

705604096_stats101a_hw3

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Question 1

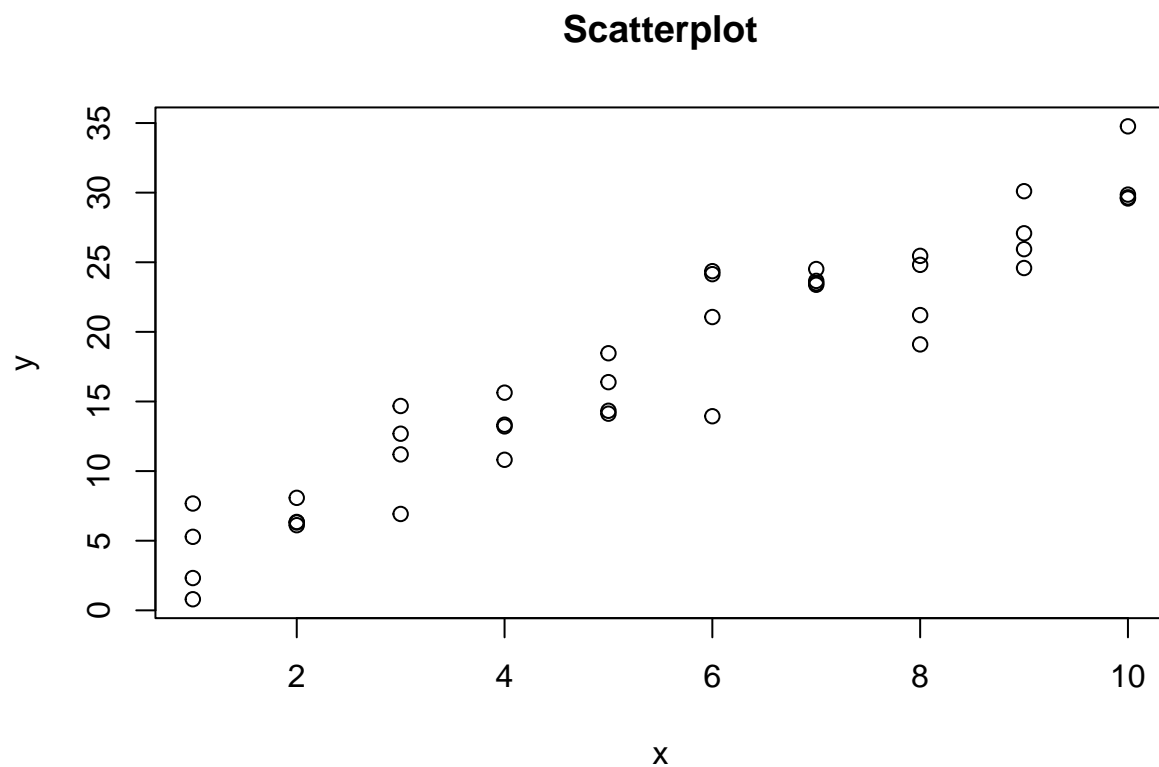
A)

a)

```
slr <- function(beta_0, beta_1, sigma, x, rns){  
  set.seed(rns)  
  eps <- rnorm(length(x), 0, sigma)  
  beta_0 + (beta_1 * x) + eps  
}
```

b)

```
x <- rep(1:10, by = .1, 4)  
y <- slr(1, 3, 3, x, 123)  
plot(x, y, main = "Scatterplot")
```



B)

```
cor(x, y)
```

```
## [1] 0.9529631
```

The correlation coefficient is 0.9529631

C)

```
x2 <- rep(1:10, by = .1, 4)
y2 <- slr(1, 30, 3, x2, 123)
cor(x2, y2)
```

```
## [1] 0.9995272
```

