## Baseball Salary Data Study

1. # (1) Data = read.csv('baseball.csv') baseball=Data[,1:17] # This takes data from x1..x16. fit = lm(salary ~., data=baseball) # use the data frame baseball # salary on the left of ~ has R use salary column of baseball as response  $\sharp$  . on right of  $\sim$  tells R to use all other columns of baseball as independent variables > summary(fit) Call: lm(formula = salary ~ ., data = baseball) Residuals: 10 Median Min 30 10.9 340.7 3181.7 -1908.3 -463.0 Coefficients: Estimate Std. Error t value Pr(>|t|) 223.115 332.717 0.671 0.502970 (Intercept) 2712.536 1.122 0.262746 3043.192 batting.average 2376.084 -1.485 0.138581 -3528.013 on.base.percent 7.100 5.643 1.258 0.209259 3.312 -0.815 0.415788 hits -2.698 doubles 1.368 8.611 0.159 0.873846 -17.922 21.647 -0.828 0.408339 triples 12.583 1.548 0.122506 home.runs 19.483 3.436 0.000668 \*\*\* 17.415 rbi 5.068 1.285 0.199548 walks 5.815 4.523 -9.586 2.151 -4.457 1.15e-05 \*\*\* strike.outs 13.044 4.714 2.767 0.005988 \*\* stolen.bases -9.553 7.500 -1.274 0.203693 errors free.agent.eligible 1372.886 108.594 12.642 < 2e-16 \*\*\* free.agent -280.790 137.640 -2.040 0.042168 \* arbitration.eligible 783.592 118.289 6.624 1.48e-10 \*\*\* 352.114 241.829 1.456 0.146361 Signif. codes: 0 \\*\*\*' 0.001 \\*\*' 0.01 \\*' 0.05 \.' 0.1 \' 1 Residual standard error: 694.3 on 320 degrees of freedom Multiple R-squared: 0.7014, Adjusted R-squared: 0.6865

F-statistic: 46.99 on 16 and 320 DF, p-value: < 2.2e-16

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# (2)
# coefficient of independent variable "hits": -2.698
# according to this, getting 10 extra hits means a salary DECREASE of 10*2.7 = $27.
# This does not make sense. By my intuition, one would expect that more hits means better payday.
# There should be a positive coefficient for hits, not negative.

variation_percentage = summary(lm(salary ~.,data=baseball))$r.squared
# 70.14% of variation in salaries explained by linear model
> variation_percentage = summary(lm(salary ~.,data=baseball))$r.squared
> variation_percentage
[1] 0.7014386
```

3. Based on the summary of the fit in problem 1, we found that the p-value when testing the independent variables against the response (salary) is 2.2 \* 10^-16. This is much less than the level of significance 0.05 and would be lower than most level of significances anyway. Even when we tested individual response variables, the largest p-value attained was 0.02, still less than 0.05. The linear model's variables are all at least somewhat related to the salary.

```
# (3)
baseball none = Data[,c(1,13)] \# testing free agent status vs salary had the highest p-value
# But this p-value of 0.02 was still < 0.05.
fit none = lm(salary ~., data=baseball none) # use data from the eleven columns above
> baseball none = Data[,c(1,13)]
> fit none = lm(salary ~.,data=baseball none) # use data from the eleven columns above
> summary(fit none)
Call:
lm(formula = salary ~ ., data = baseball none)
Residuals:
   Min 1Q Median 3Q
                                 Max
-1455.6 -969.1 -528.8 821.2 4644.4
Coefficients:
      Estimate Std. Error t value Pr(>|t|)
(Intercept) 1078.11 102.05 10.565 <2e-16 ***
errors 25.17
                       11.35 2.218 0.0272 *
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 1233 on 335 degrees of freedom
Multiple R-squared: 0.01447, Adjusted R-squared: 0.01153
F-statistic: 4.919 on 1 and 335 DF, p-value: 0.02723
```

4. From the summary of the fitted model below using the 11 chosen independent variables, the p value attained = 2.2e^-16 which is much less than the level of significance of 0.05. Based on this, we can agree with the null hypothesis and say that batting average, OBP, hits, doubles, and triples are not needed in the same model with the other 11 variables. This result is very surprising because you would expect the best players (the ones with the higher salaries) to have many more hits and a better batting average. We take these pieces away but still get a relatively strong model even with the other 11 variables like home runs, walks, strikeouts, etc.

