

Q1

In the last homework assignment, the relationship between weight and height of 18 year old girls was examined. For this assignment, you will examine additional variables collected in the Berkeley Guidance Study. The data are posted in the file BGSgirls2.txt. There is one line of data for each of 70 girls with the variables appearing in the following order:

- ID: Girl identification number
- WT2: Weight (kg) at 2 years
- HT2: Height (cm) at 2 years
- WT9: Weight (kg) at 9 years
- HT9: Height (cm) at 9 years
- LG9: Leg circumference (cm) at 9 years
- ST9: Strength (kg) at 9 years
- WT18: Weight (kg) at 18 years
- HT18: Height (cm) at 18 years
- LG18: Leg circumference (cm) at 18 years
- ST18: Strength (kg) at 18 years
- BMI: Body Mass Index at 18 years
- SOMA: Somatotype (SOMA), on a scale from 1, very thin, to 7, very obese

The goal of this exercise is to determine how well the measurements at ages 2 and 9 can predict BMI at age 18.

Use R to complete the following exercises:

(a)

Compute the sample correlations between BMI at age 18 and each of the following explanatory variables HT2, HT9, WT2, WT9, and ST9. Which of these explanatory variables have significant correlations with BMI at age 18?

```
data <- read.table("BGSgirls2.txt", header = F)

colnames(data) <- c(
  "ID",      # Girl identification number
  "WT2",     # Weight (kg) at 2 years
  "HT2",     # Height (cm) at 2 years
  "WT9",     # Weight (kg) at 9 years
  "HT9",     # Height (cm) at 9 years
  "LG9",     # Leg circumference (cm) at 9 years
  "ST9",     # Strength (kg) at 9 years
  "WT18",    # Weight (kg) at 18 years
  "HT18",    # Height (cm) at 18 years
  "LG18",    # Leg circumference (cm) at 18 years
  "ST18",    # Strength (kg) at 18 years
```

```

"BMI",      # Body Mass Index at 18 years
"SOMA"      # Somatotype (SOMA), on a scale from 1, very thin, to 7, very obese
)

response_var <- "BMI"
explanatory_vars <- c("HT2", "HT9", "WT2", "WT9", "ST9")

correlations <- sapply(X = explanatory_vars,
                      FUN = function(var) {
                        cor(data[[var]], data[[response_var]])
                      })
correlations

```

```

##           HT2           HT9           WT2           WT9           ST9
## 0.042573733 0.236907969 0.190947873 0.545925753 0.005603061

```

```

correlation_results <- sapply(X = explanatory_vars,
                             FUN = function(var) {
                               test <- cor.test(data[[var]], data[[response_var]])
                               c("correlation" = round(test$estimate, 4), "p-value" = round(test$p.value, 4))
                             })

correlation_results_df <- as.data.frame(t(correlation_results))
correlation_results_df <- cbind(variable = explanatory_vars, correlation_results_df)
correlation_results_df

```

```

##      variable correlation.cor p-value
## HT2      HT2          0.0426 0.7264
## HT9      HT9          0.2369 0.0483
## WT2      WT2          0.1909 0.1133
## WT9      WT9          0.5459 0.0000
## ST9      ST9          0.0056 0.9633

```

Using `cor.test` we are able to extract both the correlation coefficient as well as its associated p-value for determining significance. For something to be “significant”, we evaluate it at the $\alpha = 0.05$ level, such that we are testing whether the explanatory variable in question has a correlation coefficient with the response variable (BMI) significantly different from 0 (null hypothesis, alternative being the correlation is not zero). That being said:

Significant Findings

The only significant explanatory variables are for WT9 and HT9 at the level specified above are:

WT9: Weight (kg) at 9 years - Correlation: 0.5459 - p-value: $1.018837e-6$ - Significant: The correlation is strong, greater in magnitude to 0.5, and the p-value is smaller than our reference threshold.

HT9: Height (cm) at 9 years - Correlation: 0.2369 - p-value: 0.0483 - Significant: The correlation is not especially strong, but the p-value is below the 0.05 threshold.

Not Significant Findings

The other explanatory variables considered had small (close to zero estimated sample) correlations, and as such are found not to meet the significance threshold.

ST9: - Correlation: 0.0056 - p -value: 0.9633 - Not Significant: The correlation is near zero and the p -value is nearly 1.

HT2: - Correlation: 0.0426 - p -value: 0.7264 - Not Significant: The correlation is near zero and the p -value is rather large.