With parameters

$$\beta_0 = 1$$

$$\beta_1 = 30$$

$$\sigma = 3$$

we get a correlation of 0.9995272

Question 2

```
my_data <- read_csv('armspans2022_gender.csv')

## Rows: 46 Columns: 5
## -- Column specification -----
## Delimiter: ","
## chr (2): compmother, compfather
## dbl (3): height, armspan, is.female
##
## i Use 'spec()' to retrieve the full column specification for this data.
## i Specify the column types or set 'show_col_types = FALSE' to quiet this message.

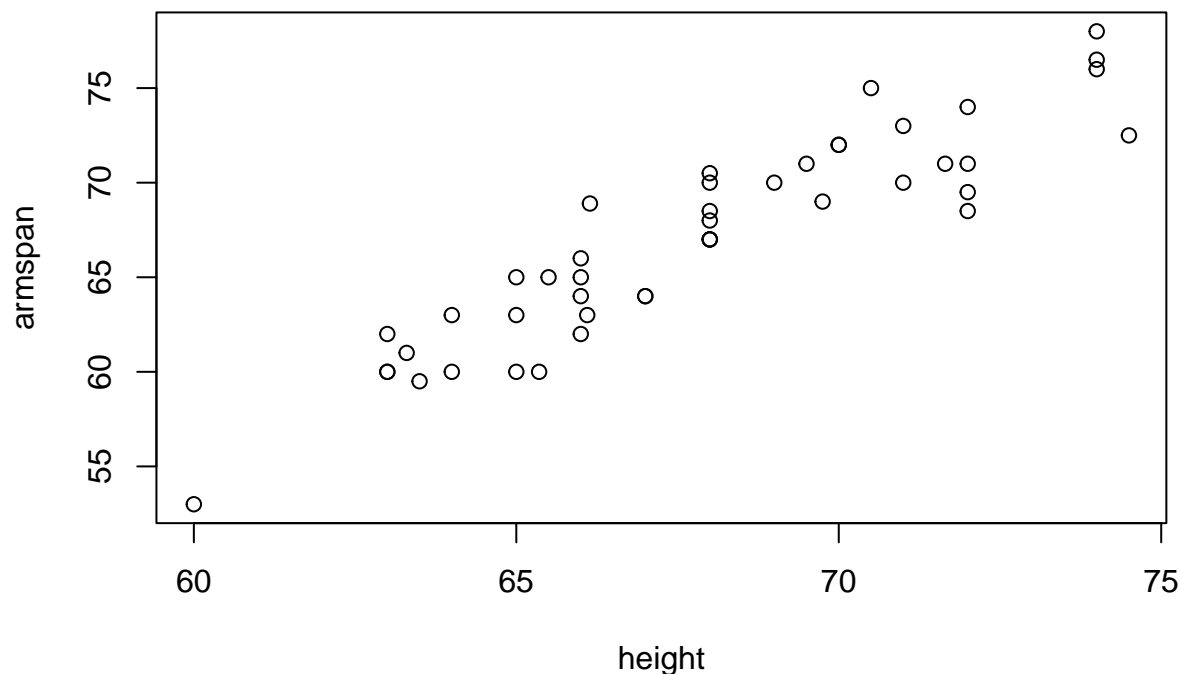
my_data['compmother'] <- tolower(my_data$compmother)
my_data['compfather'] <- tolower(my_data$compfather)
my_data$compmother <- str_replace_all(my_data$compmother, c(" " = "_", "' " = ""))
my_data$compfather <- str_replace_all(my_data$compfather, c(" " = "_", "' " = ""))
my_data

## # A tibble: 46 x 5
##   height armspan is.female compmother    compfather
##   <dbl>   <dbl>   <dbl> <chr>         <chr>
## 1    74     76         0 taller        taller
## 2    65     65         0 taller        about_the_same
## 3    60     53         1 shorter       shorter
## 4   69.8     69         0 taller        about_the_same
## 5    70     72         0 taller        about_the_same
## 6    68    70.5         0 taller        shorter
## 7    64     60         0 taller        taller
## 8    68     67         0 taller        about_the_same
## 9    68     67         1 taller        shorter
## 10   63     60         1 about_the_same shorter
## # ... with 36 more rows
```

When cleaning this data, I decided to make all of the elements in the columns lower case to make it easier to manipulate in the future. I also replaced spaces in between words with underscores for the same reason.

a)

```
plot(armspan ~ height, data = my_data )
```



We can observe a positive linear trend in this scatter plot. We can describe the strength as in between moderate and strong between the observations, since there is a clear linear pattern that is noticeable. There are not any unusual features we can discern from this scatter plot, except for the lowest value being farther from the bulk of the data but still in line with our linear pattern.

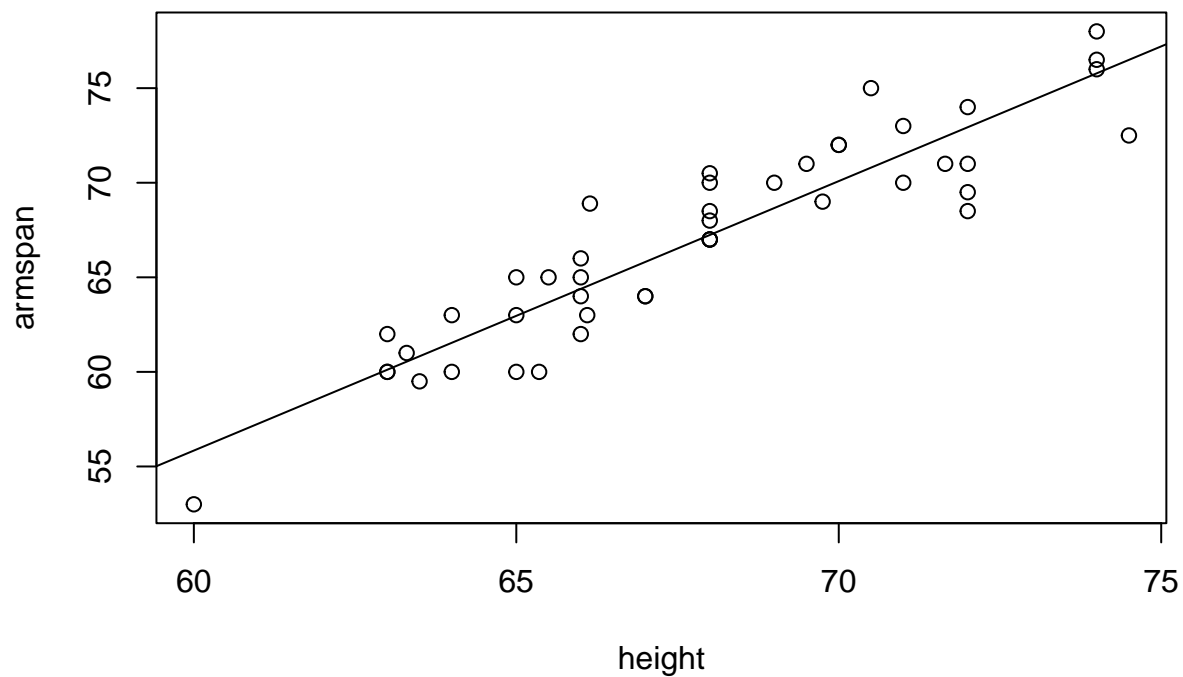
b)

```
plot(armspan ~ height, data = my_data )
my_lm <- lm(armspan ~ height, data = my_data)
summary(my_lm)
```

```
##
## Call:
## lm(formula = armspan ~ height, data = my_data)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -4.4353 -1.5302  0.0369  1.4893  4.3080
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -29.63530     6.22105  -4.764 2.19e-05 ***
## height       1.42459     0.09158  15.555 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
##
## Residual standard error: 2.135 on 43 degrees of freedom
## (1 observation deleted due to missingness)
## Multiple R-squared: 0.8491, Adjusted R-squared: 0.8456
## F-statistic: 242 on 1 and 43 DF, p-value: < 2.2e-16
```

```
abline(my_lm)
```



