STAT 423 Homework 1

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Question 1

```
a library(alr4)
```

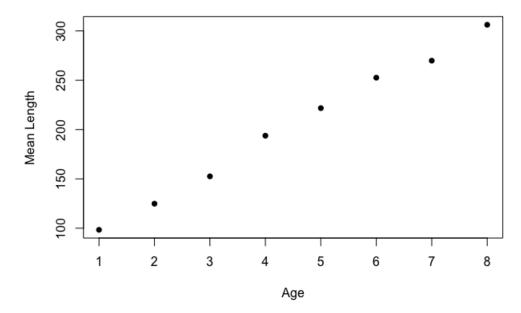
age_dt <- sort(unique(wblake\$Age))</pre>

```
leng_mean_dt <- numeric(length(age_dt))
for (i in age_dt) {
  leng_mean_dt[i] <- mean(wblake$Length[wblake$Age == i])
}</pre>
```

}
age_leng_mean_df <- data.frame(age_dt, leng_mean_dt)</pre>

plot(age_leng_mean_df, pch=16, ylab='Mean Length', xlab='Age', main='Age vs Mean of Length')

Age vs Mean of Length



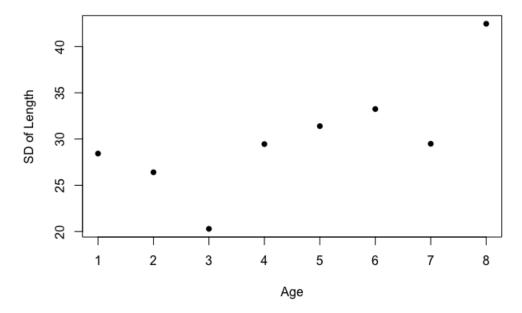
```
> age_leng_mean_df
  age_dt leng_mean_dt
1
       1
              98.34211
2
       2
             124.84722
3
       3
             152.56383
4
       4
             193.80000
            221.72059
5
       5
6
       6
            252.59770
7
       7
            269.86885
             306.25000
8
       8
```

The pattern of scatterplot shows possible linear relation between age and its corresponding mean length.

b

```
leng_sd_dt <- numeric(length(age_dt))
for (i in age_dt) {
   leng_sd_dt[i] <- sd(wblake$Length[wblake$Age == i])
}
age_leng_sd_df <- data.frame(age_dt, leng_sd_dt)
plot(age_leng_sd_df, pch=16, ylab='SD of Length', xlab='Age', main='Age vs SD of Length')</pre>
```

