

violin-plots

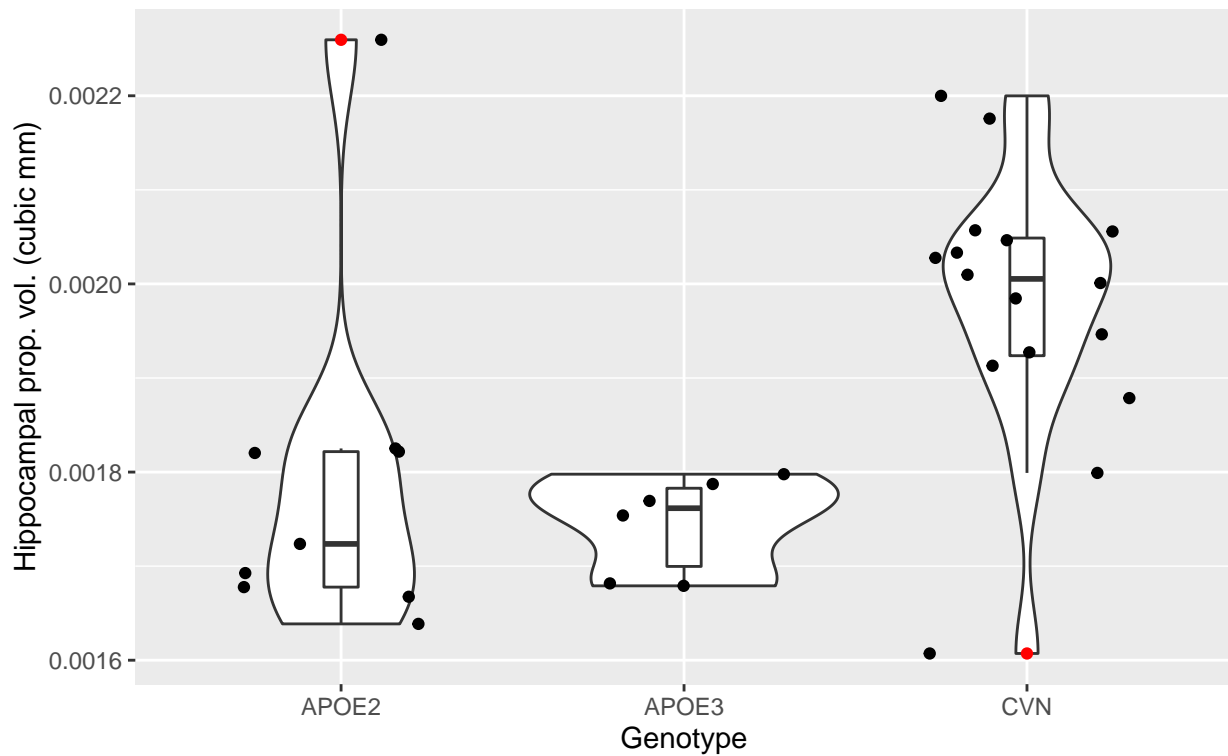
Anna MacFarlane

11/14/2020

Violin plots

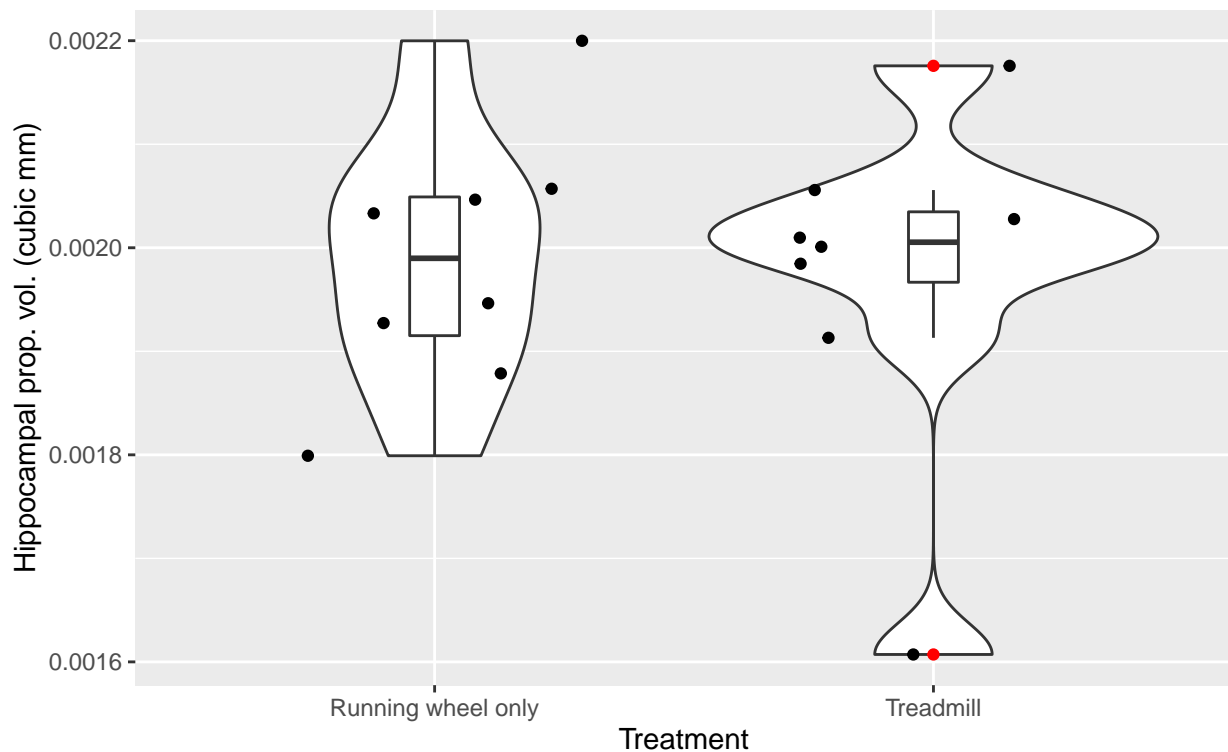
Distributions of hippocampal proportional volumes by genotype

Red points denoting outliers



Distributions of hippocampal proportional volumes by treatment

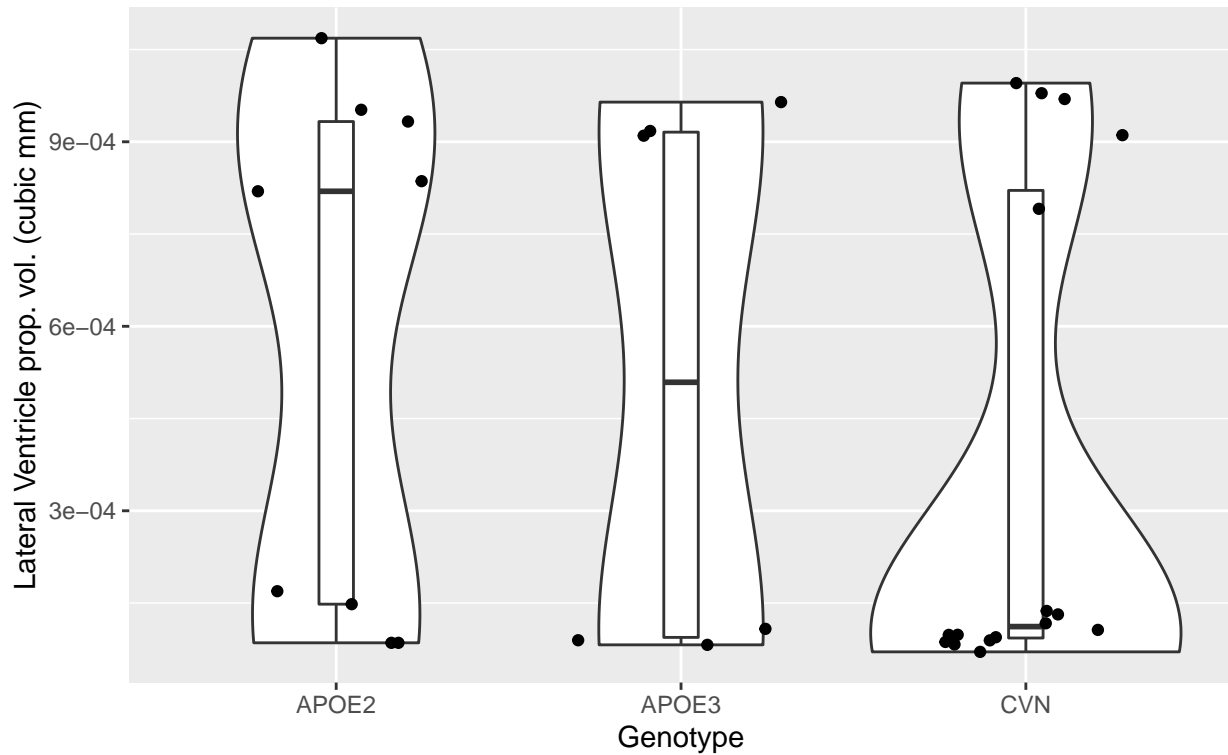
In CVN genotype with red points denoting outliers



```
##
## Welch Two Sample t-test
##
## data:  rw and tm
## t = 0.19398, df = 13.009, p-value = 0.8492
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.0001436222  0.0001719597
## sample estimates:
## mean of x mean of y
## 0.001986000 0.001971831
```