The equation for our estimated line is

$$\hat{y}_i = -29.63530 + 1.42459x_i$$

c)

```
my_func <- function(x){
  -29.6353 + 1.42459 * x
}
my_func(63)
```

```
## [1] 60.11387
```

```
residual <- 61 - my_func(63)
residual
```

```
## [1] 0.88613
```

Based on our model, the predicted arm span for my height is 60.11387 inches. The residual is 0.88613 inches.

d)

```
my_func(76)
```

```
## [1] 78.63354
```

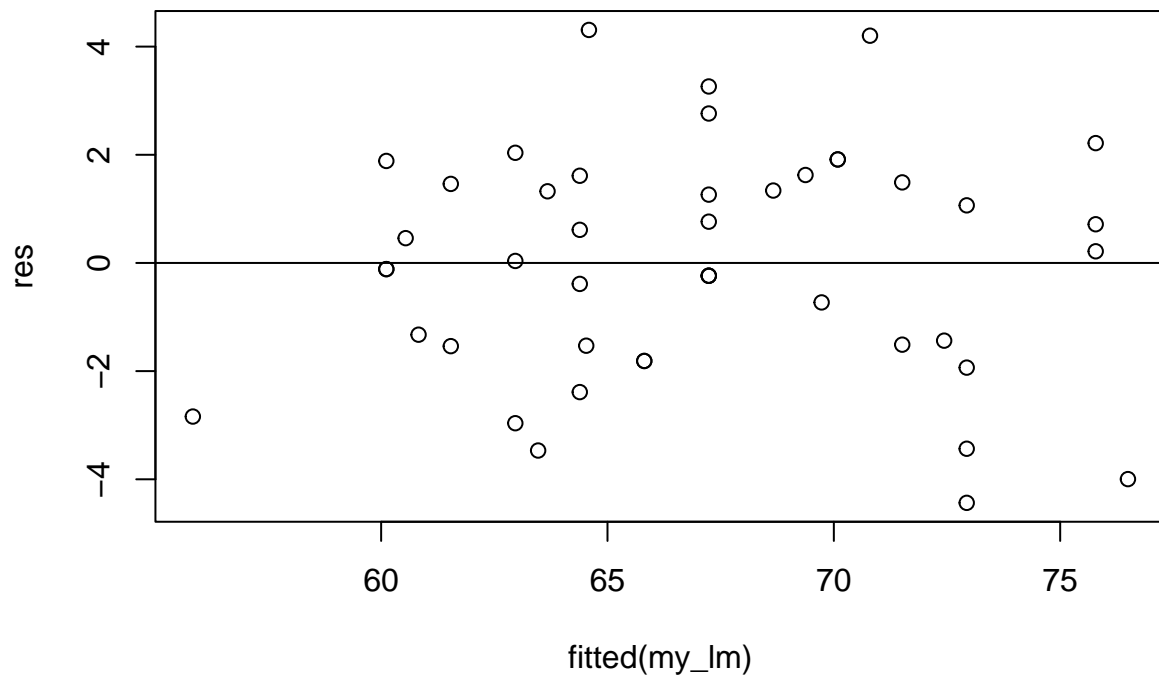
```
residual2 <- 79 - my_func(76)  
residual2
```

```
## [1] 0.36646
```

No, this does not seem unusual as the residual is relatively small.

