

Q5

**Comments**

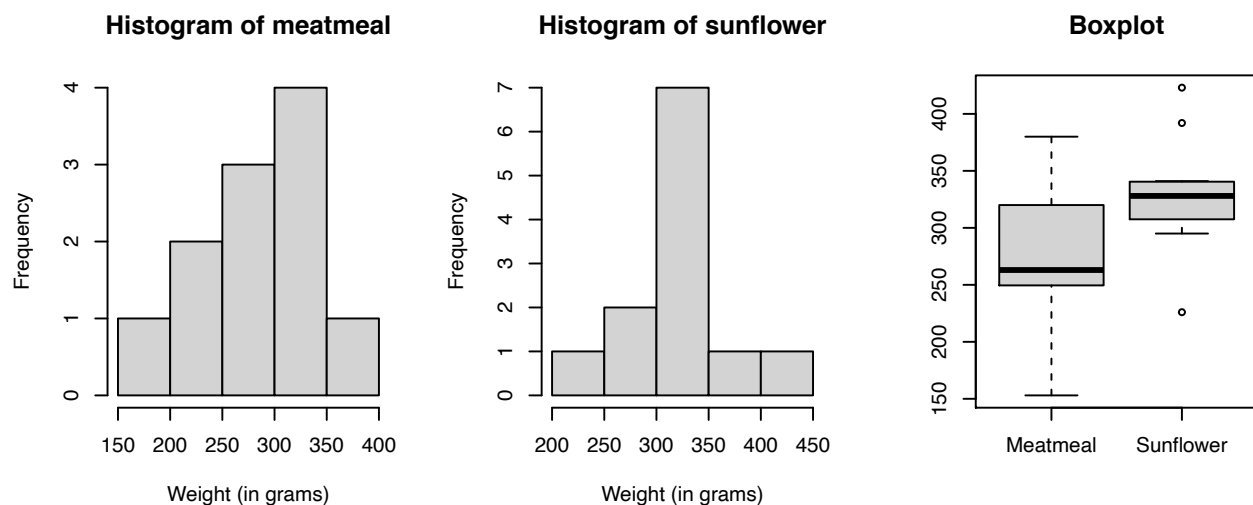
a)correct.

b)correct.

c)Very good for also checking the variance even not necessary here.

d)correct

2 / 2 pts

**Exercise 5 - Chick Weights****a) Meatmeal and Sunflower**

As shown in the boxplot the chicks that get the sunflower ( $M=329$ ) supplement seem to weigh more than the chicks that get the meatmeal ( $M=277$ ) supplement. To test whether this result is significant a t-test, Mann-Whitney and Kolmogorov-Smirnov test are executed. Since the chicks are divided between groups and the difference between the groups is tested, a two independent samples t-test will be used. The t-test showed that there is a difference between the `meatmeal` and the `sunflower` groups ( $t(18) = -2.2$ ,  $p < .05$ ).

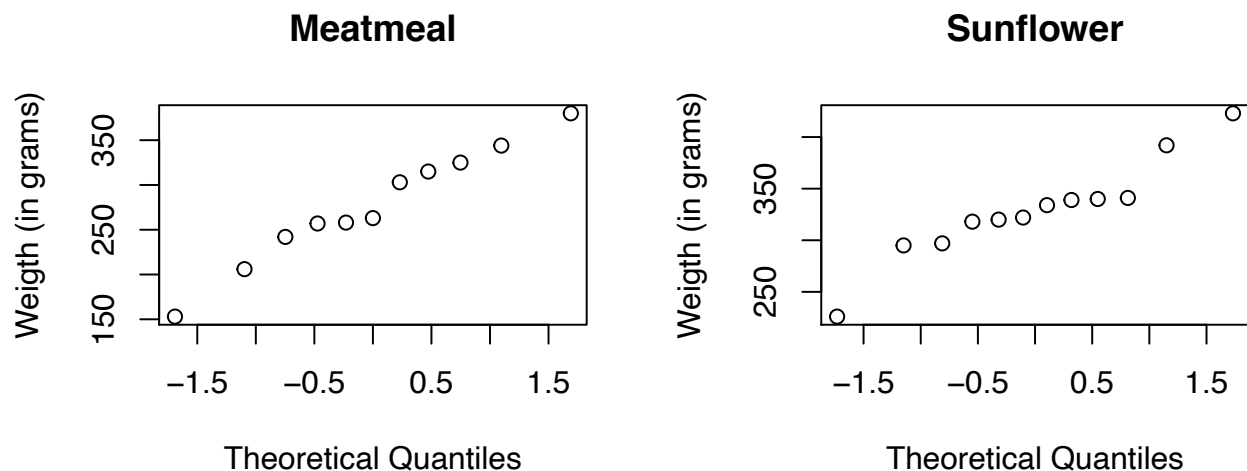
```
t.test(meatmeal, sunflower)
##
## Welch Two Sample t-test
##
## data: meatmeal and sunflower
## t = -2, df = 19, p-value = 0.04
```

```
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -102.57 -1.44
## sample estimates:
## mean of x mean of y
##      277      329
```

Whether we can trust this result depends on whether the assumption of normality is met. The Mann-Whitney and Kolmogorov-Smirnov test do not assume normality of the data and both the Mann-Whitney ( $W = 36$ ,  $p > .05$ ) and Kolmogorov-Smirnov ( $D = 0.5$ ,  $p > .05$ ) test show no difference in distribution between the groups.

```
wilcox.test(meatmeal, sunflower)
##
## Wilcoxon rank sum exact test
##
## data: meatmeal and sunflower
## W = 36, p-value = 0.07
## alternative hypothesis: true location shift is not equal to 0
ks.test(meatmeal, sunflower)
##
## Two-sample Kolmogorov-Smirnov test
##
## data: meatmeal and sunflower
## D = 0.5, p-value = 0.1
## alternative hypothesis: two-sided
```

Looking at the histograms and qq-plots we cannot assume normality and we should therefore conclude that there is no difference between the two groups.



