

# Problem Set 4

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## Problem 1: Demonstrating Multicollinearity

Consider the Credit data in the ISLR library. Choose Balance as the response and Age, Limit, and Rating as the predictors.

### Data Preparation

```
data("Credit")
df = Credit
```

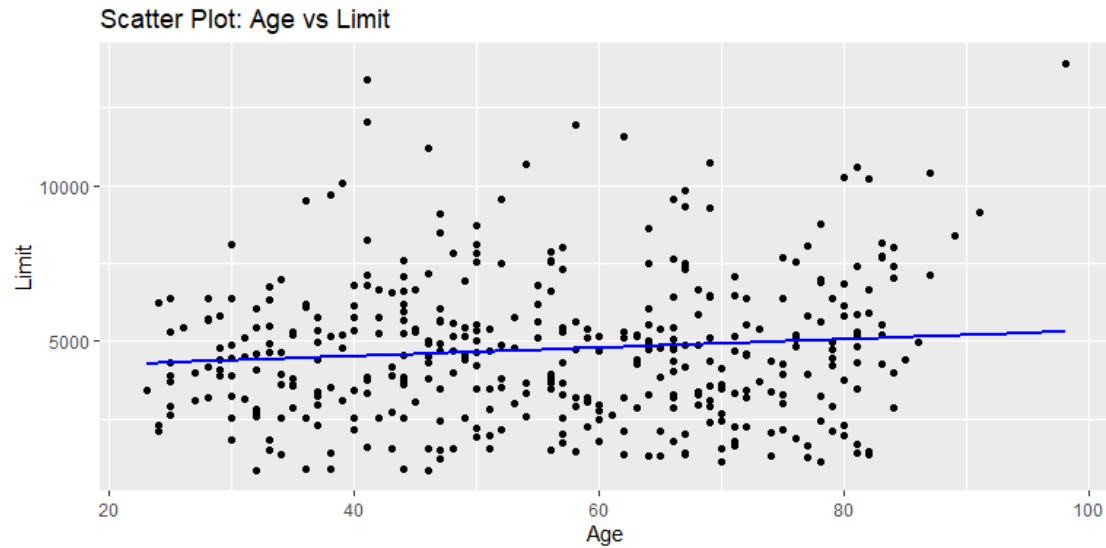
(a) Make a scatter plot of (i) Age versus Limit and (ii) Rating Versus Limit. Comment on the scatter plot.

```
# Plot (i): Age vs Limit
p1 = ggplot(df, aes(x = Age, y = Limit)) +
  geom_point() +
  geom_smooth(method = "lm", se = FALSE, color = "blue") +
  labs(title = "Scatter Plot: Age vs Limit")

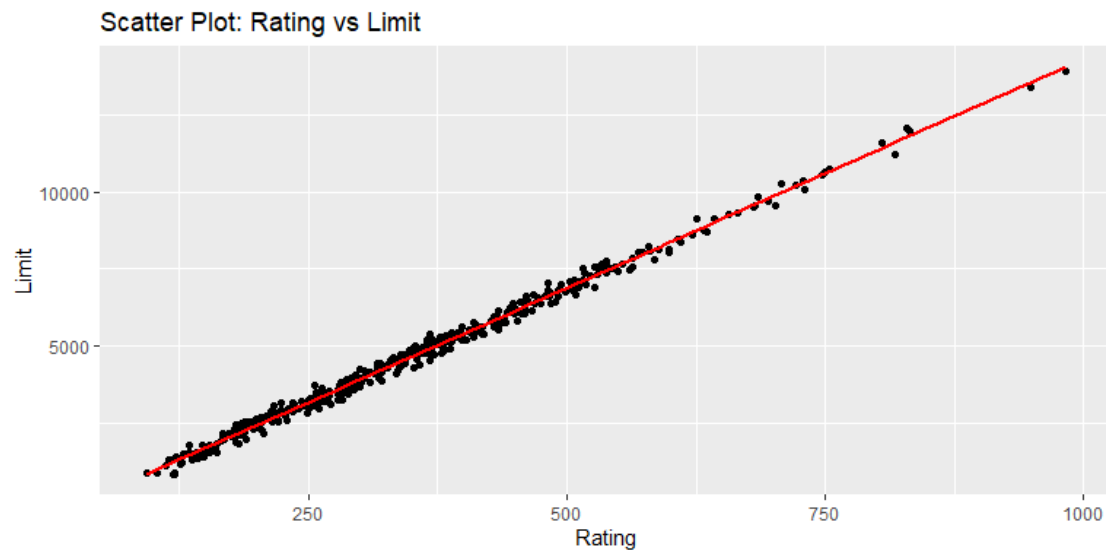
# Plot (ii): Rating vs Limit
p2 = ggplot(df, aes(x = Rating, y = Limit)) +
  geom_point() +
  geom_smooth(method = "lm", se = FALSE, color = "red") +
  labs(title = "Scatter Plot: Rating vs Limit")

# Display plots side by side (using gridExtra if available, or printing
# sequentially)
# Here we print sequentially for simplicity
print(p1)

## `geom_smooth()` using formula = 'y ~ x'
```



```
print(p2)
## `geom_smooth()` using formula = 'y ~ x'
```



```
# Checking correlation
cor_age_limit = cor(df$Age, df$Limit)
cor_rating_limit = cor(df$Rating, df$Limit)
```