e)

```
res <- resid(my_lm)  
plot(fitted(my_lm), res)  
abline(0,0)
```

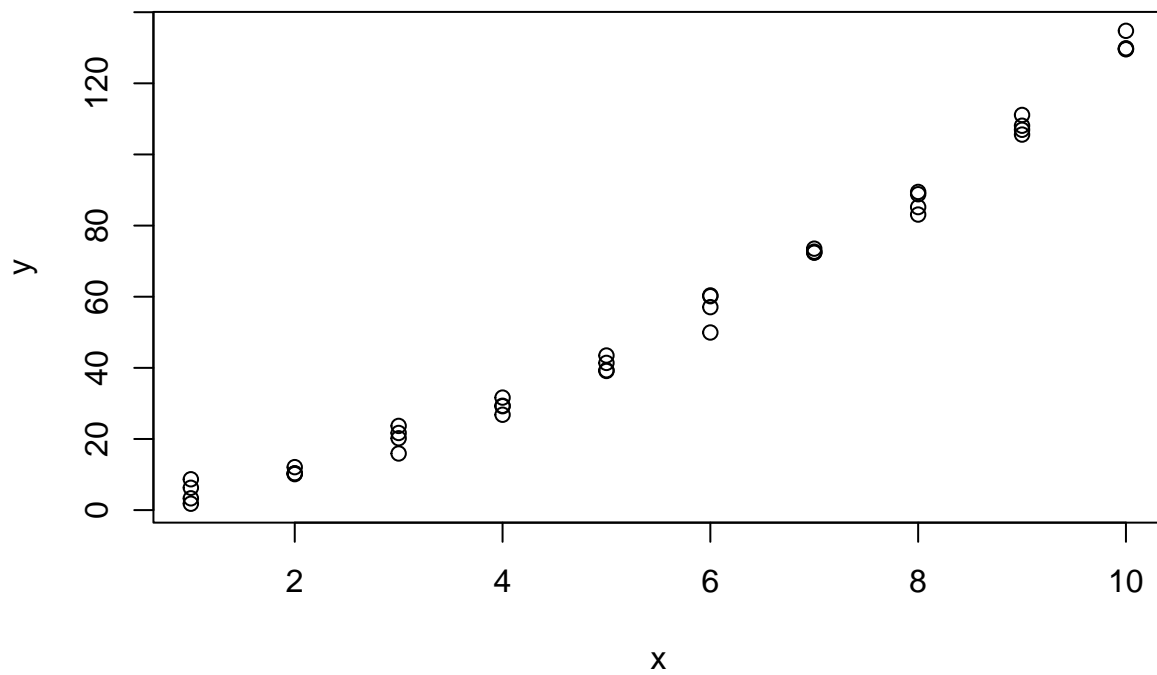


Because there does not seem to be any pattern in the residual points with respect to the line at $x = 0$, we can conclude that the linear model is the right fit for this data. We can also note that there is a fan shape that appears in our residual plot, with exception to the outlier as previously mentioned, that would indicate that the standard deviation increases as x increases.

Question 3

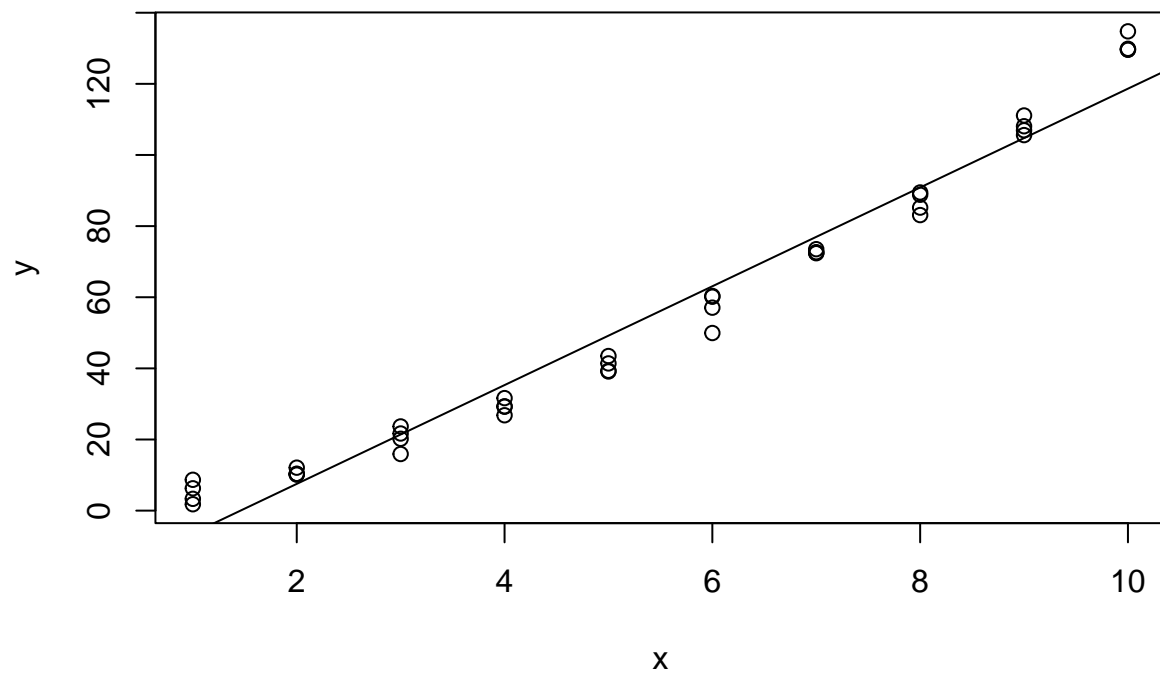
a)

```
quad_func <- function(a, b, c, sigma, x = rep(1:10,by=.1,4), random.seed){  
  set.seed(random.seed)  
  a + (b * x) + (c * (x^2)) + rnorm(length(x), 0, sigma)  
}  
x <- rep(1:10,by=.1,4)  
y <- quad_func(1, 3, 1, 3, x, 123)  
plot(x, y)
```

