DiZhen_1717719

dizhen

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Set directory

```
Data

WELL CT 0(22 62 50 01 70 22 46 57 20 42 74 72 43 76 72 61 63 77 25 21 70 02 76 01 72 103 00
```

```
KBI <- c(28,68,59,91,70,38,46,57,89,48,74,78,43,76,72,61,63,77,85,31,79,92,76,91,78,103,99,73,88,64,52,74)

ADL <- c(39,52,89,57,28,34,42,52,88,90,38,83,30,45,47,90,63,34,76,26,68,85,22,82,80,80,81,30,27,72,46,6)

MEM <- c(4,33,17,31,35,3,16,6,41,24,22,41,9,33,36,17,14,35,33,13,34,28,12,57,51,20,20,7,27,9,15,52,26,5)

COG <- c(18,9,3,7,19,25,17,26,13,3,13,11,24,14,18,0,16,22,23,18,26,10,16,3,3,18,1,17,27,0,22,13,18,0,19)

mydata <- data.frame(KBI,ADL,MEM,COG)

str(mydata)
```

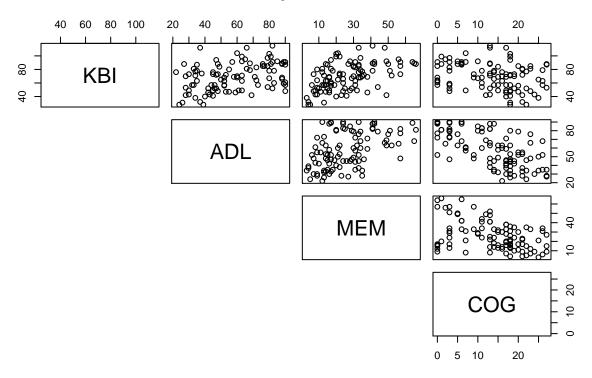
```
## 'data.frame': 100 obs. of 4 variables:
## $ KBI: num 28 68 59 91 70 38 46 57 89 48 ...
## $ ADL: num 39 52 89 57 28 34 42 52 88 90 ...
## $ MEM: num 4 33 17 31 35 3 16 6 41 24 ...
## $ COG: num 18 9 3 7 19 25 17 26 13 3 ...
```

head(mydata)

```
## KBI ADL MEM COG
## 1 28 39 4 18
## 2 68 52 33 9
## 3 59 89 17 3
## 4 91 57 31 7
## 5 70 28 35 19
## 6 38 34 3 25
```

Visualization

Scatterplot Matrix



Question 1

```
mydata.LM1 <- lm(KBI ~ ADL + MEM + COG, data = mydata)
summary(mydata.LM1)
##
## Call:</pre>
```

```
## lm(formula = KBI ~ ADL + MEM + COG, data = mydata)
##
## Residuals:
##
       Min
                10 Median
                                3Q
                                       Max
   -42.037 -10.535 -1.503
                             9.213 43.151
##
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
##
                                     4.008 0.000121 ***
## (Intercept) 40.4908
                           10.1030
## ADL
                 0.2162
                            0.1168
                                     1.851 0.067273 .
## MEM
                 0.5547
                            0.1300
                                     4.267 4.65e-05 ***
## COG
                 0.1210
                            0.3003
                                     0.403 0.687978
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 17.26 on 96 degrees of freedom
```

```
## Multiple R-squared: 0.282, Adjusted R-squared: 0.2596
## F-statistic: 12.57 on 3 and 96 DF, p-value: 5.315e-07
```

The multiple regression equition is:

```
KBI = 40.5 + 0.2 \times ADL + 0.6 \times MEM + 0.1 \times COG
```

The mean KBI when ADL, MEM and COG are zero is 40.5. For every extra unit of ADL, the expected KBI increases by 0.2 points, holding other variables constant. For every extra unit of MEM, the mean KBI increases by 0.6 points, holding other variables constant. For every extra unit of COG, the mean KBI increases by 0.1 points, holding other variables constant.

Question 2

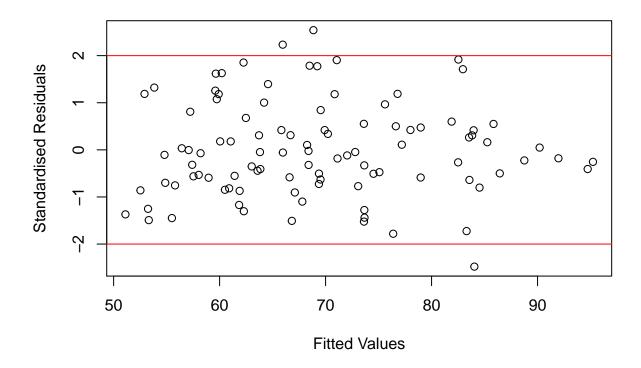
Evaluate the presence of outliers in the dependent variable using standardised residuals.

