# RADPAD Mixed Model

#### Mixed Model

I used a log transform on the response to improve the model fit (by inspection of residual plots). The random effect is procedure ID.

```
m <- lmer(log(Dose + 1) ~ RADPAD*Occupation + (1 | ID), data = radpad)</pre>
```

#### ANOVA for Fixed Effects

```
anova(m)
```

```
## Type III Analysis of Variance Table with Satterthwaite's method
## Sum Sq Mean Sq NumDF DenDF F value Pr(>F)

## RADPAD 34.34 34.340 1 203.49 55.213 2.932e-12 ***

## Occupation 403.91 80.783 5 867.39 129.886 < 2.2e-16 ***

## RADPAD:Occupation 50.45 10.090 5 867.39 16.224 2.565e-15 ***

## ---

## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

## LS Means for RADPAD by Occupation

LS means have been back-transformed from log scale.

```
emmeans(m, ~ RADPAD | Occupation, type = "response")
```

```
## Occupation = Anesthesia:
  RADPAD response
                       SE
                            df lower.CL upper.CL
             4.2981 0.3851 892
                                            5.110
##
                                  3.5937
                                            0.410
##
            0.1002 0.1391 884 -0.1416
##
## Occupation = Faculty:
   RADPAD response
                        SE
                            df lower.CL upper.CL
##
             1.9012 0.2096 887
                                  1.5177
                                            2.343
##
   Y
             0.7126 0.2166
                           884
                                  0.3362
                                            1.195
##
## Occupation = Resident 1:
   RADPAD response
                        SE
                             df lower.CL upper.CL
            15.6726 1.2010 884 13.4744
##
                                           18.205
##
   Y
            5.6412 0.8398
                           884
                                  4.1815
                                            7.512
##
## Occupation = Resident 2:
## RADPAD response
                        SE
                             df lower.CL upper.CL
```

```
1.8890 0.2234 938
                                  1.4821
                                            2.362
##
             0.9189 0.2449 892
                                  0.4937
                                            1.465
##
## Occupation = Tech 1:
## RADPAD response
                        SE
                             df lower.CL upper.CL
             0.1181 0.0828 906
                                -0.0332
##
             0.0589 0.1339
                           884 -0.1738
                                            0.357
##
## Occupation = TEE:
  RADPAD response
                        SE
                             df lower.CL upper.CL
             2.5156 0.3708 1045
                                  1.8583
                                            3.324
                                  0.7105
                                            2.180
## Y
             1.3321 0.3684 1017
## Degrees-of-freedom method: kenward-roger
## Confidence level used: 0.95
## Intervals are back-transformed from the log(mu + 1) scale
```

### Pairwise contrasts for RADPAD by Occupation

Contrasts have been back-transformed from log scale.

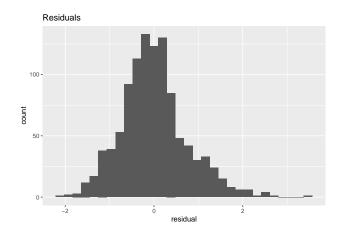
rg <- ref\_grid(m)

## Confidence level used: 0.95

```
em <- emmeans(regrid(rg, transform = "response"), ~ RADPAD | Occupation)</pre>
c <- contrast(em, interaction = "pairwise")</pre>
summary(c, infer = c(TRUE, TRUE))
## Occupation = Anesthesia:
## RADPAD pairwise estimate
                                SE
                                     df lower.CL upper.CL t.ratio p.value
## N - Y
                      4.1979 0.409
                                    884
                                           3.394
                                                    5.001 10.253 <.0001
##
## Occupation = Faculty:
## RADPAD_pairwise estimate
                                SE
                                     df lower.CL upper.CL t.ratio p.value
## N - Y
                      1.1886 0.301
                                           0.597
                                                    1.780 3.944 0.0001
                                    884
##
## Occupation = Resident 1:
  RADPAD_pairwise estimate
                                SE
                                     df lower.CL upper.CL t.ratio p.value
##
  N - Y
                     10.0314 1.466
                                           7.155
                                                   12.908
                                                          6.845 <.0001
                                    884
## Occupation = Resident 2:
   RADPAD_pairwise estimate
                                SE
                                     df lower.CL upper.CL t.ratio p.value
                      0.9701 0.332
                                           0.319
##
   N - Y
                                    892
                                                    1.621
                                                            2.926 0.0035
##
## Occupation = Tech 1:
## RADPAD_pairwise estimate
                                SE
                                     df lower.CL upper.CL t.ratio p.value
## N - Y
                      0.0592 0.157
                                    884
                                          -0.250
                                                    0.368
                                                          0.376 0.7072
##
## Occupation = TEE:
## RADPAD_pairwise estimate
                                     df lower.CL upper.CL t.ratio p.value
                                SE
## N - Y
                      1.1834 0.523 1017
                                           0.158
                                                    2.209
                                                            2.264 0.0238
##
## Degrees-of-freedom method: inherited from kenward-roger when re-gridding
```

## Residual Analysis

```
mdat <- data.frame(predicted = predict(m), residual = residuals(m))
ggplot(mdat, aes(x = residual)) + geom_histogram() +
labs(title = "Residuals")</pre>
```



```
ggplot(mdat,aes(sample=residual)) + stat_qq() + stat_qq_line() +
labs(title = "QQ Plot")
```

