

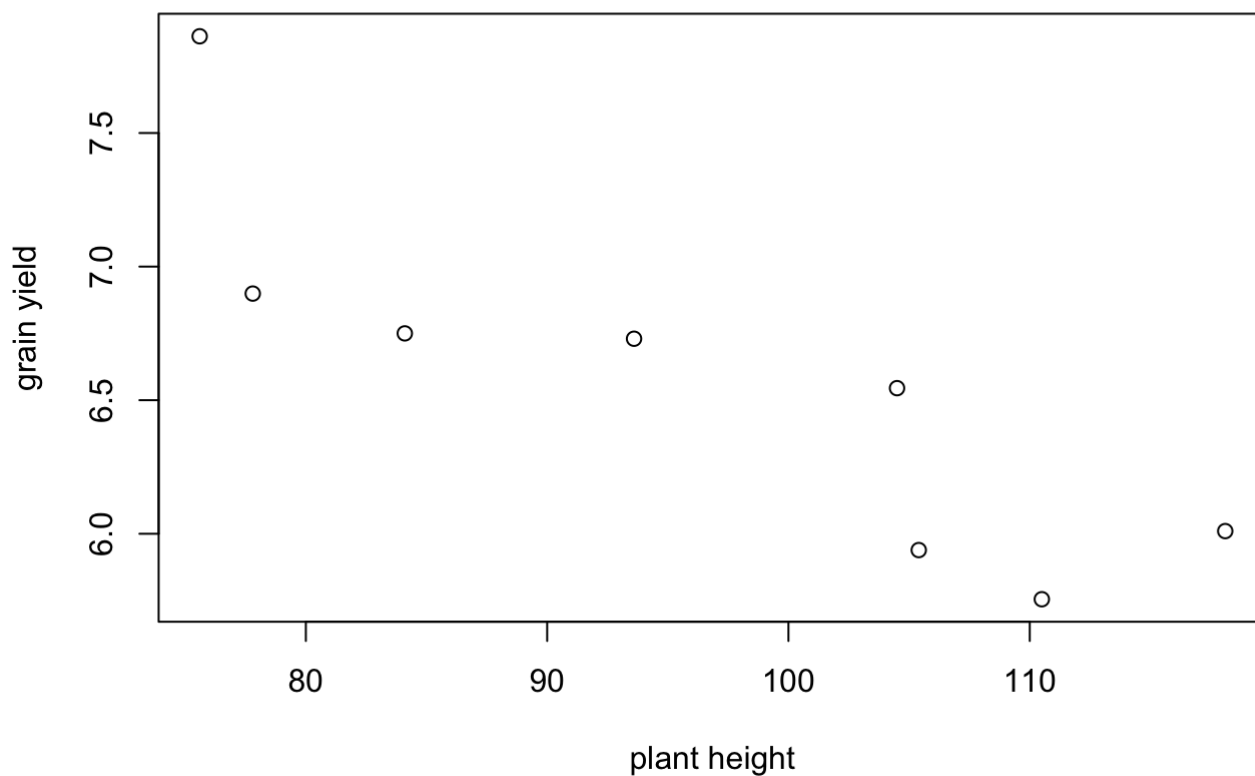
Hmw_06

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Q1

```
d = data.frame(x = c(110.5, 105.4, 118.1, 104.5, 93.6, 84.1, 77.8, 75.6),  
y = c(5.755, 5.939, 6.010, 6.545, 6.730, 6.750, 6.899, 7.862))  
plot(d$x, d$y, xlab = "plant height", ylab = "grain yield")
```



Answer a

```
cor(d$x, d$y)
```

```
## [1] -0.868707
```

```
fit_d = lm(y ~ x, data = d)  
summary(fit_d)
```

```
##
## Call:
## lm(formula = y ~ x, data = d)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.34626 -0.27605 -0.09448  0.27023  0.53495
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 10.137455   0.842265  12.036   2e-05 ***
## x           -0.037175   0.008653  -4.296   0.00512 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.3624 on 6 degrees of freedom
## Multiple R-squared:  0.7547, Adjusted R-squared:  0.7138
## F-statistic: 18.46 on 1 and 6 DF,  p-value: 0.005116
```

```
#beta 0
coef(fit_d)[1]
```