WT2: - Correlation: 0.1909 - p -value: 0.1133 - Not Significant: While the correlation is not especially strong, and despite being a somewhat small p -value it is nonetheless larger than the comparison value of 0.05.

(b)

Compute the sample correlations among the five explanatory variables HT2, HT9, WT2, WT9, and ST9. Which of these explanatory variables have significant correlations with other explanatory variables?

```
explanatory_vars <- c("HT2", "HT9", "WT2", "WT9", "ST9")
explanatory_data <- data[, explanatory_vars]
cor_matrix <- cor(explanatory_data)
n <- nrow(explanatory_data)
p_values <- matrix(NA, nrow = length(explanatory_vars), ncol = length(explanatory_vars))
rownames(p_values) <- colnames(p_values) <- explanatory_vars

# For loops are allowed
# Yipee!
for (i in 1:length(explanatory_vars)) {
  for (j in 1:length(explanatory_vars)) {
    if (i != j) {
      test <- cor.test(explanatory_data[[i]], explanatory_data[[j]])
      p_values[i, j] <- test$p.value
    }
  }
}

significant_correlations <- which(p_values < 0.05, arr.ind = TRUE)
significant_results <- data.frame(
  Var1 = rownames(p_values)[significant_correlations[, 1]],
  Var2 = colnames(p_values)[significant_correlations[, 2]],
  "correlation" = round(cor_matrix[significant_correlations], 4),
  "p-Value" = round(p_values[significant_correlations], 4)
)

significant_results <- significant_results[!duplicated(t(apply(significant_results, 1, sort))), ]
significant_results
```

##	Var1	Var2	correlation	p.Value
## 1	HT9	HT2	0.7384	0.0000
## 2	WT2	HT2	0.6445	0.0000
## 3	WT9	HT2	0.5229	0.0000
## 4	ST9	HT2	0.3617	0.0021
## 6	WT2	HT9	0.6071	0.0000
## 7	WT9	HT9	0.7276	0.0000
## 8	ST9	HT9	0.6034	0.0000
## 11	WT9	WT2	0.6925	0.0000
## 12	ST9	WT2	0.4516	0.0001
## 16	ST9	WT9	0.4530	0.0001

HT9, WT9, WT2, and ST9 are strongly or moderately correlated with multiple other variables, where “strongly” or “moderately” is whether the magnitude of correlation ($|\text{Rho}|$) is greater than or equal to 0.35 (for moderate, 0.50 for strong). Pairwise comparisons are listed below for variables with correlations stronger in magnitude than 0.05, which is all of them. Additionally, all of the pairwise comparisons considered had significant p-values, or p-values whose values are below the $\alpha = 0.05$ threshold.

Please pardon the redundancies as $\text{Cor}(\text{HT9}, \text{HT2}) = \text{Cor}(\text{HT2}, \text{HT9})$:

HT2

- HT9 $r = 0.7384$, $p \approx 3.00e - 13$
- WT2 $r = 0.6445$, $p \approx 1.73e - 9$
- WT9 $r = 0.5229$, $p \approx 3.42e - 6$
- ST9 $r = 0.3617$, $p \approx 2.09e - 3$

HT9

- HT2 $r = 0.7384$, $p \approx 3.00e - 13$
- WT9 $r = 0.7276$, $p \approx 9.69e - 13$
- WT2 $r = 0.6071$, $p \approx 2.52e - 8$
- ST9 $r = 0.6034$, $p \approx 3.23e - 8$

WT2

- WT9 $r = 0.6925$, $p \approx 3.11e - 11$
- HT2 $r = 0.6445$, $p \approx 1.73e - 9$
- HT9 $r = 0.6071$, $p \approx 2.52e - 8$
- ST9 $r = 0.4516$, $p \approx 8.71e - 5$

WT9

- HT9 $r = 0.7276$, $p \approx 9.69e - 13$
- WT2 $r = 0.6925$, $p \approx 3.11e - 11$
- HT2 $r = 0.5229$, $p \approx 3.42e - 6$
- ST9 $r = 0.4530$, $p \approx 8.22e - 5$

ST9

- HT9 $r = 0.6034$, $p \approx 3.23e - 8$
- WT2 $r = 0.4516$, $p \approx 8.71e - 5$
- WT9 $r = 0.4530$, $p \approx 8.22e - 5$
- HT2 $r = 0.3617$, $p \approx 2.09e - 3$

(c)

Find least squares estimates of the parameters in the regression of BMI at age 18 on strength at age 9,

