Project #1

Singh, Gurjeet

This report contains analysis done while building regression models for the professional baseball team to predict the number of wins for the team.

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Introduction:

The purpose of this project is to build OLS ("Linear") Regression models using given statistics of the professional baseball team to predict the number of wins for each team.

For our purposes, we use the data set that contains performance information of the team for the given year. Each record in the data represents a professional baseball team from the years 1871 to 2006 inclusive. The performance statistics have been adjusted to match the performance of a 162 games season. The training dataset contains 2276 observations and 17 explanatory variables. The test dataset contains 259 observations and 16 explanatory variables. In the test dataset, there is no "Target Wins" information. We will be using our selected model to score the test data file to predict the number of wins.

Figure 1 gives the basic descriptions of each field and how those affect the winning chances of a team.

Figure 1: Statistical Values

VARIABLE NAME	DEFINITION	THEORETICAL EFFECT
INDEX	Identification Variable	None
TARGET_WINS		
TEAM_BATTING_H	Base Hits by batters (1B,2B,3B,HR)	Positive Impact on Wins
TEAM_BATTING_2B	Doubles by batters (2B)	Positive Impact on Wins
TEAM_BATTING_3B	Triples by batters (3B)	Positive Impact on Wins
TEAM_BATTING_HR	Homeruns by batters (4B)	Positive Impact on Wins
TEAM_BATTING_BB	Walks by batters	Positive Impact on Wins
TEAM_BATTING_HBP	Batters hit by pitch (get a free base)	Positive Impact on Wins
TEAM_BATTING_SO	Strikeouts by batters	Negative Impact on Wins
TEAM_BASERUN_SB	Stolen bases	Positive Impact on Wins
TEAM_BASERUN_CS	Caught stealing	Negative Impact on Wins
TEAM_FIELDING_E	Errors	Negative Impact on Wins
TEAM_FIELDING_DP	Double Plays	Positive Impact on Wins
TEAM_PITCHING_BB	Walks allowed	Negative Impact on Wins
TEAM_PITCHING_H	Hits allowed	Negative Impact on Wins
TEAM_PITCHING_HR	Homeruns allowed	Negative Impact on Wins
TEAM_PITCHING_SO	Strikeouts by pitchers	Positive Impact on Wins

Now that we have some context for our analysis and dataset, let's look at the results in the next section.

Section 1: Data Exploration

The first step towards any modeling project is the Data Exploration i.e. Exploratory Data Analysis (EDA). This helps us to understand and analyze the data set to summarize the main characteristics of variables. For this purposes, we will look into the basic statistics to understand the data, examine the distribution, and examine relationships.

Section 1.1: Statistics

Figure 2 shows the statistical values of the Moneyball training data set. The number of observations in the data is 2276 records. We can clearly see that there are missing values (NAs) in the data for variables, TEAM_BATTING_SO, TEAM_BASERUN_SB, TEAM_BASERUN_CS, TEAM_BATTING_HBP, TEAM_PITCHING_H, TEAM_PITCHING_SO, and TEAM_FIELDING_DP. For TEAM_BATTING_HBP, we can see that more than ninety percent (90%) of the values are missing. Hence, we will not be using this variable in any of our models. In Figure 2, we can also see outliers in various variables in the data. For instance, variables like TEAM_PITCHING_H and TEAM_PITCHING_SO have some extreme outliers. TEAM_PITCHING_H has a maximum value of 30,132. But the median value is only 1518. Therefore, we will need to fix the missing values and outliers prior to building our models.

We explained how we fixed the missing values and outliers in Section 2: Data Preparation.

Variable Names Minimum Maximum 1. Quartile Mean Median Skewness **Kurtosis** INDEX 2276 2535 630.75 1915.5 1268.46353 1270.5 736.34904 0.004215 -1.216756 TARGET WINS 2276 80.79086 15.75215 1.027476 146 -0.3987231537.25 TEAM BATTING H 2276 0 891 2554 1383 1469.26977 1454 144.59119 1.571333 7.278526 TEAM_BATTING_2B 0 458 46.80141 2276 69 208 241.24692 238 0.215102 0.006161 273 TEAM_BATTING_3B 2276 ol 0 223 34 72 55.25 47 27.93856 1.109465 1.503242 TEAM_BATTING_HR 0 42 147 99.61204 102 60.54687 0.186042 -0.963119 2276 264 TEAM_BATTING_BB 2276 0 0 878 451 580 501.55888 512 122.67086 -1.02576 2.182854 TEAM BATTING SO 2276 102 0 1399 548 930 735.60534 750 248.52642 -0.2978 -0.320799 TEAM BASERUN SB 2276 131 0 697 66 156 124.76177 101 87.79117 1.972414 5.489675 TEAM_BASERUN_CS 2276 772 38 52.80386 22.95634 7.620382 201 62 49 1.976218 TEAM_BATTING_HBP 2276 2085 29 95 50.5 67 59.35602 58 12.96712 0.318575 -0.111983 1682.5 TEAM_PITCHING_H 1137 30132 1419 1779.21046 1406.84293 10.329511 141.839699 2276 1518 TEAM_PITCHING_HR 2276 0 0 343 50 105.69859 107 61.29875 0.287788 -0.604631 TEAM PITCHING BB 476 96.96764 2276 0 3645 553.00791 536.5 166.35736 6.743899 611 TEAM PITCHING SO 2276 102 0 19278 615 968 817.73045 813.5 553.08503 22.174554 671.189129 TEAM_FIELDING_E 227.77097 2276 65 1898 127 249.25 246.48067 159 2.990466 10.970272 TEAM_FIELDING_DP 2276 52 228 131 146.38794 149 26.22639 -0.388939 0.18174

Figure 2: Statistical Values

Section 1.2: Examining Distributions

Figure 3 shows the histogram for each variable to help us understand the distribution of the variables. We can see that there are extreme outliers in the variables, TEAM_PITCHING_H, TEAM_PITCHING_BB, TEAM_PITCHING_SO, and TEAM_FIELDING_E. Variables like TEAM_BATTING_3B, TEAM_BASERUN_CS, and TEAM_FIELDING_E seems to be skewed right. Variables like TEAM_BATTING_BB and TEAM_FIELDING_DP seems to be skewed left. Both, right and left skewed are telling us that there are some outliers in the variables which need to be fixed. Variables like TEAM_BATTING_HR, TEAM_BATTING_SO, and TEAM_PITCHING_HR seem to bimodal shape.

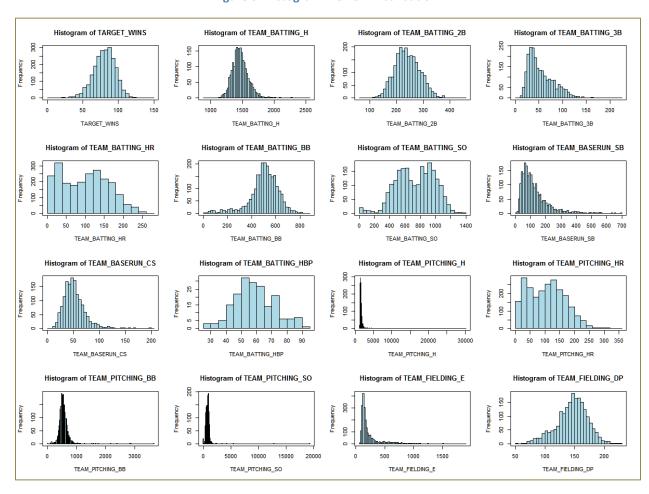


Figure 3: Histogram: Review Distribution

Figure 4 shows the QQ-plots to help us further review the distribution. These variables are selected based on some baseball knowledge I possess. Figure 4 confirms the present of outliers and the evidence of non-normality in the data.

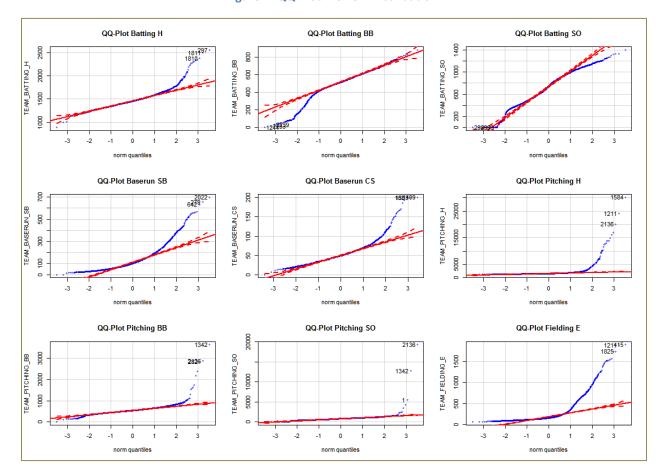


Figure 4: QQ-Plot: Review Distribution

Section 1.3: Examine Relationships

Figure 4 shows the relationship between TARGET_WINS and other variables. Based on the result, it does not appear that there's a linear positive or negative relationship. These could be due to outliers and missing values. Once we clean the data with missing values and outliers, we will do the sanity check and see if there's any difference in the relationship with the transformed data. Checking the correlation would help us determine if there's any relationship.

