

HOMEWORK 3

Spring 2025

Instructions

- The homework is due on **Mar. 7, 2025**, 11:59 pm.
 - The homework should be submitted electronically via Canvas.
 - Please upload a single PDF containing the solutions in the correct order. If you include scanned images, make sure that they are organized and easy to read.
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1. (12*6=72 points) Using the 'Baseball Salary Data' in canvas, which includes the data (a `.csv` file) and a `.txt` file with some background information on the data and the variables, do the following:
 - (a) Fit a linear regression model with **"salary"** as the response, and the other 16 variables (excluding **"names"**) as the predictors.
 - (b) What percentage of the variation in salaries is explained by the linear model above?
 - (c) Comment on the coefficient of the predictor **"hits"**. Is this coefficient consistent with what your intuition says should be the relationship between number of **hits** and **salary**? Why or why not?
 - (d) Test the null hypothesis (using level of significance $\alpha = 0.05$) that none of the 16 predictors is related to **salary**. What is the proper conclusion about the linear model above and its utility?
 - (e) Test the null hypothesis (using level of significance $\alpha = 0.05$) that the variables **"batting average"**, **"on base percentage"**, **"hits"**, **"doubles"** and **"triples"** are **not** needed in the same model with the other 11 predictors. Is the result surprising? Give a possible explanation for the result.
 - (f) What percentage of the variation in salaries is explained by the linear model containing the 11 variables not named in part (e) above?
 - (g) Obtain residuals from the linear model fitted in part (f) above, and produce the following three plots: (i) the residuals versus the predicted values, (ii) a kernel density estimate of the residuals, and (iii) a Normal probability (or Q-Q) plot of the *standardized* residuals. Comment on the plots.
 - (h) Use the command `'leaps'` in the R package **"leaps"** along with the strategy discussed in class to choose a good subset of the 16 predictors to include in a linear model with **"salary"** as the response. **Describe fully the rationale you use** in choosing your model.
 - (i) Plot the *standardized* residuals from your chosen model versus an index running from 1 to 337. Identify any players who have standardized residuals that are larger in absolute value than 3. Are these players different in any important way from most of the other players?
 - (j) Provide a plot of the standardized residuals versus the predicted values and comment on the plot.
 - (k) Provide a Normal probability (or Q-Q) plot of the standardized residuals and comment on the plot.
 - (l) Provide a plot of Cook's D values. Do any data points seem to be influential? Why or why not?
2. (Problems 3, on page 359 of textbook, Section 10.2). (9+9+10=28 points) Four different concentrations of ethanol are compared at level $\alpha = 0.05$ for their effect on sleep time. Each concentration was given to a sample of 5 rats and the REM (rapid eye movement) sleep time for each rat was recorded (SleepRem.txt). Do the four concentrations differ in terms of their effect on REM sleep time?
 - (a) State the relevant null and alternative hypotheses for answering this question, and use hand calculations to conduct the ANOVA F test at level of significance 0.05. State any assumptions needed for the validity of this test procedure. (Hint. You may use the summary statistics $\bar{X}_1 = 79.28$, $\bar{X}_2 = 61.54$, $\bar{X}_3 = 47.92$, $\bar{X}_4 = 32.76$, and $MSE = S_p^2 = 92.95$.)

- (b) Import the data into the R data frame `sl`, and use the R command `anova(aov(sl$values sl$ind))` to conduct the ANOVA F test. Give the ANOVA table, stating the p-value, and the outcome of the test at level of significance 0.05.
- (c) Use R commands to test the assumptions of equal variances and normality. Report the p-values from the two tests and the conclusions reached. Next, construct a boxplot and the normal Q-Q plot for the residuals, and comment on the validity of the normality assumption on the basis of these plots.