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# Model Adequecy Check
### B Aditi \|\| MM19B022
## Problem 4.2
```{r}
#Getting table from 3.1
tb1 <- read.csv('Table_B1.csv',header=TRUE,
 stringsAsFactors=FALSE)
print(tb1)
```{r}
#model fittin as per 3.1
model2 < -lm(Y \sim X2 + X7 + X8, data = tb1)
summary(model2)
#### Part a: QQ plot of residuals
```{r}
#probability plot
stdres2 = rstandard(model2)
qqnorm(stdres2,
 ylab="Standardized Residuals",
 xlab="Normal Scores",
 main="Residual distribution plot")
qqline(stdres2)
From the above graph we can conclude that there is a slight issue with the normality of
the residuals.
Part b: Residual vs Prediction
```{r}
plot(model2, which = c(1,1))
This plot looks balanced.
#### Part c: Residuals vs Regressor plots
```{r}
install.packages("ggplot2")
```{r}
library(ggplot2)
```{r}
res2 <- resid(model2)</pre>
```{r}
ggplot(tb1, aes(x = X2, y = res2)) +
  geom_point() +
  geom_hline(yintercept = 0, linetype = "dashed") +
 xlab("Predictor") +
ylab("Residuals") +
 ggtitle("Residuals vs. X2 Plot")
. . .
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```{r}
ggplot(tb1, aes(x = X7, y = res2)) +
 geom point() +
 geom hline(yintercept = 0, linetype = "dashed") +
 xlab("Predictor") +
 ylab("Residuals") +
ggtitle("Residuals vs. X7 Plot")
```{r}
ggplot(tb1, aes(x = X8, y = res2)) +
  geom point() +
  geom hline(yintercept = 0, linetype = "dashed") +
 xlab("Predictor") +
 ylab("Residuals") +
ggtitle("Residuals vs. X8 Plot")
While the plot for X8 shows somewhat constant variance, the plot for X7 shows non-
constant variance and the plot for X2 is partially constant.
#### Part d: Partial regressor plots
```{r}
library(car)
```{r}
#create partial residual plots
crPlots(model2)
#### Part e: Studentised Residuals
```{r}
library(MASS)
```{r}
stud resids2 <- studres(model2)</pre>
print(stud_resids2)
These can be used fo find outliers in the model.
## Problem 4.5
```{r}
#Dataset from 3.7
tb4 <- read.csv('Table_B4.csv',header=TRUE,
 stringsAsFactors=FALSE)
print(tb4)
```{r}
#fitting model
model5 = lm(X \sim X1 + X2 + X3 + X4 + X5 + X6 + X7 + X8 + X9, data = tb4)
summary(model5)
#### Part a: QQ plot of residuals
```{r}
plot(model5, which = c(1,2))
Based on the plot, the assumption of normality of residuals seems valid without any
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problems.

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Part b: Residual vs Predicted values
```{r}
plot(model5, which = c(1,1))
While the variance is fairly uniform, there is a slight upward shift in the graph.
#### part c: Partial regression plots
```{r}
library(gridExtra)
```{r}
plot5 = crPlots(model5)
From these graphs we can see that pther than X1 rest deviate from the values and are
thus not necessary.
#### Part d: studentised and R-studentised residuals
```{r}
Compute the studentized residuals
studentized_res5 <- rstandard(model5)</pre>
Compute the R-student residuals
rstudent res5 <- rstudent(model5)</pre>
```{r}
print(studentized_res5)
```{r}
print(rstudent_res5)
Problem 4.9
```{r}
#inputting dataset
tb9 <- read.csv('table_2_ozone.csv',header=TRUE,</pre>
                      stringsAsFactors=FALSE)
print(tb9)
```{r}
#Fitting model
model9 = lm(Days ~ Index, data = tb9)
summary(model9)
Part a: QQ plot for Residuals
```{r}
plot(model9, which = c(1,2))
The normal assumptions of residuals seem to be valid.
