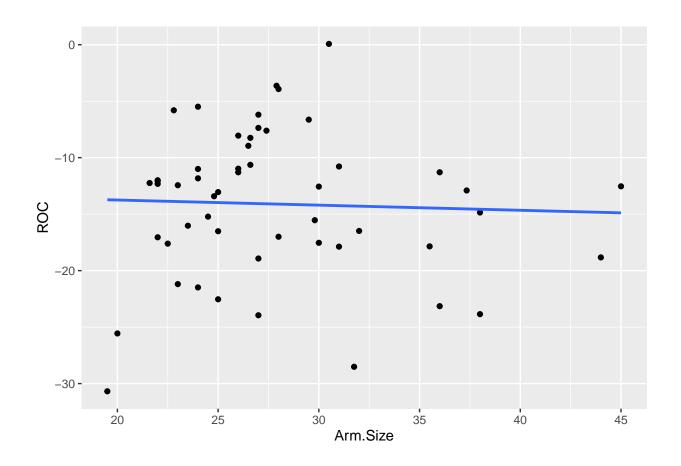
prelim

Hewlett Madrid

2023-04-20

```
library(tidyverse)
library(ggpubr)
library(broom)
library(car)
library(rstatix)
```

```
roc <- read.csv("C:/Users/Xyrine/Documents/School Stuff/BS BIO 4th Year/1st Semester/BIO 118/Module 1/M
colnames(roc) <- c("Arm.Size", "ROC")
ggplot(roc, aes(Arm.Size, ROC)) + geom_point() +
   geom_smooth(method = "lm", se = FALSE)</pre>
```

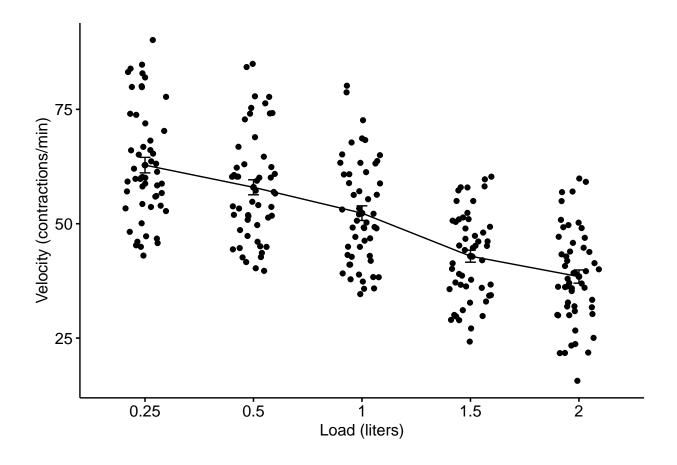


```
roc.lm <- lm(ROC ~ Arm.Size, roc)</pre>
summary(roc.lm)
##
## Call:
## lm(formula = ROC ~ Arm.Size, data = roc)
##
## Residuals:
      Min
##
               1Q Median
                              3Q
                                     Max
## -16.956 -3.458
                  1.485
                          3.428 14.299
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## Arm.Size
             -0.04541
                          0.16037 -0.283 0.77824
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 6.549 on 50 degrees of freedom
## Multiple R-squared: 0.001601, Adjusted R-squared: -0.01837
## F-statistic: 0.08017 on 1 and 50 DF, p-value: 0.7782
cor.test(roc$Arm.Size, roc$ROC)
##
## Pearson's product-moment correlation
##
## data: roc$Arm.Size and roc$ROC
## t = -0.28314, df = 50, p-value = 0.7782
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## -0.3095303 0.2354616
## sample estimates:
##
          cor
## -0.04000959
# imports pooled data into R
msc <- read.csv("C:/Users/Xyrine/Documents/School Stuff/BS BIO 4th Year/1st Semester/BIO 118/Module 1/M
# isolates columns for arm size and lengths, and velocity
msc.cont <- msc %>%
 select(Upper.Arm.Size, Upper.Arm.Length, Forearm.Length,
        X0.25, X0.5, X1, X1.5, X2) %>%
 drop na() %>%
 gather(Load, Velocity, X0.25, X0.5, X1, X1.5, X2)
```