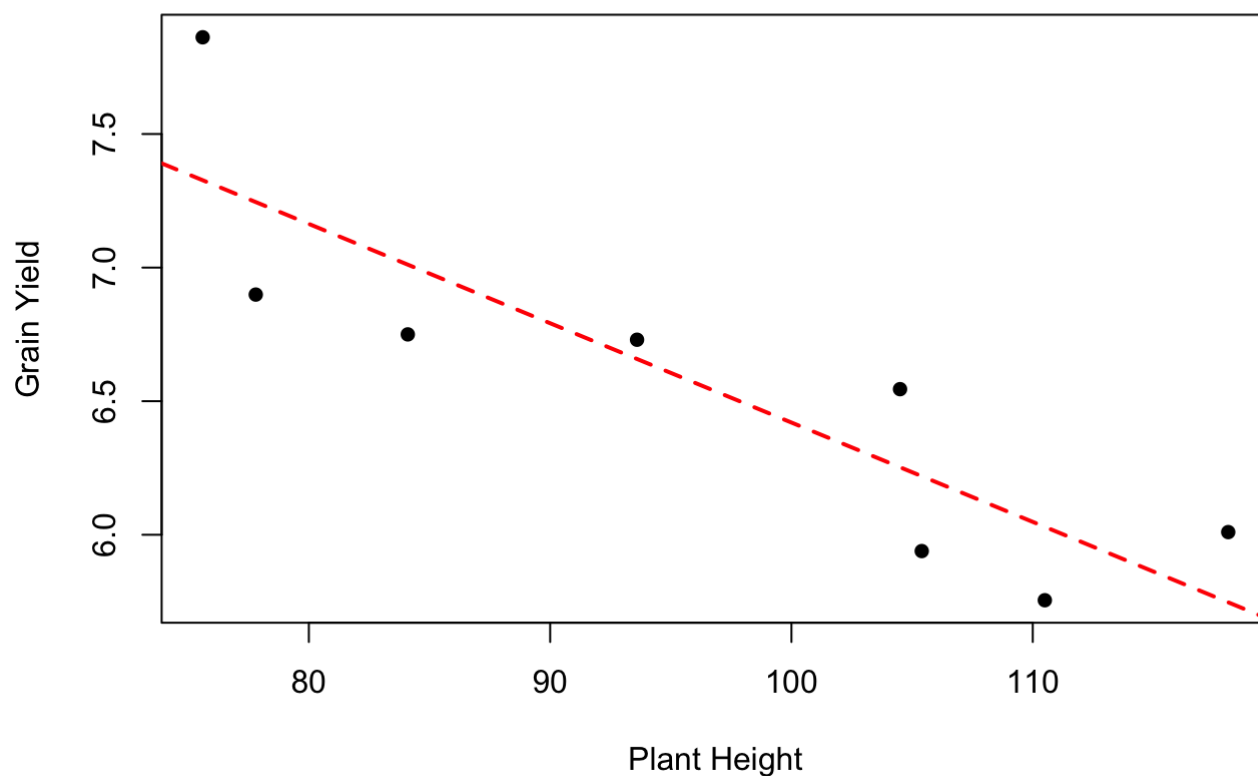
```
## (Intercept)
##      10.13746
```

```
# beta 1 Units??
coef(fit_d)["x"]
```

```
##              x
## -0.03717469
```

```
plot(d$x, d$y, pch=16,
      xlab = "Plant Height",
      ylab = "Grain Yield",
      main = "Scatterplot of Rice Plant Growth vs Grain Yield") +
abline(coefficients(fit_d), lwd = 2, lty = 2,
        col = "red")
```

Scatterplot of Rice Plant Growth vs Grain Yield



```
## integer(0)
```

the estimate tells us that the grain yield will decrease as the plant gets taller. The regression line indicates that yield will likely fall below 6.0 once plant height is more than about 105