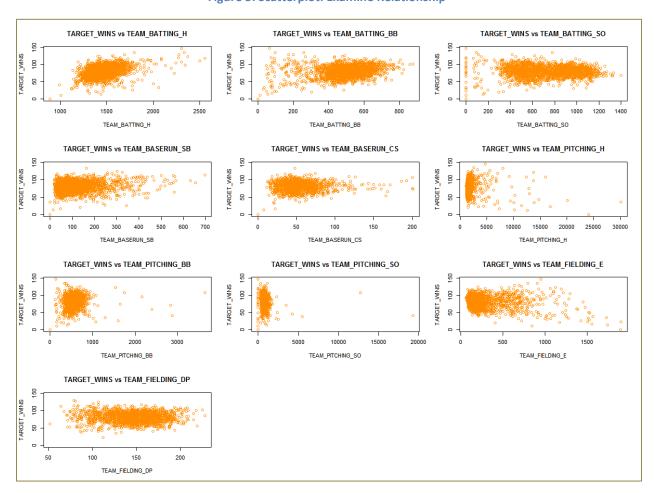


Figure 5: Scatterplot: Examine Relationship

Figure 6 shows the correlation matrix after fixing the missing values. As we saw in Figure 4, Figure 5 also confirms that there was no strong positive or negative linear relationship between the response variable i.e. TARGET_WINS and other predictor variables. In Figure 6, we can also see the relationship of predictor variables with other predictor variables. Here we notice that TEAM_FIELDING_E is negatively related to TEAM_BATTING_BB and IMP_TEAM_BATTING_SO. We will use this information when building models so we can test for multicollinearity.

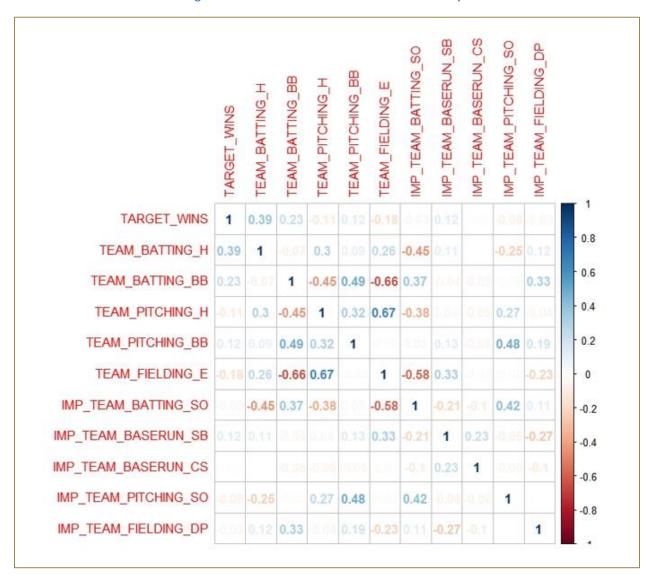


Figure 6: Correlation Matrix: Examine Relationship

Section 2: Data Preparation

In Section 1: Data Exploration, we noticed that we had some missing values and outliers in the data. This section explains the methodology we took to fix the missing values and transform data to eliminate outliers. First, we fixed the missing values by using the Median values for some variables. For variables, IMP_TEAM_BATTING_SO: we added two (2) additional Strike Out by Batters than the Median value; and IMP_TEAM_PITCHING_SO: we added one (1) additional Strike Out by Pitchers than the Median value. These changes will not impact a lot since the values are pretty close to the Median value. There are five (5) flag variables (indicator/dummy) for variables that had a missing value. These flag variables contain values zeros (0) and ones (1). Zero (0) for original missing values and one (1) for new values. At the bottom of Figure 6, we can see the new values used for the missing values (highlighted in green).

Next, we cleaned the data to fix extreme outliers. For this, we started with our approach to caps the lower and higher values with 1st percentile and 99th percentile respectively. However, after the detailed review of the source data, we used the values that made sense for each variable. For instance, 99th percentile for the variable, IMP_TEAM_PITCHING_H, is 7054 whereas we used the max value of 2200. Using 7000 as the max value did not help much with the distribution. We noticed a histogram plot to a right-skewed. Hence, after going back and forth for a while, we cap it at 2200. Likewise, the values for other variables were entered in a similar approach. Figure 7 also shows the new min and max values (highlighted in green) for each variable to fix the outliers.

Figure 7: Transforming Data: Fix Missing values and Outliers

Fix Outliers								
Predictor Variable	dictor Variable Old MIN Value Old MAX Value		1-Percentile	99-Percentile	New MIN Value	New MAX Value		
IMP_TEAM_BATTING_H	891	2554	1193.25	1945.5	1100	2000		
IMP_TEAM_BATTING_BB	0	878	79	752.75	150	800		
IMP_TEAM_BATTING_SO	0	1399	72	1191.25	72	1350		
IMP_TEAM_BASERUN_SB	0	697	24	434.25	14	500		
IMP_TEAM_BASERUN_CS	0	201	18.75	123.5	10	123		
IMP_TEAM_PITCHING_H	1137	30132	1244	7054	1200	2200		
IMP_TEAM_PITCHING_HR	0	343	8	244	8	260		
IMP_TEAM_PITCHING_BB	0	3645	240	921	119	1000		
IMP_TEAM_PITCHING_SO	0	19278	241	1461.75	241	1700		
IMP_TEAM_FIELDING_E	65	1898	86	1228	65	500		
IMP_TEAM_FIELDING_DP	52	228	80	202	71	220		

Fix Missing Values					
Predictor Variable	New Value				
IMP_TEAM_BATTING_SO	752				
IMP_TEAM_BASERUN_SB	101				
IMP_TEAM_BASERUN_CS	49				
IMP_TEAM_PITCHING_SO	814				
IMP TEAM FIELDING DP	149				

Section 3: Model Building

For the model building, I took the automated variable selections approach as well manually selected the variables to include in the model. After trying different combinations (of course, combinations that made sense), we selected three models that gave us decent metrics and also logically made sense.

This section explains the three (3) models (Model 1, Model 2, and Model 3) that we have created to predict the number of wins for each team. Out of these models, we have selected one (1) that fits the best and also logically correct to use for predicting the number of wins using test data.

Section 3.1: Model 1

Figure 8 shows the output of the Model 1. Residuals in Figure 8 are essentially the difference between the actual observed response values and the response values that the model predicted. The Residuals section of the model help us assess how well the model fits the data, we should look for a symmetrical distribution across these points on the mean value zero (0). In our case, we can see that the distribution of the residuals appears to be symmetrical since the Median value is very close to zero (0).

The coefficients of each parameter are behaving correctly (as per the theoretical effects in Data Dictionary). For instance, while holding the other predictors constant, if there is an increase of one (1) base hit by the batter (IMP_TEAM_BATTING_H), the winning chances increase by 0.0583 which is roughly 5.83% (approx.). Similarly, if a pitcher allowed one (1) additional hit (IMP_TEAM_PITCHING_H), the winning chances decrease by 0.007589 which is roughly 0.76% (approx.).

*** Significance stars in coefficients in Figure 8 shows that the predictor variables are statistically significant and it's unlikely that no relationship exists between the TARGET WINS and predictor variables.

Figure 8: Model 1: Output

```
Call:
lm(formula = TARGET_WINS ~ IMP_TEAM_BATTING_H + IMP_TEAM_BATTING_BB +
    IMP_TEAM_BASERUN_SB + M_TEAM_BASERUN_SB + IMP_TEAM_FIELDING_E +
   IMP_TEAM_PITCHING_BB + IMP_TEAM_PITCHING_H, data = MoneyBall)
Residuals:
            1Q Median
   Min
                            3Q
                                   Max
-55.473 -8.584 -0.326
                        8.337 52.355
Coefficients:
                     Estimate Std. Error t value Pr(>|t|)
(Intercept)
                    -5.833726
                                3.407116 -1.712
                                                 0.0870
                                0.003275 17.796 < 2e-16 ***
IMP_TEAM_BATTING_H
                     0.058274
                                          6.068 1.51e-09 ***
IMP_TEAM_BATTING_BB
                    0.034757
                                0.005728
                                0.004675 15.915 < 2e-16 ***
IMP_TEAM_BASERUN_SB
                    0.074402
                    28.650995
                                1.933428 14.819 < 2e-16 ***
M_TEAM_BASERUN_SB
IMP_TEAM_FIELDING E -0.055349
                                0.004245 -13.038
                                                 < 2e-16 ***
IMP_TEAM_PITCHING_BB -0.005889
                                0.004937
                                         -1.193
                                                  0.2331
                                                  0.0073 **
IMP_TEAM_PITCHING_H -0.007589
                                0.002826 -2.685
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 13.04 on 2268 degrees of freedom
Multiple R-squared: 0.3172,
                              Adjusted R-squared: 0.3151
F-statistic: 150.5 on 7 and 2268 DF, p-value: < 2.2e-16
```

In Figure 8, we also look at the Adjusted R-Squared statistic to measure how well our model is fitting the actual data. The R-Squared is a measure of how close the data are to the fitted regression line. In general, the higher the R-Squared, the better the model fits the data. However, when dealing to predict human behavior, it is expected that the R-Squared value will be low. The Adjusted R-Squared we got is **0.3151** which is roughly 31% (approx.) of the variance found in the response variable (TARGET_WINS) can be explained by the predictor variables. We notice that even though our Adjusted R-Squared value is below 50%, the predictor variables are statistically significant. Hence, this model was selected as one of the candidates for final selection.

Figure 9 shows the distribution using Q-Q plot and histogram of residuals and we see some evidence of non-normality towards the upper and lower end. The histogram shows that the residuals do not form a bell-shaped which usually present a normally distributed. In my opinion, this distribution is a slightly bimodal shape.

