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Question 1

plot_residuals <- function(model) {</pre>

```
Part 1
```

a)

```
par(mfrow=c(2, 2))
    plot(model)
    par(mfrow=c(1, 1))
raw_model <- lm(mpg~disp)</pre>
raw_model_summary <- summary(raw_model)</pre>
log_model <- lm(log.mpg~log.disp)</pre>
log_model_summary <- summary(log_model)</pre>
plot_residuals(raw_model)
plot_residuals(log_model)
The raw model is better from an R^2 perspective, as well as a model
assumptions perspective. The log model suffers from significant
shape in the Residuals vs Fitted plot which indicates a
violation of the linearity assumption.
The log model also significantly deviates from the normal QQ line,
violating the normality assumption.
Part 2
b)
log_model_summary$fstatistic
# H_0: B_{\log.disp} = 0
# H_1: B_{log.disp} != 0
# F statistic 31.64008
# Null distribution: F(1, 27)
\# p\text{-value}: 5.715 * 10^{-6}
With a 1% level of significance, we can reject the null hypothesis.
This means that the model with log.disp is better than the
intercept only model.
c)
log_model_summary$coefficients["log.disp", "Estimate"]
# -0.241509
```

```
d)
confint(log_model, level=0.99)
# (-0.3604691, -0.12225489)
e)
predict(log_model, data.frame(log.disp=log(240)), interval="confidence", level=0.99)
# (2.769225, 3.017715)
Part 3
a)
carsdat$dummy \leftarrow c(rep(0, 5), 1, rep(0, 23))
part_three_model <- lm(log.mpg~log.disp+dummy, data=carsdat)</pre>
# a)
summary(part_three_model)
b)
With a t statistic of 0.972, a t distribution of t(26) and
a p-value of 0.34, we cannot reject the null hypothesis, as
0.34 > 0.05.
Question 2
Part 1
a)
# 1 predictor: cyl
# 2 predictors: wt + cyl
# 3 predictors: wt + cyl + disp
b)
subset_summary$adjr2
# R^2: 0.5996692, 0.6091114, 0.6976197
PRESS <- function(model) {</pre>
    p_res <- residuals(model) / (1 - hatvalues(model))</pre>
    return(sum(p_res^2))
}
PRESS(lm(mpg~cyl))
PRESS(lm(mpg~wt+cyl))
PRESS(lm(mpg~wt+cyl+disp))
The best model is the 3 predictor model, with the highest
adjusted R^2; which indicates that the regression
component explains more of the total variation, and
the lowest PRESS, which indicates the best out of sample
prediction.
```

```
c)
PRESS is important to consider the out-of-sample prediction.
Using residuals to consider model prediction is a poor
test as the model is fit for the very sample. PRESS residuals
allow for the testing of out-of-sample prediction, and thus
can show relative predictive performance.
Part 2
d)
# Hypotheses:
# H_0 : wt = cyl = disp = hp = 0
# H_1 : Not all betas are 0
# F-statistic: 20.51267
# Null distribution: F(4, 27)
\# p\text{-value}: 7.294 * 10^{-8}
With a p-value < 0.05, we can reject the null hypothesis,
and conclude that the full model is better than the
intercept only model.
e)
e_test <- anova(lm(mpg~wt), full_model)</pre>
# Hypotheses:
# H_0: cyl=disp=hp=0
# H_1: Not all betas are 0
# F statistic: 9.6683
# Null distribution: F(3, 27)
\# p\text{-value} = 0.0001668
With a 5% level of significance, we have enough evidence
to reject the null hypothesis. This means that the additional
predictors do have a statistically significant positive
influence on the model.
f)
```

predict(full_model, data.frame(wt=3.6, cyl=6, disp=220, hp=160),

interval="prediction", level=0.98)

(9.123288, 26.53452)

QUESTION 3

$$5_{w}(B) = \sum_{i=1}^{n} v_{i}(y_{i} - Bx_{i})^{2}$$

$$\frac{\partial S}{\partial B} = -2 \sum_{i=1}^{N} r(x_i) (y_i - Bx_i)$$

$$=\sum_{i=1}^{n} r_i \times i \vee i - \omega_i \sum_{i=1}^{n} r_i \times i = 0.$$

$$p^{n} = \sum_{i=1}^{N} u_{i} \times \sum_{i=1}^{N} u_{i$$

$$Vor\left(\frac{\sum_{i=1}^{n} r_i x_i y_i}{\sum_{i=1}^{n} r_i x_i^2}\right) = \frac{1}{\left(\sum_{i=1}^{n} r_i x_i^2\right)^2} \sum_{i=1}^{n} r_i^2 x_i^2 Vor(y_i)$$

$$\begin{array}{l} \varepsilon(x) = \frac{1}{2} \left(\frac{1}{2}$$

e)
$$S(B) = \sum_{i=1}^{N} (J_{N_i} y_i - J_{N_i} B_{N_i})^2 = \sum_{i=1}^{N} v_i (y_i - B_{N_i})^2$$

$$= S_{ave} \quad \text{objective} \quad \text{furction}.$$

.: Consider.

If the variance ishas the relationship Var(b_) = xi262, the transformation seen in model (2) stabilises the variance

QUESTION 4.

PAIRT 1.

b) First, recall trad:

$$D_{i} = \left(\frac{(x + x)^{-1} + i + i}{1 - h_{ii}} \right)^{T} (x + x)^{-1} \left(\frac{(x + x)^{-1} + i + i}{1 - h_{ii}} \right)$$

$$= \left(\frac{(x + x)^{-1} + i + i}{1 - h_{ii}} \right)^{T} (x + x)^{-1} \left(\frac{(x + x)^{-1} + i + i}{1 - h_{ii}} \right)$$

$$= \frac{e_i^2 \times_i^T (\times TX)^T (\times TX)^T (\times TX)^T \times_i^2}{3^2 p(1-h_{ii})^2}$$

$$= \frac{e_i^2 \times_i^T (\times TX)^T \times_i^2}{3^2 p(1-h_{ii})^2}$$

PART 2.

- c) Vi,-i is the value fifted of the i-h observation with parameters littled WITHOUT to i-har observation
 - d) We assume gitte then combreto of the predictor and their betas. Thereby, define xi = (xi, xi2 -- xin)T. We can then define:

where bis is fitted astrond to inter observetor,

$$2) \quad \widehat{v_i} - \widehat{v_{i,-i}} = \lambda_i^T b - \kappa_i^T b - k_i^T b - k_i^T$$

+iT(XTX) x; = Nic, Os H= X(XTX) XT.