

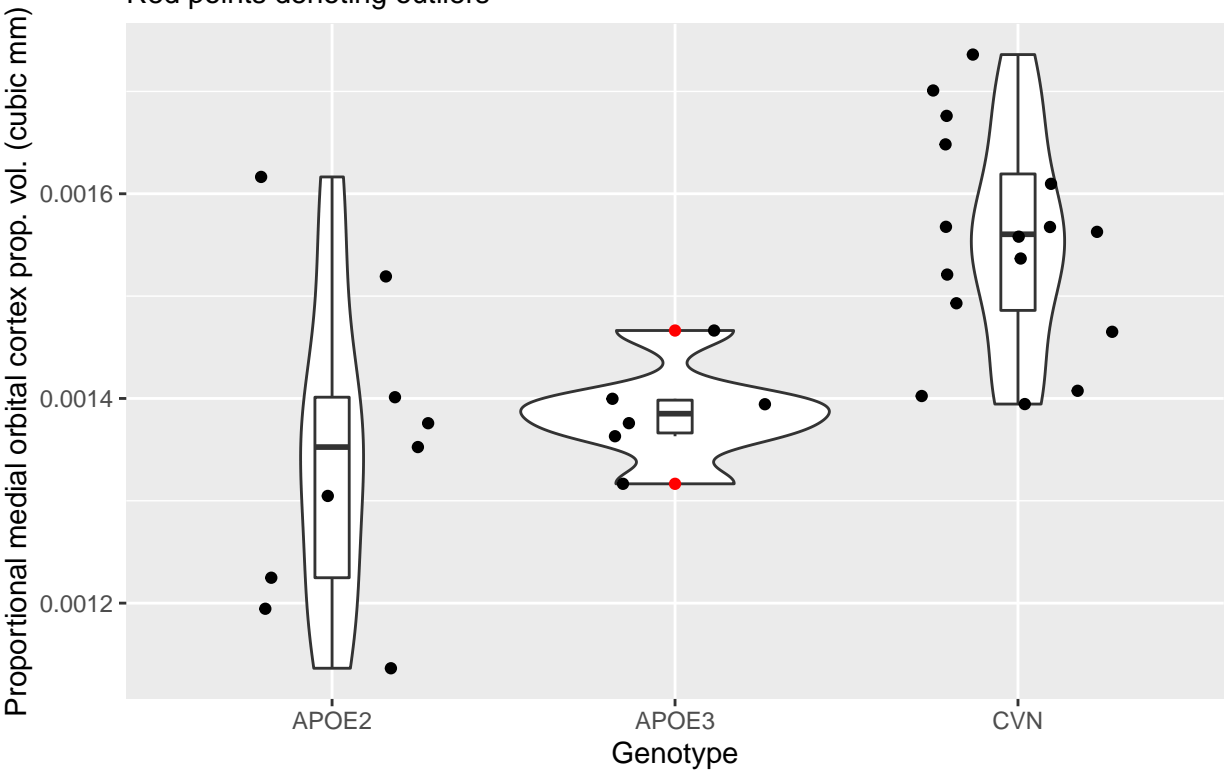
# Proportional Volume Distributions by Region

Anna MacFarlane

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## Violin plots

Distributions of medial orbital cortex proportional volumes by genotype  
Red points denoting outliers



Distributions of secondary motor cortex proportional volumes by genotyp  
Red points denoting outliers