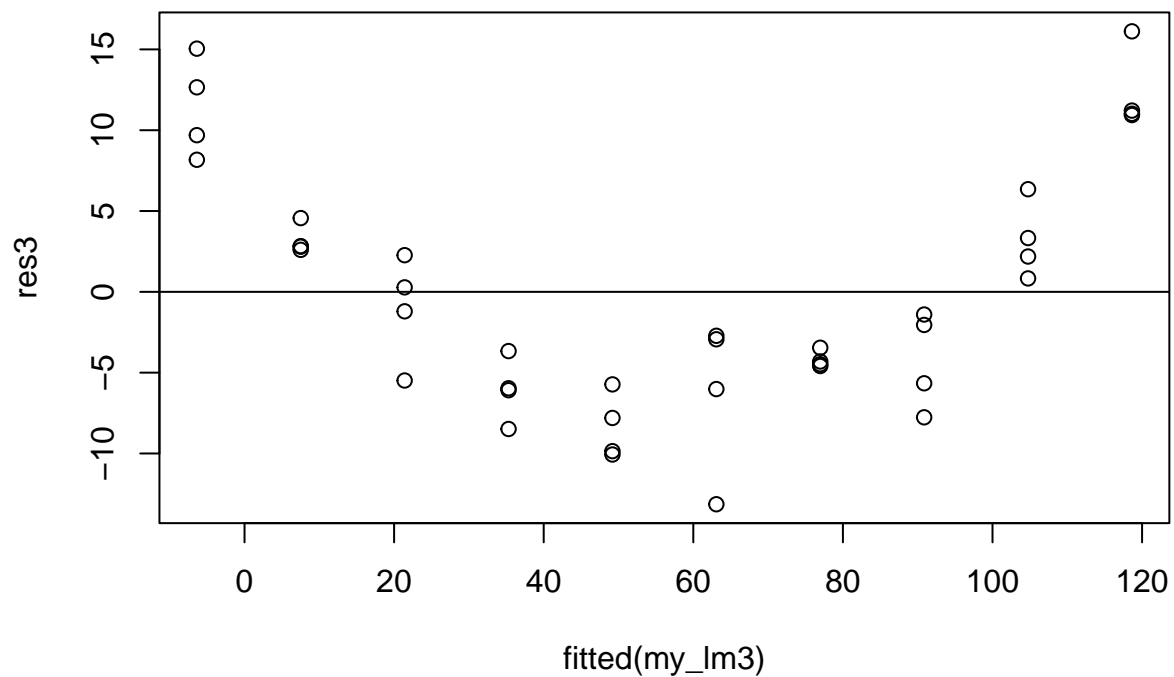


b)

```
plot(x, y)  
my_lm3 <- lm(y ~ x)  
abline(my_lm3)
```



```
res3 <- resid(my_lm3)
plot(fitted(my_lm3), res3)
abline(0,0)
```

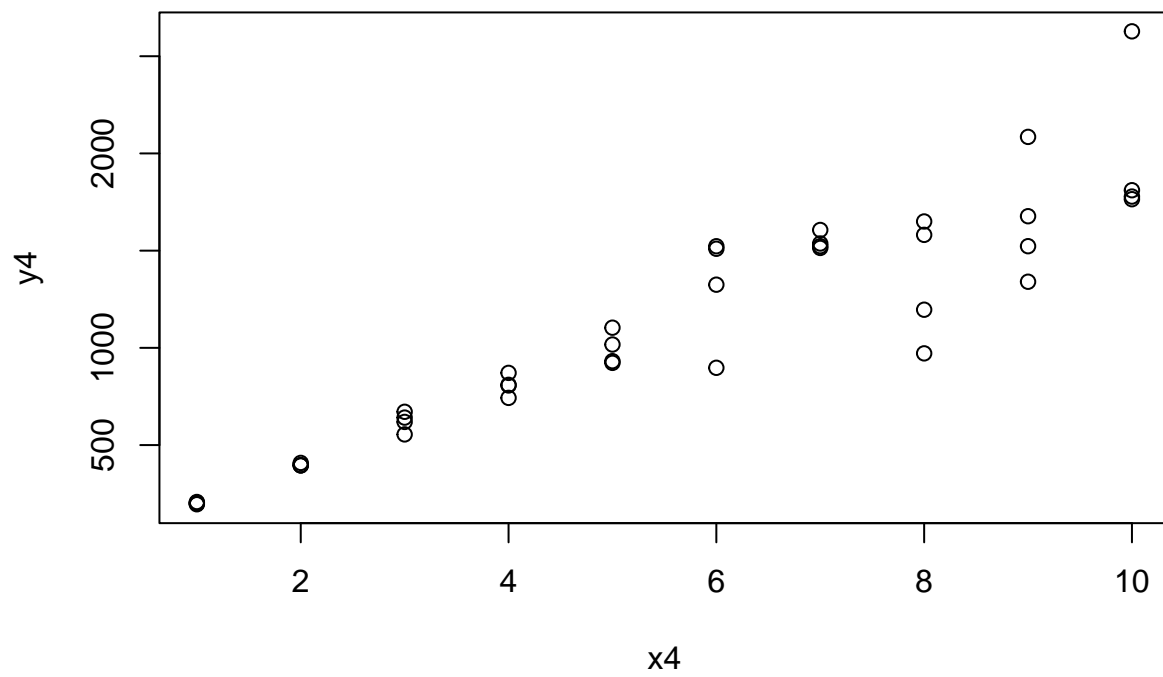



In this residual plot, we can observe a quadratic shape or a parabola appear.

c) If the residual plot shows features like parabolic shapes, we can conclude that the trend is non-linear.