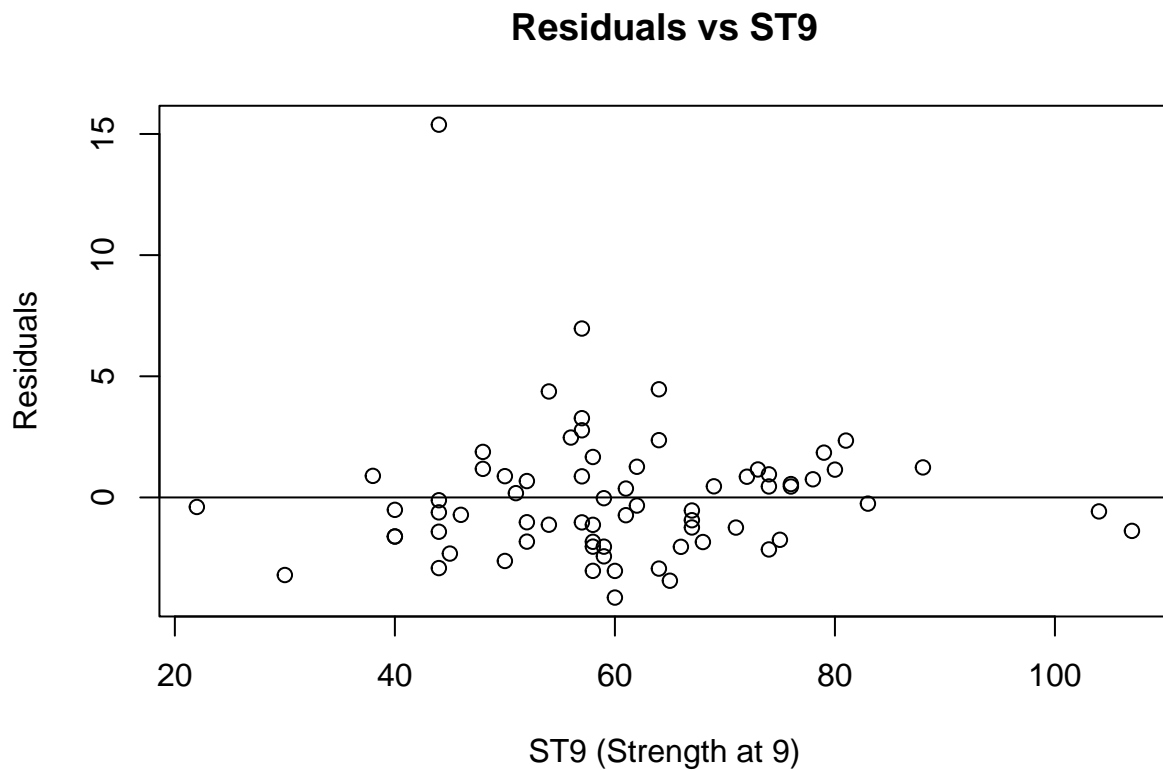
$$BMI_i = \beta_0 + \beta_1 ST9_i + \epsilon_i, \text{ for } i = 1, \dots, 70$$

Is the slope significantly different from zero? What do the residual plots reveal?