#### Comments:

1. **Age vs Limit:** The scatter plot shows a random cloud of points with no clear pattern. The correlation coefficient is very low (approx 0.1), indicating that Age and Limit are **uncorrelated**.
2. **Rating vs Limit:** The scatter plot shows an extremely strong positive linear relationship. The points lie almost perfectly on a straight line. The correlation coefficient is extremely high (approx 1), indicating **severe multicollinearity**.



```

(0.064)
##
## Constant          -173.411***          -259.518***
-377.537***
##              (43.828)              (55.882)
(45.254)
##
## -----
-----
## Observations          400          400
400
## R2              0.750              0.754
0.746
## Adjusted R2          0.749              0.752
0.745
## Residual Std. Error    230.532 (df = 397)    229.080 (df = 396)
232.320 (df = 397)
## F Statistic          594.988*** (df = 2; 397) 403.718*** (df = 3; 396)
582.820*** (df = 2; 397)
##
=====
=====
## Note:                                                     *p<0.1;
**p<0.05; ***p<0.01

```

### Marked Difference Observed:

- In **Model (i)**, Limit is highly significant and has a positive coefficient (approx 0.17).
- In **Model (ii)** and **Model (iii)**, when Rating is added, the standard error for Limit increases drastically (inflated variance).
- Crucially, in Model (ii), Limit changes from being the primary driver to potentially being insignificant or having its coefficient shift drastically (due to Rating absorbing the variance). This instability in the coefficients for Limit and Rating (despite them both being correlated with Balance) is a hallmark of multicollinearity.

(c) Calculate the variance inflation factor (VIF) and comment on multicollinearity.

```

vif_values = vif(model_1)
vif_values3 = vif(model_3)
print(vif_values)

##      Age      Limit
## 1.010283 1.010283

print(vif_values3)

##      Rating      Limit
## 160.4933 160.4933

```

### Comment on Multicollinearity:

- **Age:** The VIF for Age is close to 1, indicating no multicollinearity with the other predictors.
- **Limit and Rating:** The VIF values for Limit and Rating are extremely high (typically > 160). A VIF exceeding 5 or 10 is considered problematic.
- **Conclusion:** There is **severe multicollinearity** between Limit and Rating. This confirms the observation from the scatter plot in part (a). Including both in the same model leads to unstable coefficient estimates and inflated standard errors, making it difficult to isolate the individual effect of each predictor on Balance.