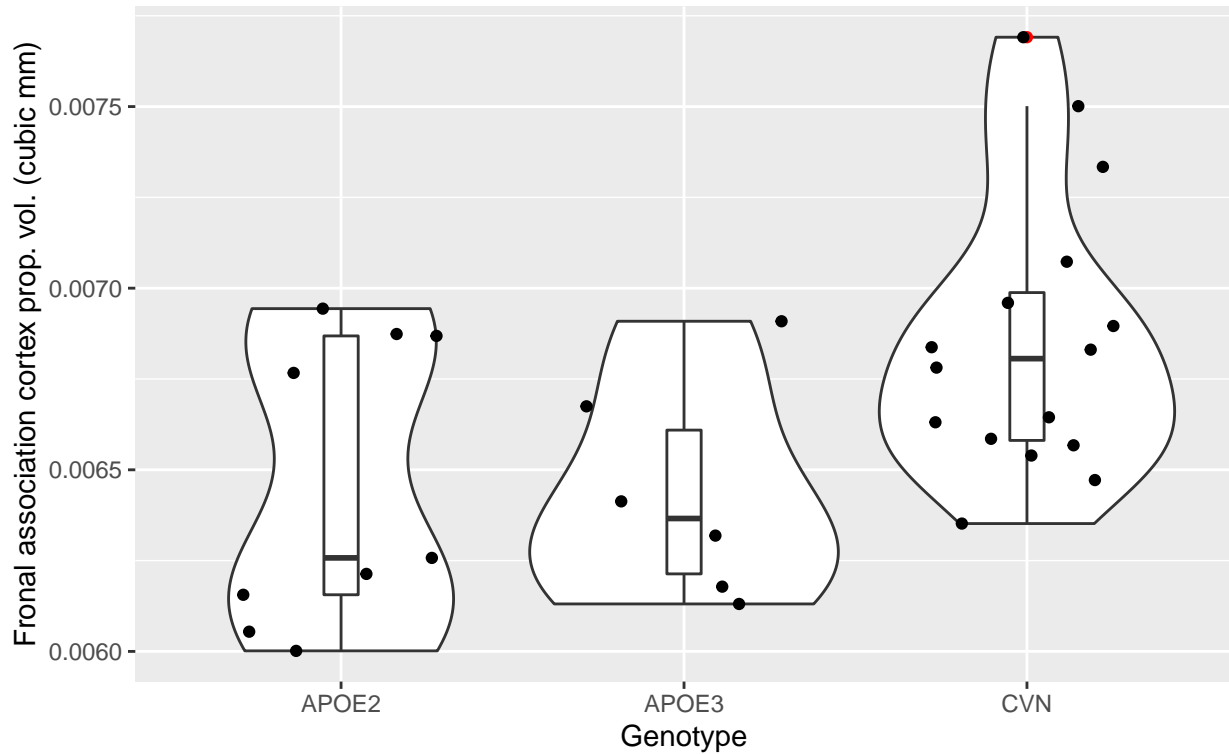


Distributions of lateral parietal association cortex proportional volumes by genotype

Red points denoting outliers



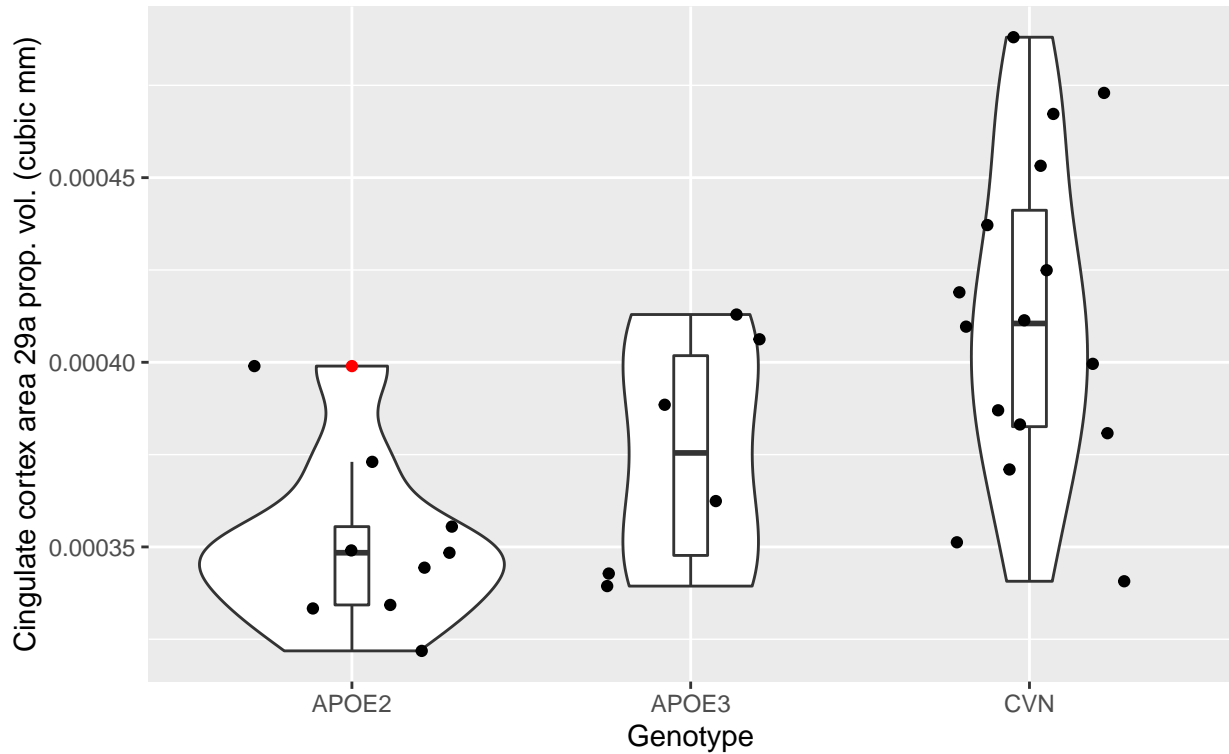
Distributions of frontal association cortex proportional volumes by genoty  
Red points denoting outliers