Answer b

```
anova(fit_d)
```

```
## Analysis of Variance Table
##
## Response: y
##          Df Sum Sq Mean Sq F value    Pr(>F)
## x          1  2.42357    2.42357   18.455 0.005116 **
## Residuals   6  0.78794    0.13132
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
summary(fit_d)
```

```
##
## Call:
## lm(formula = y ~ x, data = d)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.34626 -0.27605 -0.09448  0.27023  0.53495
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 10.137455   0.842265  12.036   2e-05 ***
## x           -0.037175   0.008653  -4.296   0.00512 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.3624 on 6 degrees of freedom
## Multiple R-squared:  0.7547, Adjusted R-squared:  0.7138
## F-statistic: 18.46 on 1 and 6 DF,  p-value: 0.005116
```

Both the F test and T test gave a p-value of less than 5%, allowing us to reject a null hypothesis of the slope value = 0

Answer C

```
qt(0.05/2, 8-2)
```

```
## [1] -2.446912
```

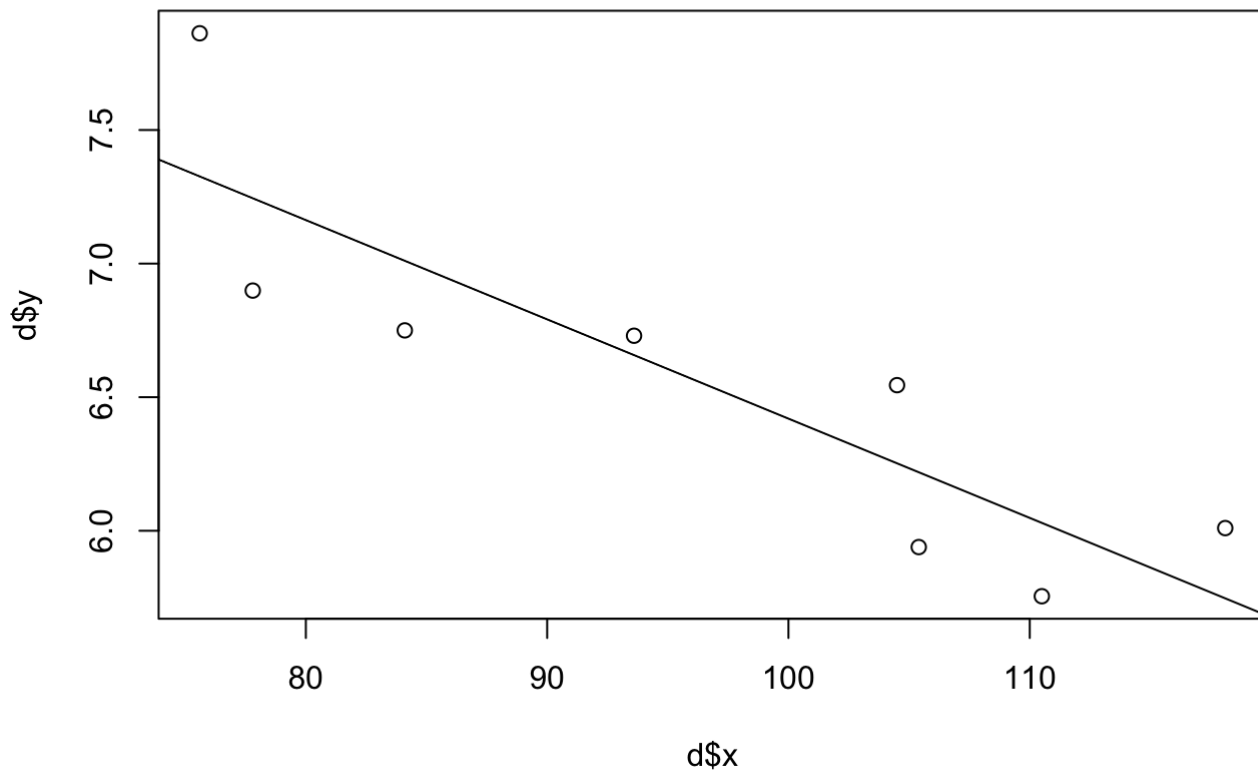
95% of the time that the confidence interval is calculate it will overlap with the true value of the slope.

```
confint(fit_d)
```