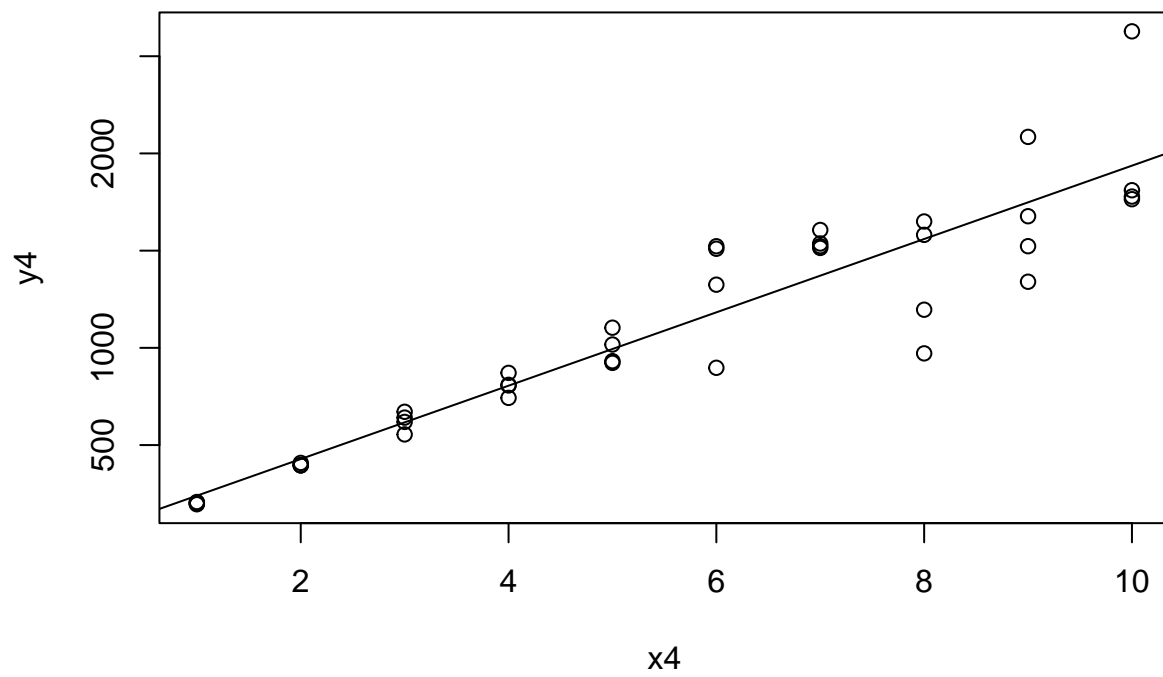
d)

```
my_nl_func <- function(a, b, x, sigma, random.seed){
  set.seed(random.seed)
  a + (b * x) + rnorm(length(x), 0, (sigma * (x^2)))
}
x4 <- rep(1:10, by = .1, 4)
y4 <- my_nl_func(1, 200, x4, 5, 123)
plot(y4 ~ x4)
```

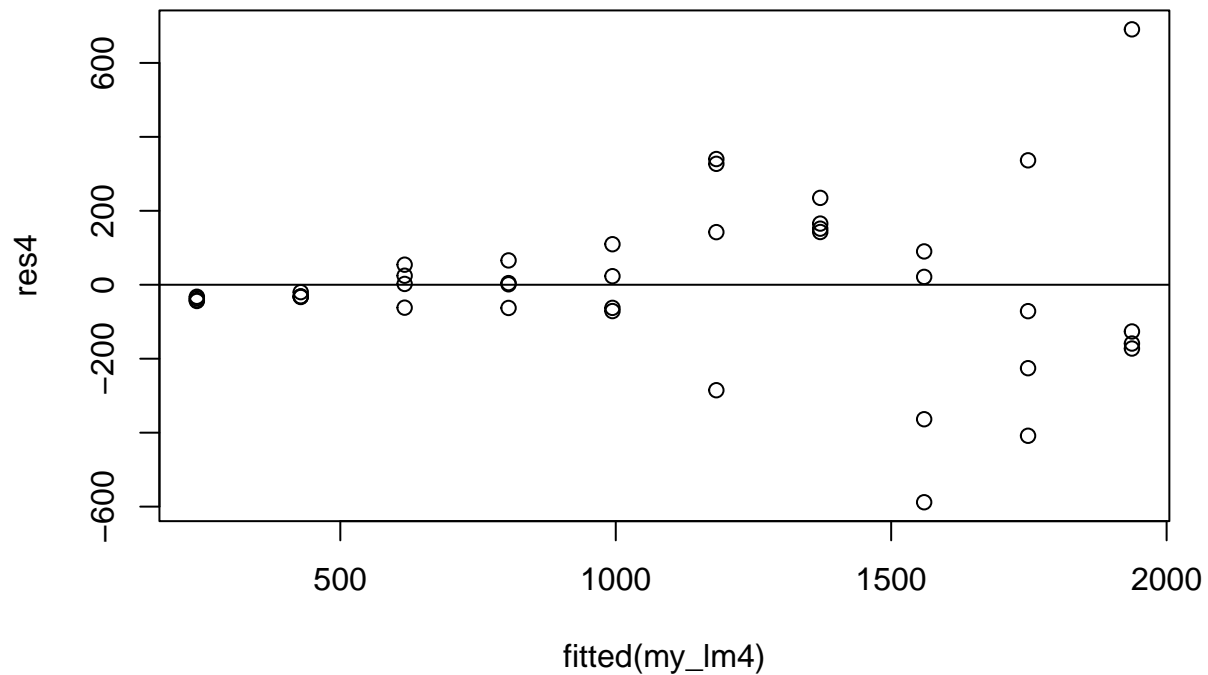


e)

```
plot(x4, y4)
my_lm4 <- lm(y4 ~ x4)
abline(my_lm4)
```



```
res4 <- resid(my_lm4)
plot(fitted(my_lm4), res4)
abline(0,0)
```



The cone shape of the residual plot indicates that the standard deviation increases as x increases, and therefore the constant standard deviation assumption is violated.