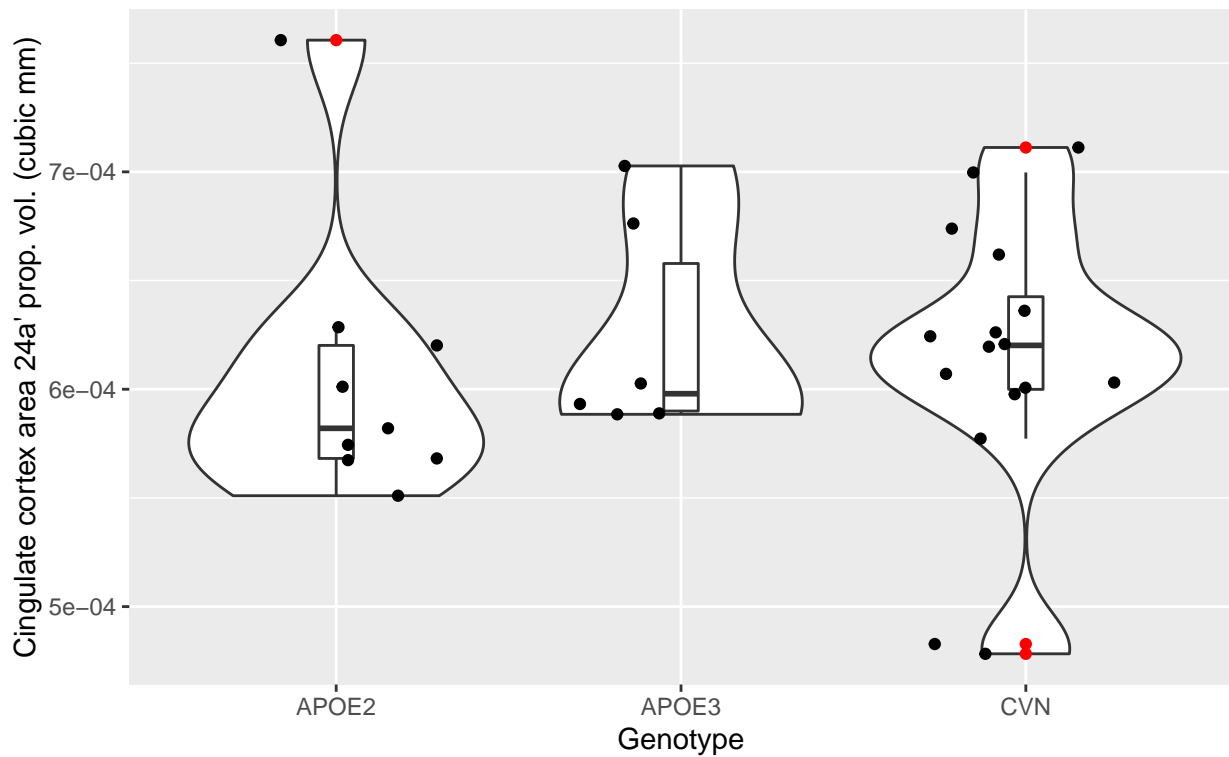
Distributions of cingulate cortex area 29c proportional volumes by genoty  
Red points denoting outliers