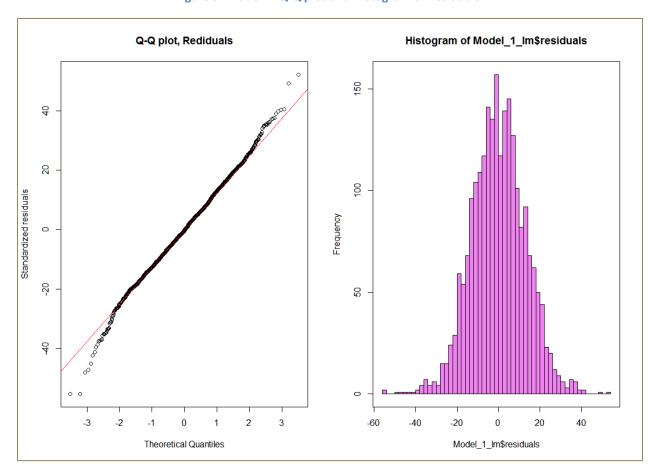


Figure 9: Model 1: Q-Q plot and Histogram of Residuals

Figure 10 shows the structure of residuals against each predictor variables using scatterplot. We notice that there's no structure in these plots. Hence, we validate the homoscedasticity assumption meaning that the variance of the error term (Residuals) is constant for all combination of independent (predictor) variables.

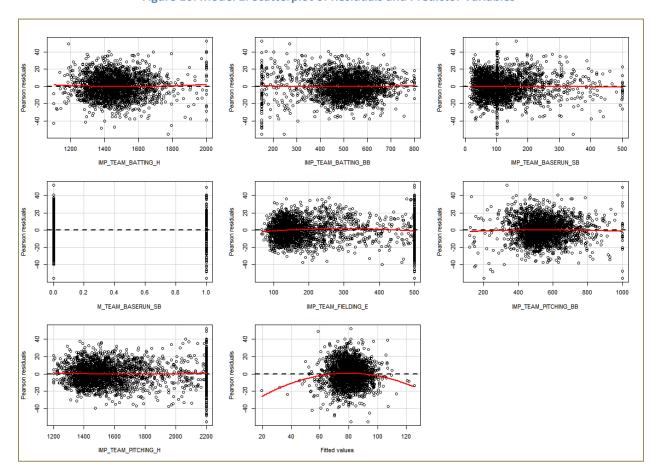


Figure 10: Model 1: Scatterplot of Residuals and Predictor Variables

Section 3.2: Model 2

Figure 11 shows the output of the Model 2. Residuals in Figure 11 appears to be symmetrical since the Median value is very close to zero (0). We would further investigate this by plotting the residuals in Q-Q plot and histogram to see whether this is normally distributed.

In Model 2, the coefficients of each parameter are also behaving correctly (as per the theoretical effects in Data Dictionary). For instance, while holding the other predictors constant, if there is an increase of one (1) base hit by the batter (IMP_TEAM_BATTING_H), the winning chances increase by 0.0584 which is roughly 5.8% (approx.). Similarly, if a pitcher allowed one (1) additional hit (IMP_TEAM_PITCHING_H), the winning chances decrease by 0.007517 which is roughly 0.75% (approx.). Lastly, if the base is stolen (IMP_TEAM_BASERUN_SB), the winning chances increase by 0.0743 which is roughly 7.43% (approx.).

*** Significance stars in coefficients in Figure 11 shows that the predictor variables are statistically significant and it's unlikely that no relationship exists between the TARGET WINS and predictor variables.

Figure 11: Model 2: Output

```
Call:
lm(formula = TARGET_WINS ~ IMP_TEAM_BATTING_H + IMP_TEAM_BATTING_BB +
   IMP_TEAM_BASERUN_SB + M_TEAM_BASERUN_SB + IMP_TEAM_FIELDING_E +
   IMP_TEAM_PITCHING_BB + IMP_TEAM_PITCHING_H + IMP_TEAM_PITCHING_HR,
   data = MoneyBall)
Residuals:
            1Q Median
   Min
                           3Q
                                  Max
-55.344 -8.578 -0.282
                         8.335 52.521
Coefficients:
                    Estimate Std. Error t value Pr(>|t|)
(Intercept)
                    -5.872491 3.409313 -1.722 0.08512
IMP_TEAM_BATTING_H 0.058421 0.003298 17.713 < 2e-16 ***
IMP_TEAM_BATTING_BB 0.034810 0.005730
                                         6.075 1.45e-09 ***
IMP_TEAM_BASERUN_SB
                    0.074371
                               0.004677
                                         15.903 < 2e-16 ***
                               1.955806 14.706 < 2e-16 ***
M_TEAM_BASERUN_SB
                    28.761232
IMP_TEAM_FIELDING_E -0.056268  0.004897 -11.490 < 2e-16 ***</pre>
IMP_TEAM_PITCHING_BB -0.005635
                               0.004984 -1.131 0.25836
                               0.002833 -2.654 0.00802 **
IMP_TEAM_PITCHING_H -0.007517
IMP_TEAM_PITCHING_HR -0.002472
                               0.006562 -0.377 0.70640
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 13.04 on 2267 degrees of freedom
Multiple R-squared: 0.3172, Adjusted R-squared: 0.3148
F-statistic: 131.6 on 8 and 2267 DF, p-value: < 2.2e-16
```

In Figure 11, we again look at the Adjusted R-Squared statistic to measure how well our model is fitting the actual data. The Adjusted R-Squared we got is **0.3148** which is roughly 31% (approx.) of the variance found in the response variable (TARGET_WINS) can be explained by the predictor variables. We notice that even though our Adjusted R-Squared value is below 50%, the predictor variables are statistically significant. Hence, this model was also selected as one of the candidates for final selection. In Model 2, adding IMP_TEAM_PITCHING_HR seems to slightly decrease the Adjusted R-Squared.

Figure 12 shows the distribution using Q-Q plot and histogram of residuals and we see some evidence of non-normality towards the upper and lower end. The histogram shows that the residuals from Model 2 also does not form a bell-shaped which usually represent a normal distribution. In my opinion, this distribution is a slightly bimodal shape. Also, the distribution seems to be fairly similar to Model 1.

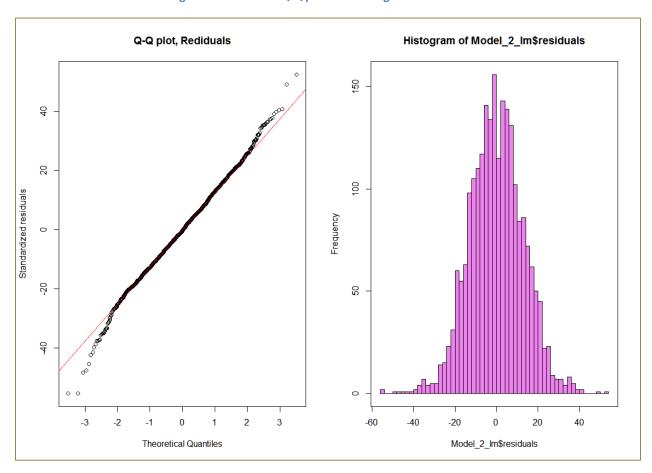


Figure 12: Model 2: Q-Q plot and Histogram of Residuals

Figure 13 shows the structure of residuals against each predictor variables using scatterplot. We notice that there's no structure in these plots except in one. The plot with IMP_TEAM_PITCHING_HR shows some form of structure. Due to this, we believe that we could not validate the homoscedasticity assumption meaning that the variance of the error term (Residuals) is not constant for all combination of independent (predictor) variables. Now, we can either transform this IMP_TEAM_PITCHING_HR variable or eliminate it from the mode completely and try to refit the model again.

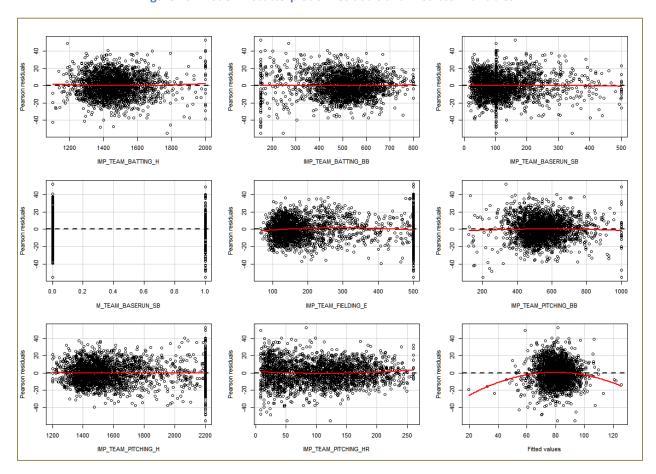


Figure 13: Model 2: Scatterplot of Residuals and Predictor Variables

Section 3.3: Model 3

Figure 14 shows the output of the Model 3. Residuals in Figure 14 appears to be symmetrical since the Median value is very close to zero (0). We would further investigate this by plotting the residuals in Q-Q plot and histogram to see whether this has a normal distribution.

