WT Validation Cohort

Before CombatSeq

Mucosal Dataset

```
Permutation: free
Number of permutations: 10000
Terms added sequentially (first to last)
              Df SumsOfSqs MeanSqs F.Model R2 Pr(>F)
Sequencing_Run 1
                   0.943 0.943 0.681 0.00338
                                                0.5159
Sex
              1
                  4.211 4.211 3.042 0.01508
                                                 0.0459 *
Site_General 1 92.763 92.763 67.012 0.33218 9.999e-05 ***
Residuals
          131 181.339 1.384
                                       0.64937
Total
            134 279.256
                                       1.00000
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
```

```
Permutation: free
Number of permutations: 10000
Terms added sequentially (first to last)
               Df SumsOfSqs MeanSqs F.Model R2 Pr(>F)
               1 0.943 0.943 0.678 0.00338
Sequencing_Run
                                                  0.5087
                    4.211 4.211 3.029 0.01508
Sex
                1
                                                  0.0470 *
Site_General 1 92.763 92.763 66.731 0.33218 9.999e-05 ***
Sex:Site_General 1
                    0.626 0.626 0.450 0.00224
                                                 0.6451
Residuals 130 180.713 1.390
                                        0.64712
Total
             134 279.256
                                        1.00000
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
```

Mucosal Colon

```
Terms added sequentially (first to last)

Df SumsOfSqs MeanSqs F.Model R2 Pr(>F)
```

```
Sequencing_Run 1 35.075 35.075 35.472 0.25510 9.999e-05 ***
Sex
             1
                  0.387 0.387 0.391 0.00281
                                               0.6833
Site
             2 39.741 19.871 20.095 0.28903 9.999e-05 ***
Residuals
           63 62.296 0.989
                                       0.45306
Total
           67 137.499
                                       1.00000
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Permutation: free
Number of permutations: 10000
Terms added sequentially (first to last)
             Df SumsOfSqs MeanSqs F.Model R2
                                                 Pr(>F)
Sequencing_Run 1
                  35.075 35.075 34.989 0.25510 9.999e-05 ***
                  0.387 0.387 0.386 0.00281
Sex
             1
Site
              2 39.741 19.871 19.822 0.28903 9.999e-05 ***
           2 1.146 0.573 0.572 0.00833 0.6843
Sex:Site
Residuals
           61 61.150 1.002
                                       0.44473
           67 137.499
Total
                                       1.00000
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Mucosal SI

```
Permutation: free
Number of permutations: 10000
Terms added sequentially (first to last)
             Df SumsOfSqs MeanSqs F.Model R2
                                                  Pr(>F)
Sequencing_Run 1 1.545 1.5445 0.8068 0.01091 0.4535546
                   3.093 3.0931 1.6157 0.02185 0.1976802
Sex
             1
Site
              2
                  18.201 9.1004 4.7536 0.12860 0.0008999 ***
            62 118.694 1.9144
Residuals
                                        0.83864
         66 141.533
Total
                                        1.00000
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

After CombatSeq

Mucosal

```
Permutation: free
Number of permutations: 10000
Terms added sequentially (first to last)
              Df SumsOfSqs MeanSqs F.Model R2
                                                 Pr(>F)
Sequencing_Run
               1
                  5.499 5.499 3.970 0.01953 0.0163 *
                    4.435 4.435 3.201 0.01575
Sex
               1
                                                 0.0387 *
Site_General 1 88.816 88.816 64.110 0.31538 9.999e-05 ***
Residuals
          132 182.868 1.385
                                       0.64935
Total
          135 281.618
                                        1.00000
___
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Mucosal Colon

```
Permutation: free
Number of permutations: 10000
Terms added sequentially (first to last)
             Df SumsOfSqs MeanSqs F.Model R2
                                                 Pr(>F)
Sequencing_Run 1 61.042 61.042 79.796 0.44347 9.999e-05 ***
Sex
             1
                  0.065 0.065 0.085 0.00047
                                                 0.9293
Site
             2 28.345 14.172 18.526 0.20592 9.999e-05 ***
Residuals 63 48.194 0.765
                                      0.35013
Total
           67 137.646
                                       1.00000
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Mucosal SI

```
Permutation: free
Number of permutations: 10000
Terms added sequentially (first to last)
             Df SumsOfSqs MeanSqs F.Model R2 Pr(>F)
Sequencing_Run 1
                  1.986 1.9857 1.0291 0.01380 0.35976
Sex
             1
                   4.566 4.5661 2.3663 0.03172 0.09069 .
Site
             2
                  15.813 7.9063 4.0974 0.10986 0.00280 **
Residuals 63 121.564 1.9296
                                        0.84461
Total
                  143.928
                                        1,00000
             67
```

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

Permutation: free
Number of permutations: 10000

Terms added sequentially (first to last)

Df SumsOfSqs MeanSqs F.Model R2 Pr(>F)
Sequencing_Run 1 1.986 1.9857 1.0402 0.01380 0.36236
Sex 1 4.566 4.5661 2.3920 0.03172 0.09289 .
Site 2 15.813 7.9063 4.1418 0.10986 0.00200 **
Sex:Site 2 5.120 2.5600 1.3411 0.03557 0.25347
Residuals 61 116.444 1.9089 0.80904
Total 67 143.928 1.00000

--Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1