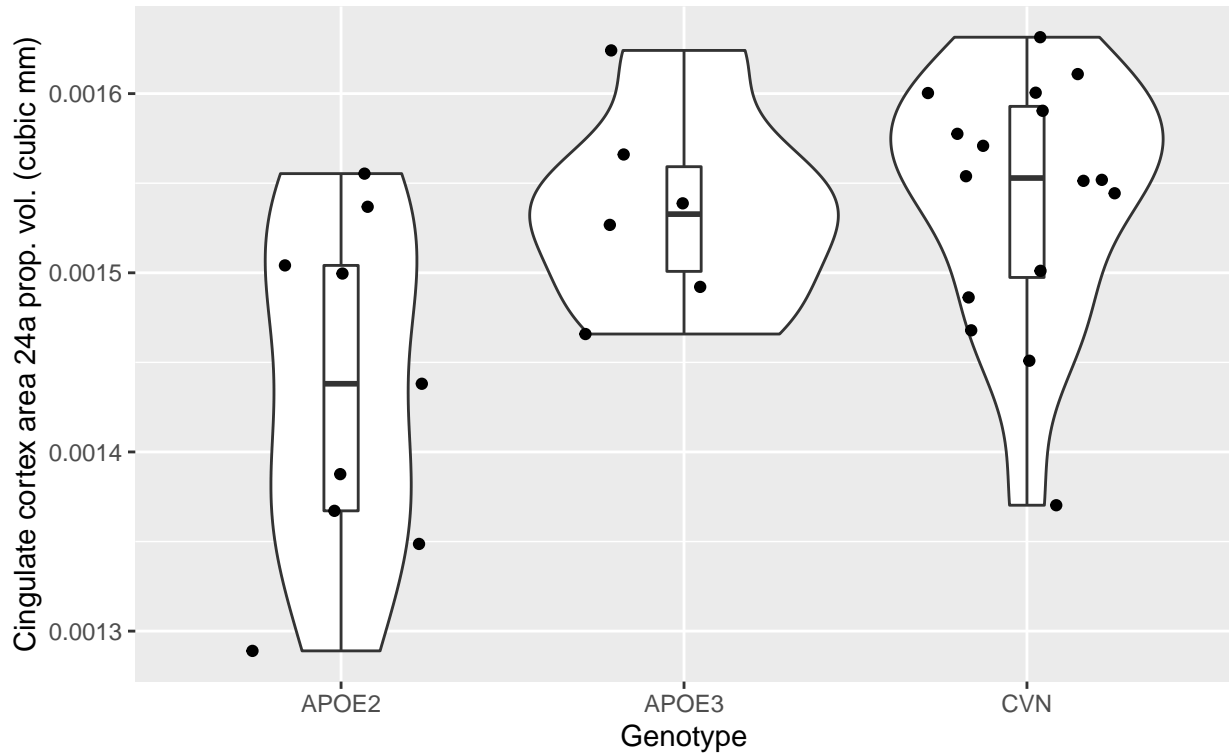
Distributions of cingulate cortex area 29a proportional volumes by genot  
Red points denoting outliers



