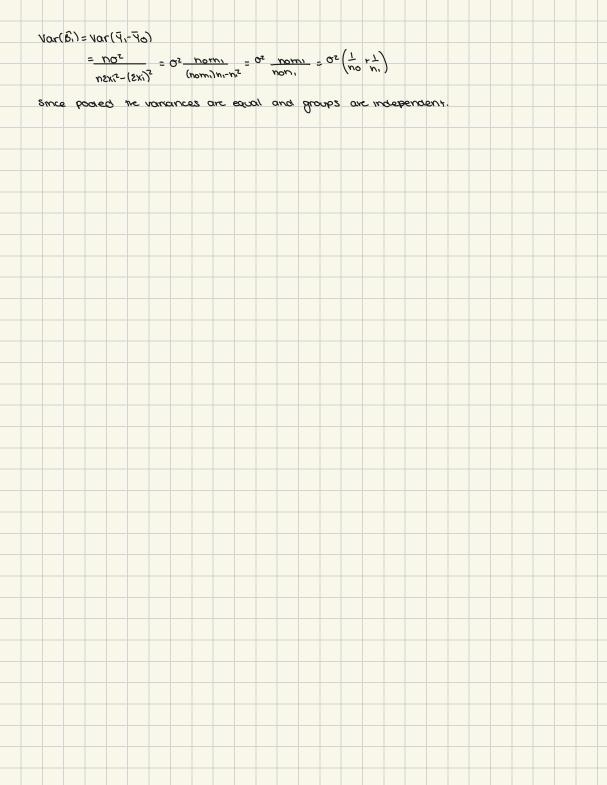
```
Saran Cooper
Week 11 Question 4
 4. ANOVA and Lack of Fit
                                                                   -equal vomance ANOVA
    a) Show that the pooled two-sample t test is a special case of simple regression on an indicator
      variable. For example, test whether goalkeepers and other players have the same mean height
      or weight. Show that the regression output for the t-test of \beta_1 = 0 exactly matches the pooled
      2-sample t test.
a) Show that pooled 2 sample t test is a special case of simple regression
   Group 1(G1) = goalies
    Group 2 (62) = field players
   Test: Are mean height and veight the same across two groups
                                                                           (equal variance)
    5M=1M2
                  Ha: MI = MZ
 Poded Z-sample Test
      t= x1-x2
                        XI and Xz are sample group means
                                                                   Sp2 = pooled sample variance
                       n, and nz are sample sizes
                                  df=n1+n2-2
                                Si and si are sample variances
   Simple Linear Regression
                                                            Y
     Y= response variable (height or veight)
     X=O goalles
                            $ Indicator
     X=1 for Reld players
   YL=BO+BIXL+EL
                                                                                          X
    Yi = neight or velght of player i
    XI = Indicator variable
    Bo = intercept, mean height (or reignt) for goalies (x=0)
                                                                                             Vo at x=0
                                                                                              and 7, as
    B1 = 810pe, difference in near height (or veight) blue two groups = Uz-Wi
                                                                                                X=1
    Fi= even ferm
    HO: BI=0 same as uz=u, Ha: BI=0
     YO = BOY EL -> BO = E(YO) = MO
                                           MLE: BO=YO
                                                                     * E(y,-w) = E(Y;- Y)
     Y1= BO+ B1+Ei -> B1= E(Y1)-E(Y0)= M-MO ME: B. = Y1-Y0
     Thus Bi= 71-90
   Comparing t-stats:
                                                                       *both denominators are the
                                                                         SE of difference between
                                                                         means
        Bi = corresponds directly to difference in sample means
        The t-stats are mathematically equivalent since born estimating for difference blum groups
```



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Part A)
soccer player data
<-read.table("C:\\Users\\scoop\\OneDrive\\SWAT\\soccerplayer data.txt", header</pre>
= TRUE, sep = ",")
>
> # Check the first few rows and structure of the data
> head(soccer player data)
 Division Pos GK Weight Height
        1 F N 158
        1 M N
                             71
2
                    145
        1 M N
3
                    150
                            67
        1 D N
                    147
4
                            68
5
        1
            F N
                    160
                            68
           M N
6
        1
                    150
                            68
> str(soccer_player data)
'data.frame': 1040 obs. of 5 variables:
 $ Division: int 1 1 1 1 1 1 1 1 1 ...