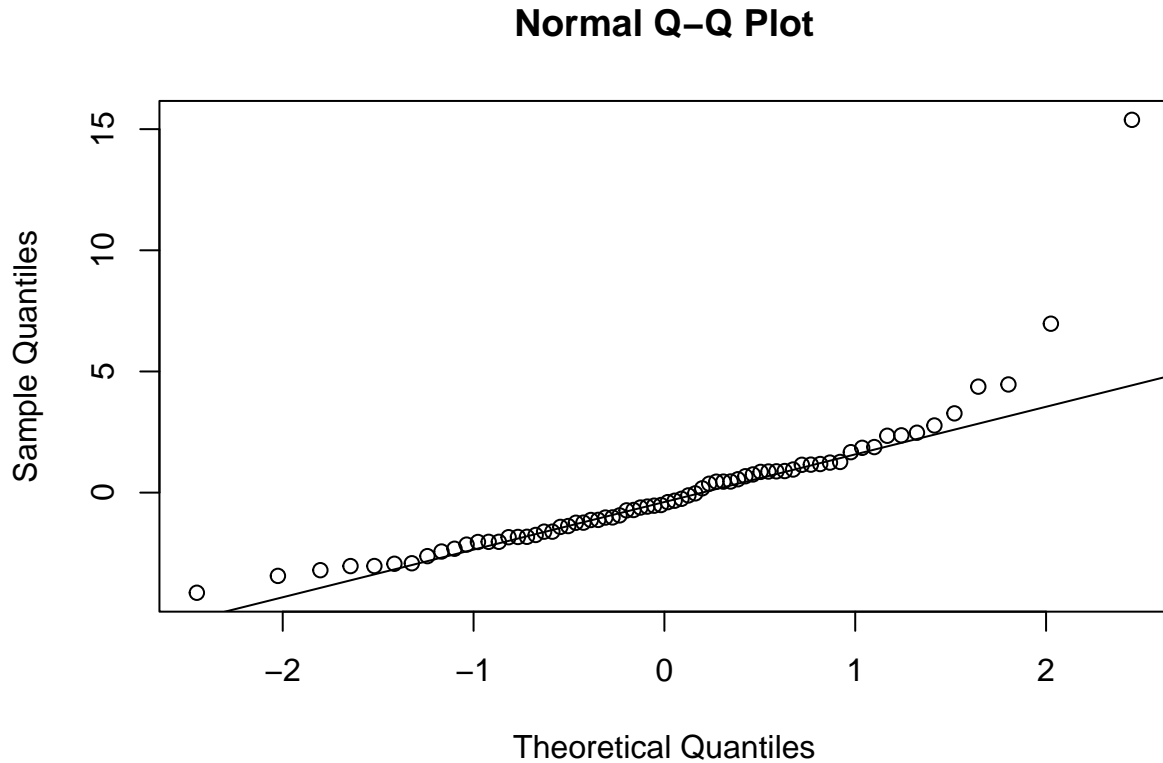
```
model <- lm(BMI ~ ST9, data = data)
summary(model)
```

```
##
## Call:
## lm(formula = BMI ~ ST9, data = data)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -4.1324 -1.7138 -0.4527  0.9375 15.3840
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 21.471009   1.379242  15.567  <2e-16 ***
## ST9          0.001023   0.022141   0.046    0.963
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.782 on 68 degrees of freedom
## Multiple R-squared:  3.139e-05, Adjusted R-squared:  -0.01467
## F-statistic: 0.002135 on 1 and 68 DF,  p-value: 0.9633
```

```
plot(data$ST9, resid(model), main = "Residuals vs ST9", xlab = "ST9 (Strength at 9)", ylab = "Residuals",
      abline(h = 0, col = "black"))
```



```
qqnorm(resid(model))
qqline(resid(model), col = "black")
```



Least Squares Formula:

$$\hat{Y}_i = b_0 + b_1 x_{(i, ST9)}$$

$$\hat{Y}_i = 21.471009 + 0.001023 x_{(i, ST9)}$$

So our estimated intercept of the least squares model, b_0 is 21.471009, and for b_1 we have 0.001023.

Significance of Slope:

- $H_0: \beta_1 = 0$
- $H_a: \beta_1 \neq 0$
- t -value for the slope: $t = 0.046$.
- p -value for the slope: $p = 0.963$.

For $p = 0.963$, we do not meet the threshold of $\alpha = 0.05$, such that we fail to reject the null hypothesis. This means we do not have evidence to support the slope of the linear regression to be significantly different from zero. (We do not have evidence to reject that strength at age 9 does not significantly predict BMI at age 18.)

Residual Plots:

Residuals Plot:

- The residuals are scattered randomly around 0, which indicates no clear pattern and suggests that the assumption of linearity is reasonable.
- The aforementioned point is conditional on the removal of outliers. There are a few points that could be outliers in the data, such that their removal would provide a residual plot that looks good, i.e. conforms to our expectations were our assumptions (of linearity and of equal variances) not being violated.

Normal Q-Q Plot:

- The residuals generally follow the reference line for the first few theoretical quantiles, though we do observe some deviations from the reference line at the tails of the distribution. Generally, it appears that our assumption of normality of errors is not being violated.
- Again, we do observe some points that could be considered outliers, such that their possible removal would help to further validate our assumptions are not being violated.

A Quick Note on Model Fit: $R^2 = 0.00003139$: This indicates that only 0.003% of the variation in BMI at age 18 is explained by strength at age 9. And Adj. $R^2 = -0.01467$ is negative. So this really is not an especially great model fit.