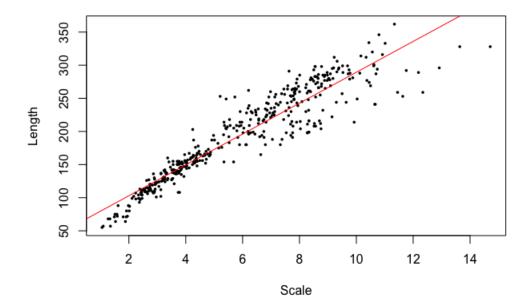
Age vs SD of Length



 \mathbf{c}

The variance of standard deviation seem to distribute around 30, except for standard deviation for age 8, which is slightly higher than others.

```
lm1 <- lm(wblake$Length~wblake$Scale)</pre>
plot(wblake$Scale, wblake$Length, pch=16, cex=0.5,
     xlab='Scale', ylab='Length')
abline(lm1, col=2)
summary(lm1)
Call:
lm(formula = wblake$Length ~ wblake$Scale)
Residuals:
   Min
             1Q Median
                             ЗQ
                                    Max
-84.896 -9.643 -0.021 14.651 75.290
Coefficients:
             Estimate Std. Error t value Pr(>|t|)
(Intercept)
              56.2986
                          2.6423
                                   21.31
                                           <2e-16 ***
wblake$Scale 23.3068
                          0.4096
                                   56.90
                                           <2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```



A simple linear regression line fits the data very well when scale is low, however, when scale is high, the residual increases.

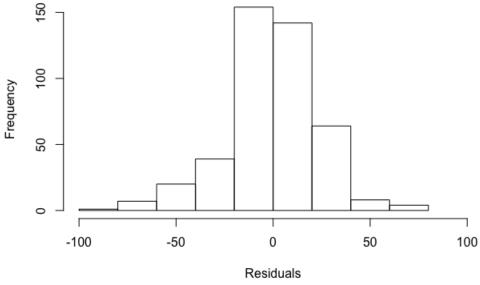
Base on the summary of linear regression, P-value of both intercept and coefficient of scale are much less than zero, indicating that both beta zero and beta one are unlikely to be zero.

\mathbf{d}

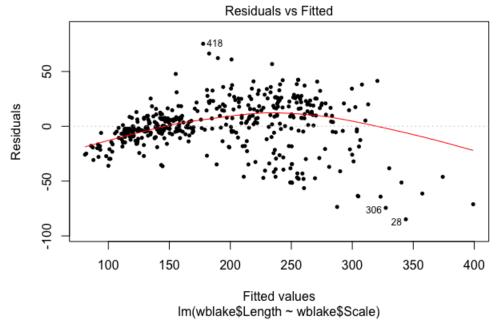
```
resid1 <- summary(lm1)$residual
hist(resid1, xlab='Residuals', main='Histogram of Residuals', xlim=c(-100, 100))
plot(lm(wblake$Length~wblake$Scale), which = 1, pch=16)</pre>
```

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The histogram has a symmetric bell shape and centered around zero, indicating the distribution of residuals is approximately normal. The normality assumption is not violated.



The pattern of TA plot shows that residuals randomly distributed around zero when fitted value is small. However, at the high end of fitted value, residuals tend to be farther away from zero. The relatively high curvature of line also reflects the unconstant variance within residuals. Therefore the constant variance assumption is violated.

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```
new_pt <- data.frame(Scale=200)</pre>
fit <- lm(Length~Scale, data=wblake)</pre>
pred_200 <- predict(fit, new_pt, se.fit = T, )$fit</pre>
pred_200_CI <- predict(fit, new_pt, interval = 'confidence', level=0.95)</pre>
pred_200_PI <- predict(fit, new_pt, interval = 'prediction', level=0.95)</pre>
> pred_200
       1
4717.665
> pred_200_CI
       fit
                 lwr
                           upr
1 4717.665 4561.348 4873.983
> pred_200_PI
       fit
                lwr
                          upr
1 4717.665 4554.91 4880.421
```

The fitted value \hat{y} for length given that scale equals 200 is 4717.665.

The 95 % confidence interval for fitted value is $4561.348 < \hat{y} < 4873.983$.

The 95 % prediction interval is [4554.91, 4880.421].

 \mathbf{e}

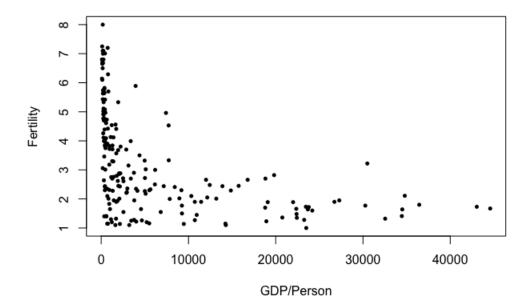
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Question 2

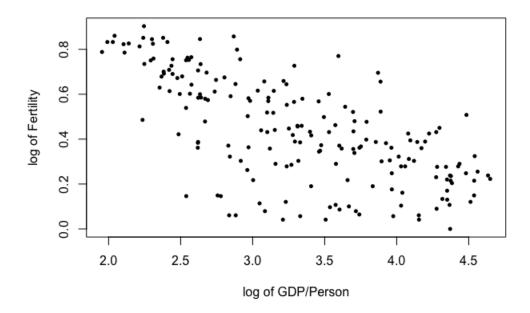
 \mathbf{a}

Using this data set, we can analyze whether the average gdp of a country affects fertility rate. For this scenario, the predictor is gdp per person, and the response is fertility rate.