In Model 3, the coefficients of each parameter are also behaving correctly (as per the theoretical effects in Data Dictionary). For instance, while holding the other predictors constant, if there is an increase of one (1) base hit by the batter (IMP_TEAM_BATTING_H), the winning chances increase by 0.06074 which is roughly 6.07% (approx.). Similarly, if a pitcher allowed one (1) additional hit (IMP_TEAM_PITCHING_H), the winning chances decrease by 0.00930 which is roughly 0.93% (approx.). Since we have included another predictor variable (IMP_TEAM_BASERUN_CS) in this mode, let's explain how that variable behaves in this model. While holding all the other variables constant, if the batter is caught stealing (IMP_TEAM_BASERUN_CS) once, the winning chances decrease by 0.03456 which is roughly 3.46% (approx.). Lastly, if the base is stolen (IMP_TEAM_BASERUN_SB), the winning chances increase by 0.07521 which is roughly 7.52% (approx.).

As you may have noticed, we have removed IMP_TEAM_PITCHING_HR predictor since it wasn't helping in Model 2. Instead, we added another predictor variable IMP_TEAM_BASERUN_CS and corresponding flag variable.

*** Significance stars in coefficients in Figure 14 shows that the predictor variables are statistically significant.

Figure 14: Model 3: Output

```
Call:
lm(formula = TARGET_WINS ~ IMP_TEAM_BATTING_H + IMP_TEAM_BATTING_BB +
   IMP_TEAM_BASERUN_SB + M_TEAM_BASERUN_SB + IMP_TEAM_BASERUN_CS +
   M_TEAM_BASERUN_CS + IMP_TEAM_PITCHING_H + IMP_TEAM_PITCHING_BB +
   IMP_TEAM_FIELDING_E, data = MoneyBall)
Residuals:
   Min
            1Q Median
                            3Q
                                   Max
-56.979 -8.556 -0.275
                         8.213 59.265
Coefficients:
                     Estimate Std. Error t value Pr(>|t|)
(Intercept)
                    -3.960326
                                3.509798 -1.128 0.259286
IMP_TEAM_BATTING_H
                     0.060744
                                0.003242 18.737
                                                 < 2e-16 ***
IMP_TEAM_BATTING_BB
                    0.031502
                                0.005672
                                          5.554 3.11e-08 ***
                                          15.280 < 2e-16 ***
IMP_TEAM_BASERUN_SB
                    0.075213
                                0.004922
                                1.917537
                                          15.792 < 2e-16 ***
                    30.282484
M_TEAM_BASERUN_SB
                                         -1.987 0.047058 *
IMP_TEAM_BASERUN_CS -0.034563
                                0.017396
                                0.861711
                                          7.460 1.23e-13 ***
M_TEAM_BASERUN_CS
                    6.428316
IMP_TEAM_PITCHING_H -0.009303
                                0.002793
                                         -3.330 0.000882 ***
IMP_TEAM_PITCHING_BB -0.001774
                                0.004897 -0.362 0.717179
IMP_TEAM_FIELDING_E -0.073967
                                0.004755 -15.555 < 2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 12.85 on 2266 degrees of freedom
Multiple R-squared: 0.3371,
                              Adjusted R-squared: 0.3344
F-statistic:
              128 on 9 and 2266 DF, p-value: < 2.2e-16
```

In Figure 14, we look at the Adjusted R-Squared statistic to measure how well our model is fitting the actual data. The Adjusted R-Squared we got is **0.3344** which is roughly **33%** (approx.) of the variance found in the response variable (TARGET_WINS) can be explained by the predictor variables. This model by far performs the best. Removing IMP_TEAM_PITCHING_HR and adding IMP_TEAM_BASERUN_CS seems to increase the Adjusted R-Squared value. Earlier in Figure 14, we noticed that coefficients improved as well.

Figure 15 shows the distribution using Q-Q plot and histogram of residuals and we see some evidence of non-normality towards the upper and lower end. The histogram shows that the residuals from Model 3 form a bell-shaped which usually represent a normally distribution. Again, this by far better than both the previous models we have seen. Overall, the residuals look pretty normal distributed.

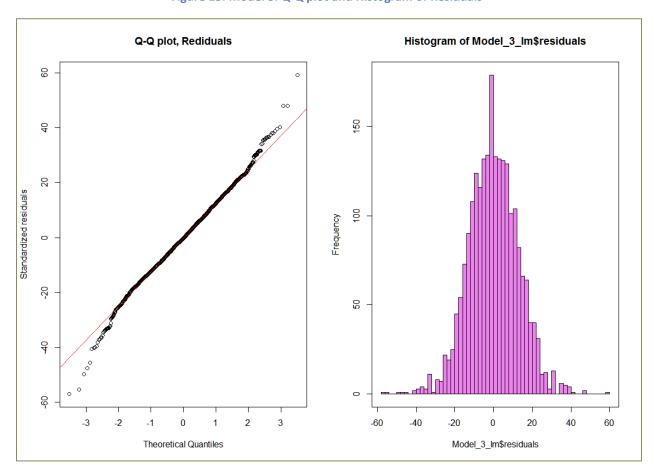


Figure 15: Model 3: Q-Q plot and Histogram of Residuals

Figure 16 shows the structure of residuals against each predictor variables using scatterplot. We notice that there's no structure in these plots. We validate the homoscedasticity assumption meaning that the variance of the error term (Residuals) is constant for all combination of independent (predictor) variables.

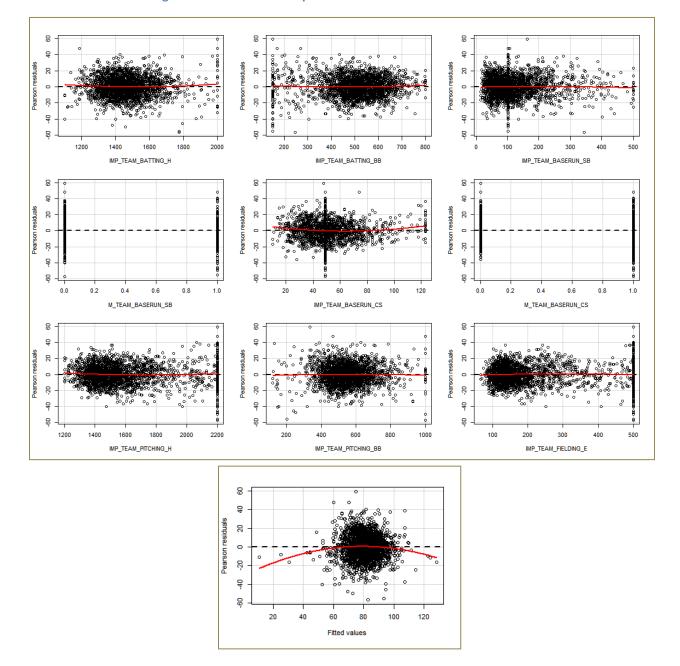


Figure 16: Model 3: Scatterplot of Residuals and Predictor Variables

Section 4: Model Comparison and Selection

Figure 17 shows each of our three models i.e. Model 1, Model 2, and Model 3. Difference between these models is that Model 2 has all the variables from Model 1 and includes an additional variable, IMP_TEAM_PITCHING_HR. Model 3 also has all the variables from Model 1 but includes two additional variables, IMP_TEAM_BASERUN_CS and M_TEAM_BASERUN_CS (flag variable). Model 3 does not include IMP_TEAM_PITCHING_HR.

When comparing our models against Adjusted R-Squared, Model 3 here outperformed all the other models. However, there's no guarantee that Model 3 will be predictive. Hence, we check for the multicollinearity for each to the models to make sure the predictor variables used in these models have no high correlation among themselves. We will also calculate other measures to compare and based on the results, we will select our final model.

Figure 17: Models: Model 1, Model 2, and Model 3

Figure 18 shows the Variance Inflation Factor (VIF) values of all the models for the predictor variables. The VIF values of the predictor variables indicate the strength of the linear relationship between the variable and remaining predictor variables. A good rule of thumb is that VIF values greater than 10 give some cause for concern. A low VIF value means that there are no high correlations among some or all predictor variables. In Figure 14, we see that all the VIF values are below 7 for Model 1, Model 2, and Model 3. Therefore, multicollinearity is not a problem for these models.

Figure 18: VIF Values: Model 1, Model 2, and Model 3

Predictor Variables	Model 1 VIF Values	Model 2 VIF Values	Model 3 VIF Values
IMP_TEAM_PITCHING_H	6.722779	6.753197	6.760962
IMP_TEAM_BATTING_BB	6.107077	6.110652	6.162474
IMP_TEAM_PITCHING_BB	4.729828	4.818159	4.788496
IMP_TEAM_FIELDING_E	3.962756	5.271085	5.117097
M_TEAM_BASERUN_SB	2.715479	2.777652	2.748805
IMP_TEAM_BATTING_H	2.598941	2.635439	2.621315
IMP_TEAM_BASERUN_SB	2.029005	2.029641	2.314802
IMP_TEAM_PITCHING_HR		2.124017	
M_TEAM_BASERUN_CS			2.293748
IMP_TEAM_BASERUN_CS			1.179531

Figure 19 shows the adjusted R-Squared, AIC, BIC, mean squared error (MSE), and the mean absolute error (MSE) for each of these models i.e. Model 1, Model 2, and Model 3. Based on the results, Model 3 performs better than all the other models by every measure. Model 3 has Adjusted R-Squared is higher, low AIC and BIC values, low Mean Squared Error (MSE), and low Mean Absolute Error (MAE). Hence, we will use Model 3 as our final model to predict the number of wins in the test data.