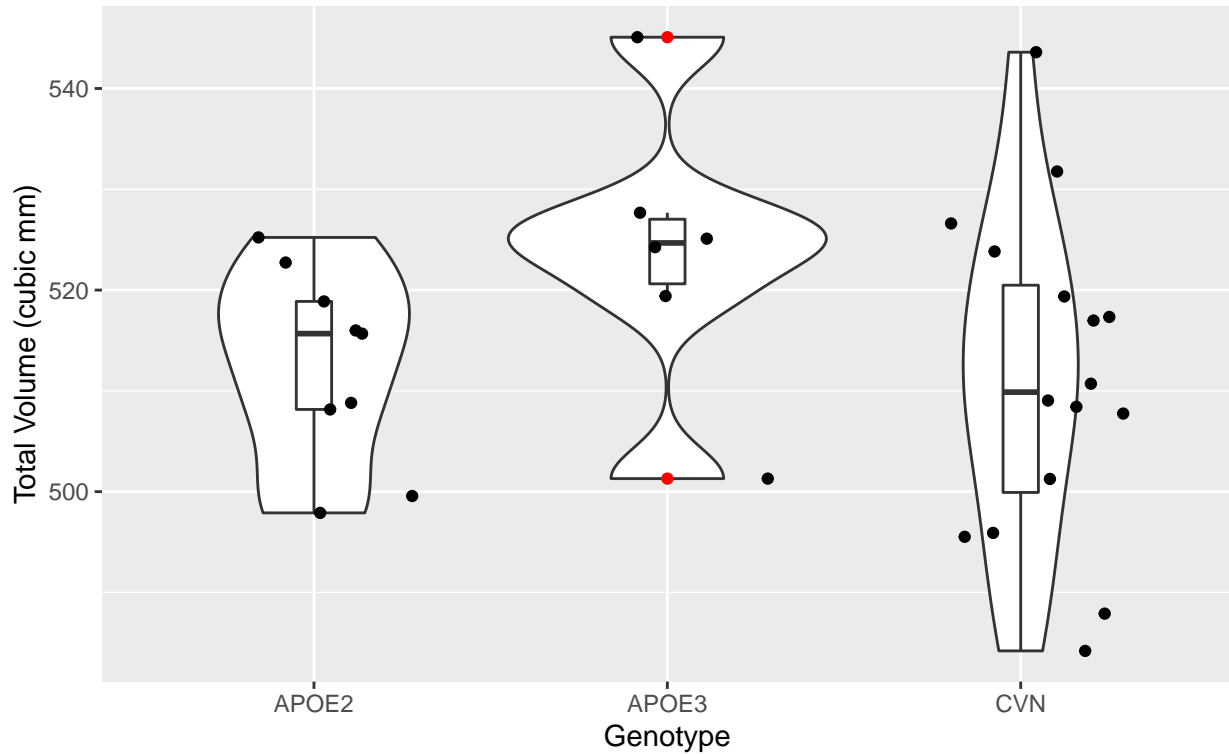
Distributions of lateral ventricle proportional volumes by genotype

