



CMP4011 Big Data and Cloud Computing

Lab 5 Linear Regression

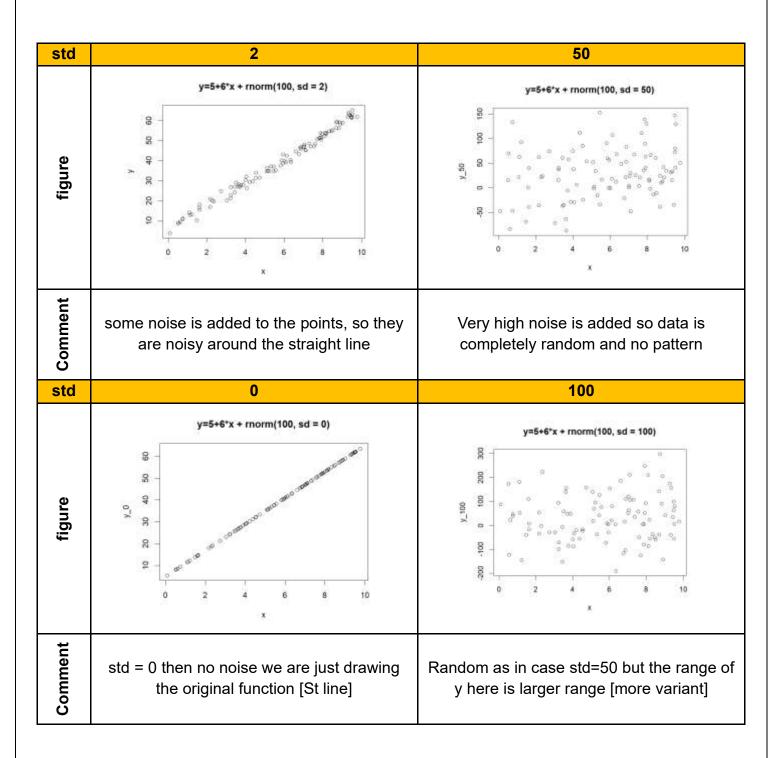
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Page 1 of 15 Linear Regression

Part (1)

(Q1) Try changing the value of standard deviation (sd) in the next command. How do the data points change for different values of standard deviation?

Gold: y=5+6x. Y is a linear transformation of input feature x.

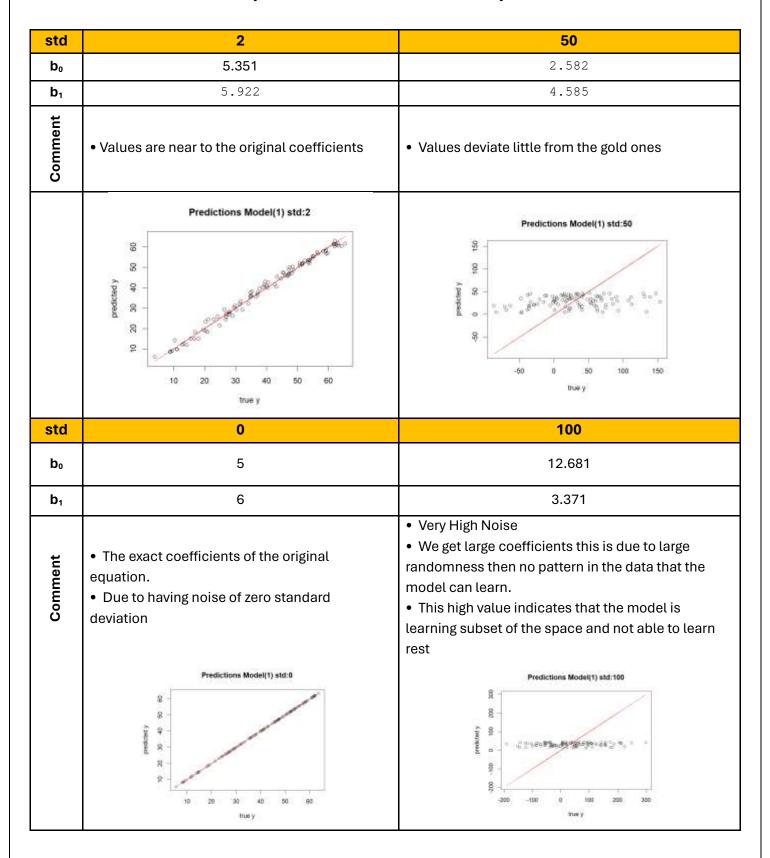


Page 2 of 15 Linear Regression

(Q2) How are the coefficients of the linear model affected by changing the value of standard deviation in Q1?

y=b0+b1x

Gold: y=5+6x.



