

Lab 4

Karen Lopez

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Note: the content of this lab is on the midterm exam (March 5) even though the lab itself is due after the midterm exam.

We now move on to simple linear modeling using the ordinary least squares algorithm.

Let's quickly recreate the sample data set from practice lecture 7:

```
n = 20
x = runif(n)
beta_0 = 3
beta_1 = -2
y = beta_0 + beta_1 * x + rnorm(n, mean = 0, sd = 0.33)
```

Solve for the least squares line by computing b_0 and b_1 *without* using the functions `mean`, `cor`, `cov`, `var`, `sd` but instead computing it from the x and y quantities manually using base function such as `sum` and other basic operators. See the class notes.

```
mean_x = sum(x) / n
mean_y = sum(y) / n
b_1 = (sum((x * y)) - n * mean_x * mean_y) / (sum(x^2) - n*mean_x^2)
b_0 = mean_y - b_1 * mean_x
```

Verify your computations are correct using the `lm` function in R:

```
lm_mod = lm(y~x)
b_vec = coef(lm_mod)
pacman::p_load(testthat)
expect_equal(b_0, as.numeric(b_vec[1]), tol = 1e-4)
expect_equal(b_1, as.numeric(b_vec[2]), tol = 1e-4)
```

6. We are now going to repeat one of the first linear model building exercises in history — that of Sir Francis Galton in 1886. First load up package `HistData`.

```
pacman::p_load(HistData)
```

In it, there is a dataset called `Galton`. Load it up.

```
data("Galton")
```

You now should have a data frame in your workspace called `Galton`. Summarize this data frame and write a few sentences about what you see. Make sure you report n , p and a bit about what the columns represent and how the data was measured. See the help file `?Galton`.

```
summary(Galton)
```

##	parent	child
##	Min. :64.00	Min. :61.70
##	1st Qu.:67.50	1st Qu.:66.20
##	Median :68.50	Median :68.20
##	Mean :68.31	Mean :68.09
##	3rd Qu.:69.50	3rd Qu.:70.20
##	Max. :73.00	Max. :73.70

```
str(Galton)
```

```
## 'data.frame':   928 obs. of  2 variables:
## $ parent: num  70.5 68.5 65.5 64.5 64 67.5 67.5 67.5 66.5 66.5 ...
## $ child : num  61.7 61.7 61.7 61.7 61.7 62.2 62.2 62.2 62.2 62.2 ...
```

```
?Galton
```

n = 928 observations p = 2 variables - represented in two columns one named parent which represents the average height of the father and mother and the second column named chil which is the height of the child

Find the average height (include both parents and children in this computation).

```
avg_height = (sum(Galton$parent+Galton$child))/(nrow(Galton)*2)
```

```
mean(c(Galton$parent,Galton$child))
```

```
## [1] 68.19833
```

```
(sum(Galton$parent)+sum(Galton$child))/(928*2)
```

```
## [1] 68.19833
```

If you were to use the null model, what would the RMSE be of this model be?

```
y_hat=rep(avg_height,1856) #null model
y_vec=c(Galton$parent,Galton$child)
SSE=sum((y_vec-y_hat)^2)
MSE=SSE/1854
RMSE=sqrt(MSE)
```

```
RMSE
```

```
## [1] 2.186179
```

Note that in Math 241 you learned that the sample average is an estimate of the “mean”, the population expected value of height. We will call the average the “mean” going forward since it is probably correct to the nearest tenth of an inch with this amount of data.

Run a linear model attempting to explain the childrens’ height using the parents’ height. Use `lm` and use the R formula notation. Compute and report b_0 , b_1 , RMSE and R^2 . Use the correct units to report these quantities.

```
ols1 = lm(child~parent,Galton)
summary(ols1)
```

```
##
## Call:
## lm(formula = child ~ parent, data = Galton)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -7.8050 -1.3661  0.0487  1.6339  5.9264
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  23.94153    2.81088   8.517  <2e-16 ***
## parent        0.64629    0.04114  15.711  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
##
## Residual standard error: 2.239 on 926 degrees of freedom
## Multiple R-squared:  0.2105, Adjusted R-squared:  0.2096
## F-statistic: 246.8 on 1 and 926 DF,  p-value: < 2.2e-16
```

```
length(ols1$residuals)
```

```
## [1] 928
```

```
sse = sum(ols1$residuals^2) #units= inches squared
mse = sse / (928 - 2) #units = inches squared
rmse = sqrt(mse) #units = inches
rmse #
```

```
## [1] 2.238547
```

Interpret all four quantities: b_0 , b_1 , RMSE and R^2 .