Red points denoting outliers



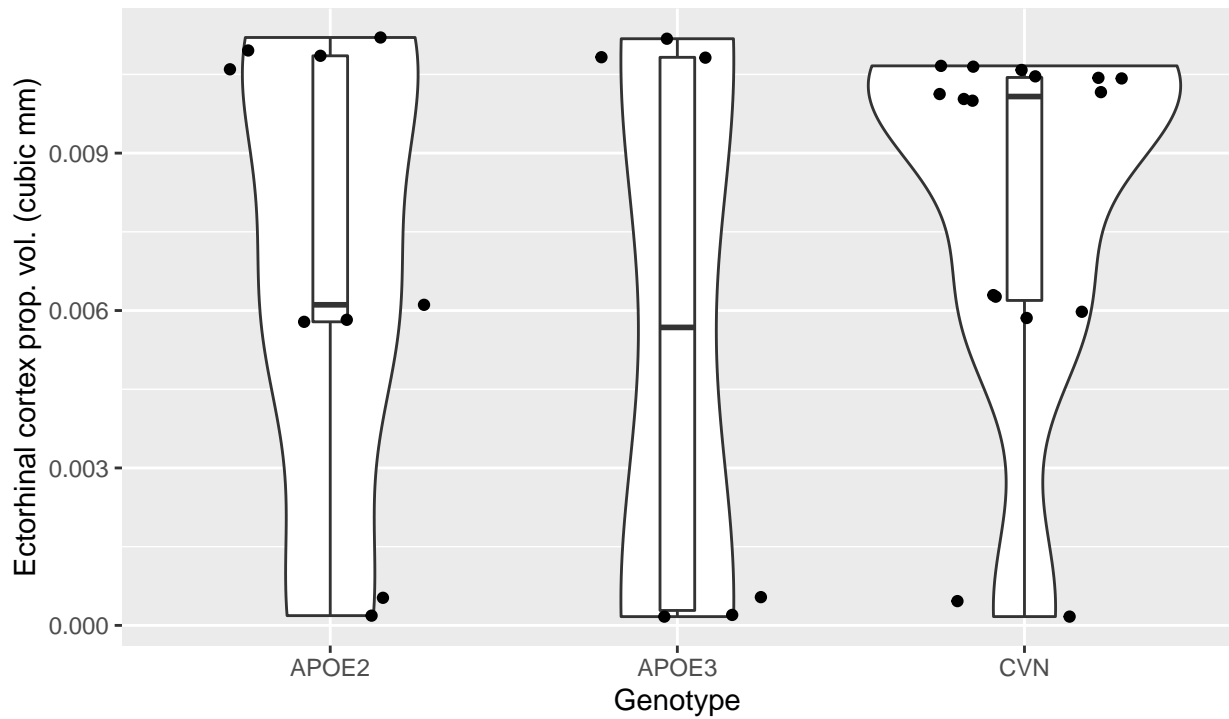
Distributions of total brain volumes by genotype