(d)

Now compute the multiple regression of the body mass index at age 18 on both weight at age 9 and strength at age 9, i.e. fit the model

$$BMI_i = \beta_0 + \beta_1 ST9_i + \beta_2 WT9_i + \epsilon_i, \text{ for } i = 1, \dots, 70$$

Is the estimate of β_1 for this model, the coefficient for strength at age 9, the same as the estimate of β_1 for the model in part (c)? Did you expect the estimates to be different? Explain. Is the effect of strength at age 9 significant in this model?

```
multi_model <- lm(BMI ~ ST9 + WT9, data = data)
summary(multi_model)
```

```
##
## Call:
## lm(formula = BMI ~ ST9 + WT9, data = data)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -4.1736 -1.2146 -0.2474  1.1231 11.2834
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 14.63878    1.54661   9.465 5.66e-14 ***
## ST9         -0.05552    0.01983  -2.799 0.00668 **
## WT9          0.32418    0.05151   6.293 2.72e-08 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
##
## Residual standard error: 2.222 on 67 degrees of freedom
## Multiple R-squared:  0.3715, Adjusted R-squared:  0.3528
## F-statistic: 19.8 on 2 and 67 DF,  p-value: 1.747e-07
```

Estimate of β_1 (Effect of Strength at Age 9):

- From the multiple regression model: $\hat{\beta}_1 = -0.05552$
- From the simple regression model in part (c): $\hat{\beta}_1 = 0.001023$

Comparison:

- Between the two models, the estimates for β_1 differ significantly, both in sign (positive to negative), to magnitude (from decimal place of hundredths to decimal place of thousandths).
- This is somewhat understandable though, as their interpretations between models differ: In the simple regression model, the estimate reflects the total (unadjusted) association between ST9 and BMI. By contrast, in the multiple regression model, the estimate reflects the unique contribution of ST9 to BMI, after adjusting for WT9. The negative coefficient in the multiple regression suggests that, after accounting for weight at age 9, greater strength is associated with lower BMI.
- I would argue that this change is not unexpected, but instead expected, because WT9 and ST9 are likely correlated given our prior correlations (the two variables ST9 and WT9 in our multiple regression are positively correlated with one another).

Is the Effect of Strength at Age 9 Significant (In the Multiple Regression Model)?

- The t -value (test statistic) for ST9: $t = -2.799$.
- The p -value for the estimated parameter ST9: $p = 0.00668$.

Since the above p -value is smaller than the $\alpha = 0.05$ level threshold, we have evidence to reject the null hypothesis. This means we have evidence that strength at age 9 (ST9) is statistically significant in the multiple regression model estimating BMI at age 18 that includes WT9 as an explanatory variable. This indicates that strength at age 9 has a significant relationship with BMI at age 18, after adjusting for weight at age 9 (WT9) (or for a constant/fixed value of WT9).

(e)

Fit the multiple regression model

$$BMI_i = \beta_0 + \beta_1 WT2_i + \beta_2 HT2_i + \beta_3 WT9_i + \beta_4 HT9_i + \beta_5 ST9_i + \epsilon_i, \text{ for } i = 1, \dots, 70$$

Report and interpret in context the R^2 value.

```
full_model <- lm(BMI ~ WT2 + HT2 + WT9 + HT9 + ST9, data = data)
summary(full_model)
```

```
##
## Call:
## lm(formula = BMI ~ WT2 + HT2 + WT9 + HT9 + ST9, data = data)
##
## Residuals:
```



```
##      Min      1Q  Median      3Q      Max
## -5.0948 -1.2186 -0.2533  1.0090 10.4951
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 30.855335   8.781156   3.514 0.000817 ***
## WT2         -0.317779   0.278736  -1.140 0.258505
## HT2         -0.193997   0.130819  -1.483 0.142996
## WT9          0.419762   0.075211   5.581 5.2e-07 ***
## HT9          0.008057   0.096344   0.084 0.933613
## ST9         -0.044416   0.022219  -1.999 0.049853 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.14 on 64 degrees of freedom
## Multiple R-squared:  0.4431, Adjusted R-squared:  0.3996
## F-statistic: 10.19 on 5 and 64 DF,  p-value: 3.294e-07
```

$R^2 = 0.4431$:

Interpretation: 44.31% of the variability in BMI at age 18 is explained by the multiple linear regression that uses weight and height at ages 2 and 9 (WT2, HT2, WT9, HT9), and strength at age 9 (ST9) as explanatory variables. Frankly, this isn't especially great, as we can explain less than half the variability in what we are attempting to estimate.

