1.不同玉米材料农艺性状有较大差异,通过对不同自交系穗长、穗粗、穗行数、株高、穗位高、散粉期 和果穗重的测定,试写出果穗重的回归模型,诊断是否存在异常值。

a[,8]代表最后一列, 其余同理......可以看到回归方程为:

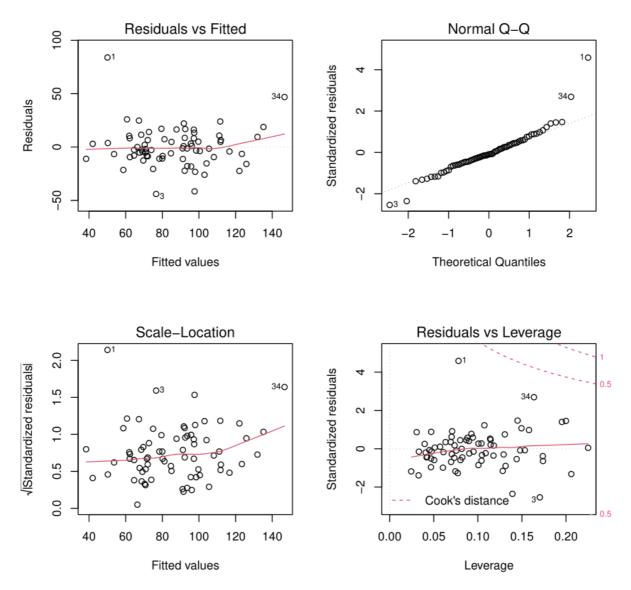
 $earweight = -66.27904 + 4.93213 \times earlong + 3.19007 \times earwidth.....$ (略)

```
Call:
lm(formula = a[, 8] \sim a[, 2] + a[, 3] + a[, 4] + a[, 5] + a[,
    6] + a[, 7], data = a)
Coefficients:
(Intercept)
                 a[, 2]
                             a[, 3]
                                          a[, 4]
                                                        a[, 5]
                                                                    a[, 6]
                                                                     0.05596
  -66.27904
                 4.93213
                              3.19007
                                         -0.33549
                                                        0.11799
     a[, 7]
   -1.07416
```

然后输入如下代码:

```
summary(lm1)
```

```
Call:
lm(formula = a[, 8] \sim a[, 2] + a[, 3] + a[, 4] + a[, 5] + a[,
   6] + a[, 7], data = a)
Residuals:
   Min
            1Q Median
                                  Max
                            3Q
-44.028 -9.464 -1.642 8.300 83.945
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) -66.27904 35.37115 -1.874 0.065452 .
                       1.30169 3.789 0.000333 ***
a[, 2]
            4.93213
a[, 3]
                      0.70855 4.502 2.86e-05 ***
            3.19007
a[, 4]
            -0.33549
                       1.40051 -0.240 0.811436
a[, 5]
                                1.017 0.313084
             0.11799
                       0.11606
            0.05596
                       0.20959 0.267 0.790318
a[, 6]
            -1.07416
a[, 7]
                       0.51075 -2.103 0.039334 *
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 19.06 on 65 degrees of freedom
Multiple R-squared: 0.6005, Adjusted R-squared: 0.5637
F-statistic: 16.29 on 6 and 65 DF, p-value: 2.423e-11
```



2.将三种不同菌型的伤寒病毒a、b、c分别接种于10,9,11只小白鼠上,观察其存活天数,问三种菌型下小白鼠的平均存活天数是否有显著性差异。 a菌株: 2,4,3,2,4,7,7,2,5,4 b菌株: 5,6,8,5,10,7,12,6,6 c菌株: 7,11,6,6,7,9,5,10,6,3,10

由于比较平均存活天数是否有显著差异,因此用t检验:

```
h1=c(2,4,3,2,4,7,7,2,5,4)

h2=c(5,6,8,5,10,7,12,6,6)

h3=c(7,11,6,6,7,9,5,10,6,3,10)

t.test(h1,h2)

t.test(h2,h3)

t.test(h1,h3)
```

结果如下:

## 第一组与第二组对比:

```
Welch Two Sample t-test

data: h1 and h2
t = -3.2413, df = 15.24, p-value = 0.005387
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
-5.338236 -1.106208
sample estimates:
mean of x mean of y
4.000000 7.222222
```

## 第二组与第三组对比:

```
Welch Two Sample t-test

data: h2 and h3
t = -0.046497, df = 17.405, p-value = 0.9634
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
-2.33812 2.23711
sample estimates:
mean of x mean of y
7.222222 7.272727
```

## 第一组与第三组对比:

```
Welch Two Sample t-test

data: h1 and h3

t = -3.4447, df = 18.527, p-value = 0.002795

alternative hypothesis: true difference in means is not equal to 0

95 percent confidence interval:

-5.264722 -1.280732

sample estimates:

mean of x mean of y

4.000000 7.272727
```

可以看到2和3之间差异较小,1和2、1和3有显著差异。