#### Part b: Residuals vs Predictions
```{r}
plot(model9, which = c(1,1))
```

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There is a clear pattern that is being captured in this plot
Part c: Residual vs time plot
```{r}
res9 = resid(model9)
time_order <- 1:nrow(tb9)</pre>
plot(time_order, res9,
     ylab="Residuals", xlab="Time order",
    main="Residual vs Time plot")
The graph shows a positive autocorrelation.
## Problem 4.16
```{r}
#inputting dataset
tb16 <- read.csv('table_B8.csv',header=TRUE,</pre>
 stringsAsFactors=FALSE)
print(tb16)
```{r}
#building model
model16 = lm(y \sim x1 + x2, data = tb16)
summary(model16)
#### Part a: QQ plot for Residuals
```{r}
stdres16 = rstandard(model16)
qqnorm(stdres16,
 ylab="Standardized Residuals",
 xlab="Normal Scores",
 main="Residual distribution plot")
qqline(stdres16)
The plot is abnormal near the tails
Part b: Residual vs Predictions plot
```{r}
plot(model16, which = c(1,1))
The fit seems to be pretty good.
#### Part c: PRESS values
```{r}
model16_2 = lm(y \sim x2, data = tb16)
summary(model16_2)
```{r}
#calculating residula
r16_2 <- resid(model16_2)
r16 = resid(model16)
. . .
```{r}
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. . .

```
pr16 <- (resid(model16)/(1 - lm.influence(model16)$hat))^2</pre>
print(sum(pr16))
```{r}
pr16 2 \leftarrow (resid(model16 2)/(1 - lm.influence(model16 2)$hat))^2
print(sum(pr16_2))
Based on the press values of both values, the model with both x1 and x2 will work
better.
## Problem 4.20
```{r}
#importing data
tb20 <- read.csv('table_4_20.csv',header=TRUE,
 stringsAsFactors=FALSE)
print(tb20)
```{r}
#fitting the model
model20 = lm(y ~ Acid.Temp. + Acid.Conc. + WaterTemp. + Sulfide.Conc., data = tb20)
summary(model20)
#### Part a: Model adequecy test
```{r}
#QQ plot
plot(model20, which = c(1,2))
Clearly, there is a problem with the normal assumption
```{r}
res20 = resid(model20)
# produce residual vs. fitted plot
plot(fitted(model20), res20)
# add a horizontal line at 0
abline(0,0)
We also note that there is non constant variance.
```{r}
plot20 = crPlots(model20)
Part b: Lack of fit test
There is no test for lack of fit since there are no replicate points. It is possible to
use the near-neighbor approach.
Problem 4.21
Please refer the pdf file with the name "Problem4_21"
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```
PROBLEM 4.21
 To find:
 E[MSpe]=?
 E [MSLOF] = ?
 IE [MSpe] = I IE [M. R. (Yi) - Yi)2 where
 Working:-
 = \sum_{i=1}^{m} \sum_{j=1}^{2} \left\{ E(y_{ij}^{2}) - 2E(y_{ij}^{2}, y_{ij}^{2}) + E(y_{ij}^{2}) \right\}
 = n e^{2} + m e^{2} - 2 = (2 = \frac{n_{i}}{2} (2 = \frac{y_{i} y_{i}}{n_{i}})
 n_{6}^{2} + m_{6}^{2} - 2 \frac{m}{l=1} \frac{n_{16}^{2}}{n_{1}^{2}} = n_{6}^{2} + m_{6}^{2} - 2m_{6}^{2}
** [SSLOF] = [SSPE] = (N-2) = 2+ = [E(yi) - B - ANI)]2
 = (m-2)e^{2} + \sum_{i=1}^{m} \left[\mathbb{E}[y_{i}] - \beta_{0} - \beta_{i} \chi_{i} \right]^{2}
\mathbb{E}\left[MS_{LOF}\right] = \mathbb{E}\left[SS_{LOF}\right] = \mathbb{E}\left[MS_{LOF}\right] = S^{-2} + \mathbb{E}\left[E(y_1) - \beta_0 - \beta_1 \mathcal{H}\right]^2
```