



UNIVERSITI
TEKNOLOGI
PETRONAS

LAB WEEK 11

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BACHELOR OF COMPUTER SCIENCE

DATA SCIENCE

TEB2164

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Code

Activity 1

```
#Load Dataset
data("ToothGrowth")

#Convert Categorical to Numerical
df <- ToothGrowth
#Assign Auto
df$supp <- as.numeric(df$supp) #VC vitamin C, OJ orange juice

#Assign Manually
df$supp <- ifelse(ToothGrowth == "OJ", 1, 0)

#Calc Corr Analysis
corr_mat <- round(cor(df), 2)

#Reshape
library(reshape2)

melted_corr <- melt(corr_mat)

#Plot
library(ggplot2)
ggplot(melted_corr, aes(x = Var1, y = Var2, fill = value)) +
  geom_tile(color = "white") +
  geom_text(aes(label = value), color = "white", size = 5) +
  scale_fill_gradient2(low = "blue", high = "red", mid = "white", midpoint = 0) +
  theme_minimal()
```

Activity 2

```
#Load Dataset
```

```
data(mtcars)
```

```
df <- mtcars
```

```
#Lib
```

```
library(caret)
```

```
#Log Transformation
```

```
df_log <- log(df)
```

```
#Standard Scaling
```

```
df_standard <- as.data.frame(scale(df))
```

```
#Min-Max Scaling
```

```
minmax <- preProcess(as.data.frame(df), method = c("range"))
```

```
df_minmax <- predict(minmax, as.data.frame(df))
```

```
#Summary
```

```
summary(df)
```

```
summary(df_log)
```

```
summary(df_standard)
```

```
summary(df_minmax)
```

```
#Plot
```

```
par(mfrow = c(2, 2)) #grid 2 by 2 for side by side
```

```
boxplot(df, main = "Original Data")
```

```
boxplot(df_log, main = "Log Transformation")
```

```
boxplot(df_standard, main = "Standard Scaling")
```

```
boxplot(df_minmax, main = "Min-Max Scaling")
```

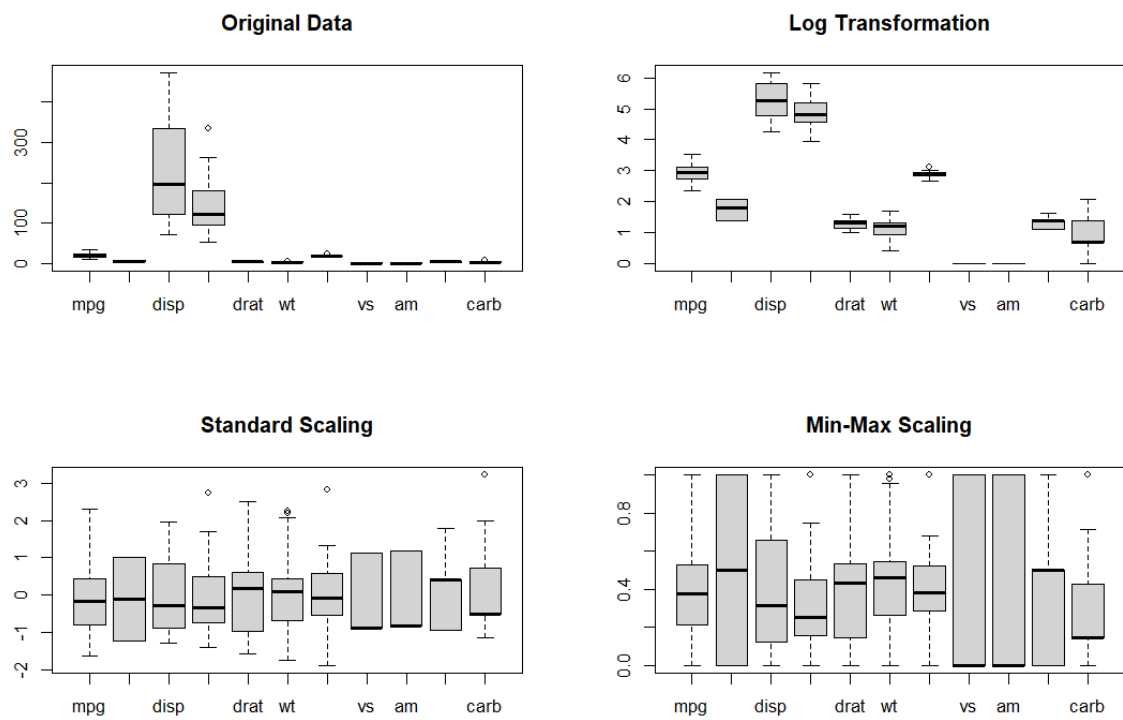
Visualisation

Activity 1



There's a very strong correlation between dose and len which account to 0.8. While dose and supp have a 0 and len with supp have a negative correlatio of -0.24.

Activity 2



Log transformation compresses the high values, standard scaling centres data around a mean of 0 and standard deviation of 1, and min-max scaling bounds values in a range of [0, 1]

