# Linear models vignette

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# 1 Purpose

This vignette aims to introduce you liner models using R.

## 2 Reference

 $Dr.\ Bharatendra\ https://www.youtube.com/watch?v=rsfV57N7Uns\&list=PL34t5iLfZddtUUABMikey6NtL05hPAp42\&index=7$ 

## 3 Libraries

## 4 Load some data

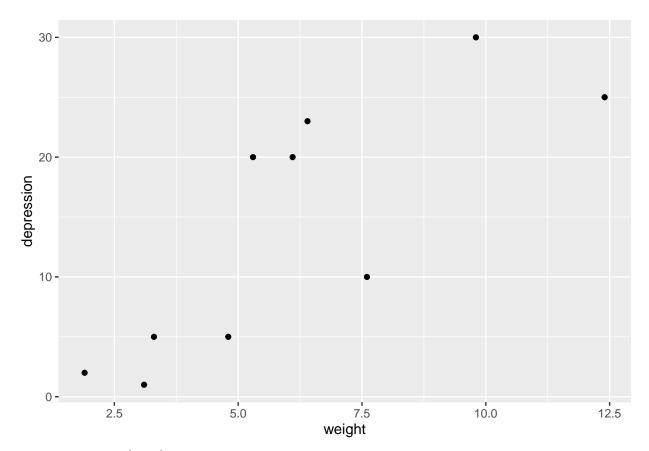
Look at 'roller' dataset from DAAG library.

```
# Example the roller dataset
str(roller)

## 'data.frame': 10 obs. of 2 variables:
## $ weight : num 1.9 3.1 3.3 4.8 5.3 6.1 6.4 7.6 9.8 12.4
## $ depression: num 2 1 5 5 20 20 23 10 30 25
```

# 5 Plot it (scatter plot)

```
ggplot(roller, aes(x=weight, y=depression)) +
  geom_point()
```



# Linear model lm{stats}

Reference: Dr. Bharatendra https://youtube.com/watch?v=utjaosw7wi0&si=EnSIkaIECMiOmarE

```
# lm model, with formula y \sim x..., and dataset name
# here dependent var will be depression, inde variable weight from dataset roller
model <- lm(depression ~ weight, roller)</pre>
# Get the summary
summary(model)
##
## Call:
## lm(formula = depression ~ weight, data = roller)
##
## Residuals:
     Min
              1Q Median
                            3Q
                                  Max
## -8.180 -5.580 -1.346 5.920 8.020
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -2.0871
                            4.7543 -0.439 0.67227
## weight
                 2.6667
                            0.7002
                                   3.808 0.00518 **
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## Residual standard error: 6.735 on 8 degrees of freedom
## Multiple R-squared: 0.6445, Adjusted R-squared: 0.6001
## F-statistic: 14.5 on 1 and 8 DF, p-value: 0.005175
```

### 5.1 Explore the model

#### 5.1.1 What the model has

```
names(model)
```

```
## [1] "coefficients" "residuals" "effects" "rank"

## [5] "fitted.values" "assign" "qr" "df.residual"

## [9] "xlevels" "call" "terms" "model"
```

### 5.1.2 Look at coefficients only

```
coef(model)
```

```
## (Intercept) weight
## -2.087148 2.666746
```

#### 5.1.3 look at the residuals

Residuals = Observed - Predicted

```
residuals(model)
```

```
## 1 2 3 4 5 6 7
## -0.9796695 -5.1797646 -1.7131138 -5.7132327 7.9533944 5.8199976 8.0199738
## 8 9 10
## -8.1801213 5.9530377 -5.9805017
```

### 5.1.4 Look at predictions (i.e. the fitted model)

In stats terms we say the 'fitted model' while in machine learning we say 'prediciton', same thing.

#### 5.1.5 What that means

## 24.046962 30.980502

Look at the prediction (fit) and calculate the residual by hand. Compare.

```
# Residual from the first prediction
r1 <- residuals(model)[1]

# Actual number
a1 <- roller[1,]

# This result must be zero (or super close to zero)
residual_test = (a1[2] - f[1]) - r1</pre>
```

# 6 Linear Model in scatter plot

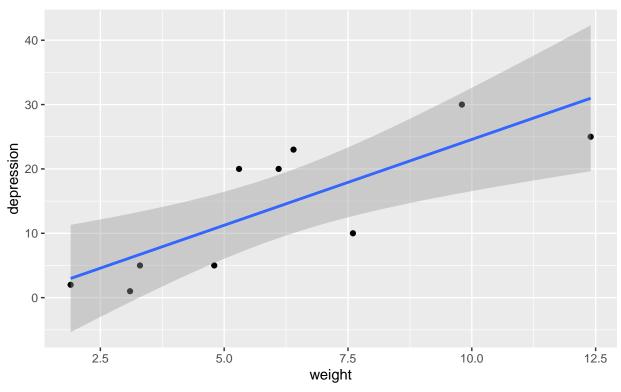
Shows the 95% confidence intervals.

```
ggplot(roller, aes(x=weight, y=depression)) +
  geom_point() +
  geom_smooth(method = 'lm') +
  ggtitle('Depression vs Weight', 'Source: Roller data from DAAG')
```