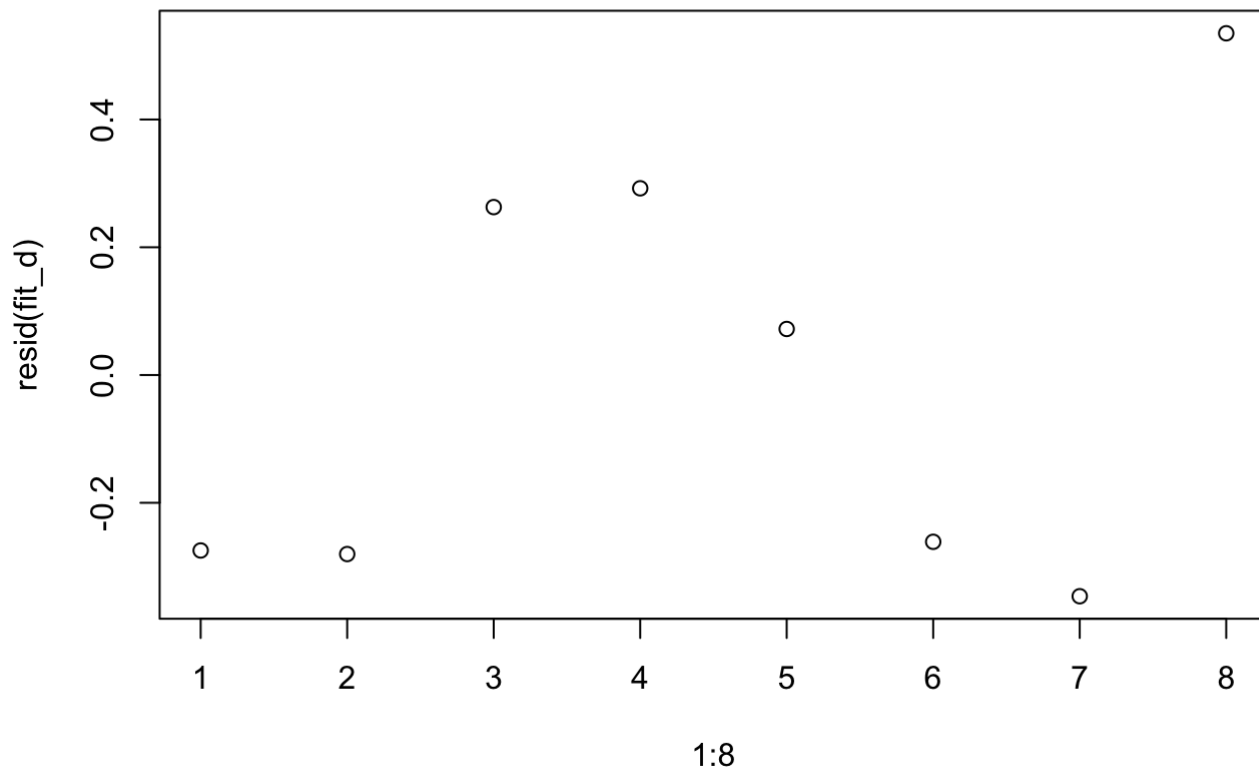
```
##              2.5 %      97.5 %
## (Intercept)  8.07650745 12.19840320
## x           -0.05834895 -0.01600043
```