Question 4

```
atus <- read.csv('atus.csv')
atus1 <- subset(atus, homework > 0)
head(atus1)
```

```
##      caseid      state age gender      citizen marital_stat
## 24 2.01201e+13   Florida  20  Male Native, Born in USA Never married
## 32 2.01201e+13 Wisconsin 67 Female Native, Born in USA Never married
## 50 2.01201e+13   Florida  18  Male Native, Born in USA Never married
## 65 2.01201e+13   New York 43 Female Native, Born in USA      Married
## 103 2.01201e+13 Pennsylvania 30  Male Native, Born in USA Never married
## 105 2.01201e+13 Connecticut 20 Female Native, Born in USA Never married
##      veteran active_armedforces      emp_status multi_jobs
## 24 Non-Veteran                No Unemployed, Looking No answer
## 32  Veteran                No Not in labor force No answer
## 50 Non-Veteran                No      Employed      No
## 65 Non-Veteran                No      Employed      No
## 103 Non-Veteran                No      Employed      No
## 105 Non-Veteran                No Not in labor force No answer
##      work_class      retired fulltime_emp hours_worked
```

```

## 24          No answer No answer    No answer    NA
## 32          No answer No answer    No answer    NA
## 50          Private, for profit No answer    Part time    15
## 65  Self-employed, unincorporated No answer    Part time    30
## 103         Private, for profit No answer    Full time    40
## 105         No answer No answer    No answer    NA
##          fam_income household_size household_kids household_child
## 24    $12,500 to $14,999             7             2             Yes
## 32    $12,500 to $14,999             1             0             No
## 50   $100,000 to $149,999            2             0             No
## 65    $60,000 to $74,999             5             3             Yes
## 103   $50,000 to $59,999             1             0             No
## 105   $150,000 and over              6             1             Yes
##          phys_challenge travel phone volunteer religion sports social food gov_civic
## 24  Has difficulty      30      0      0      120      0      590      10      0
## 32  Has difficulty      60      0      0      0      0      225      80      0
## 50  No difficulty      30      0      0      0      0      290      30      0
## 65  No difficulty     107      0      0      0      0      90      45      0
## 103 No difficulty      70     60      0      0      0      0      95      0
## 105 No difficulty       2      0      0      0      0      150     119      0
##          household pro_services purchasing education work care_nonhousehold
## 24           0           0           0           30      0           0
## 32           0           0           0          189      0           0
## 50           0           0          10          300      0           0
## 65           0          280         105           30      0           0
## 103          0           0           0          190     515           0
## 105          0           0           0          461      0           0
##          care_household household_chores personal_care sleep groom health_related
## 24           0           0           660      630      30           0
## 32           0          315           451      420      30           1
## 50           0           0           780      735      45           0
## 65           3          45           735      670      65           0
## 103          0           0           510      480      30           0
## 105          0           5           700      690      10           0
##          eating class homework socializing holiday      day year  month      date
## 24          10      0      30           0      No  Sunday 2012  January 2012-01-22
## 32          80     70     119           0      No  Friday 2012  January 2012-01-27
## 50          30      0     300           0      No  Sunday 2012  January 2012-01-29
## 65          45      0      30          82      No  Sunday 2012  January 2012-01-22
## 103         95     100      90           0      No Thursday 2012  January 2012-01-26
## 105        119      0     461           0      No  Sunday 2012  January 2012-01-29

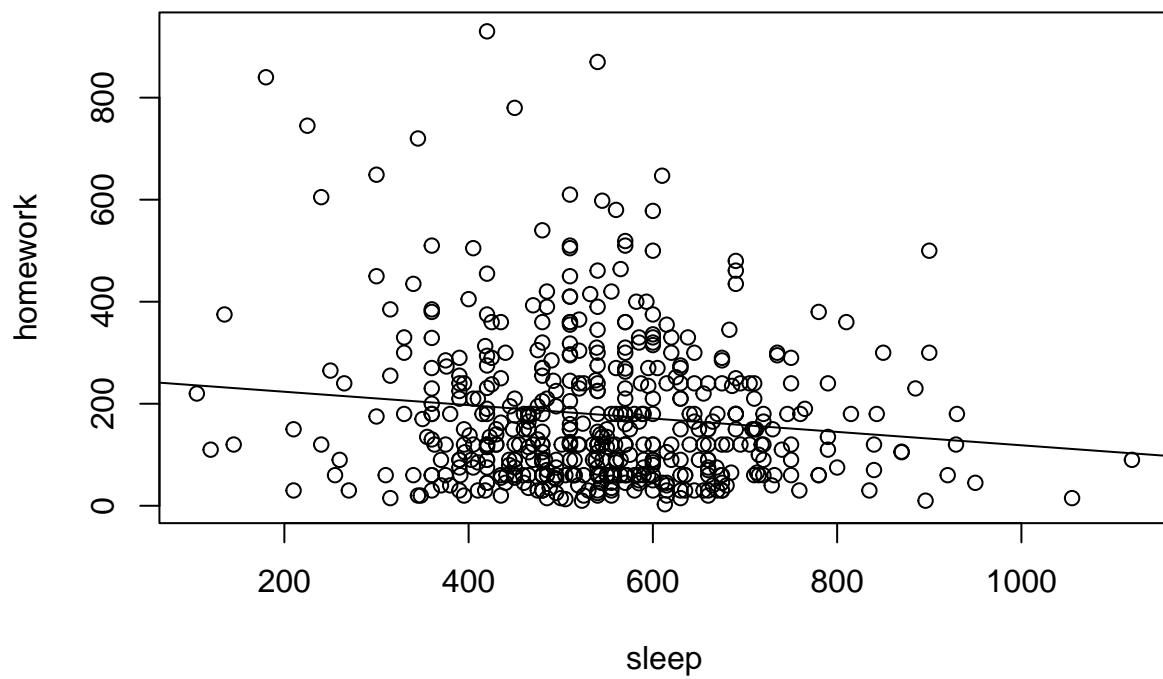
```