Page **3** of **15** Linear Regression

Note: ML is all about learning from data so nothing could be leant from a random data

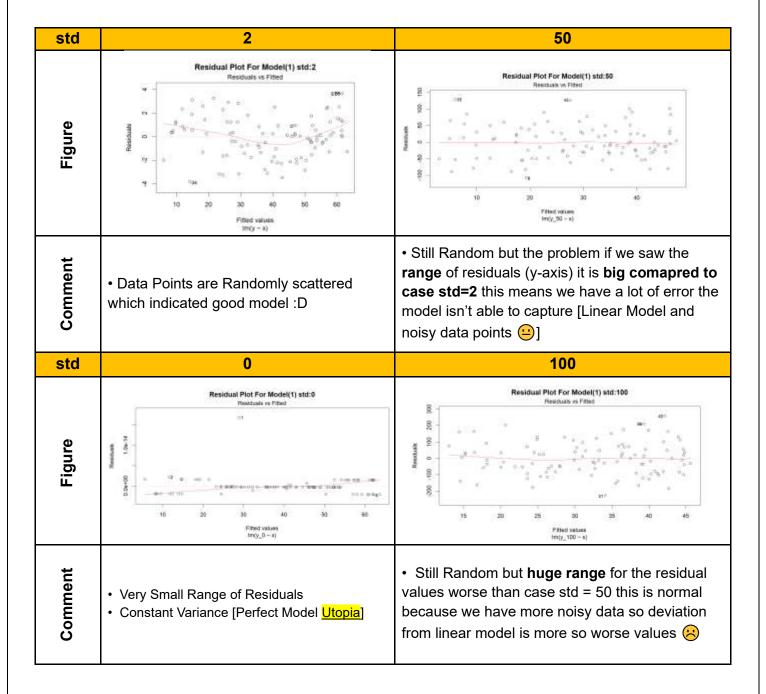
Conclusion: Higher std in noise make more noisy data so due learning this noisy data the model overfits because of trying to learn this noisy pattern this overfitting is obvious in the higher values for the coefficient as the data is more noisy

(Q3) How is the value of R-squared affected by changing the value of standard deviation in Q1?

std	2	50	
R ²	0.9899548	0.0582	
Comment	Near 1 but not perfect 1 because data has some noise values so the no line can perfectly fit the data points [Data becomes not perfectly linear]	Very Small Value due to high value noise so fitting data point with line will get bad results [High Error]	
std	0	100	
R ²	1	0.008	
Comment	• 1 Perfect © because no noise is added and also data is generated from linear line so linear regression will perfectly learn the target function achieving R ² = 1	Very Small even smaller than case of std 50 due to higher std for noise then we get more points more randomly deviated from the gold	

Page **4** of **15** Linear Regression

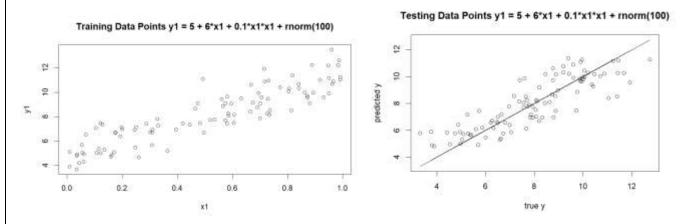
(Q4) What do you conclude about the residual plot? Is it a good residual plot?



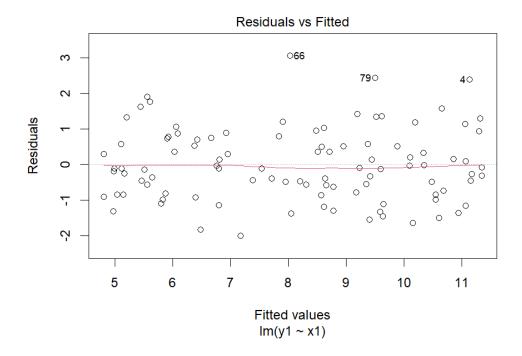
Page **5** of **15** Linear Regression

Part (2):

(Q5) What do you conclude about the residual plot? Is it a good residual plot?