Figure 19: Metrics: Model 1, Model 2, and Model 3

Models	Adjusted_R_Squared	AIC_Values	BIC_Values	MSE_Values	MAE_Values
Model_1	0.3150547	18157.48	18209.05	169.3583	10.26199
Model_2	0.3147955	18159.34	18216.64	169.3477	10.25872
Model_3	0.3344353	18094.14	18157.17	164.4212	10.02838

Section 5: Model Testing and Scoring

Next, we use our Model 3 to score the MONEYBALL Test data file ("test data"). Prior to running our model, we will clean the test data in a similar way as we cleaned our training dataset. We will fix missing values as well outliers in the data set. Note, a separate protocol document will accompany the stand-alone R program. This document will list the directions on how to import file and acquire results.

Figure 20 shows the summary statistics of the predicted wins. We notice that we have all the observations i.e. 259 observations, and there is not missing values. Our model predicted a minimum of 38 wins for a team and maximum of 111 wins. The average number of wins is about 80 wins. In my opinion, the results seem to be OK. I would have been surprised or alarmed if our model predicted more than 162 wins for any team in the 162 game seasons,

Therefore, we will accept this score data result and use it as our final output.

Figure 20: Moneyball Test Result Stats

Variable	nobs	NAs	Minimum	Maximum	1. Quartile	3. Quartile	Mean	Median	Stdev
P_TARGET_WINS	259	0	38	111	75	86	80.3475	81	9.74376

Conclusion:

In conclusion, I would like to state that all three of my models had a low Adjusted R-Squared value. However, each of them behaved logically correct meaning their coefficients in the model made sense for each parameter. We selected the Model 3 that outperformed the other two models by every measure. We assessed the Goodness-Of-Fit for these models i.e. validate the normality assumption, validate the homoscedasticity assumption (equal variance), and check for multicollinearity. Model 3 passed all the assumptions and had the highest Adjusted R-squared. Hence, it was picked as our final model for prediction.

Lastly, Model 3 states that a team would have a higher number of winning chances if they do the following:

- Bat well and get to more bases (get to 1st, 2nd, 3rd, and 4th base more)
- Get more walks by batters
- Steal more bases
- Do not get caught stealing
- Pitchers allow very minimum hits or no hits
- Pitchers allow very minimum walks or no walks
- There's minimum or no fielding errors.