Answer d

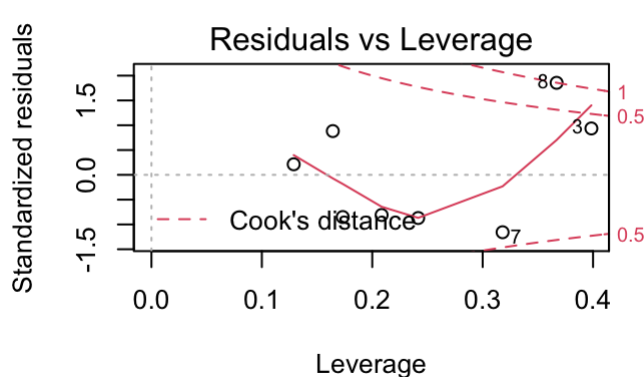
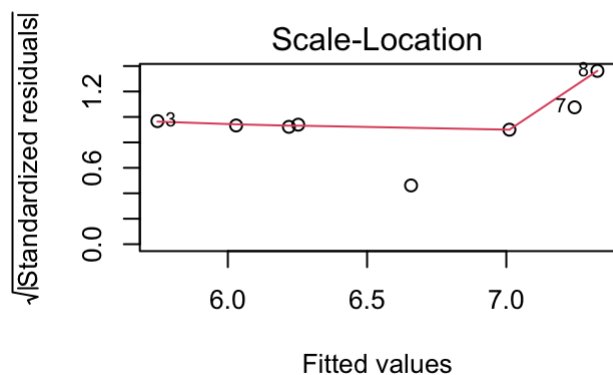
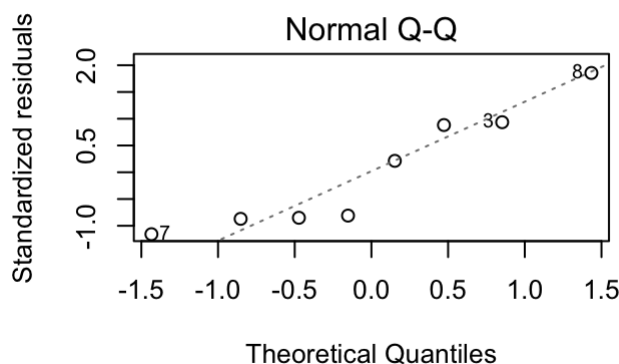
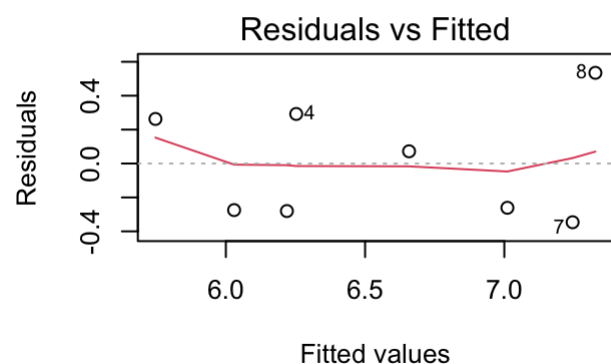
```
{plot(d$x,d$y)
abline(fit_d)}
```



```
{y=plot(resid(fit_d), x=1:8)  
abline(fit_d)}
```



```
### resid(fit_d) gives me the actual value of the residuals
### abline(fit_d) plots the slope
{par(mfrow=c(2,2)) ## plotting the residuals vs fitted lines.
plot(fit_d)}
```



Answer e

```
SSE<-sum((fitted(fit_d) - d$y)^2)

ev<-SSE/(8-2)
sum(fit_d$residuals**2)
```

```
## [1] 0.7879367
```

```
ev1<-anova(fit_d)
ev1<-ev1$`Mean Sq`[2]
```

Answer f

```
est<-predict(fit_d, newdata = data.frame(x = 100), interval = "confidence")
####int[1] its how much the 100 things is gonna yield, int[2] is the lower bound of the
ci and int[3] is the upper bound
```

Answer g

```
pred<-predict(fit_d, newdata = data.frame(x = 100), interval = "prediction")  
#####int[1] its how much the 100 things is gonna yield, int[2] is the lower bound of the  
ci and int[3] is the upper bound
```