```
# (4)
baseball eleven = Data[,c(1,8:17)] # This takes data from x8..x17 and compares with salary in x1
fit eleven = lm(salary ~., data=baseball eleven) # use data from the eleven columns above
> baseball eleven = Data[,c(1,8:17)] # This takes data from x8..x17 and compares with salary in x1
> fit_eleven = lm(salary ~.,data=baseball_eleven)
 > variation percentage eleven = summary(lm(salary ~.,data=baseball eleven))$r.squared
 > variation percentage eleven
C:[1] 0.6972869
lm(rormula = salary ~ ., data = paseball eleven)
Residuals:
           1Q Median
                            3Q
  Min
                                     Max
-1856.2 -463.6 42.9 349.0 3260.0
Coefficients:
                   Estimate Std. Error t value Pr(>|t|)
(Intercept)
                    -90.563 88.553 -1.023 0.3072
24.786 9.774 2.536 0.0117 *
18.068 3.257 5.547 6.01e-08 ***
home.runs
rbi

    walks
    3.857

    strike.outs
    -9.837

    stolen.bases
    15.196

    errors
    -8.491

                                   2.465 1.565 0.1185
                                  1.931 -5.095 5.92e-07 ***
3.666 4.145 4.33e-05 ***
7.167 -1.185 0.2370
free.agent.eligible 1367.017 104.696 13.057 < 2e-16 *** free.agent -280.462 136.714 -2.051 0.0410 *
arbitration.eligible 782.842 116.474 6.721 8.06e-11 ***
                      373.259 238.871 1.563 0.1191
arbitration
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 692.6 on 326 degrees of freedom
Multiple R-squared: 0.6973, Adjusted R-squared: 0.688
F-statistic: 75.09 on 10 and 326 DF, p-value: < 2.2e-16
  5.
# (5)
variation percentage eleven = summary(lm(salary ~.,data=baseball eleven))$r.squared
# 69.73% of variation in salaries explained by linear model of given 11 columns
```

6.

```
# (6)
resid = fit_eleven$residuals
# put predicted values into a predict vector
predict = fit$fitted.value

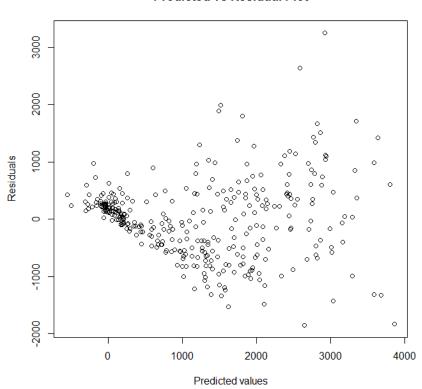
plot(predict,resid, main="Predicted vs Residual Plot", xlab="Predicted values", ylab="Residuals")
# plot predicted values vs residuals

plot(density(fit_eleven$residuals), main = "Kernel Density plot") # kernel density estimate

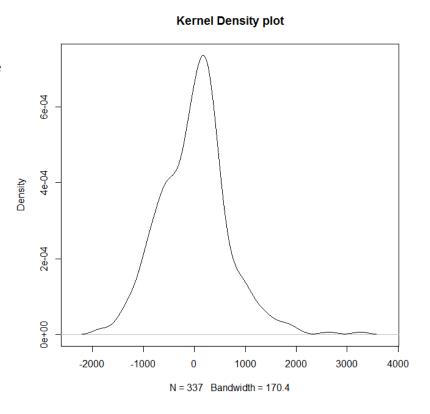
out = qqnorm((resid-mean(resid))/sd(resid)) # normal probability plot
x = range(out$x)
lines(c(x[1],x[2]),c(x[1],x[2]))
```

 In the predicted vs residual plot, as the predicted values increase, there appears to be more large residual values. This would lead us to say that the model we have fits well for predicted values less than 1000 but after 1000 there's much more fluctuation in terms of how accurately our model fits the given dataset.

## **Predicted vs Residual Plot**



The kernel density plot to the left shows that the data is distributed normally. You can see a rough normal curve on the plot with one slight dent between -1000 and 0 on the x-axis on the curve. Based on this plot, we can assume normality for the data and proceed with any calculations based on this assumption.



• The normal probability plot to the right shows a very linear relationship between the theoretical quantiles and the sample quantiles. The plotted values closely follow a 45 degree line here so just like we concluded earlier in the kernel density plot, we can safely say that the data in the baseball file are approximately normally distributed.

