

Estadística (M)

Resolución del ejercicio hecho en clase de Estadística Descriptiva

Datos: Todas las muestras son de tamaño 100.

a) >summary(DietaA)

```
Min. 1st Qu. Median Mean 3rd Qu. Max.
68.00 90.75 99.00 98.68 105.20 121.00
```

> summary(DietaB)

```
Min. 1st Qu. Median Mean 3rd Qu. Max.
68.00 79.00 94.50 94.46 110.20 122.00
```

> summary(DietaC)

```
Min. 1st Qu. Median Mean 3rd Qu. Max.
68.00 70.00 73.00 77.29 81.00 121.00
```

> mean(gluco,trim=0.1)

```
dietaa dietab dietac
98.525 94.500 75.300
```

> mean(gluco,trim=0.2)

```
dietaa dietab dietac
98.53333 94.40000 74.50000
```

b) Varianza muestral

> var(DietaA)

```
[1] 103.7147
```

> var(DietaB)

```
[1] 280.4125
```

> var(DietaC)

```
[1] 112.006
```

#Desvío Muestral: raíz cuadrada de la varianza muestral

> sqrt(var(DietaA))

```
[1] 10.18404
```

> sqrt(var(DietaB))

```
[1] 16.74552
```

> sqrt(var(DietaC))

```
[1] 10.58329
```

#Distancia Intercuartos

##¿Cómo hacemos para obtener el primer cuarto?

> median(sort(DietaA)[1:50])

```
[1] 90.5
```

> median(sort(DietaB)[1:50])

```
[1] 79
```

> median(sort(DietaC)[1:50])

```
[1] 70
```

> ##¿Y el tercer cuarto?

> median(sort(DietaA)[51:100])

```
[1] 105.5
```

> median(sort(DietaB)[51:100])

```
[1] 110.5
```

> median(sort(DietaC)[51:100])

```
[1] 81
```

> ##Luego, las distancias intercuartos de cada uno son:

> median(sort(DietaA)[51:100])-median(sort(DietaA)[1:50])

```

[1] 15
> median(sort(DietaB)[51:100])-median(sort(DietaB)[1:50])
[1] 31.5
> median(sort(DietaC)[51:100])-median(sort(DietaC)[1:50])
[1] 11
> #Mad
> mad(DietaA)
[1] 11.1195
> mad(DietaB)
[1] 22.9803
> mad(DietaC)
[1] 5.9304

c) > quantile(DietaA) #Da los percentiles 0% (mínimo), 25%, 50%, 75% y 100%(máximo)
  0%  25%  50%  75% 100%
68.00 90.75 99.00 105.25 121.00
> quantile(DietaB)
  0%  25%  50%  75% 100%
68.00 79.00 94.50 110.25 122.00
> quantile(DietaC)
  0% 25% 50% 75% 100%
 68  70  73  81 121
#Percentiles 10% y 90%
> quantile(DietaA,0.10)
10%
86
> quantile(DietaB,0.10)
10%
75.8
> quantile(DietaC,0.10)
10%
68
> quantile(DietaA,0.90)
90%
111.1
> quantile(DietaB,0.90)
90%
115
> quantile(DietaC,0.90)
90%
90
> #Observación: Para obtener las distancias intercuartiles podemos hacer:
> quantile(DietaA,0.75)-quantile(DietaA,0.25)
75%
14.5
> quantile(DietaB,0.75)-quantile(DietaB,0.25)
75%
31.25
> quantile(DietaC,0.75)-quantile(DietaC,0.25)
75%
11
> #Rango muestral
> range(DietaA)

```

```
[1] 68 121
> range(DietaB)
[1] 68 122
> range(DietaC)
[1] 68 121
```

```
d) >stem(DietaA)
```

The decimal point is 1 digit(s) to the right of the |

```
6 | 8
7 |
8 | 011456666667777888899
9 | 0001122334555556666777788999999
10 | 000001122222444445555668888999
11 | 000111234566899
12 | 1
```

```
> stem(DietaB)
```

