CBF Correlations

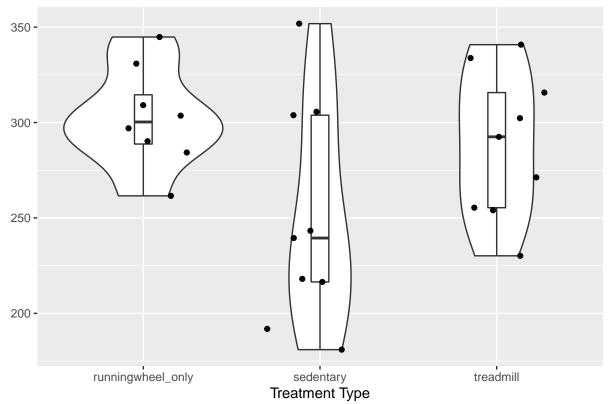
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4/7/2021

Correlations CBF and Exercise Treatment

```
ggplot(data = cbf, aes(factor(Treatment), CBF)) +
  geom_violin() +
  geom_boxplot(width = 0.1, outlier.color = "red") +
  geom_jitter(height = 0, width = 0.3) +
  labs(x = "Treatment Type",
    y = "",
    title = "CBF distrubution by treatment type")
```

CBF distrubution by treatment type



```
res.aov <- aov(CBF ~ Treatment, data = cbf)
summary(res.aov)</pre>
```

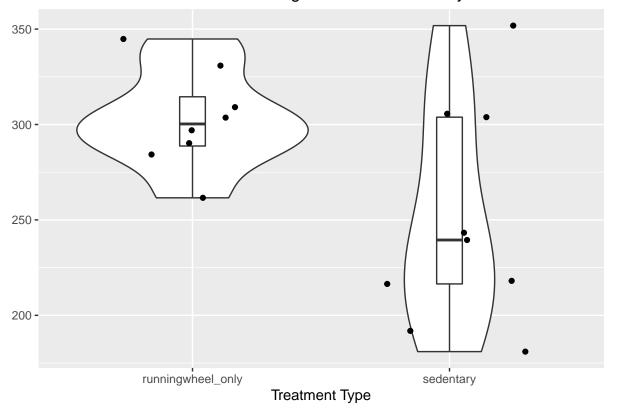
```
## Df Sum Sq Mean Sq F value Pr(>F)
## Treatment 2 12767 6384 3.386 0.0514 .
## Residuals 23 43356 1885
```

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

specific correlations

```
ggplot(data = cbf1, aes(factor(Treatment), CBF)) +
  geom_violin() +
  geom_boxplot(width = 0.1, outlier.color = "red") +
  geom_jitter(height = 0, width = 0.3) +
  labs(x = "Treatment Type",
    y = "",
    title = "CBF distrubution between running wheel and sedentary treatments")
```

CBF distrubution between running wheel and sedentary treatments

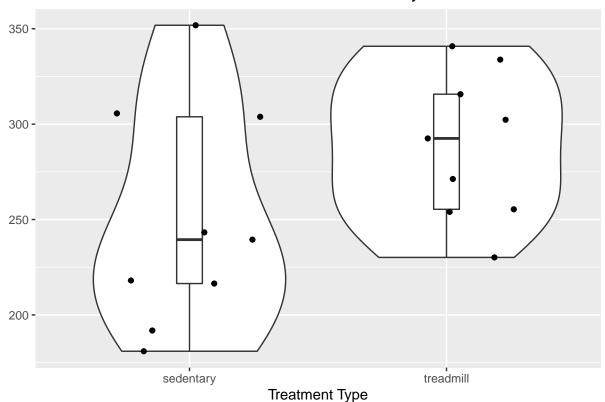


```
res.aov <- aov(CBF ~ Treatment, data = cbf1)
summary(res.aov)

### Of Sum Sa Mean Sa E welve Dr(NE)
```

```
##
              Df Sum Sq Mean Sq F value Pr(>F)
## Treatment
               1 11684
                        11684
                                  5.54 0.0326 *
              15 31635
## Residuals
                           2109
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
ggplot(data = cbf2, aes(factor(Treatment), CBF)) +
 geom_violin() +
 geom_boxplot(width = 0.1, outlier.color = "red") +
 geom_jitter(height = 0, width = 0.3) +
 labs(x = "Treatment Type",
y = "",
```

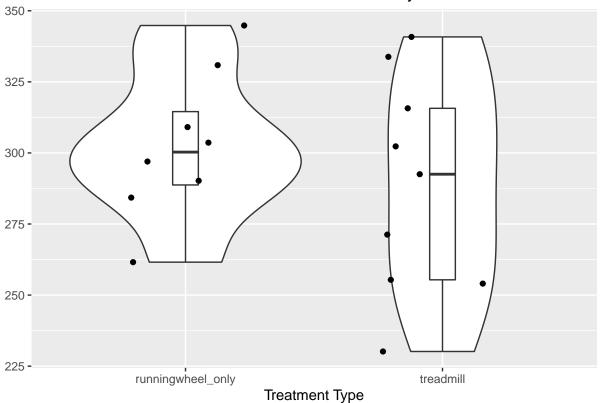
CBF distrubution between treadmill and sedentary treatments



```
res.aov <- aov(CBF ~ Treatment, data = cbf2)
summary(res.aov)</pre>
```

```
##
               Df Sum Sq Mean Sq F value Pr(>F)
## Treatment
                    6595
                            6595
                                   2.739 0.117
## Residuals
               16 38528
                            2408
ggplot(data = cbf3, aes(factor(Treatment), CBF)) +
  geom_violin() +
  geom_boxplot(width = 0.1, outlier.color = "red") +
  geom_jitter(height = 0, width = 0.3) +
 labs(x = "Treatment Type",
    y = "",
    title = "CBF distrubution between treadmill and sedentary treatments")
```

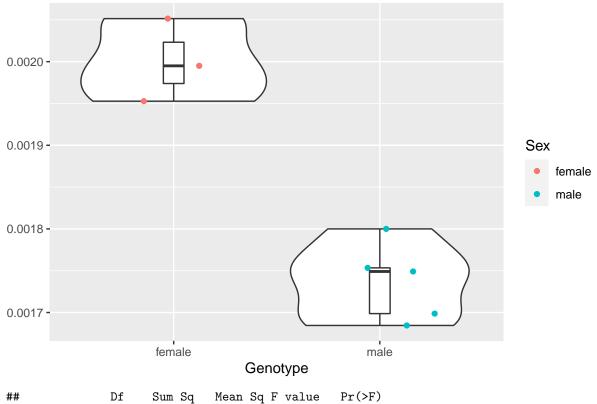




```
res.aov <- aov(CBF ~ Treatment, data = cbf3)
summary(res.aov)</pre>
```

```
## Treatment 1 859 Mean Sq F value Pr(>F)  
## Residuals 15 16548 1103.2
```

Primary Somatosensory Cortex Hindlimb Region Red points denoting outliers



```
## Sex 1 1.293e-07 1.293e-07 57.42 0.000275 ***

## Residuals 6 1.351e-08 2.250e-09

## ---

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