Appendix I: Model Development R Code

```
# Predict 411 - MONEYBALL Data Analysis Project
# Singh, Gurjeet
# 10/14/2017
#-----
library(readr)
library(car)
library(fBasics)
library(ggplot2)
library(corrplot)
## 1 - DATA EXPLORATION
colnames (MoneyBall) [1] <- "INDEX"</pre>
#Understand the stats and summary
str(MoneyBall)
summary(MoneyBall)
View(t(basicStats(MoneyBall)))
##Check the distribution of the quantitative variables
par(mfrow = c(4,4))
with (MoneyBall, hist (TARGET WINS, breaks = "FD", col = "light blue")); box();
with (MoneyBall, hist (TEAM BATTING H, breaks = "FD", col = "light blue"));
with (MoneyBall, hist (TEAM BATTING 2B, breaks = "FD", col = "light blue"));
with (MoneyBall, hist (TEAM BATTING 3B, breaks = "FD", col = "light blue"));
with (MoneyBall, hist (TEAM BATTING HR, breaks = "FD", col = "light blue"));
with (MoneyBall, hist (TEAM BATTING BB, breaks = "FD", col = "light blue"));
with (MoneyBall, hist (TEAM BATTING SO, breaks = "FD", col = "light blue"));
with (MoneyBall, hist (TEAM BASERUN SB, breaks = "FD", col = "light blue"));
with (MoneyBall, hist (TEAM BASERUN CS, breaks = "FD", col = "light blue"));
with (MoneyBall, hist (TEAM BATTING HBP, breaks = "FD", col = "light blue"));
with (MoneyBall, hist (TEAM PITCHING H, breaks = "FD", col = "light blue"));
with (MoneyBall, hist (TEAM PITCHING HR, breaks = "FD", col = "light blue"));
with (MoneyBall, hist (TEAM PITCHING BB, breaks = "FD", col = "light blue"));
with (MoneyBall, hist (TEAM PITCHING SO, breaks = "FD", col = "light blue"));
with (MoneyBall, hist (TEAM FIELDING E, breaks = "FD", col = "light blue"));
box();
```

```
with (MoneyBall, hist (TEAM FIELDING DP, breaks = "FD", col = "light blue"));
box();
par(mfrow = c(1,1))
#check the normal distribution using QQplot
par(mfrow = c(3,3))
with (MoneyBall, qqPlot(TEAM BATTING H, labels=row.names (MoneyBall),
                            main="QQ-Plot Batting H", col = "blue",id.n=3));
with (MoneyBall, qqPlot(TEAM BATTING BB, labels=row.names(MoneyBall),
                            main="QQ-Plot Batting BB", col = "blue",id.n=3));
with (MoneyBall, qqPlot(TEAM BATTING SO, labels=row.names(MoneyBall),
                            main="QQ-Plot Batting SO", col = "blue",id.n=3));
with (MoneyBall, qqPlot(TEAM BASERUN SB, labels=row.names(MoneyBall),
                            main="QQ-Plot Baserun SB", col = "blue",id.n=3));
with (MoneyBall, qqPlot(TEAM BASERUN CS, labels=row.names(MoneyBall),
                            main="QQ-Plot Baserun CS", col = "blue",id.n=3));
with (MoneyBall, qqPlot(TEAM PITCHING H, labels=row.names(MoneyBall),
                            main="QQ-Plot Pitching H", col = "blue", id.n=3));
with (MoneyBall, qqPlot(TEAM PITCHING BB, labels=row.names (MoneyBall),
                            main="QQ-Plot Pitching BB", col = "blue",
id.n=3));
with (MoneyBall, qqPlot(TEAM PITCHING SO, labels=row.names(MoneyBall)
                             , main="QQ-Plot Pitching SO", col = "blue",
id.n=3));
with (MoneyBall, qqPlot(TEAM FIELDING E, labels=row.names(MoneyBall),
                            main="QQ-Plot Fielding E",col = "blue", id.n=3));
par(mfrow = c(1,1))
##Examine Relationships
par(mfrow = c(4,3))
with (MoneyBall, plot (TEAM BATTING H , TARGET WINS,
                            main="TARGET WINS vs TEAM BATTING H",
                            col = "dark orange"))
with (MoneyBall, plot (TEAM BATTING BB , TARGET WINS,
                           main="TARGET WINS vs TEAM BATTING BB",
                           col = "dark orange"))
with (MoneyBall, plot (TEAM BATTING SO, TARGET WINS ,
                           main="TARGET WINS vs TEAM BATTING SO",
                           col = "dark orange"))
with (MoneyBall, plot (TEAM BASERUN SB, TARGET WINS ,
                           main="TARGET WINS vs TEAM BASERUN SB",
                           col = "dark orange"))
with (MoneyBall, plot (TEAM BASERUN CS, TARGET WINS ,
                           main="TARGET WINS vs TEAM BASERUN CS",
                            col = "dark orange"))
with (MoneyBall, plot (TEAM PITCHING H , TARGET WINS,
                           main="TARGET WINS vs TEAM PITCHING H",
                           col = "dark orange"))
with (MoneyBall, plot (TEAM PITCHING BB, TARGET WINS ,
                           main="TARGET_WINS vs TEAM_PITCHING_BB",
                           col = "dark orange"))
with (MoneyBall, plot (TEAM PITCHING SO, TARGET WINS,
                           main="TARGET WINS vs TEAM PITCHING SO",
                           col = "dark orange"))
with (MoneyBall, plot (TEAM FIELDING E, TARGET WINS,
                           main="TARGET WINS vs TEAM FIELDING E",
                           col = "dark orange"))
```

```
with (MoneyBall, plot (TEAM FIELDING DP , TARGET WINS,
                          main="TARGET WINS vs TEAM FIELDING DP",
                          col = "dark orange"))
par(mfrow = c(1,1))
#-----
## 2 - DATA PREPARATION
##clean missing values with median values
summary (MoneyBall)
##clean missing values with median values
MoneyBall$IMP TEAM BATTING SO <-ifelse(is.na(MoneyBall$TEAM BATTING SO),
                                         MoneyBall$TEAM BATTING SO)
MoneyBall$M TEAM BATTING SO <- ifelse(is.na(MoneyBall$TEAM BATTING SO),
MoneyBall$IMP TEAM BASERUN SB <- ifelse(is.na(MoneyBall$TEAM BASERUN SB),
                                         MoneyBall$TEAM BASERUN SB)
MoneyBall$M TEAM BASERUN SB <- ifelse(is.na(MoneyBall$TEAM BASERUN SB),
MoneyBall$IMP TEAM BASERUN CS <- ifelse(is.na(MoneyBall$TEAM BASERUN CS),
                                         MoneyBall$TEAM BASERUN CS)
MoneyBall$M TEAM BASERUN CS <- ifelse(is.na(MoneyBall$TEAM BASERUN CS),
MoneyBall$IMP TEAM PITCHING SO <- ifelse(is.na(MoneyBall$TEAM PITCHING SO),
                                         814,
                                         MoneyBall$TEAM PITCHING SO)
MoneyBall$M TEAM PITCHING SO <- ifelse(is.na(MoneyBall$TEAM PITCHING SO),
MoneyBall$IMP TEAM FIELDING DP <- ifelse(is.na(MoneyBall$TEAM FIELDING DP),
                                         MoneyBall$TEAM FIELDING DP)
MoneyBall$M TEAM FIELDING DP <- ifelse(is.na(MoneyBall$TEAM_FIELDING_DP),
                                          1, 0)
##Checking Correlation
#selectiong variables for correlation list
corr.list <- c('TARGET WINS', 'TEAM BATTING H', 'TEAM BATTING BB',
               'IMP TEAM BATTING SO', 'IMP TEAM BASERUN SB',
               'IMP TEAM BASERUN CS', 'TEAM PITCHING H',
               'TEAM PITCHING BB', 'IMP TEAM_PITCHING_SO',
               'TEAM FIELDING E', 'IMP TEAM FIELDING DP')
```

```
#Checking correlation only for the variables above.
MoneyBall.Corr <- MoneyBall[, (names (MoneyBall) %in% corr.list )]</pre>
corrplot(cor(MoneyBall.Corr), method="number")
#-----
##creating a 1st and 99th percentile
#-----
percentile.val <- matrix (with (MoneyBall,
                                c ("TEAM BATTING H",
                                      min (TEAM BATTING H),
                                      max (TEAM BATTING H),
                                      quantile (TEAM BATTING H, (0.01))[[1]],
                                       quantile (TEAM BATTING H, (0.99))[[1]]
)),
                                ncol = 5, nrow = 1)
colnames(percentile.val) <- c("FieldName", "Min", "Max", "1-Percentile",</pre>
                              "99-Percentile")
percentile.val <- rbind(percentile.val,</pre>
                        matrix(with(MoneyBall,
                                c ("TEAM BATTING 2B",
                                      min (TEAM BATTING 2B),
                                      max (TEAM BATTING 2B),
                                      quantile (TEAM BATTING 2B, (0.01))[[1]],
                                       quantile (TEAM BATTING 2B, (0.99))[[1]]
)),
                                  ncol = 5, nrow = 1))
percentile.val <- rbind (percentile.val,
                        matrix (with (MoneyBall,
                                c("TEAM BATTING 3B",
                                      min (TEAM BATTING 3B),
                                      max(TEAM BATTING 3B),
                                       quantile (TEAM BATTING_3B, (0.01))[[1]],
                                       quantile (TEAM BATTING 3B, (0.99))[[1]]
)),
                                  ncol = 5, nrow = 1))
percentile.val <- rbind (percentile.val,
                        matrix (with (MoneyBall,
                                c ("TEAM BATTING HR",
                                      min (TEAM BATTING HR),
                                      max (TEAM BATTING HR),
                                       quantile (TEAM BATTING HR, (0.01))[[1]],
                                       quantile (TEAM BATTING HR, (0.99)) [[1]]
)),
                                  ncol = 5, nrow = 1))
percentile.val <- rbind(percentile.val,
                        matrix (with (MoneyBall,
                                c("TEAM BATTING BB",
                                      min (TEAM BATTING BB),
                                      max(TEAM BATTING BB),
                                       quantile (TEAM BATTING BB, (0.01))[[1]],
```

```
quantile (TEAM BATTING BB, (0.99))[[1]]
)),
                                    ncol = 5, nrow = 1))
percentile.val <- rbind(percentile.val,</pre>
                         matrix (with (MoneyBall,
                                  c("IMP TEAM BATTING SO",
                                        min (IMP TEAM BATTING SO),
                                        max(IMP TEAM BATTING SO),
                                        quantile (IMP TEAM BATTING SO,
(0.01))[[1]],
                                        quantile (IMP TEAM BATTING SO,
(0.99))[[1]])),
                                    ncol = 5, nrow = 1))
percentile.val <- rbind(percentile.val,</pre>
                         matrix (with (MoneyBall,
                                  c("IMP TEAM BASERUN SB",
                                        min (IMP TEAM BASERUN SB),
                                        max(IMP_TEAM_BASERUN_SB),
                                        quantile (IMP_TEAM_BASERUN_SB,
(0.01))[[1]],
                                        quantile (IMP TEAM BASERUN SB,
(0.99))[[1]])),
                                    ncol = 5, nrow = 1))
percentile.val <- rbind (percentile.val,
                         matrix (with (MoneyBall,
                                  c ("IMP TEAM BASERUN CS",
                                        min (IMP TEAM BASERUN CS),
                                        max (IMP TEAM BASERUN CS),
                                        quantile (IMP TEAM BASERUN CS,
(0.01))[[1]],
                                        quantile (IMP TEAM BASERUN CS,
(0.99))[[1]])),
                                    ncol = 5, nrow = 1)
percentile.val <- rbind(percentile.val,</pre>
                         matrix (with (MoneyBall,
                                      c("TEAM PITCHING H",
                                        min (TEAM PITCHING H),
                                        max(TEAM PITCHING H),
                                        quantile (TEAM PITCHING H, (0.01))[[1]],
                                        quantile(TEAM PITCHING_H, (0.99))[[1]]
)),
                                ncol = 5, nrow = 1)
percentile.val <- rbind(percentile.val,
                         matrix (with (MoneyBall,
                                      c("TEAM PITCHING HR",
                                        min (TEAM PITCHING HR),
                                        max (TEAM PITCHING HR),
```

```
quantile (TEAM PITCHING HR,
(0.01))[[1]],
                                        quantile (TEAM PITCHING HR, (0.99)) [[1]]
)),
                                 ncol = 5, nrow = 1))
percentile.val <- rbind(percentile.val,</pre>
                          matrix (with (MoneyBall,
                                       c("TEAM PITCHING_BB",
                                         min (TEAM PITCHING BB),
                                         max(TEAM PITCHING BB),
                                         quantile (TEAM PITCHING BB,
(0.01))[[1]],
                                         quantile (TEAM PITCHING BB,
(0.99))[[1]])),
                                  ncol = 5, nrow = 1))
percentile.val <- rbind(percentile.val,</pre>
                         matrix(with(MoneyBall,
                                      c("IMP TEAM PITCHING SO",
                                        min(IMP_TEAM_PITCHING_SO),
                                        max(IMP_TEAM_PITCHING_SO),
                                        quantile (IMP TEAM PITCHING SO,
(0.01))[[1]],
                                        quantile (IMP TEAM PITCHING SO,
(0.99))[[1]])),
                                 ncol = 5, nrow = 1))
percentile.val <- rbind(percentile.val,</pre>