The decimal point is 1 digit(s) to the right of the |

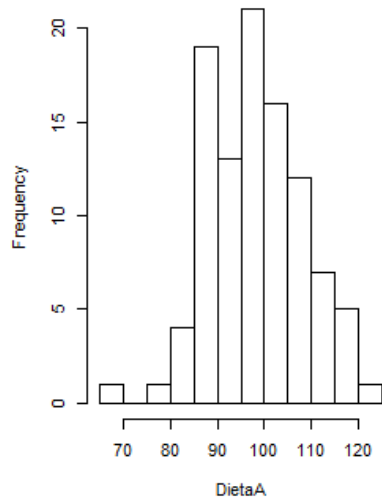
```
6 | 889
7 | 002334466666777788899999
8 | 00000001222233333346678
9 |
10 | 1124445566666678899999
11 | 00011111122223445555778899
12 | 2
```

```
> stem(DietaC)
```

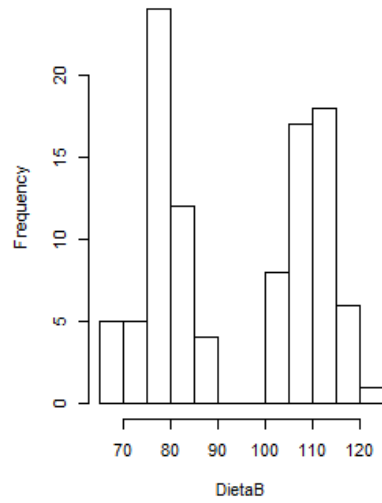
The decimal point is 1 digit(s) to the right of the |

```
6 | 88888888888999999999
7 | 000000000111111222222233333334444555567889999
8 | 0000111222344556679
9 | 000455
10 | 1379
11 | 0
12 | 1
```

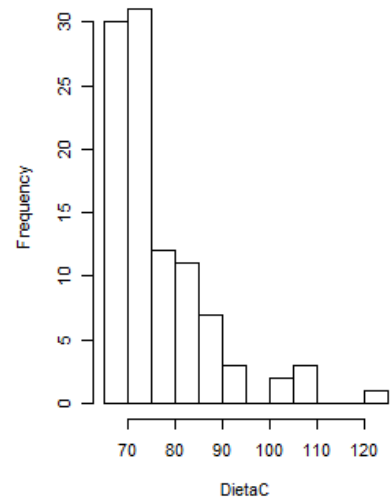
Histogram of DietaA



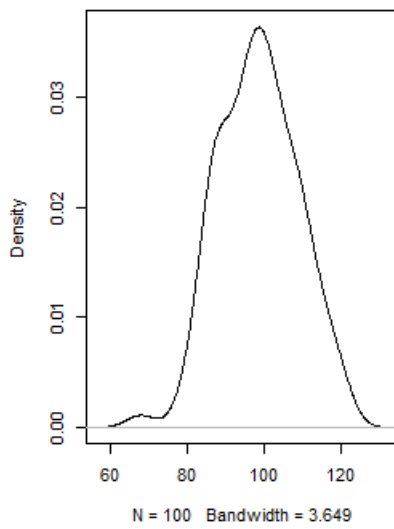
Histogram of DietaB



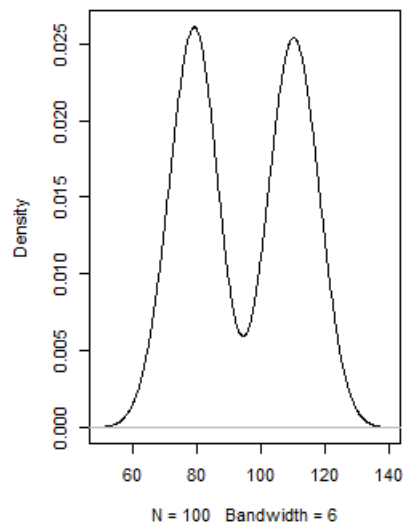
Histogram of DietaC



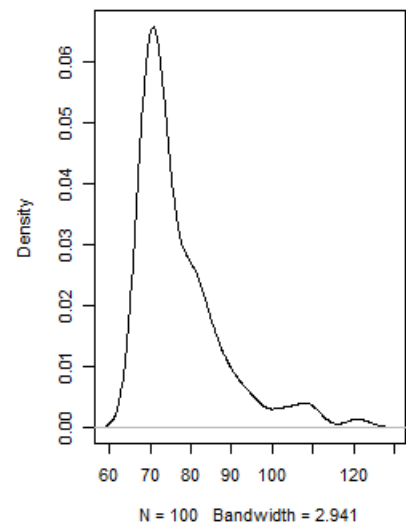
density.default(x = DietaA)