$b_0 = 23.9415$ inches (intercept) $b_1 = 0.6463$ inches (For a one inch increase in the parents height the child high increases by .6463 inches) $R^2 = 0.2105$ (21 percent) (21% of variance in y explained) $rmse = 2.238547$ inches (the model is plus or minus 2.23 inch off of the actual.. the model plus or minus 4.46 is 95% confidence set of y)

How good is this model? How well does it predict? Discuss.

Since the R^2 represents 21% of the variance in y. Also based on the RMSE your off by +/- 4 inches. Therefore, it does better than the null model but it is not a great model.

It is reasonable to assume that parents and their children have the same height? Explain why this is reasonable using basic biology and common sense.

This is reasonable to assume because of the way genetics work so it is expected that children reflect their parents because they share genes.

If they were to have the same height and any differences were just random noise with expectation 0, what would the values of β_0 and β_1 be?

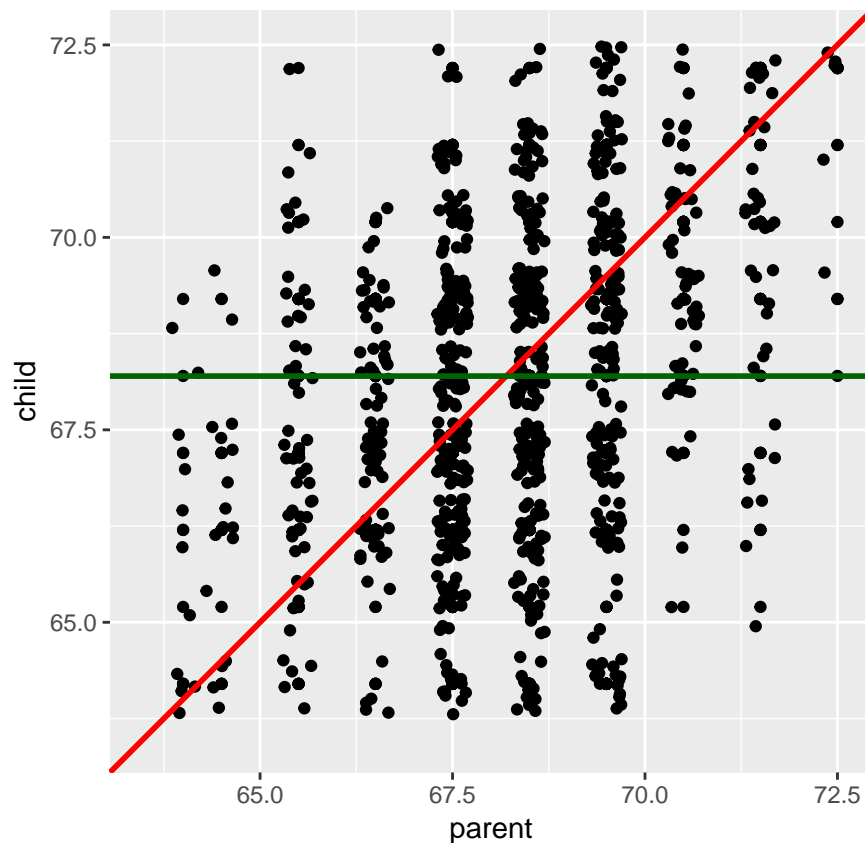
TO-DO

Let's plot (a) the data in \mathbb{D} as black dots, (b) your least squares line defined by b_0 and b_1 in blue, (c) the theoretical line β_0 and β_1 if the parent-child height equality held in red and (d) the mean height in green.

```
pacman::p_load(ggplot2)
ggplot(Galton, aes(x = parent, y = child)) +
  geom_point() +
  geom_jitter() +
  geom_abline(intercept = b_0, slope = b_1, color = "blue", size = 1) +
  geom_abline(intercept = 0, slope = 1, color = "red", size = 1) +
  geom_abline(intercept = avg_height, slope = 0, color = "darkgreen", size = 1) +
  xlim(63.5, 72.5) +
  ylim(63.5, 72.5) +
  coord_equal(ratio = 1)
```

```
## Warning: Removed 76 rows containing missing values (geom_point).
```

```
## Warning: Removed 88 rows containing missing values (geom_point).
```



Fill in the following sentence:

Children of short parents became taller on average and children of tall parents became shorter on average.