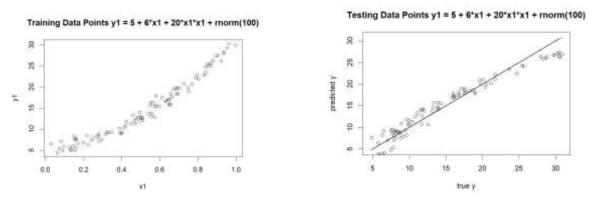
The Training & Testing Data are noisy linear [0.*x2]



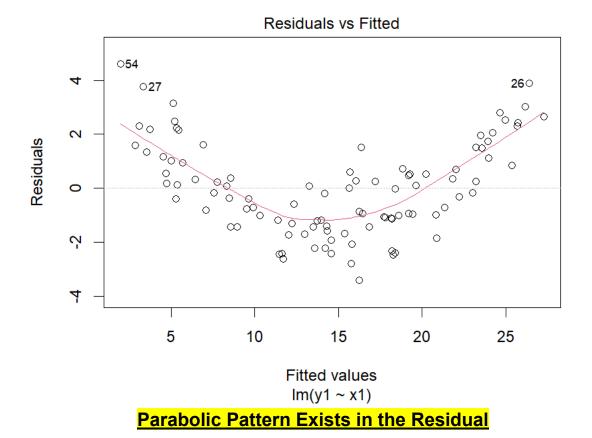
The residuals plot is random with no pattern exist which suggest it is good I think the only problem is bit large range for the residuals.

Page 6 of 15 Linear Regression

(Q6) Now, change the coefficient of the non-linear term in the original model for (A) training.



The Training & Testing Data are non-linear (Parabola like :D)



If a pattern exists in the residual plot, it suggests that the model's **predictions are not capturing all the information** present in the data, indicating potential **issues** with the model's specification or **assumption**. Which is clear than Linear Regression **assumes Linear Data** but the data we are using for test and train are from second order [Nonlinearly:D]

Page **7** of **15** Linear Regression

Part (3):

Q (7) Import the data set LungCapData.tsv. What are the variables in this dataset?

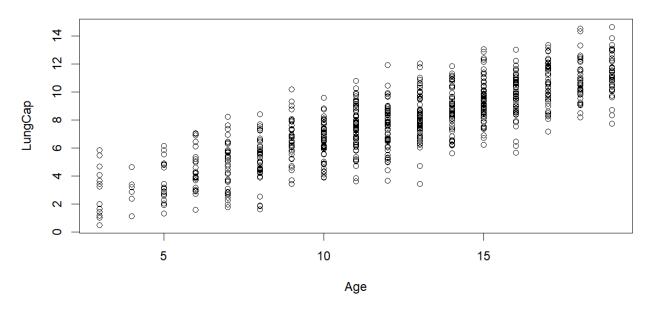
- ✓ LungCap [Continues]
- ✓ Age [Continues]
- √ Height [Continues]
- ✓ Smoke [Categorial]
- ✓ Gender [Categorial]
- ✓ Caesarean [Categorial] → [Target]

Info About each col for better understanding for me:

- LungCap→ This could represent lung capacity, which is the maximum amount of air that a
 person can inhale and exhale from their lungs.
- Caesarean → This variable likely represents whether the individuals were born via Caesarean section (C-section) or not.

Q (8) Draw a scatter plot of Age (x-axis) vs. LungCap (y-axis). Label x-axis "Age" and y-axis "LungCap"

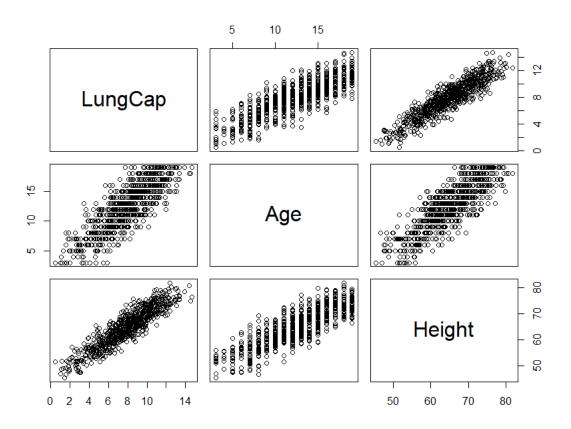
Scatter Plot of Age vs. LungCap



In general, there is increasing lung Cap as Age increases.

Page 8 of 15 Linear Regression

(Q9) Draw a pair-wise scatter plot between Lung Capacity, Age and Height.