1. Load and Velocity

1.a. Data summary and visualization

```
# prints summary of data
msc.cont %>%
 group_by(Load) %>%
 summarise(
   Mean.Velocity = mean(Velocity),
   SD.Velocity = sd(Velocity)
)
## # A tibble: 5 x 3
## Load Mean. Velocity SD. Velocity
## <chr> <dbl> <dbl>
## 1 X0.25
                62.8
                           12.2
                58.0
## 2 X0.5
                          11.8
                          11.5
## 3 X1
                52.3
## 4 X1.5
                42.9
                           9.53
## 5 X2
                 38.4
                           10.4
# visualizes data
ggline(msc.cont, x = "Load", y = "Velocity",
      add = c("mean_se", "jitter"),
      order = c("X0.25", "X0.5", "X1", "X1.5", "X2"),
      ylab = "Velocity (contractions/min)", xlab = "Load (liters)") +
  scale_x_discrete(labels=c("0.25","0.5", "1", "1.5", "2"))
```

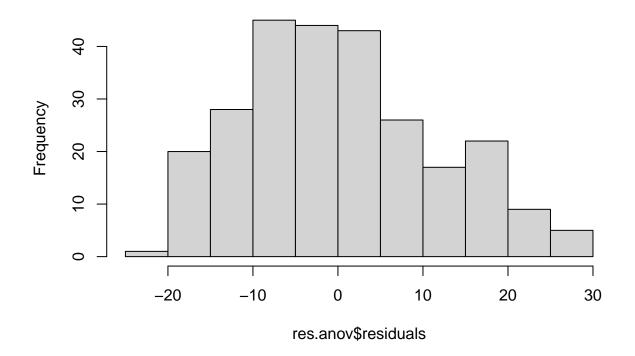


1.b. ANOVA

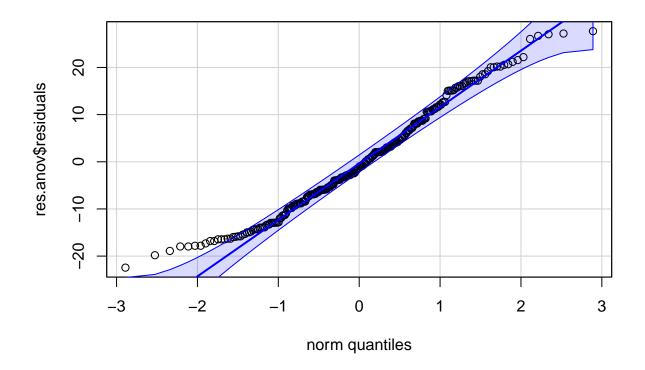
```
# Assumption: homogeneity of variances
leveneTest(Velocity ~ Load, data = msc.cont)
## Levene's Test for Homogeneity of Variance (center = median)
         Df F value Pr(>F)
## group
           4 0.6653 0.6166
##
         255
## Accept null hypothesis (p > 0.05)
## Not enough evidence suggests that differences in variances between groups
### are statistically significant
## Assume homoscedasticity
# Assumption: normality
shapiro.test(msc.cont$Velocity[msc.cont$Load == "X0.25"]) # (p < 0.05)
##
    Shapiro-Wilk normality test
##
```

```
## data: msc.cont$Velocity[msc.cont$Load == "X0.25"]
## W = 0.95087, p-value = 0.03171
shapiro.test(msc.cont$Velocity[msc.cont$Load == "X0.5"]) # (p < 0.05)
##
## Shapiro-Wilk normality test
##
## data: msc.cont$Velocity[msc.cont$Load == "X0.5"]
## W = 0.95079, p-value = 0.03147
shapiro.test(msc.cont$Velocity[msc.cont$Load == "X1"]) # (p > 0.05)
## Shapiro-Wilk normality test
##
## data: msc.cont$Velocity[msc.cont$Load == "X1"]
## W = 0.95969, p-value = 0.07586
shapiro.test(msc.cont$Velocity[msc.cont$Load == "X1.5"]) # (p > 0.05)
##
## Shapiro-Wilk normality test
##
## data: msc.cont$Velocity[msc.cont$Load == "X1.5"]
## W = 0.97211, p-value = 0.2591
shapiro.test(msc.cont$Velocity[msc.cont$Load == "X2"]) # (p > 0.05)
##
## Shapiro-Wilk normality test
## data: msc.cont$Velocity[msc.cont$Load == "X2"]
## W = 0.98488, p-value = 0.7461
res.anov <- aov(Velocity ~ Load, msc.cont)
## histogram
hist(res.anov$residuals)
```

Histogram of res.anov\$residuals



```
## qqplot
qqPlot(res.anov$residuals,
  id = FALSE # id = FALSE to remove point identification
)
```



```
## histogram shows roughly a bell curve, and points in the QQ plot nearly adhere to
### a streight line and are well within the confidence bands

## We can assume normality

# ANOVA
anov <- aov(Velocity ~ Load, msc.cont)
summary(anov)</pre>
```