

```
library(tidyverse)
f=file.choose()
fullBumpus=read_table(f)
```

1.

a)

```
> with(fullBumpus,t.test(Weight[Survive==1],
+                        Weight[Survive==0],
+                        var.equal=TRUE))
```

### Two Sample t-test

data: Weight[Survive == 1] and Weight[Survive == 0]  
 t = -2.6093, df = 134, p-value = 0.0101  
 alternative hypothesis: true difference in means is not equal to 0  
 95 percent confidence interval:  
 -1.1399459 -0.1569291  
 sample estimates:  
 mean of x mean of y  
 25.21250 25.86094

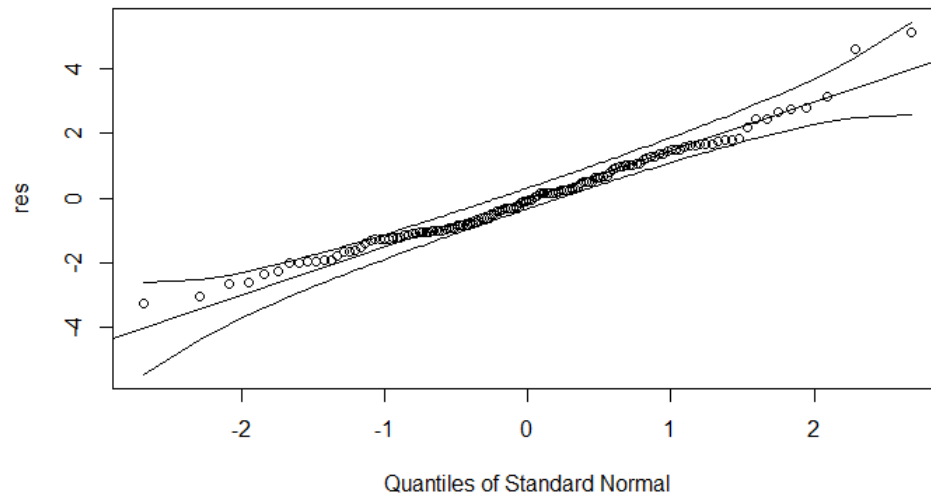
```
> with(fullBumpus,t.test(Weight[Survive==1],
+                        Weight[Survive==0],
+                        var.equal=FALSE))
```

### Welch Two Sample t-test

data: Weight[Survive == 1] and Weight[Survive == 0]  
 t = -2.5703, df = 117.95, p-value = 0.01141  
 alternative hypothesis: true difference in means is not equal to 0  
 95 percent confidence interval:  
 -1.1480287 -0.1488463  
 sample estimates:  
 mean of x mean of y  
 25.21250 25.86094

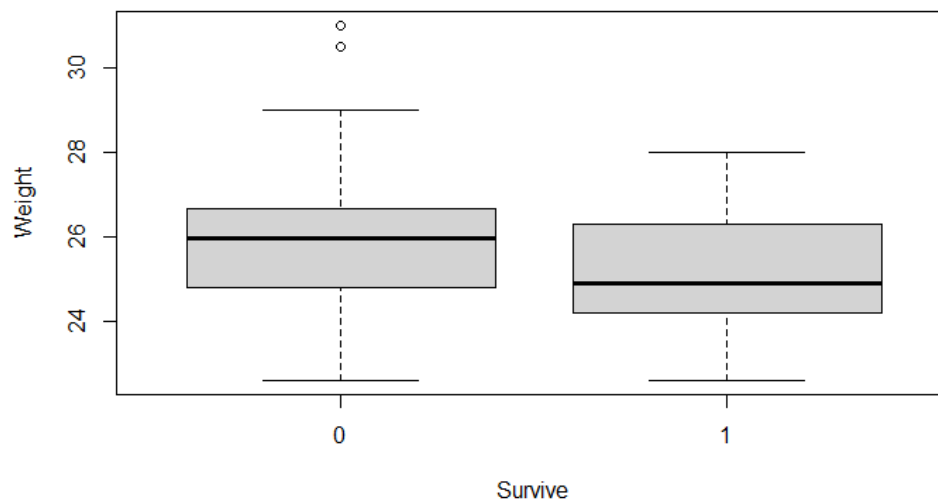
b)

```
res = resid(lm(Weight~Survive, fullBumpus))
source("http://www.stat.cmu.edu/~hseltman/files/qqn.R")
qqn(res)
```