b
plot(x=UN1\$PPgdp, y=UN1\$Fertility, xlab='GDP/Person', ylab='Fertility', pch=16, cex=0.7)



The observations at lower GDP is much more denser than high GDP. Though we may perceive that countries with high GDP generally have a low fertility rate, the unbalance in density makes linear model inappropriate here.



After applying log to both predictor and response, linear regression seems to fit the new data better. But the mean square error of linear model might be relatively high.

```
\mathbf{d}
```

```
lm2 <- lm(log10(Fertility)~log10(PPgdp), data=UN1)
abline(a=lm2$coefficients[[1]], b=lm2$coefficients[[2]], col=2, lwd=2)</pre>
```

> summary(lm2)

Call:

```
lm(formula = log10(Fertility) ~ log10(PPgdp), data = UN1)
```

Residuals:

```
Min 1Q Median 3Q Max -0.48587 -0.08148 0.03058 0.11327 0.39130
```

Coefficients:

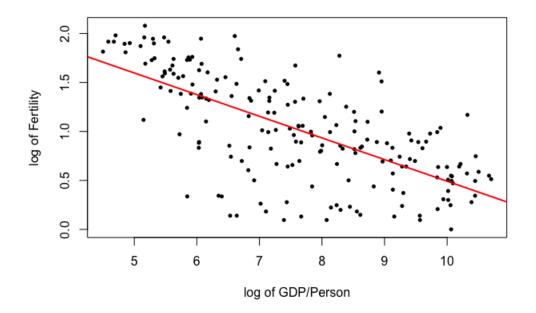
```
Estimate Std. Error t value Pr(>|t|)

(Intercept) 1.17399 0.05879 19.97 <2e-16 ***
log10(PPgdp) -0.22116 0.01737 -12.73 <2e-16 ***
---

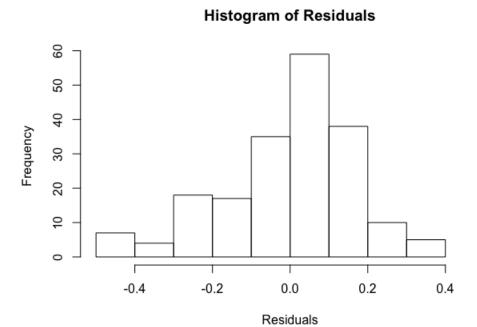
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

 \mathbf{e}

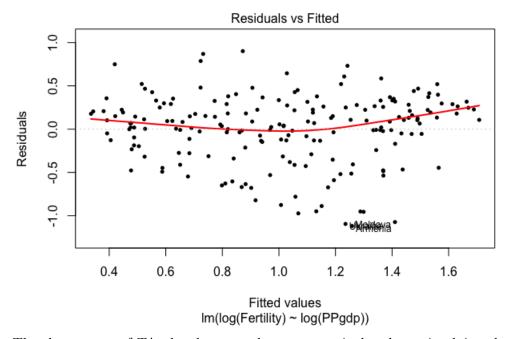
Residual standard error: 0.1721 on 191 degrees of freedom Multiple R-squared: 0.4591, Adjusted R-squared: 0.4563 F-statistic: 162.1 on 1 and 191 DF, p-value: < 2.2e-16



hist(lm2\$residuals, xlim=c(-0.5, 0.5), xlab='Residuals', main='Histogram of Residuals')
plot(lm(log(Fertility)~log(PPgdp), data=UN1), which = 1, pch=16, cex=0.7, lwd=2)



The histogram has an approximately symmetric bell shape and centered around zero, indicating the distribution of residuals is approximately normal. The normality assumption is not violated.



The the pattern of TA plot does not show any particular shape, implying that the correlation between residuals and fitted value is weak. The residuals scattered around zero randomly, therefore, the equal variance assumption is not violated

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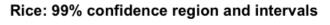
```
\mathbf{f}
```

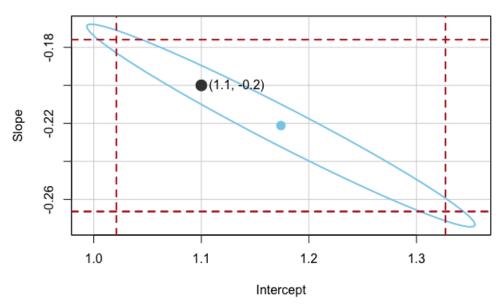
```
beta1_est <- summary(lm2)$coefficient[2,1]
beta1_se <- summary(lm2)$coefficients[2,2]
t_val <- (beta1_est - 0) / beta1_se
n <- length(UN1$Fertility)