There are two observations greater than two standard deviations larger than the mean and there are one observations less than two standard deviations below the mean. There are three outliers in the dependent variable in total.

```
mydata$FITTED <- predict(mydata.LM1, type = "response")
mydata$RESID <- resid(mydata.LM1)

mydata$RSTAND <- rstandard(mydata.LM1)
plot(RSTAND ~ FITTED, data = mydata,
         ylab = "Standardised Residuals",
         xlab = "Fitted Values",
         main = "Standardised RvF Plot")
abline(h = c(-2, 2), col = "red")</pre>
```

Standardised RvF Plot



```
mydata[abs(mydata$RSTAND) > 2,]
##
      KBI ADL MEM COG
                         FITTED
                                    RESID
                                              RSTAND
           37
               33
                    17 68.84927
                                 43.15073
                                           2.540513
                    12 84.03658 -42.03658 -2.478963
       42
           69
               49
## 92 104
           60
               21
                    7 65.95588
                                 38.04412
```

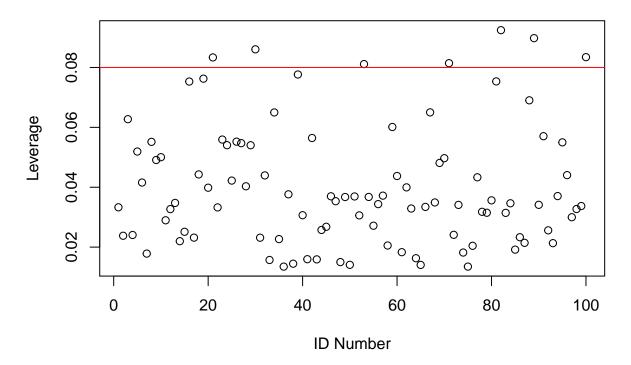
Question 3

Evaluate the presence of outliers in the independent variables using leverage values.

The plot shows 7 observations are high-leverage points. They are outliers in the independent variable.

```
mydata$HAT <- hatvalues(mydata.LM1)
HAT.CUT <- 2 * (3 + 1) / length(KBI)
ID <- seq(1,length(KBI),by = 1)
plot(HAT ~ ID, data = mydata,
        ylab = "Leverage",
        xlab = "ID Number",
        main = "Leverage by Index Plot")
abline(h = HAT.CUT, col = "red")</pre>
```

Leverage by Index Plot



```
mydata[mydata$HAT > HAT.CUT,]
```

```
##
       KBI ADL MEM COG
                                       RESID
                                                  RSTAND
                                                                 HAT
                          FITTED
## 21
        79
            68
                 34
                     26 77.19408
                                    1.805925
                                              0.1092915 0.08335487
                                              0.1789481 0.08605394
##
  30
        64
            72
                  9
                      0 61.04743
                                    2.952570
##
   53
        73
                57
                      9 83.57120
                                 -10.571198 -0.6389755 0.08112930
##
  71
        89
            68
                65
                      6 91.96908
                                   -2.969081 -0.1794948 0.08142400
##
  82
        63
            52
                15
                      0 60.05181
                                    2.948193
                                              0.1793117 0.09245325
## 89
            90
                12
                      0 66.60258
                                   -9.602580 -0.5831815 0.08978768
        57
        88
                      1 94.72923
                                   -6.729226 -0.4072611 0.08344422
## 100
                66
```

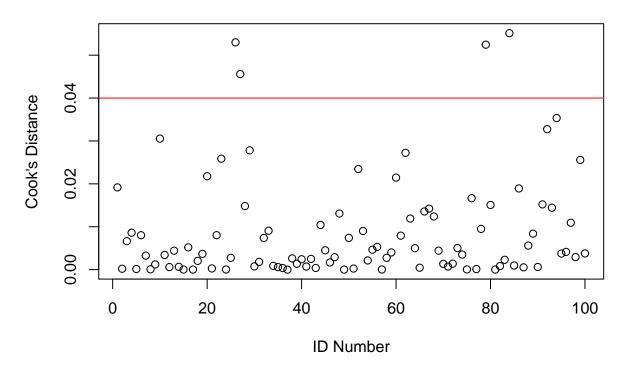
Question 4

Evaluate the presence of influential observations using Cook's distance.

The plot shows there are 4 observations with high Cook's distance values. Two of them (ID79 and ID84) are detected as having high standardised residuals.

```
mydata$COOK <- cooks.distance((mydata.LM1))
COOK.CUT <- 4/length(ID)
plot(COOK ~ ID, data = mydata,
    ylab = "Cook's Distance",
    xlab = "ID Number",
    main = "Cook's Distrance by Index Plot")
abline(h = COOK.CUT, col = "red")</pre>
```

Cook's Distrance by Index Plot



mydata[mydata\$COOK > COOK.CUT,]

```
##
      KBI ADL MEM COG
                        FITTED
                                    RESID
                                             RSTAND
                                                           HAT
                                                                      COOK
                   18 71.05536
                                31.94464
                                           1.904236 0.05522324 0.05298758
           81
               20
                    1 69.21517
                                29.78483
                                           1.775010 0.05471437 0.04559108
       99
               33
                                43.15073
                                          2.540513 0.03148128 0.05244777
  79 112
           37
                   17 68.84927
                   12 84.03658 -42.03658 -2.478963 0.03464047 0.05512833
       42
               49
```

THE END