Red points denoting outliers

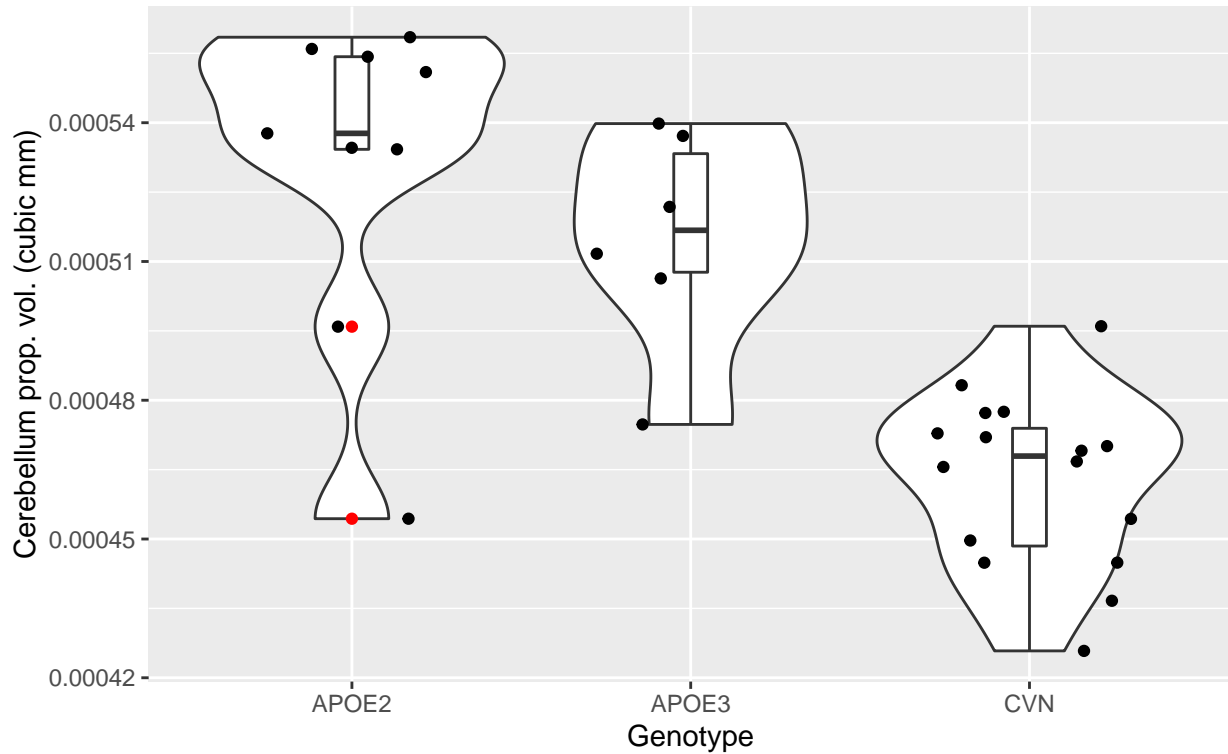


Distributions of ectorhinal cortex proportional volumes by genotype

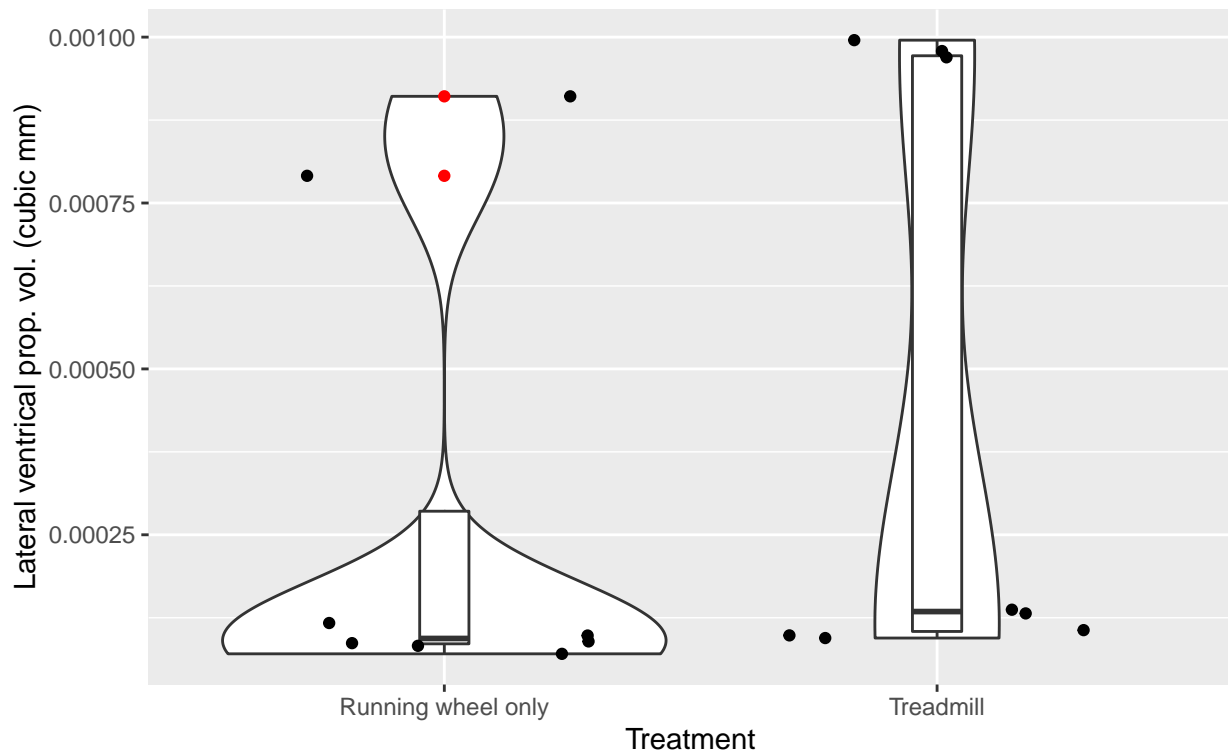
Red points denoting outliers



Distributions of cerebellum proportional volumes by genotype
Red points denoting outliers



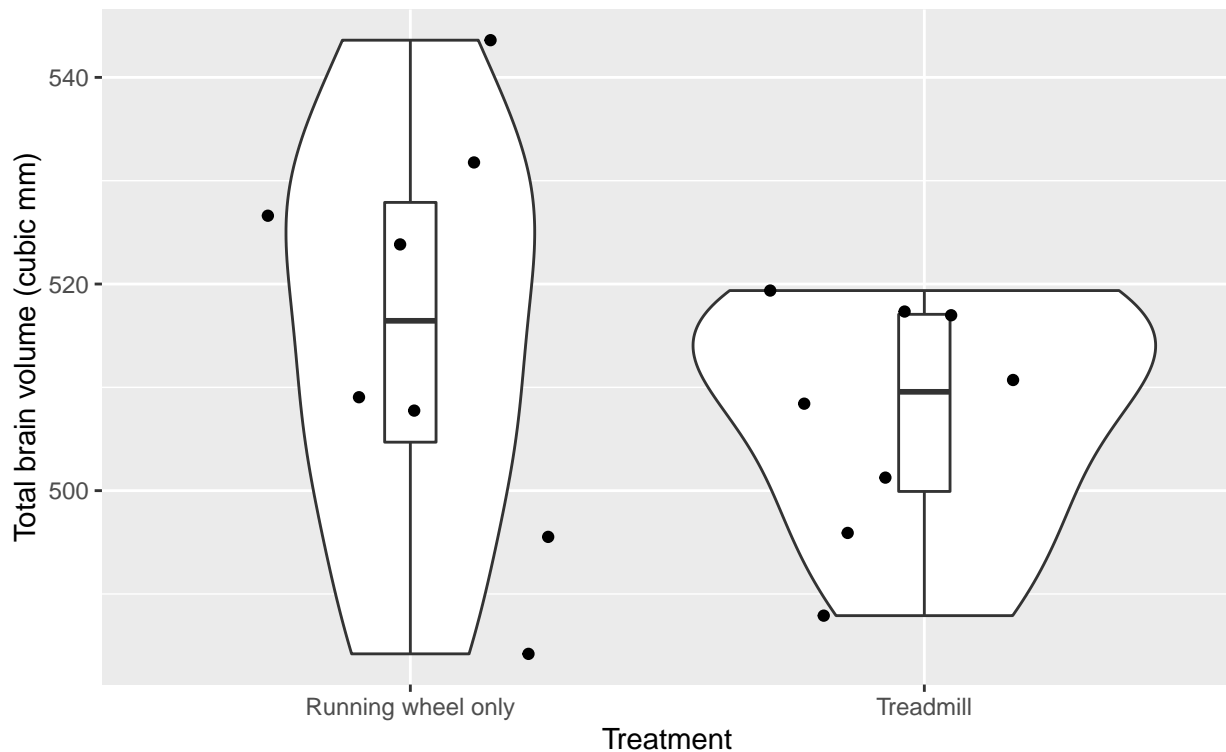
Distributions of lateral ventricular proportional volumes by treatment
In CVN genotype with red points denoting outliers



```
##
## Welch Two Sample t-test
##
## data: rw and tm
## t = -0.7824, df = 13.265, p-value = 0.4477
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.0005940676 0.0002777043
## sample estimates:
## mean of x mean of y
## 0.0002807931 0.0004389748
```