(f)

Report estimates of the six partial regression coefficients for the model in part (e), their standard errors, and the value of the corresponding t-tests and p-values (for two-sided alternatives to the null hypothesis). For each t-test, explicitly state the null hypothesis that is tested and interpret the result in context.

```
summary(full_model)
```

```
##
## Call:
## lm(formula = BMI ~ WT2 + HT2 + WT9 + HT9 + ST9, data = data)
##
## Residuals:
##      Min      1Q  Median      3Q      Max
## -5.0948 -1.2186 -0.2533  1.0090 10.4951
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 30.855335   8.781156   3.514 0.000817 ***
## WT2         -0.317779   0.278736  -1.140 0.258505
## HT2         -0.193997   0.130819  -1.483 0.142996
## WT9          0.419762   0.075211   5.581 5.2e-07 ***
## HT9          0.008057   0.096344   0.084 0.933613
## ST9         -0.044416   0.022219  -1.999 0.049853 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.14 on 64 degrees of freedom
```

Multiple R-squared: 0.4431, Adjusted R-squared: 0.3996
F-statistic: 10.19 on 5 and 64 DF, p-value: 3.294e-07

Intercept (β_0):

- H_0 : $\beta_0 = 0$ (BMI at age 18 is 0 when all predictors are 0).
- H_a : $\beta_0 \neq 0$.
- Estimate: 30.855335
- SE: 8.781156
- t-test: 3.514
- p-value: 0.000817
- Interpretation: The intercept is significant ($p = 0.0008$), indicating that the predicted baseline BMI (when all predictors are 0) is non-zero. However, the intercept may not be meaningful in this context due to the lack of real-world relevance for predictors being zero.

Weight at Age 2 (β_1):

- H_0 : $\beta_1 = 0$ (Weight at age 2 has no effect on BMI at age 18, after accounting for other predictors).
- H_a : $\beta_1 \neq 0$.
- Estimate: -0.317779
- SE: 0.278736
- t-test: -1.140
- p-value: 0.258505
- Interpretation: The coefficient for WT2 is not significant ($p = 0.2585$), suggesting that weight at age 2 does not significantly predict BMI at age 18 after adjusting for other variables.

Height at Age 2 (β_2):

- H_0 : $\beta_2 = 0$ (Height at age 2 has no effect on BMI at age 18, after accounting for other predictors).
- H_a : $\beta_2 \neq 0$.
- Estimate: -0.193997
- SE: 0.130819
- t-test: -1.483
- p-value: 0.142996
- Interpretation: The coefficient for HT2 is not significant ($p = 0.1430$), suggesting that height at age 2 does not significantly predict BMI at age 18 after adjusting for other variables.

Weight at Age 9 (β_3):

- H_0 : $\beta_3 = 0$ (Weight at age 9 has no effect on BMI at age 18, after accounting for other predictors).
- H_a : $\beta_3 \neq 0$.
- Estimate: 0.419762
- SE: 0.075211
- t-test: 5.581
- p-value: 5.2e-07
- Interpretation: The coefficient for WT9 is highly significant ($p = 5.2 \times 10^{-7}$), indicating that weight at age 9 is a strong predictor of BMI at age 18, after adjusting for other predictors.

Height at Age 9 (β_4):

- H_0 : $\beta_4 = 0$ (Height at age 9 has no effect on BMI at age 18, after accounting for other predictors).
- H_a : $\beta_4 \neq 0$.
- Estimate: 0.008057
- SE: 0.096344
- t-test: 0.084
- p-value: 0.933613
- Interpretation: The coefficient for HT9 is not significant ($p = 0.9336$), suggesting that height at age 9 does not significantly predict BMI at age 18 after adjusting for other variables.

Strength at Age 9 (β_5):

- H_0 : $\beta_5 = 0$ (Strength at age 9 has no effect on BMI at age 18, after accounting for other predictors).
- H_a : $\beta_5 \neq 0$.
- Estimate: -0.044416
- SE: 0.022219
- t-test: -1.999
- p-value: 0.049853
- Interpretation: The coefficient for ST9 is marginally significant ($p = 0.0499$), indicating that strength at age 9 has a small but significant effect on BMI at age 18, after adjusting for other predictors. The negative sign suggests that greater strength at age 9 is associated with a slightly lower BMI at age 18.

Q2

A dataset was collected from home sales in Ames, Iowa between 2006 and 2010. The variables collected are:

- Year Built: The year the house was built
- Basement Area (in sq. ft): The amount of area in the house below ground level
- Living Area (in sq. ft): The living area in the home (includes Basement Area)
- Total Room: The number of rooms in the house
- Garage Cars: The number of cars that can be placed in the garage
- Year Sold: The year the home was sold
- Sale Price: The sale price of the home (the response variable)
- Garage Size: S = Small (Garage Cars = 0,1) or L = Large (Garage Cars = 2+)
- Age (in yrs.): Age of house = Year Sold - Year Built

The data from 2,925 sales can be found in the file AmesHousing.csv posted in our course's shared folder on SAS Studio. For all parts requiring a hypothesis test, make sure to state the null and alternative hypotheses, test statistic, p-value, decision, and conclusion in context.

