ANOVA notes

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```
library(tidyverse)
df_pigs <- read_table("pig_weight.txt")</pre>
df_pigs$Drug <- as.factor(df_pigs$Drug)</pre>
pig_model1 <- lm(Pigweight ~ ., data = df_pigs)</pre>
summary(pig_model1)
##
## lm(formula = Pigweight ~ ., data = df_pigs)
##
## Residuals:
     Min
             1Q Median
                            3Q
                                 Max
## -3.905 -1.174 0.187 1.351 3.657
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 7.48163
                          9.14917 0.818 0.41628
## Drug2
              -1.60557
                          0.52788 -3.042 0.00331 **
                          0.52871 -1.333 0.18684
## Drug3
              -0.70480
## Momweight
             0.26363
                          0.04727
                                     5.578 4.28e-07 ***
## Dadweight
             0.17442
                           0.03465
                                   5.034 3.58e-06 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
## Residual standard error: 1.855 on 70 degrees of freedom
## Multiple R-squared: 0.4561, Adjusted R-squared: 0.425
## F-statistic: 14.67 on 4 and 70 DF, p-value: 9.393e-09
```

Null Model

```
null_model <- lm(Pigweight ~ NULL, data = df_pigs)</pre>
```

No Drug Model

```
nodrug_model <- lm(Pigweight ~ Momweight + Dadweight, data = df_pigs)
summary(nodrug_model)</pre>
```

```
##
## Call:
## lm(formula = Pigweight ~ Momweight + Dadweight, data = df_pigs)
## Residuals:
      Min
               1Q Median
                              3Q
## -4.4473 -1.2429 0.0714 1.2118 4.3556
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 9.48455 9.40944
                                  1.008
                                            0.317
## Momweight 0.25972
                         0.04951 5.246 1.50e-06 ***
## Dadweight 0.16183
                         0.03540 4.571 1.96e-05 ***
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## Residual standard error: 1.946 on 72 degrees of freedom
## Multiple R-squared: 0.3841, Adjusted R-squared: 0.3669
## F-statistic: 22.45 on 2 and 72 DF, p-value: 2.651e-08
```

ANOVA

```
anova(pig_model1)
## Analysis of Variance Table
## Response: Pigweight
            Df Sum Sq Mean Sq F value
## Drug
            2 23.718 11.859 3.4473 0.03733 *
## Momweight 1 91.039 91.039 26.4638 2.342e-06 ***
## Dadweight 1 87.175 87.175 25.3407 3.580e-06 ***
## Residuals 70 240.808
                       3.440
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
anova(null_model, pig_model1)
## Analysis of Variance Table
## Model 1: Pigweight ~ NULL
## Model 2: Pigweight ~ Drug + Momweight + Dadweight
## Res.Df
             RSS Df Sum of Sq F Pr(>F)
## 1
        74 442.74
## 2
        70 240.81 4 201.93 14.675 9.393e-09 ***
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
anova(nodrug_model, pig_model1)
## Analysis of Variance Table
## Model 1: Pigweight ~ Momweight + Dadweight
## Model 2: Pigweight ~ Drug + Momweight + Dadweight
## Res.Df RSS Df Sum of Sq
                               F Pr(>F)
## 1
        72 272.70
## 2
        70 240.81 2
                     31.894 4.6356 0.01286 *
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
```