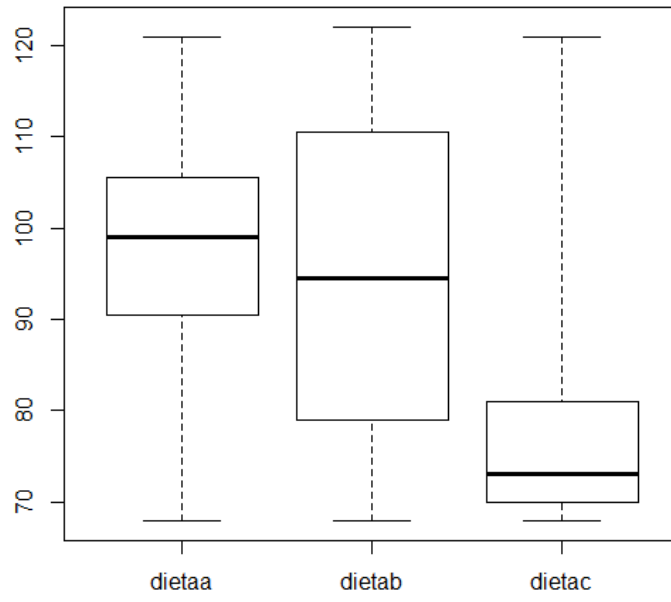
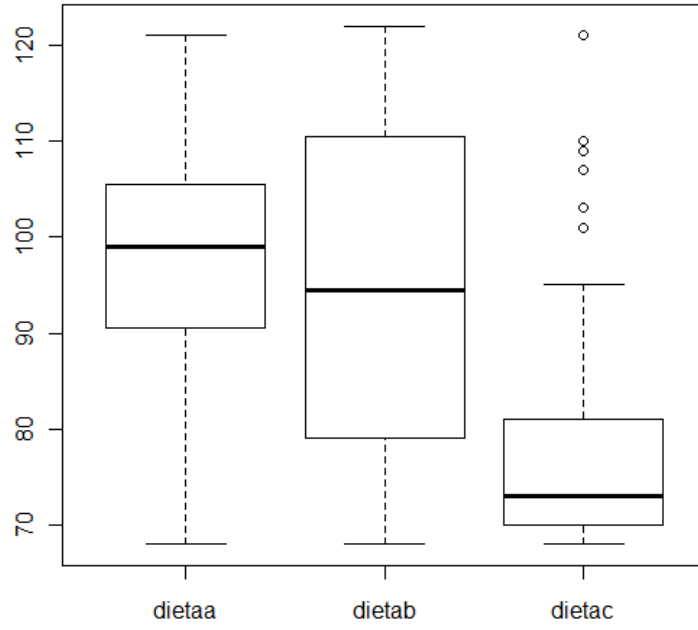


density.default(x = DietaB)

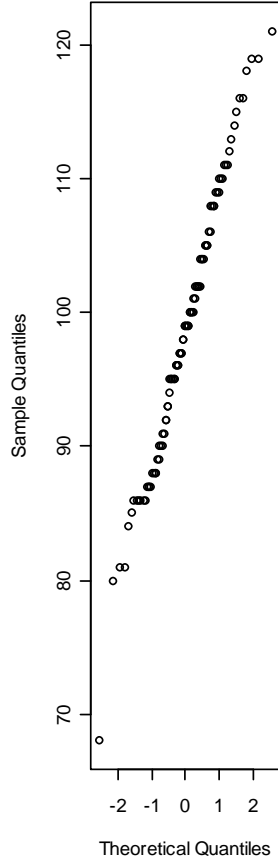


density.default(x = DietaC)

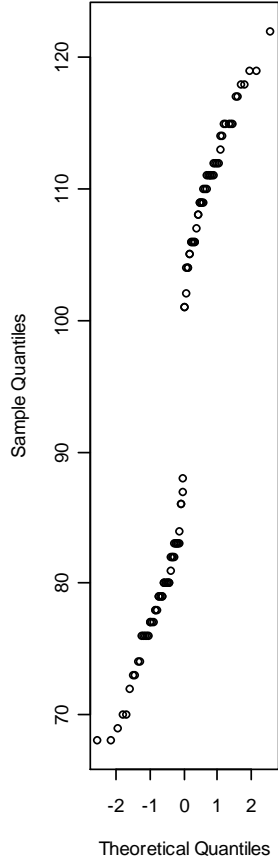




Normal Q-Q Plot



Normal Q-Q Plot



Normal Q-Q Plot

