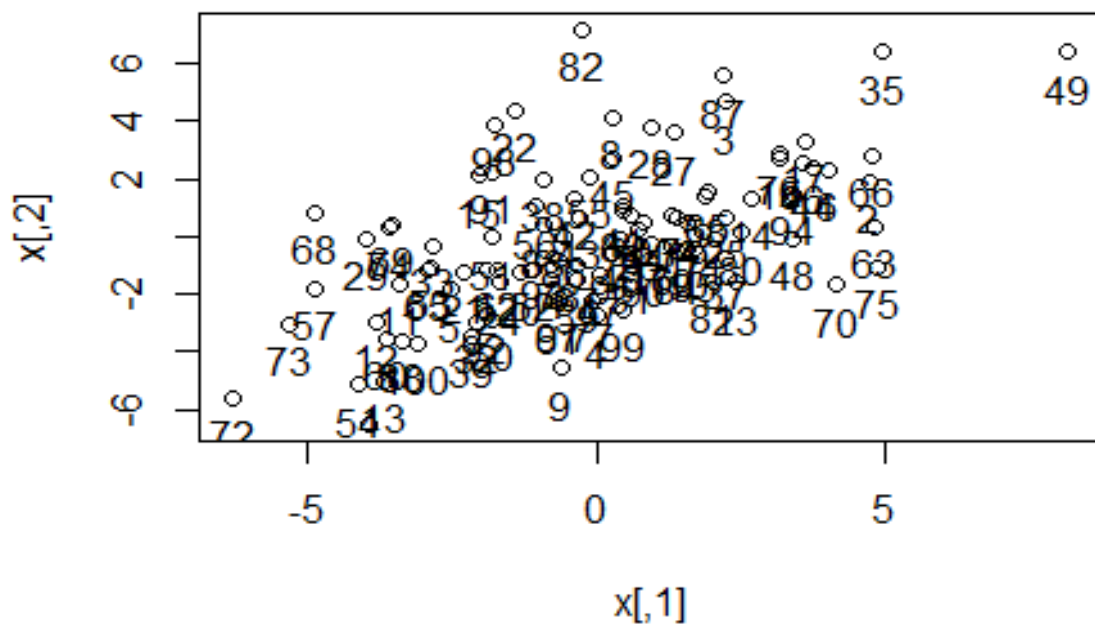


## Homework 2

### #a)

```
setwd("C:/Users/tracy/Desktop/Multivariate/作业/directory")
install.packages("mvtnorm")
library(mvtnorm)
set.seed(123)
N<-100
mean<-matrix(c(4,7),byrow=FALSE,ncol=1)
Si<-matrix(c(10,6,6,8),byrow=TRUE,ncol=2)
x <- rmvnorm(N,mean,Si)
x <- sweep(x,2,colMeans(x),"-")
plot(x)
text(x,labels=c(1:100),pos=1)
r1=range(x[,1])
r2=range(x[,2])
```



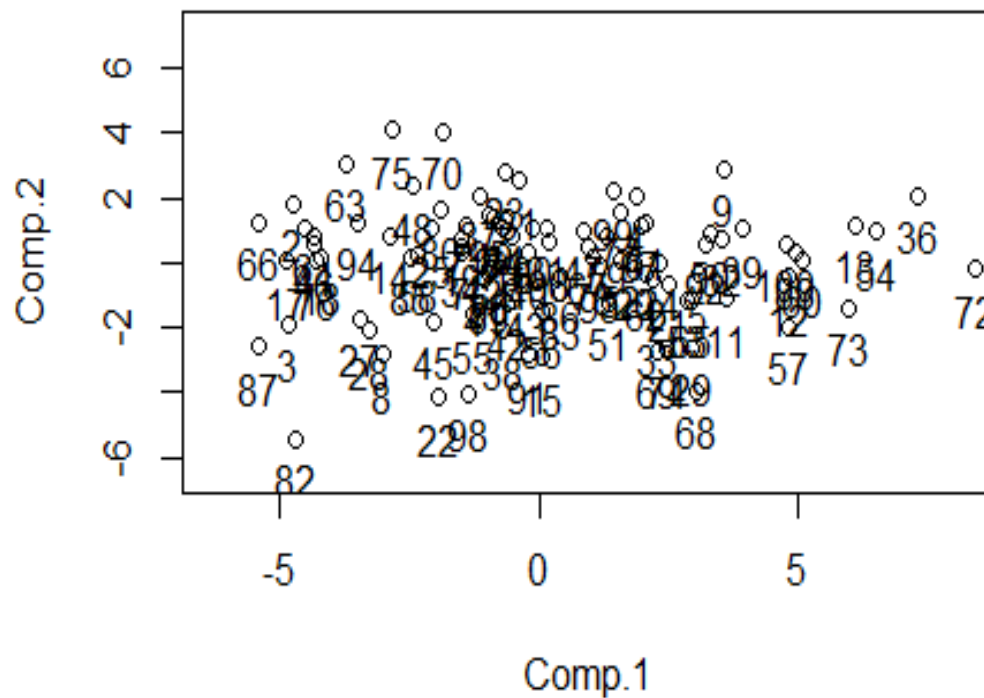
### #b)

```
x.PCA<-princomp(x,cor=FALSE)
```

### #c)

```
x.PCA$loadings
```

```
plot(x.PCA$scores,xlim=r1, ylim =r2)
text(x.PCA$scores[,1:2],labels=c(1:100),pos=1)
```



**#d)**

The range of data in Comp.2 is narrowed, while range of data in Comp.1 is not narrowed and the values show more dispersed distribution.

**#e)**

```
n <- nrow(x)
x_cov<-(n-1)/n*cov(x)
ei <- eigen(cov_x)
G <- ei$vectors
Y = x%*%G
#compare with part b
#compare G
x_cov_eval <- eigen(x_cov)$values
(x.PCA$sdev)^2
x_cov_eval-(x.PCA$sdev)^2
#compare Y
x.PCA$scores
```

x.PCA\$scores-Y

G

	[,1]	[,2]
[1,]	-0.7581476	0.6520830
[2,]	-0.6520830	-0.7581476