Distributions of total brain volumes volumes by treatment

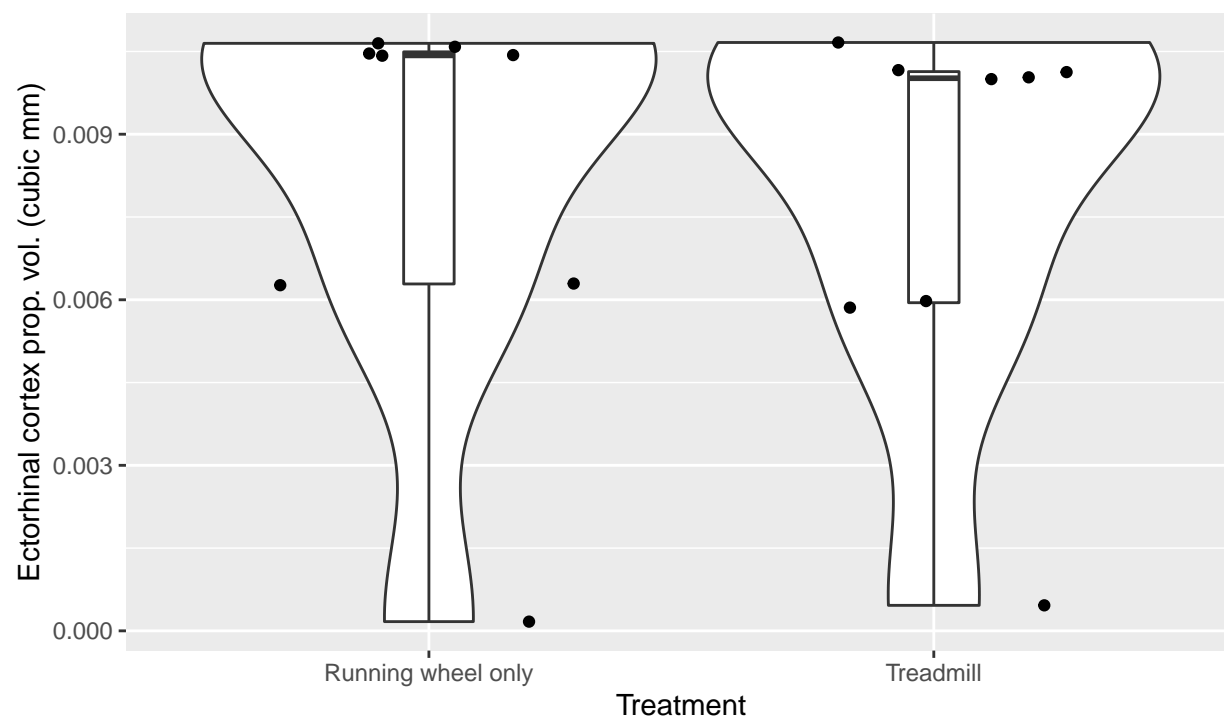
In CVN genotype with red points denoting outliers



```
##
## Welch Two Sample t-test
##
## data: rw and tm
## t = 1.0017, df = 11.149, p-value = 0.3377
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -9.617324 25.730573
## sample estimates:
## mean of x mean of y
## 515.2913 507.2346
```