> pt(t_val, df=n-2)
[1] 1.365501e-27
> t_val
[1] -12.73366
```

t-value of the slope is -12.73, the corresponding p-value is equal to $1.36 \cdot 10^{-27}$, given the null hypothesis is $\beta_1 = 0$. Since the p-value is much smaller than 0.01, we can conclude that there is strong evidence against the null hypothesis, i.e. the slope are unlikely to be zero.

\mathbf{g}





Under the confidence level of 99%, I will not reject the hypothesis that $(\beta_0, \beta_1) = (1.1, -0.2)$, because the value for null hypothesis falls in the 99% confidence region. Meanwhile, both slope and intercept of the null hypothesis fall in between the marginal confidence intervals.

Question 3

a

We have proved in class: $\hat{\beta}_1 = \sum c_i y_i$ where $c_i = \frac{x_i - \bar{x}}{SXX}$; $\hat{\beta}_0 = \bar{y} - \hat{\beta}_1 \bar{x}$

$$\hat{\beta}_0 = \bar{y} - \hat{\beta}_1 \bar{x}$$

$$= \frac{1}{n} \sum_{i=1}^{n} y_i - \sum_{i=1}^{n} c_i y_i \bar{x}$$

$$= \sum_{i=1}^{n} (\frac{1}{n} - \bar{x}c_i) y_i$$

This shows $\hat{\beta_0}$ can be written as $\sum_i^n d_i y_i$ where $d_i = \frac{1}{n} - \bar{x} c_i$ and $c_i = \frac{x_i - \bar{x}}{SXX}$.

b

We have proved in class: $\hat{\beta}_1$ is an unbiased estimator for β_1 , i.e. $E(\hat{\beta}_1|X=x)=E(\sum c_iy_i)=\beta_1$

Note that $\hat{\beta}_0 = \sum d_i y_i$, where $d_i = \frac{1}{n} \bar{x} c_i$; $y_i = \beta_0 + \beta_i x_i + \epsilon_i$, $E(y_i) = \beta_0 + \beta_i x_i$

$$E(\hat{\beta}_0|X=x) = E(\sum_{i=1}^{n} (\frac{1}{n} - \bar{x}c_i)y_i|X=x)$$

$$= E(\sum_{i=1}^{n} \frac{1}{n}y_i|X=x) - E(\bar{x}c_iy_i|X=x)$$

$$= \frac{1}{n}\sum_{i=1}^{n} E(y_i|X=x) - \bar{x}E(\sum_{i=1}^{n} c_iy_i|X=x)$$

$$= \frac{1}{n}\sum_{i=1}^{n} (\beta_0 + \beta_ix_i) - \bar{x}\beta_1$$

$$= \beta_0 + \beta_1 \frac{1}{n}\sum_{i=1}^{n} x_i - \bar{x}y_i$$

$$= \beta_0 + \beta_1 \bar{x} - \beta_1 \bar{x}$$

$$= \beta_0$$

This proves $\hat{\beta}_0$ is an unbiased estimator for β_0 .

 \mathbf{c}