                         matrix (with (MoneyBall,
                                      c ("TEAM FIELDING E",
                                        min (TEAM FIELDING E),
                                        max (TEAM FIELDING E),
                                        quantile (TEAM FIELDING E, (0.01))[[1]],
                                        quantile (TEAM FIELDING E, (0.99)) [[1]]
)),
                                 ncol = 5, nrow = 1))
percentile.val <- rbind(percentile.val,</pre>
                         matrix (with (MoneyBall,
                                      c("IMP TEAM FIELDING DP",
                                        min (IMP TEAM FIELDING DP),
                                        max(IMP TEAM FIELDING DP),
                                        quantile (IMP TEAM FIELDING DP,
(0.01))[[1]],
                                        quantile (IMP TEAM FIELDING DP,
(0.99))[[1]])),
                                 ncol = 5, nrow = 1))
View(percentile.val)
```

```
##Cleaning outliers
MoneyBall$IMP TEAM BATTING H <- with (MoneyBall,
                                           ifelse (TEAM BATTING H < 1100, 1100,
                                                ifelse (TEAM BATTING H > 2000,
2000,
                                                        TEAM BATTING H)));
#--2
MoneyBall$IMP TEAM BATTING BB <- with (MoneyBall, ifelse(TEAM BATTING BB <
                                           ifelse (TEAM BATTING BB > 800, 800,
TEAM BATTING BB)));
#--3
MoneyBall$IMP TEAM BATTING SO <- with (MoneyBall, ifelse(IMP TEAM BATTING SO <
                                           ifelse (IMP TEAM BATTING SO > 1350,
1350,
IMP TEAM BATTING SO)));
#--4
MoneyBall$IMP TEAM BASERUN SB <- with (MoneyBall, ifelse(IMP TEAM BASERUN SB <
14, 14,
                                           ifelse(IMP TEAM BASERUN SB > 500,
500,
IMP TEAM BASERUN SB)));
MoneyBall$IMP TEAM BASERUN CS <- with (MoneyBall, ifelse(IMP TEAM BASERUN CS <
10, 10,
                                           ifelse (IMP TEAM BASERUN CS > 123,
123,
IMP TEAM BASERUN_CS)));
#--6
MoneyBall$IMP TEAM PITCHING H <- with (MoneyBall, ifelse (TEAM PITCHING H <
1200, 1200,
                                           ifelse (TEAM PITCHING H > 2200,
2200,
TEAM PITCHING H)));
MoneyBall$IMP TEAM PITCHING HR <- with (MoneyBall, ifelse (TEAM PITCHING HR <
8, 8,
                                           ifelse (TEAM PITCHING HR > 260, 260,
TEAM PITCHING HR)));
#--8
```

```
MoneyBall$IMP TEAM PITCHING BB <- with (MoneyBall, ifelse (TEAM PITCHING BB <
119, 119,
                                          ifelse (TEAM PITCHING BB > 1000,
1000,
TEAM PITCHING BB)));
MoneyBall$IMP TEAM PITCHING SO <- with (MoneyBall, ifelse (IMP TEAM PITCHING SO
< 241, 241,
                                         ifelse(IMP TEAM PITCHING SO >
1700, 1700,
IMP TEAM PITCHING_SO)));
#--10
MoneyBall SIMP TEAM FIELDING E <- with (MoneyBall, ifelse (TEAM FIELDING E < 65,
                                          ifelse (TEAM FIELDING E > 500, 500,
TEAM FIELDING E)));
MoneyBall$IMP TEAM FIELDING DP <- with (MoneyBall, ifelse (IMP TEAM FIELDING DP
< 71, 71,
                                           ifelse(IMP TEAM FIELDING DP >
220, 220,
IMP TEAM FIELDING DP)));
#options(scipen = 111)
View(t(basicStats(MoneyBall)))
### Re-checking distributions and relationship after fixing outliers and
#missing values
##Check the distribution of the quantitative variables
par(mfrow = c(4,3))
with (MoneyBall, hist (TARGET WINS, breaks = "FD",
                    col = "light blue")); box();
with (MoneyBall, hist (IMP_TEAM_BATTING_H, breaks = "FD",
                    col = "light blue")); box();
with (MoneyBall, hist (IMP TEAM BATTING BB, breaks = "FD",
                    col = "light blue")); box();
with (MoneyBall, hist (IMP TEAM BATTING SO, breaks = "FD",
                    col = "light blue")); box();
with (MoneyBall, hist (IMP TEAM BASERUN SB, breaks = "FD",
                    col = "light blue")); box();
with (MoneyBall, hist (IMP TEAM BASERUN CS, breaks = "FD",
                    col = "light blue")); box();
with (MoneyBall, hist (IMP TEAM PITCHING H, breaks = "FD",
                    col = "light blue")); box();
with (MoneyBall, hist (IMP TEAM PITCHING HR, breaks = "FD",
                    col = "light blue")); box();
with (MoneyBall, hist (IMP TEAM PITCHING BB, breaks = "FD",
                    col = "light blue")); box();
```

```
with (MoneyBall, hist (IMP TEAM PITCHING SO, breaks = "FD",
                     col = "light blue")); box();
with (MoneyBall, hist (IMP TEAM FIELDING E, breaks = "FD",
                     col = "light blue")); box();
with (MoneyBall, hist (IMP TEAM FIELDING DP, breaks = "FD",
                     col = "light blue")); box();
par(mfrow = c(1,1))
##Examine Relationships
par(mfrow = c(4,3))
with (MoneyBall, plot (IMP TEAM BATTING H , TARGET WINS,
                     col = "dark orange"))
with (MoneyBall, plot (IMP TEAM BATTING BB , TARGET WINS,
                     col = "dark orange"))
with (MoneyBall, plot (IMP TEAM BATTING SO, TARGET WINS ,
                     col = "dark orange"))
with (MoneyBall, plot (IMP TEAM BASERUN SB, TARGET WINS ,
                     col = "dark orange"))
with (MoneyBall, plot (IMP TEAM BASERUN CS, TARGET WINS ,
                     col = "dark orange"))
with (MoneyBall, plot (IMP_TEAM_PITCHING_H , TARGET_WINS,
                     col = "dark orange"))
with (MoneyBall, plot (IMP TEAM PITCHING BB, TARGET WINS ,
                     col = "dark orange"))
with (MoneyBall, plot (IMP TEAM PITCHING SO, TARGET WINS ,
                     col = "dark orange"))
with (MoneyBall, plot (IMP TEAM FIELDING E, TARGET WINS,
                     col = "dark orange"))
with (MoneyBall, plot (IMP_TEAM_FIELDING DP , TARGET WINS,
                     col = "dark orange"))
par(mfrow = c(1,1))
##creating a drop list
#-----
#creating a drop list to remove not required variables.
drop.list <- c('INDEX', 'TEAM BATTING HBP', 'TEAM BATTING SO',</pre>
                'TEAM BATTING BB', 'TEAM BASERUN SB', 'TEAM BASERUN CS',
                'TEAM PITCHING SO', 'TEAM FIELDING DP', 'TEAM BATTING H',
                'TEAM_BATTING_2B', 'TEAM_BATTING_3B', 'TEAM_BATTING_HR',
                'TEAM PITCHING H', 'TEAM PITCHING HR', 'TEAM PITCHING BB'
               , 'TEAM FIELDING E')
#droping the variables
MoneyBall <- MoneyBall[,!(names(MoneyBall) %in% drop.list )]</pre>
View(t(basicStats(MoneyBall)))
```

```
## 3- Build Models
#-----
#-----
##Model 1 lm
#-----
names (MoneyBall)
##Adjusted R-squared:
Model_1_lm <- lm(TARGET WINS ~ IMP TEAM BATTING H +
                         IMP TEAM BATTING BB +
                         IMP TEAM BASERUN SB + M TEAM BASERUN SB +
                         IMP TEAM FIELDING E +
                         IMP TEAM PITCHING BB +
                         IMP TEAM PITCHING H
                       ,data=MoneyBall)
summary(Model 1 lm)
##Model 1 lm - Assessing the Goodness-Of-Fit in OLS Regression
# Validating the normality assumption:
par(mfrow = c(1,2))
#Creating 2 Q-Q plots to evaluate the distribution of
# SalePrice and L SalePrice
qqnorm (Model 1 lm$residuals, main = "Q-Q plot, Rediduals",
      xlab="Theoretical Quantiles", col = "black",
      ylab="Standardized residuals", datax=FALSE)
qqline (Model 1 lm$residuals, datax=FALSE, distribution=qnorm,
      probs=c(0.25, 0.75), qtype=7, col = "red")
hist (Model 1 lm$residuals, breaks = "FD", col = "violet"); box();
par(mfrow = c(1,1))
#Validating the homoscedasticity assumption (equal variance):
residualPlots (Model 1 lm)
   \# par(mfrow = c(1,1))
   # residualPlot(Model 1 lm)
```

```
#-----
##Model 2 lm
#-----
names (MoneyBall)
##Adjusted R-squared:
                     0.3148
Model 2 lm <- lm (TARGET WINS ~ IMP TEAM BATTING H +
                          IMP TEAM BATTING BB +
                          IMP TEAM BASERUN SB + M TEAM BASERUN SB +
                          IMP TEAM FIELDING E +
                          IMP TEAM PITCHING BB +
                          IMP TEAM PITCHING H +
                          IMP TEAM PITCHING HR
                         , data=MoneyBall)
summary(Model 2 lm)
#-----
##Model 2 lm - Assessing the Goodness-Of-Fit in OLS Regression
#-----
# Validating the normality assumption:
par(mfrow = c(1,2))
#Creating 2 Q-Q plots to evaluate the distribution of
# SalePrice and L SalePrice
qqnorm (Model 2 lm$residuals, main = "Q-Q plot, Rediduals",
      xlab="Theoretical Quantiles", col = "black",
      ylab="Standardized residuals", datax=FALSE)
qqline (Model 2 lm$residuals, datax=FALSE, distribution=qnorm,
      probs=c(0.25, 0.75), qtype=7, col = "red")
hist (Model 2 lm$residuals, breaks = "FD", col = "violet"); box();
par(mfrow = c(1,1))
#Validating the homoscedasticity assumption (equal variance):
residualPlots (Model 2 lm)
par(mfrow = c(1,1))
residualPlot (Model 2 lm)
```

```
#-----
##Model 3 lm
#-----
names (MoneyBall)
##Adjusted R-squared:
                     0.3344
Model 3 lm <- lm (TARGET WINS ~ IMP TEAM BATTING H +
                           IMP TEAM BATTING BB +
                           IMP TEAM BASERUN SB + M TEAM BASERUN SB +
                           IMP TEAM BASERUN CS + M TEAM BASERUN CS +
                           IMP TEAM PITCHING H +
                           IMP TEAM PITCHING BB +
                          IMP TEAM FIELDING E
                         , data=MoneyBall)
summary(Model 3 lm)
#-----
##Model 3 lm - Assessing the Goodness-Of-Fit in OLS Regression
#-----
# Validating the normality assumption:
par(mfrow = c(1,2))
#Creating 2 Q-Q plots to evaluate the distribution of
# SalePrice and L SalePrice
qqnorm (Model 3 lm$residuals, main = "Q-Q plot, Rediduals",
      xlab="Theoretical Quantiles", col = "black",
      ylab="Standardized residuals", datax=FALSE)
qqline (Model 3 lm$residuals, datax=FALSE, distribution=qnorm,
      probs=c(0.25, 0.75), qtype=7, col = "red")
hist (Model 3 lm$residuals, breaks = "FD", col = "violet"); box();
par(mfrow = c(1,1))
#Validating the homoscedasticity assumption (equal variance):
residualPlots (Model 3 lm)
   \# par(mfrow = c(1,1))
   # residualPlot(Model 3 lm)
```

```
## 4- SELECT MODELS - Predictive Accuracy
#extract the model information from summary output
Model_1_call <- Model_1_lm$call</pre>
Model 2 call <- Model 2 lm$call
Model 3 call <- Model 3 lm$call
#Printing the each model.
Model 1 call
Model 2 call
Model 3 call
# Compute the VIF values
library(car)
Model_1_lm.VIF <- as.matrix(sort(vif(Model_1_lm), decreasing=TRUE))</pre>
Model_2_lm.VIF <- as.matrix(sort(vif(Model_2_lm), decreasing=TRUE))</pre>
Model 3 lm.VIF <- as.matrix(sort(vif(Model 3 lm), decreasing=TRUE))</pre>
colnames (Model 1 lm.VIF) <- "VIF Values"</pre>
colnames (Model_2_lm.VIF) <- "VIF Values"</pre>
colnames(Model 3 lm.VIF) <- "VIF Values"</pre>
View (Model 1 lm.VIF)
View(Model_2 lm.VIF)
View (Model 3 lm.VIF)
##MSE
mse.Model 1 lm <- mean(Model 1 lm$residuals^2)</pre>
mse.Model 2 lm <- mean (Model 2 lm$residuals^2)
mse.Model 3 lm <- mean (Model 3 lm$residuals^2)</pre>
mae.Model 1 lm <- mean(abs(Model 1 lm$residuals))</pre>
mae.Model 2 lm <- mean(abs(Model 2 lm$residuals))</pre>
mae.Model 3 lm <- mean(abs(Model 3 lm$residuals))</pre>
##Creating a Table to include all the metrics
rsqrd.Mat <- matrix(c(summary(Model_1_lm)$adj.r.squared,</pre>
                       summary(Model 2 lm)$adj.r.squared,
                        summary(Model 3 lm)$adj.r.squared),
                     ncol = 1)
rownames(rsqrd.Mat) <- c("Model 1 lm", "Model 2 lm", "Model 3 lm")</pre>
colnames(rsqrd.Mat) <- "Adjusted R Squared"</pre>
```

```
AIC.Mat <- matrix(c(AIC(Model_1_lm),</pre>
                     AIC (Model_2_lm),
                     AIC(Model_3_lm)),
                   ncol = 1)
rownames(AIC.Mat) <- c("Model 1 lm", "Model 2 lm", "Model 3 lm")</pre>
colnames(AIC.Mat) <- "AIC Values"</pre>
BIC.Mat <- matrix(c(BIC(Model 1 lm),
                     BIC (Model 2 lm),
                     BIC (Model 3 lm)),
                   ncol = 1)
rownames(BIC.Mat) <- c("Model 1 lm", "Model 2 lm", "Model 3 lm")
colnames(BIC.Mat) <- "BIC Values"</pre>
MSE.Mat <- matrix(c(mse.Model 1 lm,</pre>
                     mse.Model 2 lm,
                     mse.Model 3 lm),
                   ncol = 1)
rownames (MSE.Mat) <- c("Model_1_lm", "Model_2_lm", "Model_3_lm")
colnames (MSE.Mat) <- "MSE Values"</pre>
MAE.Mat <- matrix(c(mae.Model 1 lm,
                     mae.Model 2 lm,
                     mae.Model_3_lm),
                   ncol = 1)
rownames(MAE.Mat) <- c("Model_1_lm", "Model 2 lm", "Model 3 lm")</pre>
colnames(MAE.Mat) <- "MAE Values"</pre>
final.table <- cbind( rsqrd.Mat,</pre>
                         AIC.Mat,
                         BIC.Mat,
                         MSE.Mat,
                         MAE.Mat)
View (final.table)
```