Why did Galton call it “Regression towards mediocrity in hereditary stature” which was later shortened to “regression to the mean”?

Galton called it “Regression towards mediocrity in hereditary stature” because essentially the regression line or relationship produced brings us towards the mediocre values essentially average or moderate values of height believed to be a hereditary construction.

Why should this effect be real?

When n is sufficiently large or we have enough observations, heights tend to go towards a central tendency.

You now have unlocked the mystery. Why is it that when modeling with y continuous, everyone calls it “regression”? Write a better, more descriptive and appropriate name for building predictive models with y continuous.

TO-DO

Create a dataset \mathbb{D} which we call Xy such that the linear model as R^2 about 50% and RMSE approximately 1.

```
x = #TO-DO
y = #TO-DO
Xy = data.frame(x = x, y = y)
```

Create a dataset \mathbb{D} which we call Xy such that the linear model as R^2 about 0% but x, y are clearly associated.

```
#circle
x = 2^2 + 3^2
```

```
y = 13
Xy = data.frame(x = x, y = y)
```

Load up the famous iris dataset and drop the data for Species “virginica”.

```
data("iris")
summary(iris)
```

```
##      Sepal.Length      Sepal.Width      Petal.Length      Petal.Width
##  Min.      :4.300    Min.      :2.000    Min.      :1.000    Min.      :0.100
##  1st Qu.:5.100    1st Qu.:2.800    1st Qu.:1.600    1st Qu.:0.300
##  Median :5.800    Median :3.000    Median :4.350    Median :1.300
##  Mean   :5.843    Mean   :3.057    Mean   :3.758    Mean   :1.199
##  3rd Qu.:6.400    3rd Qu.:3.300    3rd Qu.:5.100    3rd Qu.:1.800
##  Max.   :7.900    Max.   :4.400    Max.   :6.900    Max.   :2.500
##           Species
##  setosa      :50
##  versicolor:50
##  virginica  :50
##
##
##
```

```
iris=iris[iris$Species != "virginica",]
iris
```

```
##      Sepal.Length Sepal.Width Petal.Length Petal.Width      Species
## 1           5.1         3.5         1.4         0.2      setosa
## 2           4.9         3.0         1.4         0.2      setosa
## 3           4.7         3.2         1.3         0.2      setosa
## 4           4.6         3.1         1.5         0.2      setosa
## 5           5.0         3.6         1.4         0.2      setosa
## 6           5.4         3.9         1.7         0.4      setosa
## 7           4.6         3.4         1.4         0.3      setosa
## 8           5.0         3.4         1.5         0.2      setosa
## 9           4.4         2.9         1.4         0.2      setosa
## 10          4.9         3.1         1.5         0.1      setosa
## 11          5.4         3.7         1.5         0.2      setosa
## 12          4.8         3.4         1.6         0.2      setosa
## 13          4.8         3.0         1.4         0.1      setosa
## 14          4.3         3.0         1.1         0.1      setosa
## 15          5.8         4.0         1.2         0.2      setosa
## 16          5.7         4.4         1.5         0.4      setosa
## 17          5.4         3.9         1.3         0.4      setosa
## 18          5.1         3.5         1.4         0.3      setosa
## 19          5.7         3.8         1.7         0.3      setosa
## 20          5.1         3.8         1.5         0.3      setosa
## 21          5.4         3.4         1.7         0.2      setosa
## 22          5.1         3.7         1.5         0.4      setosa
## 23          4.6         3.6         1.0         0.2      setosa
## 24          5.1         3.3         1.7         0.5      setosa
## 25          4.8         3.4         1.9         0.2      setosa
## 26          5.0         3.0         1.6         0.2      setosa
## 27          5.0         3.4         1.6         0.4      setosa
## 28          5.2         3.5         1.5         0.2      setosa
```