Note that $\hat{\beta}_0 = \sum_i^n d_i y_i$, where $d_i = \frac{1}{n} \bar{x} c_i$; $Var(y_i | X = x) = Var(\epsilon_i) = \sigma^2$

$$Var(\hat{\beta_0}|X=x) = Var(\sum_{i=1}^{n} d_i y_i | X=x)$$

$$= \sum_{i=1}^{n} d_i^2 Var(y_i | X=x)$$

$$= \sigma^2 \sum_{i=1}^{n} (\frac{1}{n} - \bar{x}c_i)^2$$

$$= \sigma^2 \sum_{i=1}^{n} (\frac{1}{n^2} - \frac{2}{n}\bar{x}c_i + \bar{x}^2c_i^2)$$

Plug $c_i = \frac{x_i - \bar{x}}{SXX}$ into the equation,

$$Var(\hat{\beta}_0|X=x) = \sigma^2 \sum_{i}^{n} \left(\frac{1}{n^2} - \frac{2}{n} \frac{\bar{x}(x_i - \bar{x})}{SXX} + \frac{\bar{x}^2(x_i - \bar{x})^2}{SXX^2}\right)$$
$$= \sigma^2 \left(\sum_{i}^{n} \frac{1}{n^2} - \frac{2}{n} \frac{\bar{x} \sum_{i}^{n} (x_i - \bar{x})}{SXX} + \frac{\bar{x}^2 \sum_{i}^{n} (x_i - \bar{x})^2}{SXX^2}\right)$$

Note that $\sum_{i=1}^{n} (x_i - \bar{x}) = 0$, $\sum_{i=1}^{n} (x_i - \bar{x})^2 = SXX$,

$$Var(\hat{\beta}_0|X=x) = \sigma^2(\frac{1}{n} - 0 + \frac{\bar{x}^2 S X X}{S X X^2})$$
$$= \sigma^2(\frac{1}{n} - \frac{x^2}{S X X})$$

 \mathbf{d}

Note that $\hat{y}_i = \hat{\beta}_0 + \hat{\beta}_1 \bar{x}$, $\hat{\beta}_0 = \bar{y} - \hat{\beta}_1 \bar{x}$

$$\sum_{i}^{n} (y_{i} - \hat{y}_{i}) = \sum_{i}^{n} y_{i} - \sum_{i}^{n} \hat{y}_{i}$$

$$= \sum_{i}^{n} y_{i} - \sum_{i}^{n} \hat{\beta}_{0} + \hat{\beta}_{1}\bar{x}$$

$$= \sum_{i}^{n} y_{i} - \sum_{i}^{n} \bar{y} - \hat{\beta}_{1}\bar{x} + \hat{\beta}_{1}\bar{x} = \sum_{i}^{n} y_{i} - \sum_{i}^{n} \bar{y}$$

Note that $\sum_{i=1}^{n} y_i = n\bar{y}$, $\sum_{i=1}^{n} \bar{y} = n\bar{y}$ as well, the last equation shows $\sum_{i=1}^{n} (y_i - \hat{y}_i) = 0$

4

 \mathbf{a}

$$\frac{dRSS}{d\beta_1} = \sum_{i=1}^{n} -2x_i(y_i - \beta_1 x_i)$$

To minimize SSE, we want to find $\hat{\beta}_1$ such that $\frac{dRSS}{d\beta_1} = 0$.

$$\sum_{i}^{n} -2x_i(y_i - \beta_1 x_i) = 0$$
$$2\beta_1 \sum_{i}^{n} x_i^2 = 2\sum_{i}^{n} x_i y_i$$
$$\hat{\beta_1} = \frac{\sum_{i}^{n} x_i y_i}{\sum_{i}^{n} x_i^2}$$