Height and Lung Capacity are more correlated than Age and Lung Capacity

(Q10) Calculate correlation between Age and LungCap, and between Height and LungCap

```
> print(paste("Age and LungCap Correlation:", cor_age_lungcap))
[1] "Age and LungCap Correlation: 0.819674897498941"
> # Calculate correlation between Height and LungCap
> cor_height_lungcap <- cor(df$Height, df$LungCap)
> print(paste("Height and LungCap Correlation:", cor_height_lungcap))
[1] "Height and LungCap Correlation: 0.912187323133179"
> |
```

The correlation between Height & Lung Capacity is more than that between Age and Lung

Capacity which agrees with results in Q 9 (2)

Page 9 of 15 Linear Regression

(Q11) Which of the two input variables (Age, Height) are more correlated to the dependent variable (LungCap)?

Height

(Q12) Do you think the two variables (Height and LungCap) are correlated? why?

Yes, they are. In practical terms, this could imply that taller individuals tend to have larger lung capacities compared to shorter individuals. <u>However, correlation does not imply causation, so further analysis would be needed to determine the exact relationship between these variables.</u>

(Q13) Fit a liner regression model where the dependent variable is LungCap and use all other variables as the independent variables.

```
#(Q13) Fit a liner regression model where the dependent variable is LungCap
#and use all other variables as the independent variables
|m_model <- lm(LungCap - .,data = df) # the dot . represents all other variables in the dataset except the dependent
```

(Q14) Show a summary of this model.

```
R 4.3.2 · D:/Big Data Labs/Lab 5 - Predictive Analysis II/Linear Regression Requirement/
> #(Q14) Show a summary of this model
> summary(1m_model)
Call:
lm(formula = LungCap \sim ., data = df)
Residuals:
           1Q Median
                        3Q
                               Max
-3.3388 -0.7200 0.0444 0.7093 3.0172
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) -11.32249  0.47097 -24.041 < 2e-16 ***
         Age
Height
Smokeyes -0.60956 0.12598 -4.839 1.60e-06 ***
            Gendermale
Caesareanyes -0.21422 0.09074 -2.361 0.0185 *
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
Residual standard error: 1.02 on 719 degrees of freedom
Multiple R-squared: 0.8542, Adjusted R-squared: 0.8532
F-statistic: 842.8 on 5 and 719 DF, p-value: < 2.2e-16
```

Page 10 of 15 Linear Regression

(Q15) What is the R-squared value here? What does R-squared indicate?

```
Multiple R-squared: 0.8542, Adjusted R-squared: 0.8532
```

It indicates that 85.42% of the variance in the dependent variable LungCap is explained by the independent variables (Used by the model). In other words, if we draw the linear Model plane in space and the training points, we will see the points are highly around the plane.

Note: Difference between multiple vs adjusted R-squared:

Multiple R-squared (R2)

- Measures the proportion of the variance in the dependent variable that is explained by the independent variables in the model.
- Increases whenever a new predictor is added to the model, even if the predictor is not relevant.
- Does not penalize for overfitting or the inclusion of irrelevant predictors.

Adjusted R-squared(R2adj)

- Adjusts the R2 value to account for the number of predictors in the model.
- Penalizes the inclusion of irrelevant predictors by decreasing if unnecessary predictors are added to the model.
- Provides a more accurate measure of the goodness of fit when comparing models with different numbers of predictors.

Generally, **R2adj** is lower than the **R2** if unnecessary predictors are included in the **model**. The numbers above are very close to the third digit after decimal point so we will see if there are unnecessary predictors?! IThink no

Page 11 of 15 Linear Regression

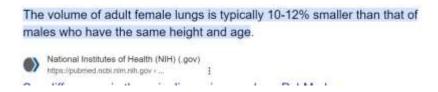
(Q16) Show the coefficients of the linear model. Do they make sense?

```
> #(Q16) Show the coefficients of the linear model. Do they make sense?
> #If not, which variables don't make sense? What should you do?
> print(lm_model$coefficients)
(Intercept) Age Height Smokeyes Gendermale Caesareanyes
-11.3224856 0.1605296 0.2641128 -0.6095592 0.3870117 -0.2142182
> |
```

Yes, they make Sense. 😉 🨉

- Age and Height have positive coefficients, which is logic becoming older means your body is growing up. Same for height taller means your larger capacity
- Smoking has a negative coefficient, yes smoking causes lung problems including cause of smaller lung capacity.
- GenderMale coefficient is 0.38701. Since this coefficient is positive, it means that there is a positive association between being male and lung capacity, holding other variables constant.