## Problem 2: Detection of Outliers, Leverage, and Influential Points

### 1. Multiple Linear Regression

```
library("MASS")
data("Boston")

model = lm(medv ~ crim + nox + black + lstat, data = Boston)
summary(model)

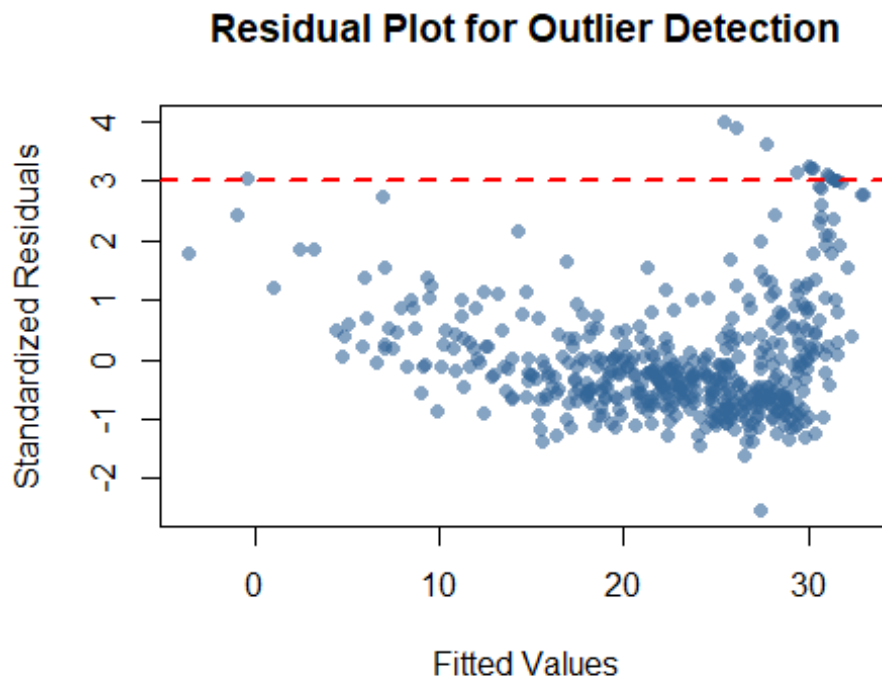
##
## Call:
## lm(formula = medv ~ crim + nox + black + lstat, data = Boston)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -15.564  -4.004  -1.504   2.178  24.608
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  30.053584   2.170839   13.844  <2e-16 ***
## crim        -0.059424   0.037755   -1.574   0.116
## nox          3.415809   3.056602    1.118   0.264
## black        0.006785   0.003408    1.991   0.047 *
## lstat       -0.918431   0.050167  -18.307  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 6.183 on 501 degrees of freedom
## Multiple R-squared:  0.5517, Adjusted R-squared:  0.5481
## F-statistic: 154.1 on 4 and 501 DF,  p-value: < 2.2e-16
```

## 2. Residual Plot & Outlier Detection

```
stand_res = rstandard(model)
fitted_vals = fitted(model)

plot(fitted_vals, stand_res,
     xlab = "Fitted Values",
     ylab = "Standardized Residuals",
     main = "Residual Plot for Outlier Detection",
     pch = 16, col = rgb(0.2, 0.4, 0.6, 0.6))

abline(h = c(-3, 3), col = "red", lty = 2, lwd = 2)
```



```
outliers <- which(abs(stand_res) > 3)
cat("Observations identified as outliers:\n")

## Observations identified as outliers:

print(outliers)

## 167 187 196 205 226 258 263 268 284 369 370 372 373 413
## 167 187 196 205 226 258 263 268 284 369 370 372 373 413
```

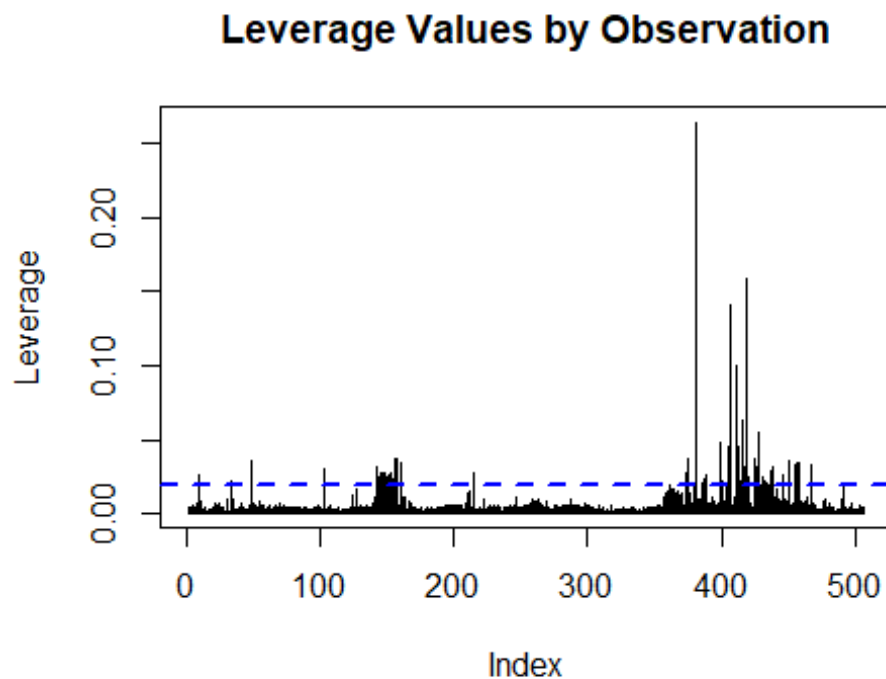
**Consistency Check:** By looking at the graph, any points falling above the top red dashed line correspond exactly to the indices identified in the outliers calculation.