## b) One-Way ANOVA

The one-way ANOVA shows that there is a difference between the means of the groups ( $F=15.37$ ,  $P < .01$ ).

```

chickwts_mod = lm(weight~feed, data=chickwts)
anova(chickwts_mod)
## Analysis of Variance Table
##
## Response: weight
##           Df Sum Sq Mean Sq F value    Pr(>F)
## feed       5 231129   46226    15.4 5.9e-10 ***
## Residuals 65 195556     3009
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

As shown in the table and boxplot below, sunflower has the highest expected mean ( $M=329$ ).

Feed Supplement	Expeted Mean
Meatmeal	276.909
Sunflower	328.917
Casein	323.583
Soybean	246.429
Horsebean	160.2
Linseed	218.75

### c) ANOVA model assumptions

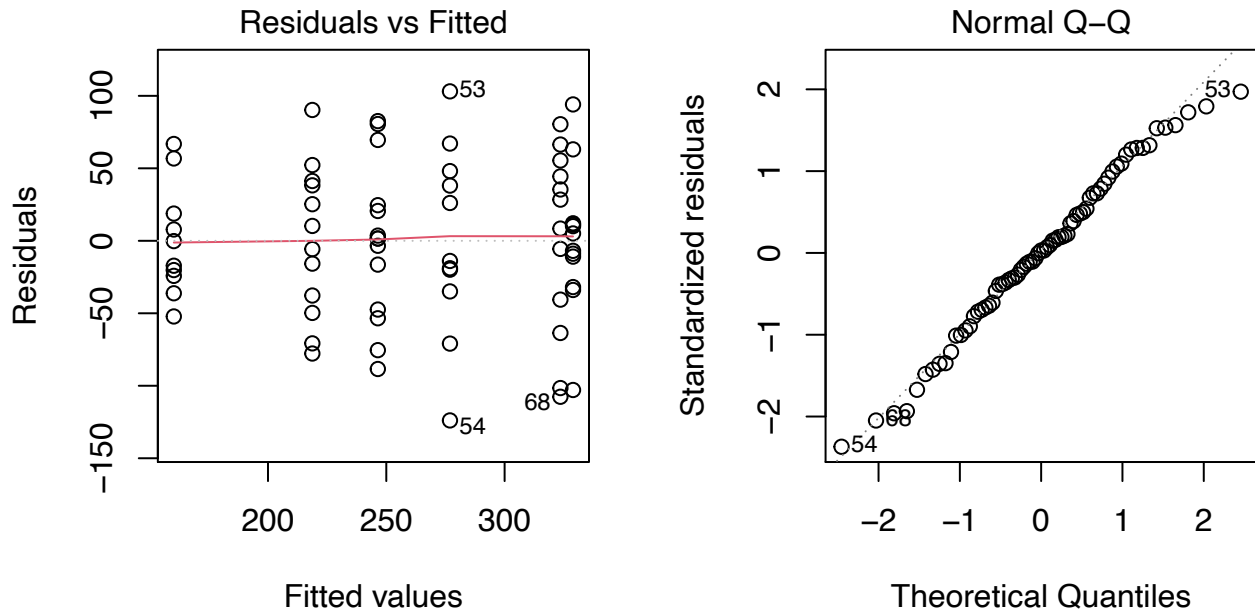
The ANOVA model assumption says, that aach group sample is drawn from a normally distributed population, and all populations have a common variance. In order to test the homogeneity of variances, we can either plot residuals or conduct a Levene test.

```

#install.packages("carData")
library(car)
leveneTest(weight ~ feed, data = chickwts)

## Levene's Test for Homogeneity of Variance (center = median)
##           Df F value Pr(>F)
## group    5      0.75  0.59
##          65

```



With the results from the Levene's test we can assume normality and homogeneity of variances. The results with  $F(5) = 0.75$  and  $p = 0.59$  suggest that the homogeneity is given. Since all points fall approximately along the reference line, we can also assume normality. This means both criteria are fulfilled.

#### d) Kruskal-Wallis vs. ANOVA

```
#Kruskal test
kruskal.test(weight~feed,data=chickwts)
```

```
##
##  Kruskal-Wallis rank sum test
##
## data:  weight by feed
## Kruskal-Wallis chi-squared = 37, df = 5, p-value = 5e-07
```

As we can see the Kruskal-Wallis test gives us a p-value of  $5.113e-07$  which means ( $p < 0.05$ ) the type of feed has an effect on the weight of the chicks. The Kruskal-Wallis test came to the same conclusion as the ANOVA test in b). However, the Kruskal-Wallis and ANOVA test could come to a different conclusion because the Kruskal-Wallis test does not rely on normality, like ANOVA does, but is based on ranks. This will lead to a different results in some cases, since Kruskal-Wallis is used when the assumptions of a one way ANOVA are not met.