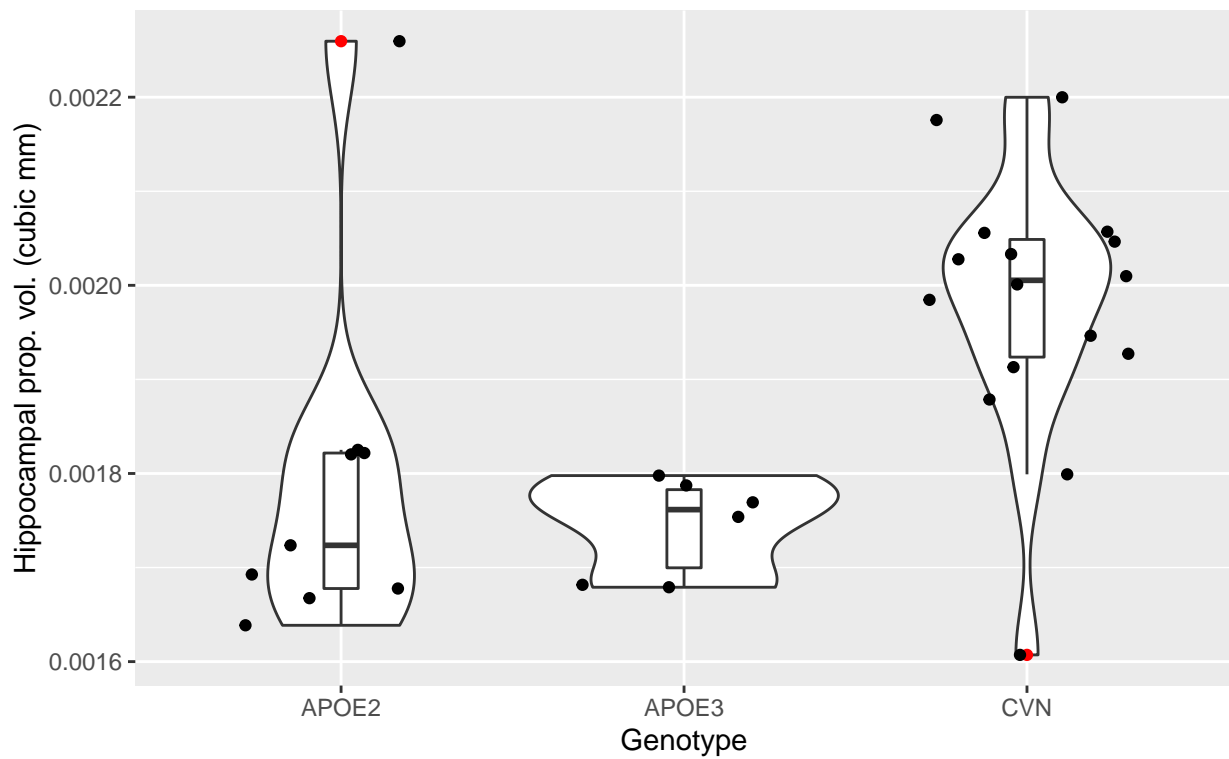
Distributions of cingulate cortex area 24a' proportional volumes by genoty  
Red points denoting outliers



Distributions of cingulate cortex area 24a proportional volumes by genoty  
Red points denoting outliers