If an individual is female, then the coefficient for Gendermale does not directly apply because Gendermale would be 0 for females. Which means Males has more lung capacity which may be biologically correct [I googled it]



➤ CaesareanYes coefficient is negative this means that individuals with a history of Caesarean birth, on average, have a lung capacity that is lower by 0.21422 units compared to individuals without a history of Caesarean birth, holding all other variables constant. Which may seem logic 😊

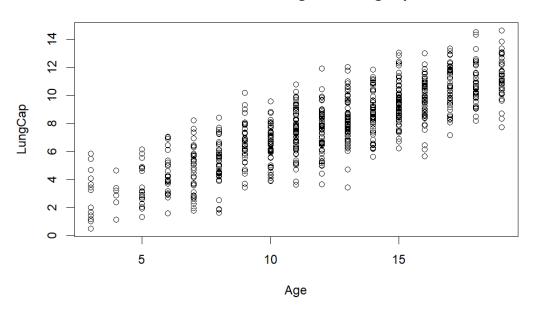
Lung function tests were carried out in the first 6 hours of 8fe on 26 babies born by vaginal delivery and 10 born by caesarean section. The babies born by caesarean section had a mean thoracic gas volume of only 19.7 milking body weight compared with 32.7 milking for the babies born vaginally.

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Page 12 of 15 Linear Regression

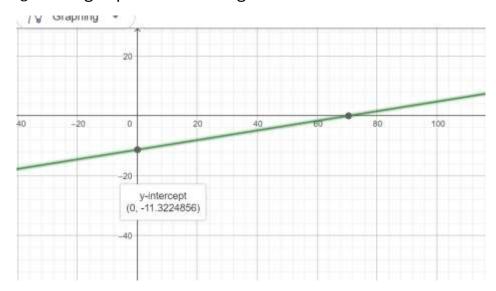
(Q17) Redraw a scatter plot between Age and LungCap. Display/Overlay the linear model (a line) over it.

Scatter Plot of Age vs. LungCap



The Linear Model Line isn't shown 😕 😕 ?

The line of the model to be drawn will have the first coefficients only which are the y-intercept and the coefficient of the Age. LungCap=0.1605296 Age - 11.3224856



The line of separator is below the graph values we have its y-intercept -11.32 and x-intercept=70 so we cannot see it here ①

Page 13 of 15 Linear Regression

(Q18) Repeat Q13 but with these variables Age, Smoke and Cesarean as the only independent variables.

```
#(Q18)Repeat Q13 but with these variables Age, Smoke and Cesarean 
1m_model_2 <- lm(LungCap ~ Age + Smoke + Caesarean,data = df)
```

(Q19) Repeat Q16, Q17 for the new model. What happened?

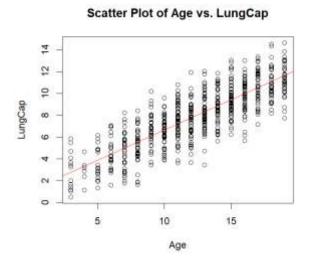
1. Show the coefficients of the linear model. Do they make sense?

```
> #(Q19)Repeat Q16, Q17 for the new model. What happened?
> print(lm_model_2$coefficients)
(Intercept) Age Smokeyes Caesareanyes
1.1086723 0.5561667 -0.6431029 -0.1460278
> print("Yes the Make Sense More Details are in the Report :D")
[1] "Yes the Make Sense More Details are in the Report :D"
```

I see the most significant difference is that y-intercept is positive and the Age coefficient is 0.55 instead of 0.16 which I think is logic we have remove other predictors so instead model positively more rely on Age

2. Redraw a scatter plot between Age and LungCap. Display/Overlay the linear model (a line) over it.

The line of the model to be drawn will have the first coefficients only which are the y-intercept and the coefficient of the Age. LungCap=0.556 Age + 1.1086



Page **14** of **15** Linear Regression

The line of separator is below the graph values we have its y-intercept -11.32 and x-intercept=70 so we cannot see it here ①

(Q20) Predict results for this regression line on the training data.

```
\#(Q20)Predict results for this regression line on the training data. predictions <- predict(lm_model_2, newdata = df)
```

predictions

Named num [1:725] 4.45 10.48 9.86 8.9 3...

```
> #(Q21)Calculate the mean squared error (MSE)of the training data.
> # Calculate Mean Squared Error (MSE)
> mse <- mean((df$LungCap - predictions)^2)
> print(paste("Mean Squared Error (MSE) of the training data:", mse))
[1] "Mean Squared Error (MSE) of the training data: 2.28016929745408"
> |
```

I Think it will be lower for Model (1)

For Model 1 we get less MSE [Better Model] This sounds logic because in Model 2 we have dropped regraters (Height, Gender) which are significantly important. This AGREES with Q (15) where R-squared is high

Page 15 of 15 Linear Regression