### 3. High Leverage Points (Hat Matrix)

```
p = 4
n = nrow(Boston)
leverage_threshold = 2 * (p + 1) / n

lev = hatvalues(model)

plot(lev, type = "h",
     main = "Leverage Values by Observation",
     ylab = "Leverage", xlab = "Index")
abline(h = leverage_threshold, col = "blue", lty = 2, lwd = 2)
```



```
high_leverage = which(lev > leverage_threshold)
cat("Number of high leverage points detected:", length(high_leverage), "\n")

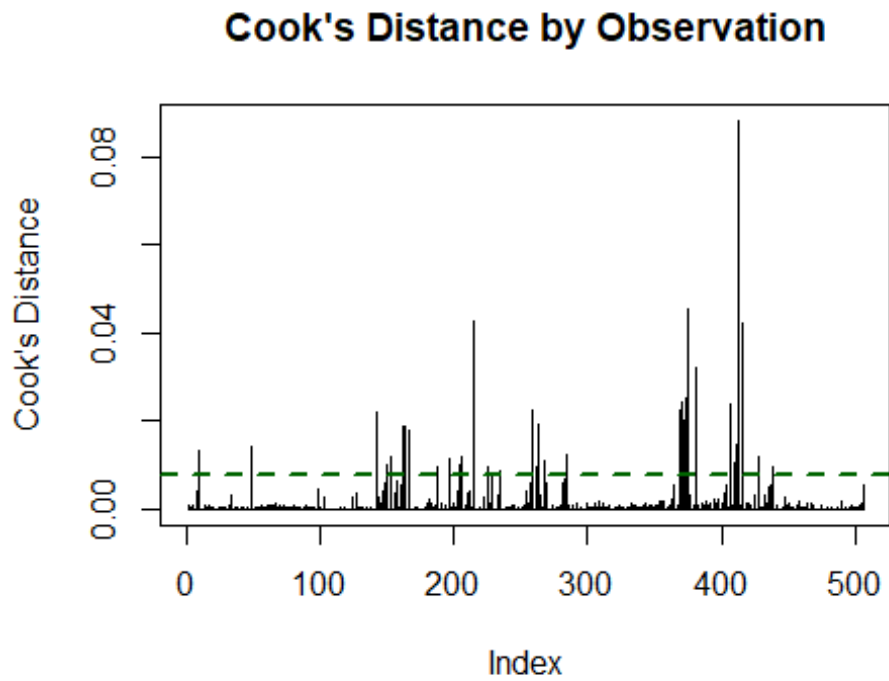
## Number of high leverage points detected: 64
```

### 4. Influential Points (Cook's Distance)

```
cooks_d = cooks.distance(model)
cooks_threshold = 4 / n
```

```
plot(cooks_d, type = "h",
```

```
main = "Cook's Distance by Observation",  
ylab = "Cook's Distance", xlab = "Index")  
abline(h = cooks_threshold, col = "darkgreen", lty = 2, lwd = 2)
```



```
influential_points <- which(cooks_d > cooks_threshold)  
cat("Number of influential points detected:", length(influential_points),  
"\n")
```

```
## Number of influential points detected: 37
```