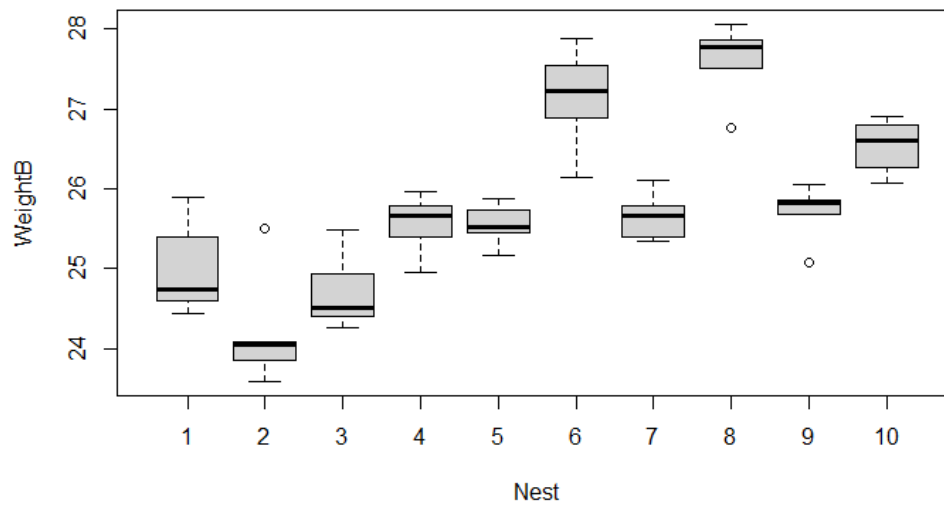
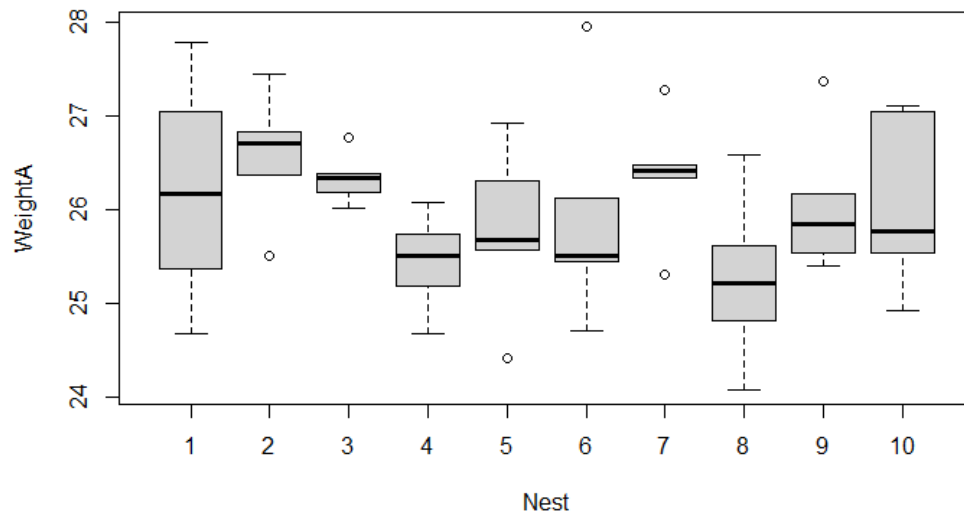
Nonnormality is not an issue here

c)  
`with(fullBumpus, boxplot(Weight~Survive))`



Ratio is pretty close to each other (1:1)

d)  
`f=file.choose()`  
`FakeCor=read_table(f)`  
`with(FakeCor, boxplot(WeightA~Nest))`  
`with(FakeCor, boxplot(WeightB~Nest))`



2.

a)

```
mdl = lm(Alar~Female+Weight, data=fullBumpus)
```

```
summary(mdl)
```

Call:

```
lm(formula = Alar ~ Female + Weight, data = fullBumpus)
```

Residuals:

	Min	1Q	Median	3Q	Max
	-13.2387	-2.6125	0.2613	2.8729	11.0747

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	202.1958	6.1318	32.975	< 2e-16 ***
Female	-4.8027	0.7271	-6.605	8.71e-10 ***
Weight	1.7553	0.2372	7.401	1.37e-11 ***

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 3.942 on 133 degrees of freedom

Multiple R-squared: 0.4961, Adjusted R-squared: 0.4885

F-statistic: 65.47 on 2 and 133 DF, p-value: < 2.2e-16

b)

> b0M

(Intercept)

202.1958

> b0F = mdl\$coefficients[1] + mdl\$coefficients[2]

> b0F

(Intercept)

197.3931

> b1 = mdl\$coefficients[3]

> b1

Weight

1.75533

c)

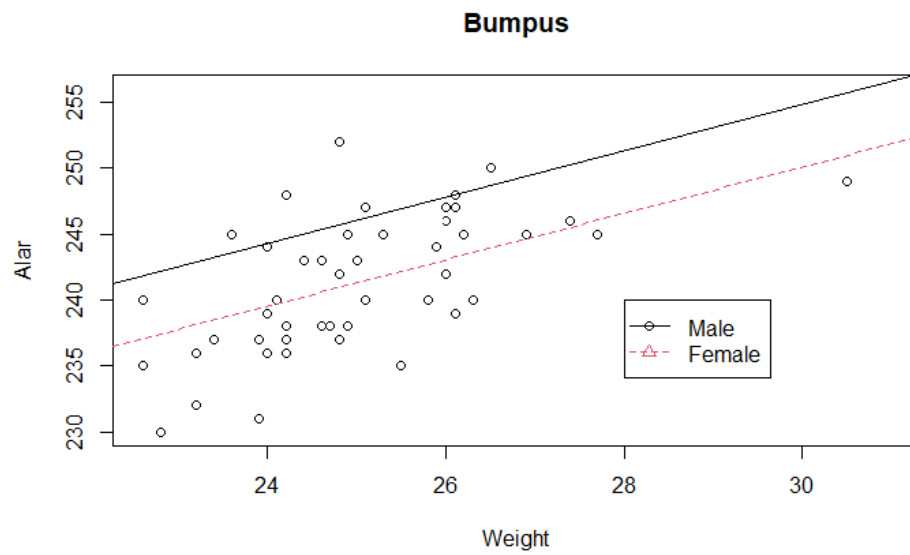
with(fullBumpus, table(Female, as.numeric(Female)))

with(fullBumpus, plot(Alar~Weight, pch=as.numeric(Female),  
col=as.numeric(Female), main="Bumpus"))

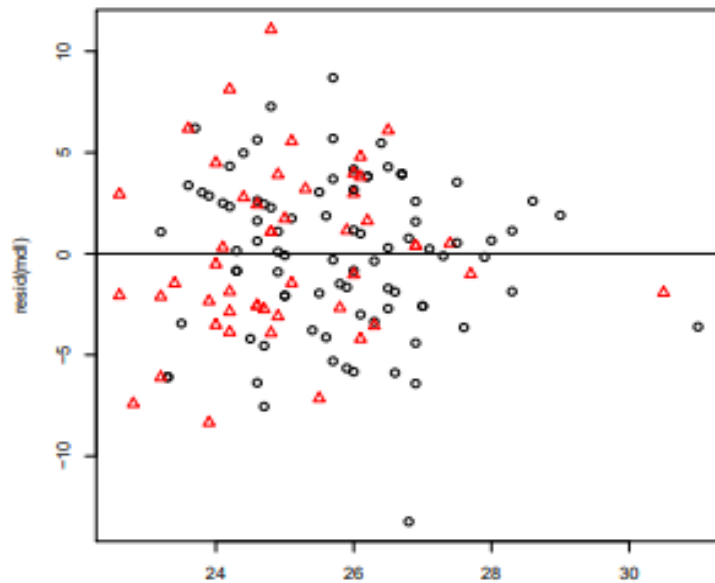
abline(b0M, b1, col=1, lty=1)

abline(b0F, b1, col=2, lty=2)

legend(28, 240, c("Male", "Female"), col=1:2, lty=1:2, pch=1:2)



```
with(fullBumpus, plot(resid(mdl)~fullBumpus$Weight, col=as.numeric(fullBumpus$Female),
  pch=as.numeric(fullBumpus$Female)))
abline(h=0)
```



d)

```
mdl2 = lm(Alar~Female*Weight, data=fullBumpus)
> summary(mdl2)
```

Call:

```
lm(formula = Alar ~ Female * Weight, data = fullBumpus)
```

Residuals:

Min	1Q	Median	3Q	Max
-13.0332	-2.5531	0.0527	2.8415	11.1550

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	207.4608	7.6956	26.958	< 2e-16 ***
Female	-18.8586	12.4580	-1.514	0.132
Weight	1.5512	0.2979	5.207	7.18e-07 ***
Female:Weight	0.5554	0.4914	1.130	0.260

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 3.938 on 132 degrees of freedom

Multiple R-squared: 0.5009, Adjusted R-squared: 0.4896

F-statistic: 44.16 on 3 and 132 DF, p-value: < 2.2e-16

```
> b0M = mdl$coefficients[1]
```

```
> b0M
```

```
(Intercept)
```

```
207.4608
```

```
> b0F = mdl$coefficients[1] + mdl$coefficients[2]
```

```
> b0F
```

```
(Intercept)
```

```
188.6022
```

```
> b1M = mdl$coefficients[3]
```

```
> b1M
```

```
Weight
```

```
1.551208
```

```
> b1F = mdl$coefficients[3] + mdl$coefficients[4]
```

```
> b1F
```

```
Weight
```

```
2.106566
```

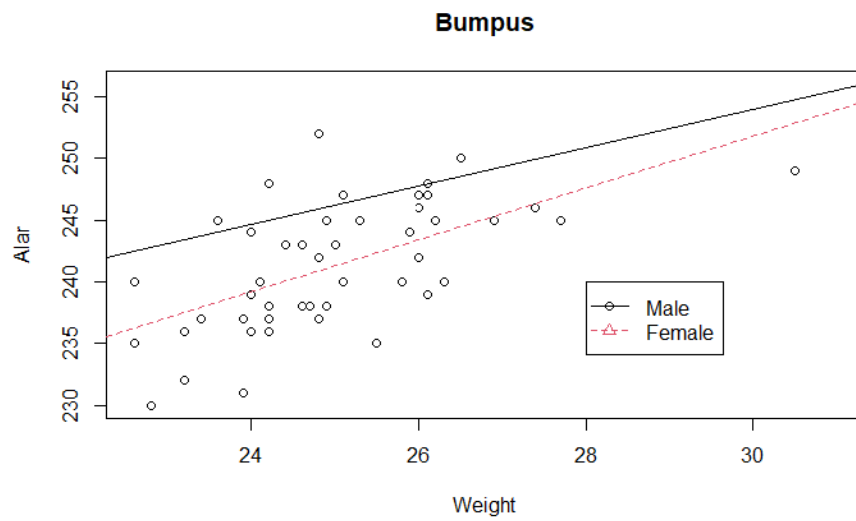
```
> with(fullBumpus, plot(Alar~Weight, pch=as.numeric(Female),
```

```
+ col=as.numeric(Female), main="Bumpus"))
```

```
> abline(b0M, b1M, col=1, lty=1)
```

```
> abline(b0F, b1F, col=2, lty=2)
```

```
> legend(28, 240, c("Male", "Female"), col=1:2, lty=1:2, pch=1:2)
```



e)

```
> anova(mdl,mdlI)
```

Analysis of Variance Table

Model 1: Alar ~ Female + Weight

Model 2: Alar ~ Female \* Weight

	Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)
1	133	2067.1				
2	132	2047.3	1	19.81	1.2773	0.2605

Do NOT have sufficient evidence

f)

```
confint(mdlI)
```

	2.5 %	97.5 %
(Intercept)	192.2381377	222.683434
Female	-43.5018377	5.784645
Weight	0.9619145	2.140502
Female:Weight	-0.4166585	1.527374