## 29	5.2	3.4	1.4	0.2	setosa
## 30	4.7	3.2	1.6	0.2	setosa
## 31	4.8	3.1	1.6	0.2	setosa
## 32	5.4	3.4	1.5	0.4	setosa
## 33	5.2	4.1	1.5	0.1	setosa
## 34	5.5	4.2	1.4	0.2	setosa
## 35	4.9	3.1	1.5	0.2	setosa
## 36	5.0	3.2	1.2	0.2	setosa
## 37	5.5	3.5	1.3	0.2	setosa
## 38	4.9	3.6	1.4	0.1	setosa
## 39	4.4	3.0	1.3	0.2	setosa
## 40	5.1	3.4	1.5	0.2	setosa
## 41	5.0	3.5	1.3	0.3	setosa
## 42	4.5	2.3	1.3	0.3	setosa
## 43	4.4	3.2	1.3	0.2	setosa
## 44	5.0	3.5	1.6	0.6	setosa
## 45	5.1	3.8	1.9	0.4	setosa
## 46	4.8	3.0	1.4	0.3	setosa
## 47	5.1	3.8	1.6	0.2	setosa
## 48	4.6	3.2	1.4	0.2	setosa
## 49	5.3	3.7	1.5	0.2	setosa
## 50	5.0	3.3	1.4	0.2	setosa
## 51	7.0	3.2	4.7	1.4	versicolor
## 52	6.4	3.2	4.5	1.5	versicolor
## 53	6.9	3.1	4.9	1.5	versicolor
## 54	5.5	2.3	4.0	1.3	versicolor
## 55	6.5	2.8	4.6	1.5	versicolor
## 56	5.7	2.8	4.5	1.3	versicolor
## 57	6.3	3.3	4.7	1.6	versicolor
## 58	4.9	2.4	3.3	1.0	versicolor
## 59	6.6	2.9	4.6	1.3	versicolor
## 60	5.2	2.7	3.9	1.4	versicolor
## 61	5.0	2.0	3.5	1.0	versicolor
## 62	5.9	3.0	4.2	1.5	versicolor
## 63	6.0	2.2	4.0	1.0	versicolor
## 64	6.1	2.9	4.7	1.4	versicolor
## 65	5.6	2.9	3.6	1.3	versicolor
## 66	6.7	3.1	4.4	1.4	versicolor
## 67	5.6	3.0	4.5	1.5	versicolor
## 68	5.8	2.7	4.1	1.0	versicolor
## 69	6.2	2.2	4.5	1.5	versicolor
## 70	5.6	2.5	3.9	1.1	versicolor
## 71	5.9	3.2	4.8	1.8	versicolor
## 72	6.1	2.8	4.0	1.3	versicolor
## 73	6.3	2.5	4.9	1.5	versicolor
## 74	6.1	2.8	4.7	1.2	versicolor
## 75	6.4	2.9	4.3	1.3	versicolor
## 76	6.6	3.0	4.4	1.4	versicolor
## 77	6.8	2.8	4.8	1.4	versicolor
## 78	6.7	3.0	5.0	1.7	versicolor
## 79	6.0	2.9	4.5	1.5	versicolor
## 80	5.7	2.6	3.5	1.0	versicolor
## 81	5.5	2.4	3.8	1.1	versicolor
## 82	5.5	2.4	3.7	1.0	versicolor

```
## 83      5.8      2.7      3.9      1.2 versicolor
## 84      6.0      2.7      5.1      1.6 versicolor
## 85      5.4      3.0      4.5      1.5 versicolor
## 86      6.0      3.4      4.5      1.6 versicolor
## 87      6.7      3.1      4.7      1.5 versicolor
## 88      6.3      2.3      4.4      1.3 versicolor
## 89      5.6      3.0      4.1      1.3 versicolor
## 90      5.5      2.5      4.0      1.3 versicolor
## 91      5.5      2.6      4.4      1.2 versicolor
## 92      6.1      3.0      4.6      1.4 versicolor
## 93      5.8      2.6      4.0      1.2 versicolor
## 94      5.0      2.3      3.3      1.0 versicolor
## 95      5.6      2.7      4.2      1.3 versicolor
## 96      5.7      3.0      4.2      1.2 versicolor
## 97      5.7      2.9      4.2      1.3 versicolor
## 98      6.2      2.9      4.3      1.3 versicolor
## 99      5.1      2.5      3.0      1.1 versicolor
## 100     5.7      2.8      4.1      1.3 versicolor
```

If the only input x is Species and you are trying to predict y which is Petal.Length, what would a reasonable, naive prediction be under both Species? Hint: it's what we did in class.

```
mean_versicolor = iris[mean(iris$Species == "versicolor")]
mean_setosa = iris[mean(iris$Species == "setosa")]
y = mean(iris$Petal.Length)
```

```
#g = mean_versicolor + (mean_versicolor - mean_setosa)
```

Prove that this is the OLS model by fitting an appropriate `lm` and then using the `predict` function to verify you get the same answers as you wrote previously.

```
#TO-DO
```