Answer h

```
sumar<-summary(fit_d)  
sumar$r.squared
```

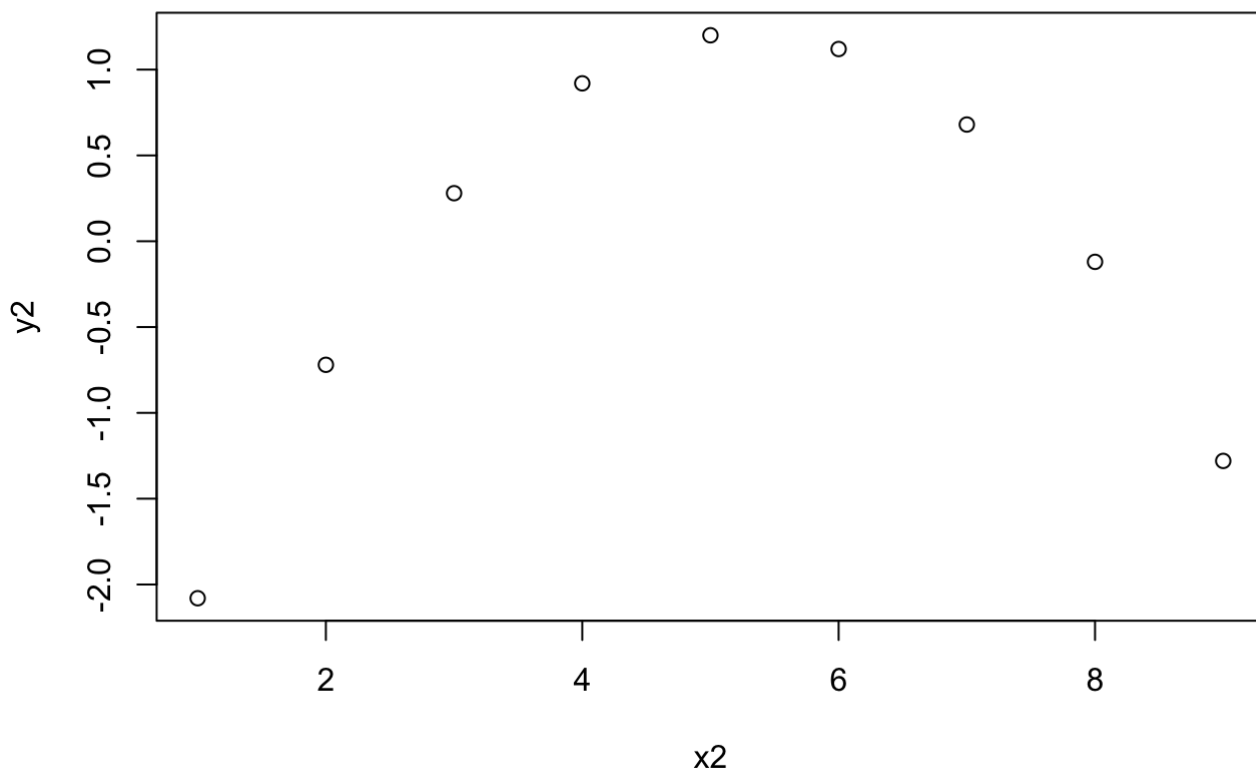
```
## [1] 0.7546518
```

```
##### This means how well the model fits the data
```

Q2

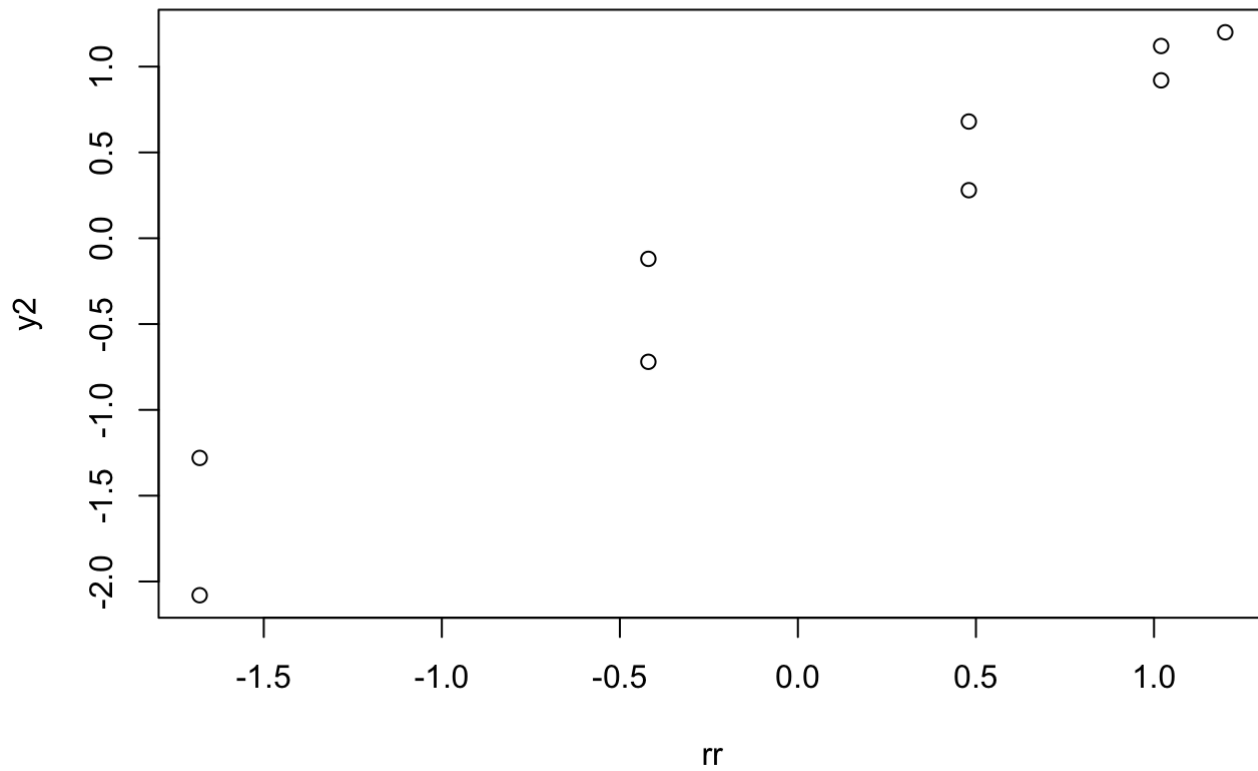
2a answer

```
x2 = c(1, 2, 3, 4, 5, 6, 7, 8, 9)  
y2 = c(-2.08, -0.72, 0.28, 0.92, 1.20, 1.12, 0.68, -0.12, -1.28)  
plot(x = x2, y = y2)
```