Appendix II: Stand-Alone R Code

```
# Predict 411 - MONEYBALL Data Analysis Project
# Singh, Gurjeet
# 10/14/2017
# Stand-Alone program
library(readr)
library(car)
library(fBasics)
library(ggplot2)
library(corrplot)
## 1 - Importing a Test File and check import
summary(MoneyBall Test)
colnames (MoneyBall Test) [1] <- "INDEX"</pre>
## 2 - DATA PREPARATION
##clean missing values with median values
MoneyBall Test$IMP TEAM BATTING SO <-
ifelse(is.na(MoneyBall Test$TEAM BATTING SO),
                                             MoneyBall Test$TEAM BATTING SO)
MoneyBall Test$M TEAM BATTING SO <-
ifelse(is.na(MoneyBall Test$TEAM BATTING SO),
MoneyBall Test$IMP TEAM BASERUN SB <-
ifelse(is.na(MoneyBall_Test$TEAM_BASERUN_SB),
                                              MoneyBall_Test$TEAM_BASERUN_SB)
MoneyBall Test$M TEAM BASERUN SB <-
ifelse(is.na(MoneyBall Test$TEAM BASERUN SB),
                                            1, 0)
MoneyBall Test$IMP TEAM BASERUN CS <-
ifelse(is.na(MoneyBall Test$TEAM BASERUN CS),
                                              MoneyBall Test$TEAM BASERUN CS)
MoneyBall Test$M TEAM BASERUN CS <-
ifelse(is.na(MoneyBall Test$TEAM BASERUN CS),
```

```
MoneyBall Test$IMP TEAM PITCHING SO <-
ifelse(is.na(MoneyBall Test$TEAM PITCHING SO),
MoneyBall Test$TEAM PITCHING SO)
MoneyBall Test$M TEAM PITCHING SO <-
ifelse(is.na(MoneyBall Test$TEAM PITCHING SO),
MoneyBall Test$IMP TEAM FIELDING DP <-
ifelse(is.na(MoneyBall_Test$TEAM_FIELDING_DP),
MoneyBall Test$TEAM FIELDING DP)
MoneyBall Test$M TEAM FIELDING DP <-
ifelse(is.na(MoneyBall Test$TEAM FIELDING DP),
#-----
##Cleaning outliers
#-----
#--1
MoneyBall Test$IMP TEAM BATTING H <- with (MoneyBall Test,
                                 ifelse (TEAM BATTING H < 1100, 1100,
                                        ifelse (TEAM BATTING H > 2000, 2000,
                                               TEAM BATTING H)));
MoneyBall Test$IMP TEAM BATTING BB <- with (MoneyBall Test,
ifelse (TEAM BATTING BB < 150, 150,
                                                  ifelse(TEAM BATTING BB >
800, 800,
TEAM BATTING BB)));
#--3
MoneyBall Test$IMP TEAM BATTING SO <- with (MoneyBall Test,
ifelse (IMP TEAM BATTING SO < 72, 72,
                                                 ifelse(IMP TEAM BATTING SO
> 1350, 1350,
IMP TEAM BATTING SO)));
#--4
MoneyBall Test$IMP TEAM BASERUN SB <- with (MoneyBall Test,
ifelse (IMP TEAM BASERUN SB < 14, 14,
                                                 ifelse (IMP TEAM BASERUN SB
> 500, 500,
IMP TEAM BASERUN SB)));
MoneyBall Test$IMP TEAM BASERUN CS <- with (MoneyBall Test,
ifelse (IMP TEAM BASERUN CS < 10, 10,
ifelse (IMP TEAM BASERUN CS > 123, 123,
```

```
IMP TEAM BASERUN CS)));
#--6
MoneyBall Test$IMP TEAM PITCHING H <- with (MoneyBall Test,
ifelse (TEAM PITCHING H < 1200, 1200,
                                                         ifelse (TEAM PITCHING H
> 2200, 2200,
TEAM PITCHING H)));
MoneyBall Test$IMP TEAM PITCHING HR <- with (MoneyBall Test,
ifelse(TEAM PITCHING_HR < 8, 8,
ifelse (TEAM PITCHING HR > 260, 260,
TEAM PITCHING HR)));
MoneyBall Test$IMP TEAM PITCHING BB <- with (MoneyBall Test,
ifelse (TEAM_PITCHING BB < 119, 119,
ifelse (TEAM PITCHING BB > 1000, 1000,
TEAM PITCHING BB)));
#--9
MoneyBall Test$IMP TEAM PITCHING SO <- with (MoneyBall Test,
ifelse (IMP TEAM PITCHING SO < 241, 241,
ifelse (IMP TEAM PITCHING SO > 1700, 1700,
IMP TEAM PITCHING SO)));
#--10
MoneyBall Test$IMP TEAM FIELDING E <- with (MoneyBall Test,
ifelse (TEAM FIELDING E < 65, 65,
                                                        ifelse (TEAM FIELDING E
> 500, 500,
                                                          TEAM FIELDING E)));
#--11
MoneyBall Test$IMP TEAM FIELDING DP <- with (MoneyBall Test,
ifelse (IMP TEAM FIELDING DP < 71, 71,
ifelse (IMP TEAM FIELDING DP > 220, 220,
IMP TEAM FIELDING DP)));
```

```
## 3- MODEL Deployment
View(t(basicStats(MoneyBall Test)))
P TARGET WINS <- with (MoneyBall Test, - 3.960326
                      + 0.060744 * IMP TEAM BATTING H
                      + 0.031502 * IMP TEAM BATTING BB
                      + 0.075213 * IMP TEAM BASERUN SB
                      + 30.282484 * M TEAM BASERUN SB
                      - 0.034563 * IMP TEAM BASERUN CS
                      + 6.428316 * M TEAM BASERUN CS
                      - 0.009303 * IMP TEAM PITCHING H
                      - 0.001774 * IMP TEAM PITCHING BB
                      - 0.073967 * IMP TEAM FIELDING E)
FINAL Submission <- with (MoneyBall Test, cbind.data.frame(INDEX,
round(P TARGET WINS)))
colnames(FINAL_Submission) <- c("INDEX", "P_TARGET_WINS")</pre>
View(FINAL_Submission)
View(t(basicStats(FINAL_Submission[-1])))
write.csv(FINAL Submission, "Singh Gurjeet MoneyBall Test Score.csv")
```