## HW4 - Simple Linear Regression

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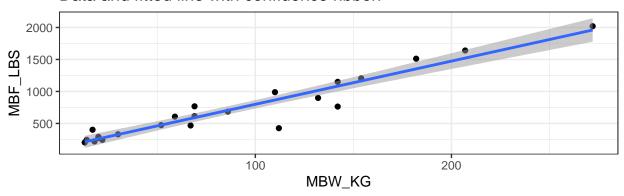
```
B <- read.csv("BodySize BiteForce2.csv",header=TRUE)
head(B)
    N MBW_KG MBF_LBS
## 1 2
       132 898.98
## 2 1
         182 1512.00
## 3 5
         20 290.44
          69 615.05
## 4 1
## 5 1
        110 988.90
## 6 2
        86 684.07
tail(B)
     N MBW KG MBF LBS
           20 273.13
## 18 4
## 19 5
           30 329.78
## 20 5
          18 218.28
## 21 3
           59 606.06
## 22 3
           13 202.32
## 23 3
           22 243.23
lm1 <- lm(MBF_LBS~MBW_KG,data=B)</pre>
summary(lm1)
##
## Call:
## lm(formula = MBF_LBS ~ MBW_KG, data = B)
## Residuals:
               10 Median
                               30
      Min
                                      Max
## -454.32 -24.25
                   24.69 75.46 177.54
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
##
                        49.914
## (Intercept) 124.518
                                   2.495
                                             0.021 *
## MBW KG
                 6.748
                            0.447 15.098 9.49e-13 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 149.1 on 21 degrees of freedom
## Multiple R-squared: 0.9156, Adjusted R-squared: 0.9116
## F-statistic: 227.9 on 1 and 21 DF, p-value: 9.49e-13
p1 <- ggplot(B, aes(x = MBW_KG, y = MBF_LBS)) +
     geom_point() +
     geom_smooth(method = "lm", alpha = .5)+
     ggtitle("Data and fitted line with confidence ribbon")+
     theme_bw()
```

```
p2 <- ggplot(B, aes(x = MBW_KG, y = MBF_LBS) ) +
    geom_point() +
    geom_smooth(method = "lm", se = FALSE)+
    ggtitle("Data and fitted line")+
    theme_bw()</pre>
```

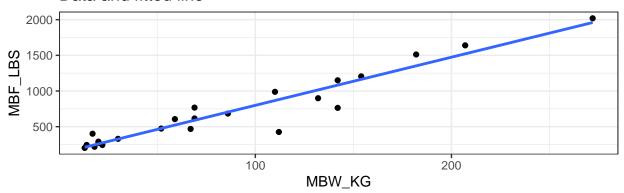
```
grid.arrange(p1,p2,nrow=2)
```

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

## Data and fitted line with confidence ribbon



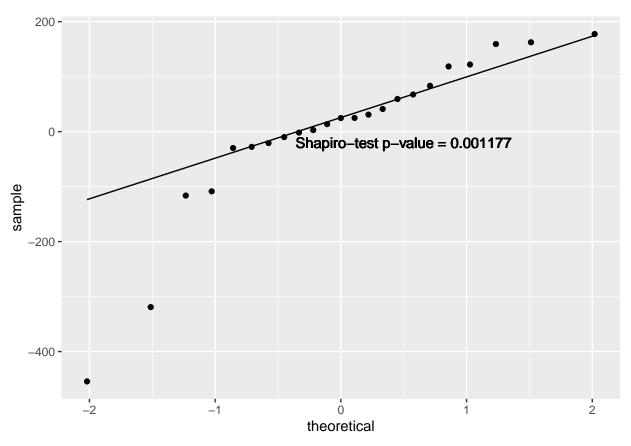
## Data and fitted line



```
#qq plot with normal line (normality test)
df <- as.data.frame(lm1$residuals)
colnames(df)[1] <- "residuals"
shapiro.test(df$residuals)</pre>
```

```
##
## Shapiro-Wilk normality test
##
## data: df$residuals
## W = 0.82939, p-value = 0.001177

ggplot(df)+stat_qq(aes(sample=residuals)) +
    geom_qq_line(aes(sample=residuals))+
    geom_text(aes(x=0.5, y=-20, label="Shapiro-test p-value = 0.001177"))
```



There is no way for us to know if this Covariance is large or small. We need a better measure for the strength of the relationship.

cov(B\$MBW\_KG,B\$MBF\_LBS)

## [1] 34152.74