 $ Pos : chr "F" "M" "M" "D" ...
          : chr "N" "N" "N" "N" ...
 $ Weight : int 158 145 150 147 160 150 150 160 175 180 ...
 $ Height : int 71 71 67 68 68 68 69 73 73 73 ...
> soccer player data$Pos <- as.factor(soccer player data$Pos)</pre>
> soccer player data$GK <- as.factor(soccer player data$GK)</pre>
> # Convert GK to a binary variable (1 if "Y", 0 if "N")
> soccer player data$GK_binary <- as.integer(soccer_player_data$GK == "Y")</pre>
> # Perform a pooled 2-sample t-test for height
> t test height <- t.test(Height ~ GK, data = soccer player data, var.equal =
TRUE)
> print(t test height)
      Two Sample t-test
data: Height by GK
t = -9.382, df = 1038, p-value < 2.2e-16
alternative hypothesis: true difference in means between group N and group Y
is not equal to 0
95 percent confidence interval:
-2.488077 -1.627333
sample estimates:
mean in group N mean in group Y
       70.71781
                      72.77551
> # Perform a pooled 2-sample t-test for weight
> t test weight <- t.test(Weight ~ GK, data = soccer player data, var.equal =
TRUE)
> print(t test weight)
      Two Sample t-test
data: Weight by GK
```

```
t = -9.9516, df = 1038, p-value < 2.2e-16
alternative hypothesis: true difference in means between group N and group Y
is not equal to 0
95 percent confidence interval:
-15.65731 -10.49967
sample estimates:
mean in group N mean in group Y
      164.3841
                  177.4626
> # Simple regression for height
> lm height <- lm(Height ~ GK, data = soccer player data)</pre>
> summary(lm height)
Call.
lm(formula = Height ~ GK, data = soccer player data)
Residuals:
           1Q Median
                            3Q
   Min
                                  Max
-7.7178 -1.7178 0.2822 1.2822 7.2822
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 70.71781 0.08246 857.625 <2e-16 ***
GKY
           2.05771
                      0.21933 9.382 <2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1
Residual standard error: 2.464 on 1038 degrees of freedom
Multiple R-squared: 0.07817, Adjusted R-squared: 0.07728
F-statistic: 88.02 on 1 and 1038 DF, p-value: < 2.2e-16
> # Simple regression for weight
> lm weight <- lm(Weight ~ GK, data = soccer player data)
> summary(lm weight)
Call:
lm(formula = Weight ~ GK, data = soccer player data)
Residuals:
           1Q Median
                         30
   Min
-37.463 -9.384 0.616 10.616 51.616
Coefficients:
          Estimate Std. Error t value Pr(>|t|)
(Intercept) 164.3841 0.4941 332.699 <2e-16 ***
GKY
           13.0785
                       1.3142 9.952 <2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1
Residual standard error: 14.77 on 1038 degrees of freedom
Multiple R-squared: 0.0871, Adjusted R-squared: 0.08622
F-statistic: 99.03 on 1 and 1038 DF, p-value: < 2.2e-16
```

Recognize that for both Height and Weight the p-values reflect that we should reject the null hypothesis. This suggests that the mean height and weight of goalkeepers and field players are not the same.