First, we will use SAS to explore predicting the Sale Price of a house from two explanatory variables: Living Area and Age. The SAS code that generated the output below is included in Canvas in the housing solution.sas file for your reference. Use the output shown on the next page to complete the exercises that follow.

(a)

The REG Procedure
Model: MODEL1
Dependent Variable: SalePrice

Number of Observations Read	2925
Number of Observations Used	2925

Analysis of Variance					
Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	2	1.283227E13	6.416137E12	3199.94	<.0001
Error	2922	5.85885E12	2005082271		
Corrected Total	2924	1.869112E13			

Root MSE	44778	R-Square	0.6865
Dependent Mean	180786	Adj R-Sq	0.6863
Coeff Var	24.76860		

Parameter Estimates					
Variable	DF	Parameter Estimate	Standard Error	t Value	Pr > t
Intercept	1	66464	3134.28250	21.21	<.0001
LivingArea	1	102.62349	1.74084	58.95	<.0001
Age	1	-1072.88363	28.21248	-38.03	<.0001

Figure 1: CocoMelon

Give a description of the parameters (β 's) for Living Area and Age in the multiple linear regression model.

Intercept (b_0): The intercept represents the expected Sale Price of a house when both Living Area (X_1) and Age (X_2) are 0. From the output, we estimate $b_0 = 66,464$. However, I cannot reasonably say that having Living Area and Age both set to input 0 makes sense in interpretation.

Living Area (b_1): The parameter $b_1 = 102.62349$ represents the expected change in Sale Price for each additional square foot of Living Area, holding Age constant. This suggests that, on average, a one-square-foot increase in Living Area is associated with an increase of approximately \$102.62 in Sale Price when holding Age constant.

Age (b_2): The parameter $b_2 = -1072.88363$ represents the expected change in Sale Price for each additional year of Age, holding Living Area constant. This indicates that, on average, each additional year in the house's age is associated with a decrease of approximately \$1,072.88 in Sale Price when holding Living Area constant.

(b)

What is the value of R^2 and its interpretation for the model including Living Area and Age?

An R^2 of 0.6865 means that approximately 68.65% of the variability in Sale Price is explained by the linear regression model that includes Living Area and Age as explanatory variables (while also including an intercept term).

(c)

Using the ANOVA table, conduct the F-test for the overall significance of the model. Report the null and alternative hypotheses, test statistic and p-value, and interpret the result in context.

$$H_0 : \beta_1 = \beta_2 = 0$$

$$H_a : \text{At least one } \beta_i \neq 0 \text{ (for } i = 1, 2\text{)}.$$

From the ANOVA table: - Model Sum of Squares (SSM): 1.283227e13 - Error Sum of Squares (SSE): 5.85885e12 - Total Sum of Squares (SST): 1.869112e13 - Degrees of Freedom for the Model: 2 - Degrees of Freedom for Error: 2922

The F-test statistic is given by:

$$F = \frac{\text{MSM}}{\text{MSE}}$$

$$\text{MSM} = \frac{\text{SSM}}{\text{DF}_{\text{Model}}}, \quad \text{MSE} = \frac{\text{SSE}}{\text{DF}_{\text{Error}}}$$

$$\text{MSM} = \frac{1.283227 \times 10^{13}}{2} = 6.416137 \times 10^{12} \quad \text{MSE} = \frac{5.85885 \times 10^{12}}{2922} = 2005082271$$

$$F = \frac{6.416137 \times 10^{12}}{2005082271} \approx 3199.94$$

From the output, the p-value is reported as < 0.0001 . This indicates overwhelming evidence in favor of rejecting the null hypothesis, which is evidence in favor of the alternative hypothesis (as stated above). Interpretation: The F-test interpretation stated previously is evidence that at least one of the predictors

(Living Area and Age) are significantly different from 0 in their estimated slopes, which is to say that at least one of the explanatory variables is able to improve the amount of variability we are able to explain in Sales Price by using a multiple linear regression.

(d)

Give the t-test for the significance of each explanatory variable in the model. Report the null and alternative hypotheses, test statistic and p-value, and interpret the result in context.

i.