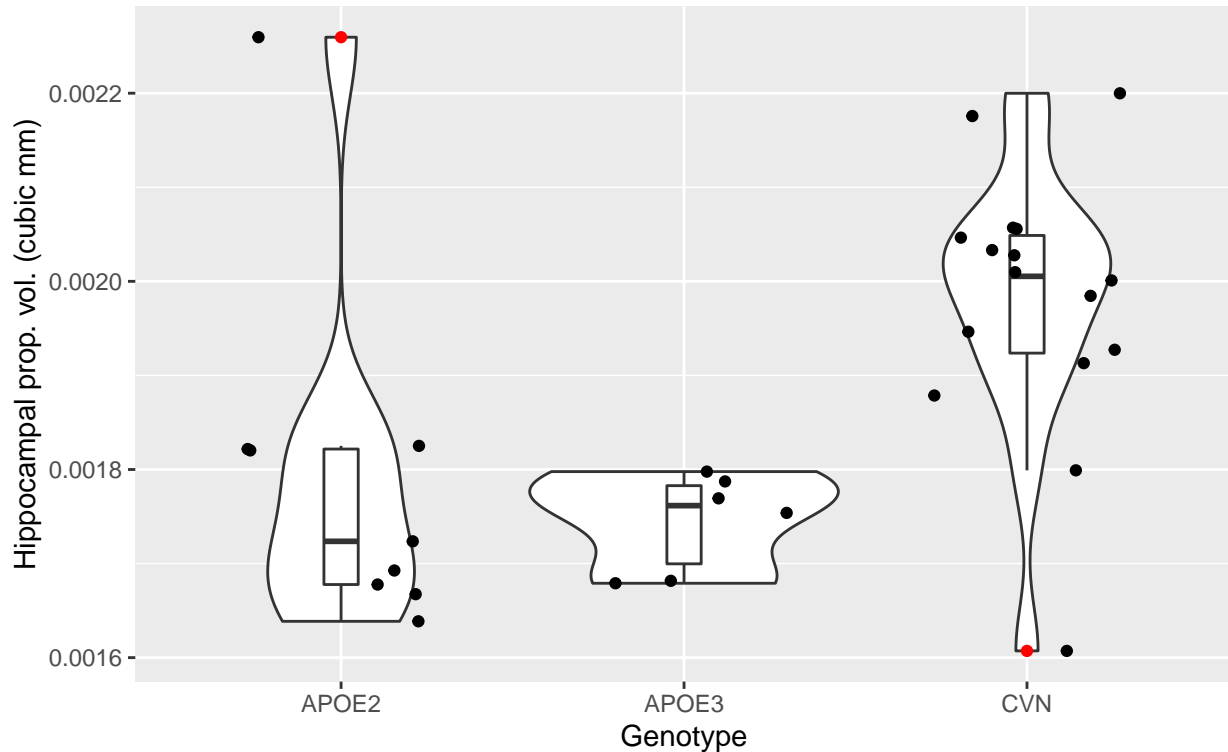


Distributions of hippocampal proportional volumes by genotype  
Red points denoting outliers



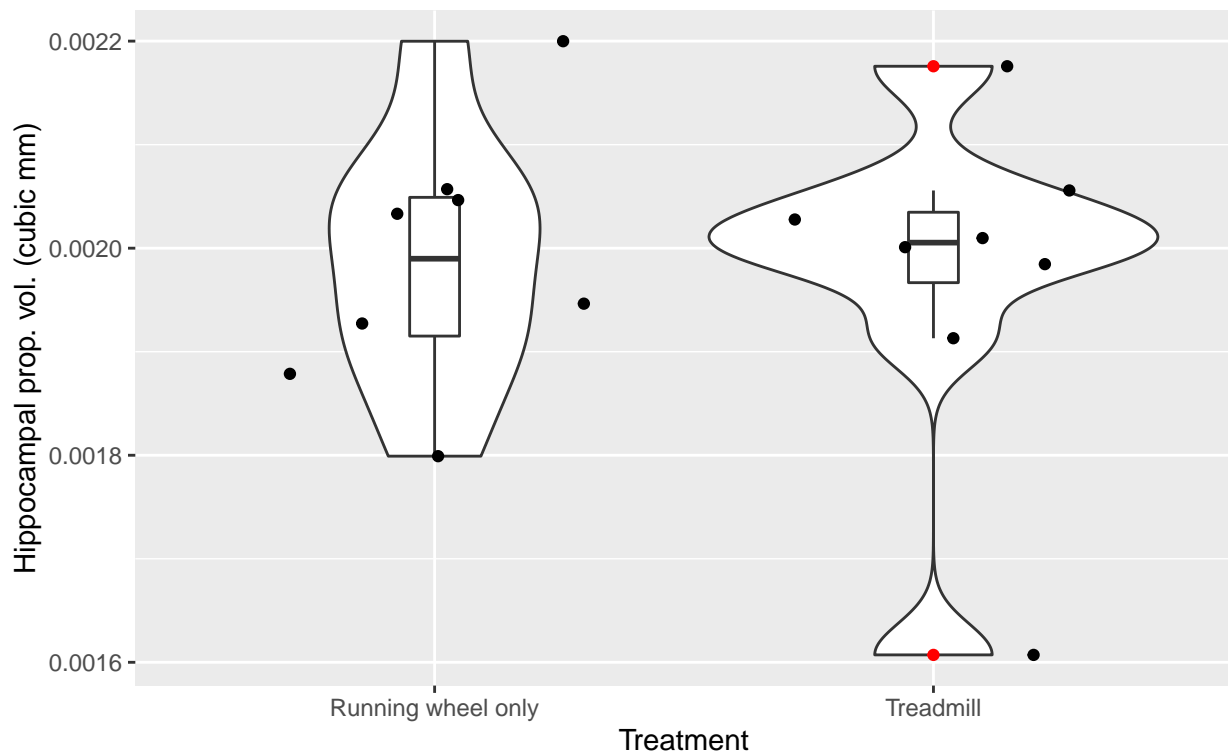
## Distributions of hippocampal proportional volumes by genotype