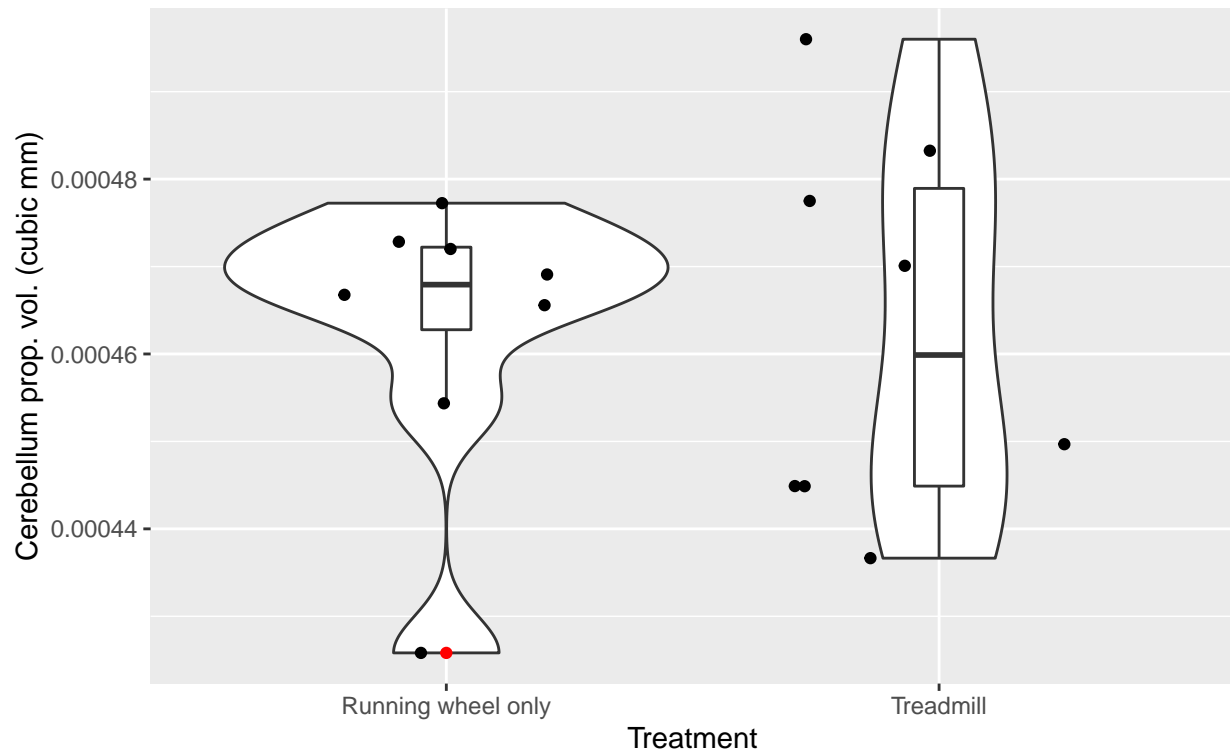
Distributions of ectorhinal cortex proportional volumes by treatment

In CVN genotype with red points denoting outliers



```
##
## Welch Two Sample t-test
##
## data:  rw and tm
## t = 0.13634, df = 13.97, p-value = 0.8935
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.003685234  0.004185445
## sample estimates:
##  mean of x  mean of y
## 0.008159700 0.007909594
```

Distributions of cerebellum proportional volumes by treatment
In CVN genotype with red points denoting outliers



```
##
## Welch Two Sample t-test
##
## data:  rw and tm
## t = 0.0099646, df = 13.06, p-value = 0.9922
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -2.068061e-05 2.087236e-05
## sample estimates:
## mean of x mean of y
## 0.0004629630 0.0004628671
```