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Part B)

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# Convert 'Pos' to a factor and set 'GK' as the baseline
(reference level)
> soccer player data$Pos <-</pre>
factor(soccer player data$Pos, levels = c('GK', 'F',
'M', 'D'))  # 'GK' as baseline
> # Perform ANOVA for height based on position
> anova height <- aov(Height ~ Pos, data =
soccer player data)
> summary(anova height)
              Df Sum Sq Mean Sq F value Pr(>F)
Pos
                   819 273.14 47.03 <2e-16 ***
Residuals
          1036 6017
                          5.81
Signif. codes:
0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> # Perform ANOVA for weight based on position
> anova weight <- aov(Weight ~ Pos, data =
soccer player data)
> summary(anova weight)
              Df Sum Sq Mean Sq F value Pr(>F)
               3 34603 11534 56.03 <2e-16 ***
Pos
          1036 213277
                            206
Residuals
Signif. codes:
0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

>

c) Explain how a simple linear regression model becomes an ANOVA model if you treat distinct x values as categories, rather than as a continuous predictor. Note how you need repeated data values with the same x values in order to fit such a model. Describe the Lack of Fit test for regression and how this saturated model is used to find an error estimate that does not depend on the linearity assumption. Demonstrate the lack of fit test for several data sets. Note how failing to reject does not imply linearity, or any other regression assumptions. C) Simple Linear Regression vs ANOVA Ti= average y value for SLR: relationary blum continuous dep. variable y and cont. indep. X. X) 7=1...K yi=Bot BixitEi, Ein N 6,02) - Normal Ineax model SSpure error = = (Yij - Vi)2 ANOVA: X as calegorical, testing differences between means of groups yi= g(xi) +Ei (g(xi) may be lirear) - you need repealed data points for each value of X. Each callegory x should have multiple ys wout it you cannot estimate variability within each group Lack of fix Test! - used to determine whether a chosen model is incidenciate (company's errors of models) Salurated Model: one parameter for each data point. Has enough parameters to 19th the data perfectly, with no residual error. Then error estimate is due to "random noise". Perfect fit. compares 1855 from chosen model to 1855 from saturated. The F-stat determines whether the complexity of saturated model is better fit that chosen model has lack of fit, more complex is reeded Ho: enosen model works SS pure error = \(\frac{\xi}{2\in (\xi)-\sqrt{1}} \) dt=n-k & variation within groups at some x 41= 9(xi) +Ei, Eit N(0.02) 88 Lock of #4 = \$ nj(7;-9;) } difference in mean model and group means

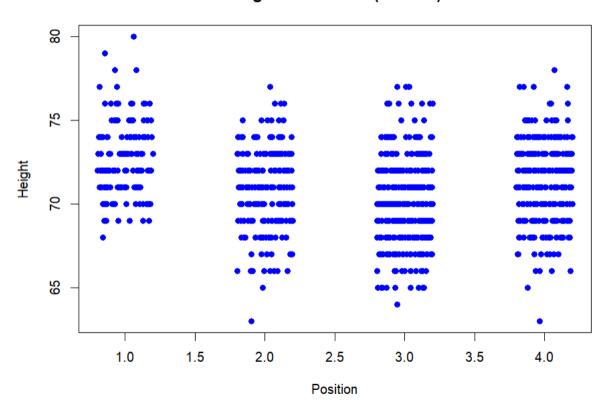
SS mear made = = (xij - Pj)2 = 55 pure error + S5 lack of Fit

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Part C)

```
# 1. Fit a linear regression model (using Pos as the predictor for Height)
> lm model <- lm(Weight ~ Height, data = soccer player data)</pre>
> print(anova(lm model))
Analysis of Variance Table
Response: Weight
            Df Sum Sq Mean Sq F value Pr(>F)
            1 133284 133284 1207.3 < 2.2e-16 ***
Residuals 1038 114596
                      110
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> # 2. Fit a saturated model (each observation gets its own level in the
> saturated model <- lm(Weight ~ factor(Height), data = soccer player data)
> print(anova(saturated model))
Analysis of Variance Table
Response: Weight
                Df Sum Sq Mean Sq F value Pr(>F)
factor(Height) 17 135080 7945.9 71.993 < 2.2e-16 ***
Residuals 1022 112799 110.4
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
> # 3. Perform an ANOVA to compare the two models
> anova result <- anova(lm model, saturated model)</pre>
> # Print the ANOVA result to check for lack of fit
> print(anova result)
Analysis of Variance Table
Model 1: Weight ~ Height
Model 2: Weight ~ factor(Height)
 Res.Df RSS Df Sum of Sq F Pr(>F)
1 1038 114596
2 1022 112799 16 1796.9 1.0175 0.4348
```

Height vs Position (Jittered)



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