Living Area

$$H_0 : \beta_1 = 0$$

$$H_a : \beta_1 \neq 0$$

Parameter Estimate ($\hat{\beta}_1$): 102.62349 Standard Error (SE): 1.74084

$$\text{Test Statistic} = t = \frac{\hat{\beta}_1}{\text{SE}} = \frac{102.62349}{1.74084} \approx 58.95$$

From the output, the p-value < 0.0001 for Living Area, indicating overwhelming evidence in favor of rejecting the null hypothesis. Interpretation: The t-test for Living Area indicates that it is a statistically significant predictor of Sale Price (we estimate the slope of the Living Area parameter to be not equal to zero). This is to say that, after controlling for Age, there is strong evidence that Living Area is associated with changes in Sale Price, specifically that for each additional square foot of Living Area, the Sale Price is expected to increase by approximately \$102.62.

ii.

Age

$$H_0 : \beta_2 = 0$$

$$H_a : \beta_2 \neq 0$$

Parameter Estimate ($\hat{\beta}_2$): -1072.88363 Standard Error (SE): 28.21248

$$\text{Test Statistic} = t = \frac{\hat{\beta}_2}{\text{SE}} = \frac{-1072.88363}{28.21248} \approx -38.03$$

From the output, the p-value < 0.0001 for Age, indicating overwhelming evidence in favor of rejecting the null hypothesis. Interpretation: The t-test for Age indicates that it is a statistically significant predictor of Sale Price (we estimate the slope of the Age parameter to be not equal to zero). This is to say that, after controlling for Living Area, there is strong evidence that Age is associated with changes in Sale Price, specifically that for each additional year of Age, the Sale Price is expected to decrease by approximately \$1,072.88.

(e)

In addition to Living Area and Age, add two additional explanatory variables Basement Area and Total Room into the multiple linear regression model. The SAS output is shown below.

The REG Procedure
Model: MODEL1
Dependent Variable: SalePrice

Number of Observations Read	2925
Number of Observations Used	2924
Number of Observations with Missing Values	1

Analysis of Variance					
Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	4	1.431999E13	3.579997E12	2396.37	<.0001
Error	2919	4.360771E12	1493926329		
Corrected Total	2923	1.868076E13			

Root MSE	38651	R-Square	0.7666
Dependent Mean	180821	Adj R-Sq	0.7662
Coeff Var	21.37550		

Parameter Estimates					
Variable	DF	Parameter Estimate	Standard Error	t Value	Pr > t
Intercept	1	32208	3719.97876	8.66	<.0001
LivingArea	1	100.79550	2.65342	37.99	<.0001
Age	1	-766.35988	26.21189	-29.24	<.0001
BasementArea	1	59.90295	1.99057	30.09	<.0001
TotalRoom	1	-5741.69710	790.86924	-7.26	<.0001

Figure 2: CocoMelon

i.

How much is the reduction to the Sums of Squares for Error for adding Basement Area and Total Room to the model with Living Area and Age?

SSE for the initial model (Living Area and Age): $5.85885e12$ SSE for the updated model (Living Area, Age, Basement Area, Total Room): $4.36077e12$

$$\text{Reduction in SSE} = \text{SSE (initial model)} - \text{SSE (updated model)} = 5.85885 \times 10^{12} - 4.36077 \times 10^{12} = 1.49808 \times 10^{12}$$

By adding Basement Area and Total Room to the model, the Sum of Squares for Error was reduced by $1.49808e12$.

ii.

Provide the partial F-test for the significance of Basement Area and Total Room in the model with Living Area and Age. Report the null and alternative hypotheses, test statistic and p-value, and interpret the result in context.

$$H_0 : \beta_{\text{BasementArea}} = \beta_{\text{TotalRoom}} = 0$$

$$H_a : \text{At least one of } \beta_{\text{BasementArea}} \text{ or } \beta_{\text{TotalRoom}} \neq 0$$

Our “m” is 2 as we add 2 additional explanatory variables. Reduction in SSE: $1.49808e12$ (calculated in part i) Mean Squared Error (MSE) of the “full” model: 1493926329

$$F = \frac{\frac{SSE_{\text{reduction}}}{m}}{MSE_{\text{full}}} = \frac{\left(\frac{1.49808 \times 10^{12}}{2} \right)}{1493926329} \approx 501.3902$$

Using the output and the extremely large F-value of 501.3902, the p-value is reported as < 0.0001 . This indicates overwhelming evidence in favor of rejecting the null hypothesis. Interpretation: The partial F-test shows that adding Basement Area and Total Room significantly improves the model for predicting Sale Price, beyond what is explained by Living Area and Age alone.