This shows the least square estimation for β_1 , $\hat{\beta}_1 = \frac{\sum_{i=1}^{n} x_i y_i}{\sum_{i=1}^{n} x_i^2}$.

b

$$E(\hat{\beta}_1|X=x) = E(\frac{\sum_{i=1}^{n} x_i y_i}{\sum_{i=1}^{n} x_i^2} | X=x)$$

$$= E(\frac{\sum_{i=1}^{n} x_i (\beta_1 x_i + \epsilon_i)}{\sum_{i=1}^{n} x_i^2} | X=x)$$

$$= \beta_1 E(\frac{\sum_{i=1}^{n} x_i^2}{\sum_{i=1}^{n} x_i^2} | X=x) + E(\frac{\sum_{i=1}^{n} x_i \epsilon_i}{\sum_{i=1}^{n} x_i^2} | X=x)$$

$$= \beta_1 + \frac{1}{\sum_{i=1}^{n} x_i^2} \sum_{i=1}^{n} E(x_i \epsilon_i | X=x)$$

Since x_i and ϵ_i are independent, $E(x_i\epsilon_i) = E(x_i)E(\epsilon_i) = 0$.

Therefore, the last equation becomes $E(\hat{\beta}_1|X=x) = \beta_1 + 0 = \beta_1$.

 \mathbf{c}

$$Var(\hat{\beta}_{1}|X = x) = Var(\frac{\sum_{i}^{n} x_{i} y_{i}}{\sum_{i}^{n} x_{i}^{2}} | X = x)$$

$$= \frac{1}{(\sum_{i}^{n} x_{i}^{2})^{2}} Var(\sum_{i}^{n} x_{i} y_{i} | X = x)$$

$$Var(\sum_{i}^{n} x_{i}y_{i}|X = x) = E((\sum_{i}^{n} x_{i}y_{i})^{2}|X = x) - E(\sum_{i}^{n} x_{i}y_{i}|X = x)^{2}$$

$$= E(\sum_{i}^{n} \sum_{j}^{n} (x_{i}y_{i})(x_{j}y_{j})|X = x) - E(\sum_{i}^{n} x_{i}y_{i}|X = x)E(\sum_{j}^{n} x_{j}y_{j}|X = x)$$

$$= \sum_{i}^{n} \sum_{j}^{n} [E((x_{i}y_{i})(x_{j}y_{j})|X = x) - E(x_{i}y_{i}|X = x)E(x_{j}y_{j}|X = x)]$$

$$= \sum_{i}^{n} \sum_{j}^{n} Cov(x_{i}y_{i}, x_{j}y_{j}|X = x)$$

$$= \sum_{i}^{n} \sum_{j}^{n} x_{i}x_{j}Cov(y_{i}, y_{j}|X = x)$$

Note that when $i \neq j$, $Cov(y_i, y_j) = 0$ since they are independent to each other.

Therefore, we get $\sum_{i=1}^{n} \sum_{j=1}^{n} x_i x_j Cov(y_i, y_j | X = x) = \sum_{i=1}^{n} x_i^2 Var(y_i)$.

$$Var(\hat{\beta}_1|X=x) = \frac{1}{(\sum_{i=1}^{n} x_i^2)^2} \sum_{i=1}^{n} x_i^2 Var(y_i|X=x)$$

Note that $Var(y_i|X=x) = Var(\beta_i x_i + \epsilon_i|X=x)$ where β_1 and x_i are constant. Thus $Var(y_i|X=x) = Var(\epsilon_i) = \sigma^2$.

$$Var(\hat{\beta}_1|X=x) = \frac{1}{(\sum_i^n x_i^2)^2} \sum_i^n x_i^2 \sigma^2$$
$$= \frac{\sigma^2}{\sum_i^n x_i^2}$$