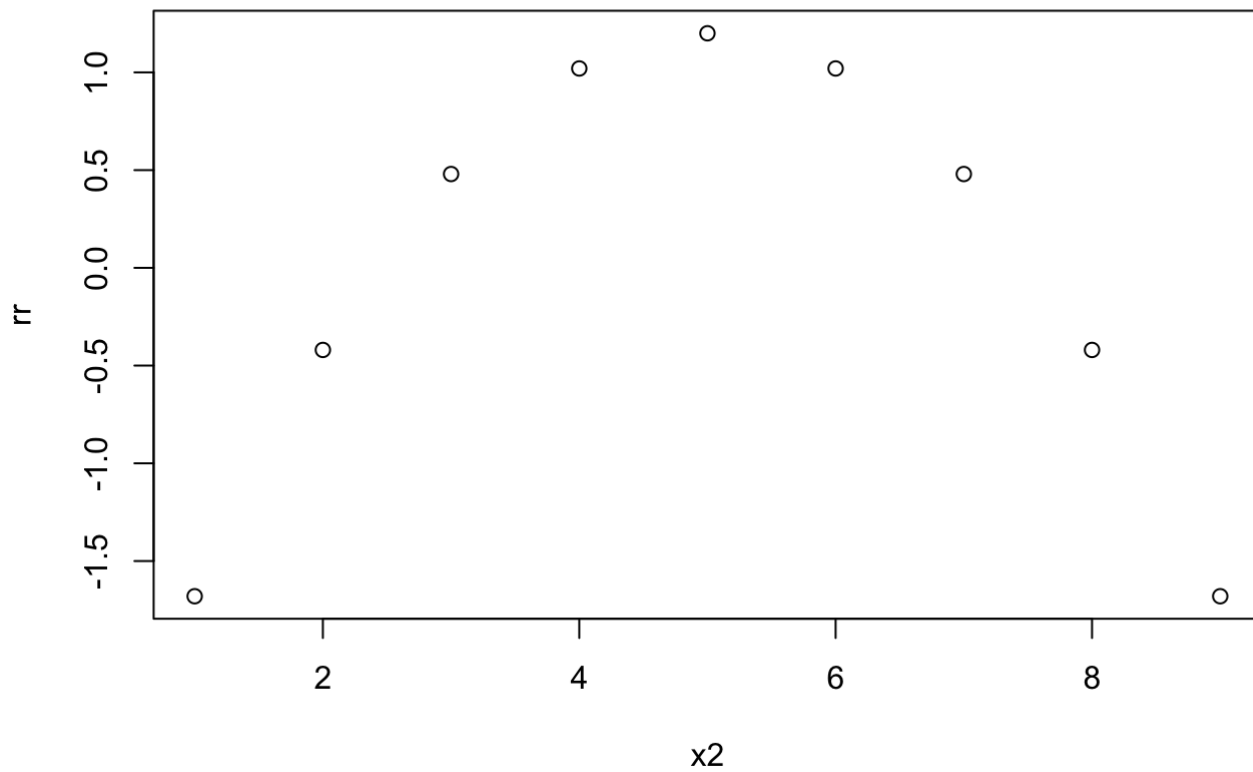
2b answer

```
modelo<-lm(y2~x2)
rr<-resid(modelo)
plot(x = rr, y = y2)
```