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
```

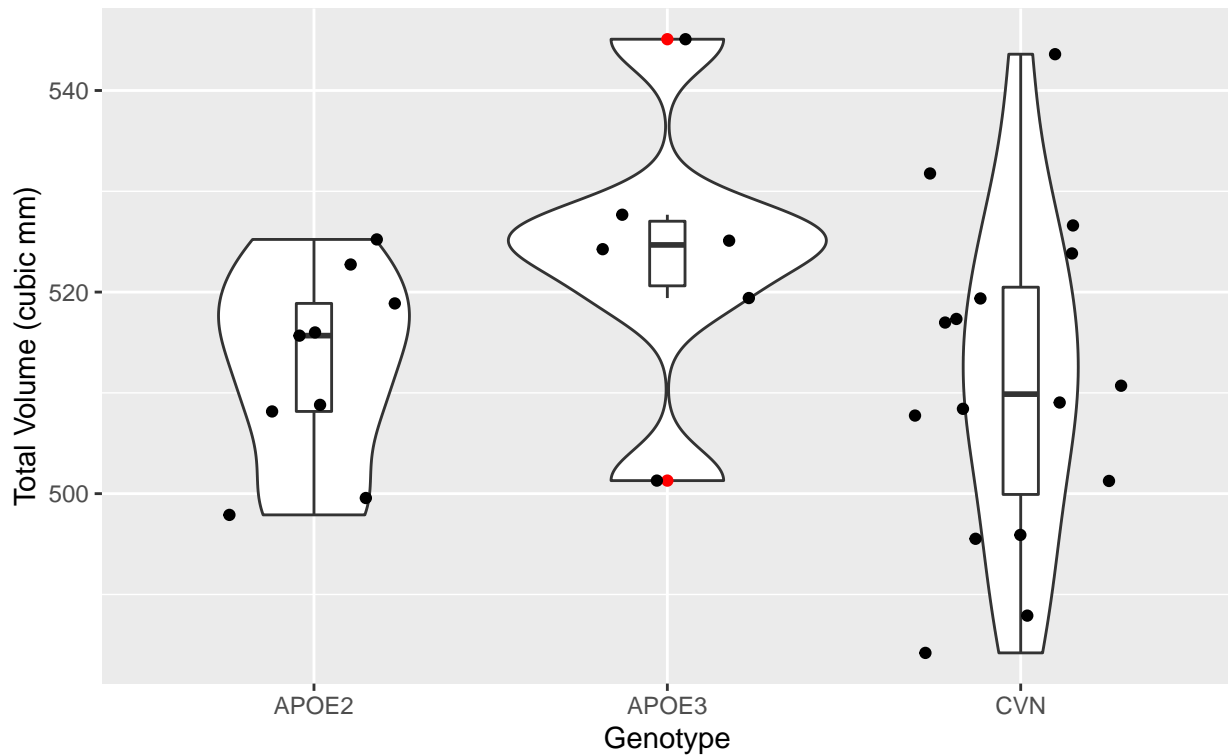
## Distributions of lateral ventricle proportional volumes by genotype

Red points denoting outliers



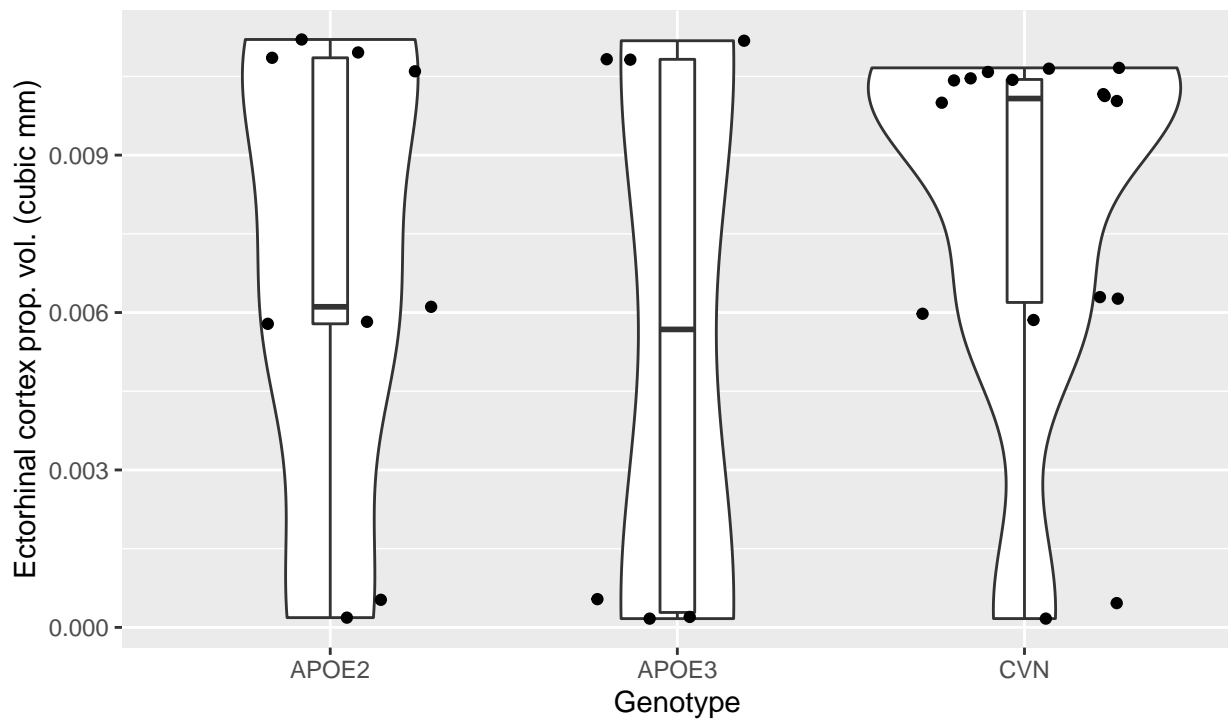