a)

```

plot(homework ~ sleep, data = atus1)
my_lm_2 <- lm(homework ~ sleep, data = atus1)
abline(my_lm_2)

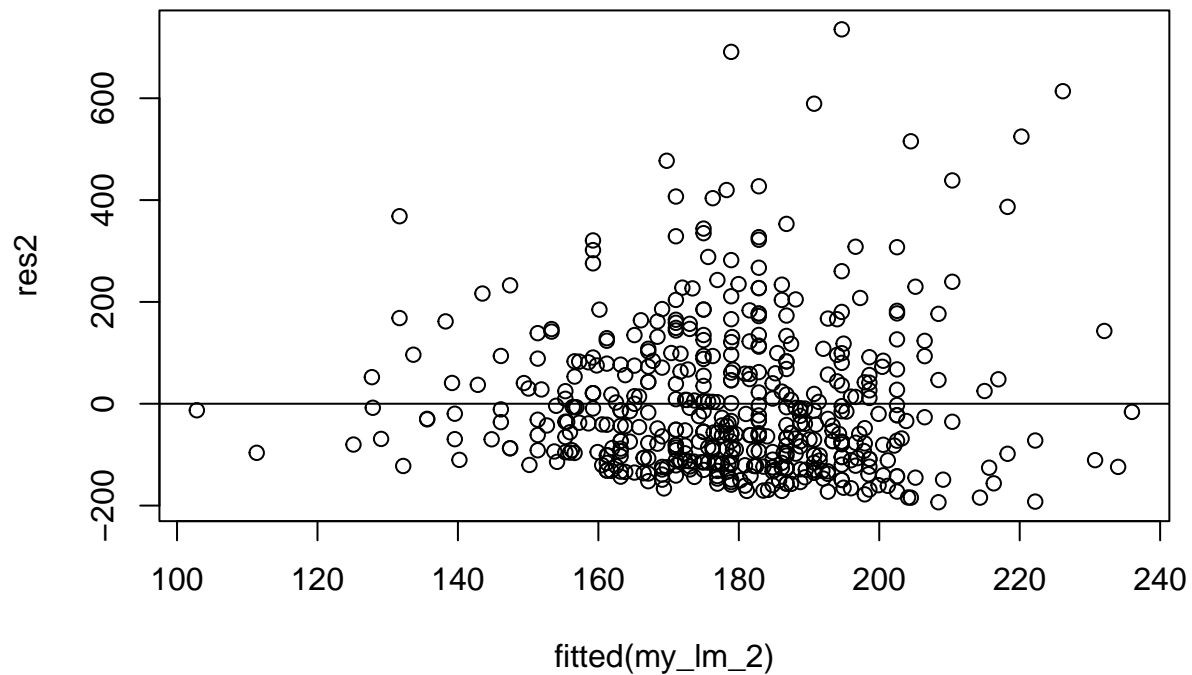
```



The linear model is not an appropriate model to describe the relationship between sleep times and homework times.

b)

```
res2 <- resid(my_lm_2)
plot(fitted(my_lm_2), res2)
abline(0,0)
```



From the residual plot, we can tell that this data is positively correlated. We can also tell that since it is not evenly distributed, a linear model would not be the best model to approximate this data.

Question 5

a)

```
t.test(atus1$household_chores ~ atus1$gender, alternative = "greater", conf.level = 0.95)
```

```
##
##  Welch Two Sample t-test
##
## data:  atus1$household_chores by atus1$gender
## t = 6.3978, df = 446.68, p-value = 1.993e-10
## alternative hypothesis: true difference in means between group Female and group Male is greater than
## 95 percent confidence interval:
##  34.62905      Inf
## sample estimates:
## mean in group Female    mean in group Male
##      77.17730           30.53052
```

Let

$$\bar{x}_f$$

represent the average time persons identifying as female spent doing chores Let

$$\bar{x}_m$$

represent the average time persons identifying as male spent doing chores

$$H_0 : \bar{x}_f - \bar{x}_m = 0$$

$$H_a : \bar{x}_f - \bar{x}_m \neq 0$$

The test statistic is 6.3978. the observed value of the statistic is 446.68. The p-value is 1.993e-10. With a 5% significance level, we reject the null hypothesis.

- b) We must assume the population distribution is normal or our sample size is sufficiently large to provide us with a good approximation. We can assume that these conditions are met because we have a large sample size observed in our data frame, which is sufficient.