2c answer

```
plot(x = x2, y = rr)
```

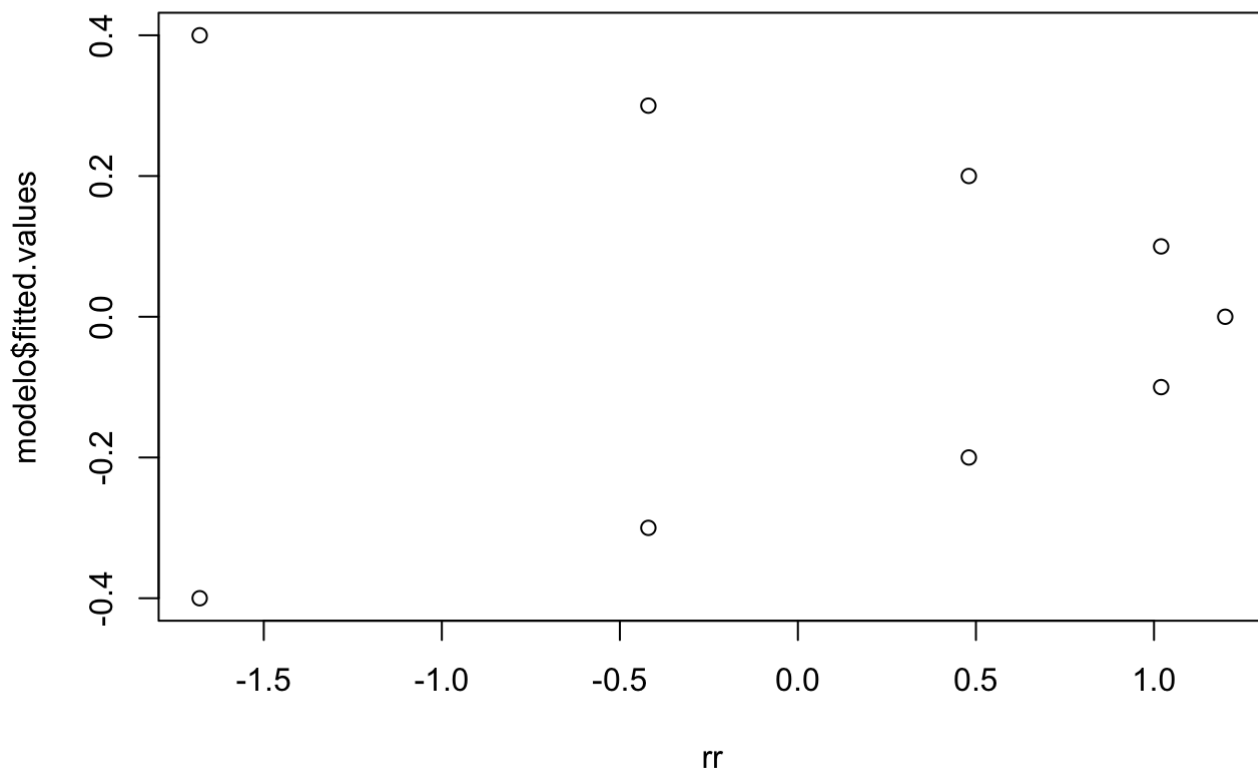


2d answer

```
modelo$fitted.values
```

```
##           1           2           3           4           5
## -4.000000e-01 -3.000000e-01 -2.000000e-01 -1.000000e-01 -2.220446e-16
##           6           7           8           9
##  1.000000e-01  2.000000e-01  3.000000e-01  4.000000e-01
```

```
plot(x = rr, y = modelo$fitted.values)
```



2e answer

Yes, there is a meaningful difference between plot c and d. In plot d, rr value maximises when fitted value is at zero. In plot c, rr value does not respond this way, instead maximizing around five.

Plot b is a better indication of lack of fit, because the points are close fit to a line of regression.