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

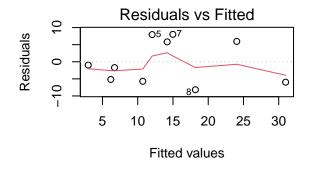
# Depression vs Weight

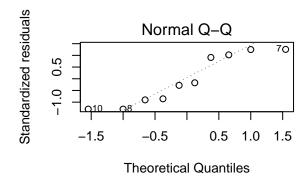
Source: Roller data from DAAG

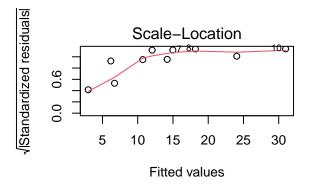


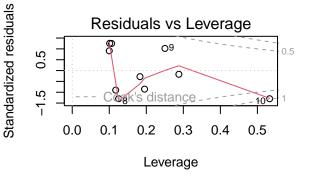
# 7 Diagnostics

```
par(mfrow=c(2, 2))
plot(model)
```









## 8 Predictions

## 8.1 One prediction

```
# Here the independent variable is 'weight'
predict(model, data.frame(weight=7))
## 1
```

### 8.2 Many predictions

## 16.58007

```
predict(model, data.frame(weight=c(7, 8, 9)))
## 1 2 3
## 16.58007 19.24682 21.91357
```

#### 8.2.1 Confidence intervals

#### 8.2.2 Prediction intervals

```
## fit lwr upr
## 1 16.58007 0.2208492 32.93930
## 2 19.24682 2.6612368 35.83240
## 3 21.91357 4.9502581 38.87687
```

Some time confidence intervals are called narrow intervals, while prediction intervals are called wider intervals.

You use it depending on the context.

If the context is about a single value then use prediciton interval, and if the context is about a wider average then use confidence interval.

### 8.3 Plots for confidence and prediction intervals

#### 8.3.1 Plot for prediction interval

```
# First make a dataset

p <- predict(model, interval = 'prediction')</pre>
```

### 8.3.1.1 make a data.frame

## Warning in predict.lm(model, interval = "prediction"): predictions on current data refer to \_future\_

```
# combine data
data <- cbind(roller, p)
str(data)</pre>
```

```
## 'data.frame': 10 obs. of 5 variables:
## $ weight : num 1.9 3.1 3.3 4.8 5.3 6.1 6.4 7.6 9.8 12.4
## $ depression: num 2 1 5 5 20 20 23 10 30 25
## $ fit : num 2.98 6.18 6.71 10.71 12.05 ...
## $ lwr : num -14.65 -10.8 -10.18 -5.71 -4.29 ...
## $ upr : num 20.6 23.2 23.6 27.1 28.4 ...
```

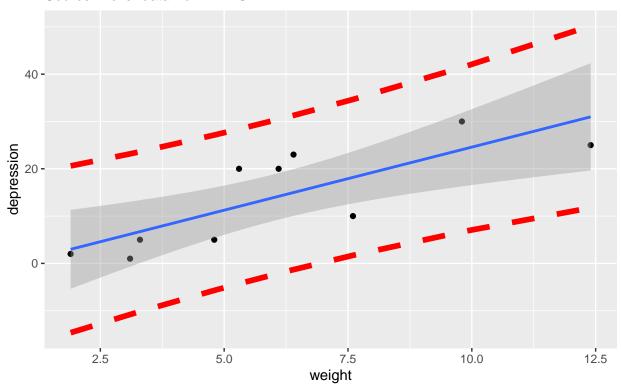
```
# Add the prediction intervals with geom_line()
ggplot(data, aes(x=weight, y=depression)) +
  geom_point() +
  geom_smooth(method = 'lm') +
  ggtitle('Depression vs Weight', 'Source: Roller data from DAAG') +
  geom_line(aes(y=lwr), color='red', linetype='dashed', lwd=2) +
  geom_line(aes(y=upr), color='red', linetype='dashed', lwd=2)
```

### 8.3.1.2 Plot it

## 'geom\_smooth()' using formula = 'y ~ x'

### Depression vs Weight

Source: Roller data from DAAG



# 9 Quadratic term

A way to help the model

```
model1 <- lm(depression ~ weight + I(weight^2), roller)
summary(model1)</pre>
```

```
##
## Call:
## lm(formula = depression ~ weight + I(weight^2), data = roller)
## Residuals:
      Min
               1Q Median
                              ЗQ
                                    Max
## -10.699 -3.192
                  1.244
                          4.792
                                   6.163
##
## Coefficients:
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) -12.1247
                       9.3821 -1.292
                                          0.2373
## weight
                        2.9822 2.090 0.0749 .
          6.2337
## I(weight^2) -0.2519 0.2051 -1.228 0.2590
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
## Residual standard error: 6.531 on 7 degrees of freedom
## Multiple R-squared: 0.7075, Adjusted R-squared: 0.624
## F-statistic: 8.467 on 2 and 7 DF, p-value: 0.01353
```