## Distributions of total brain volumes by genotype

Red points denoting outliers



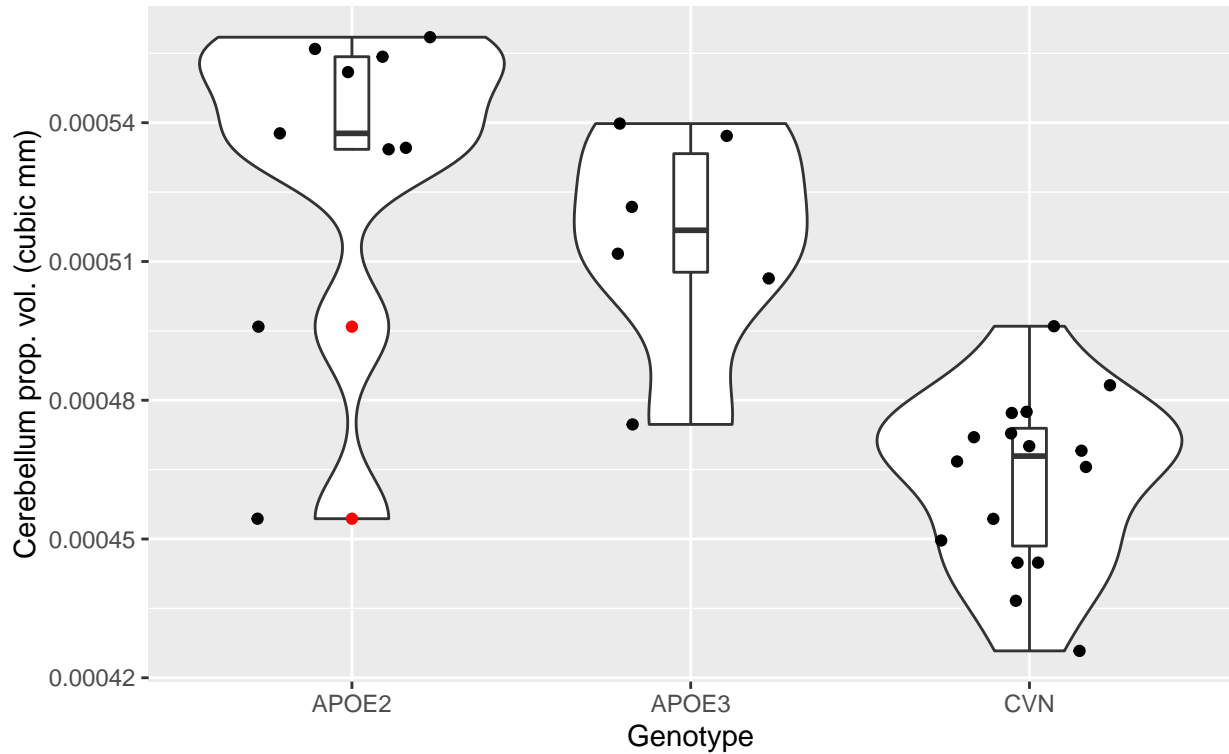
## Distributions of ectothal cortex proportional volumes by genotype

Red points denoting outliers

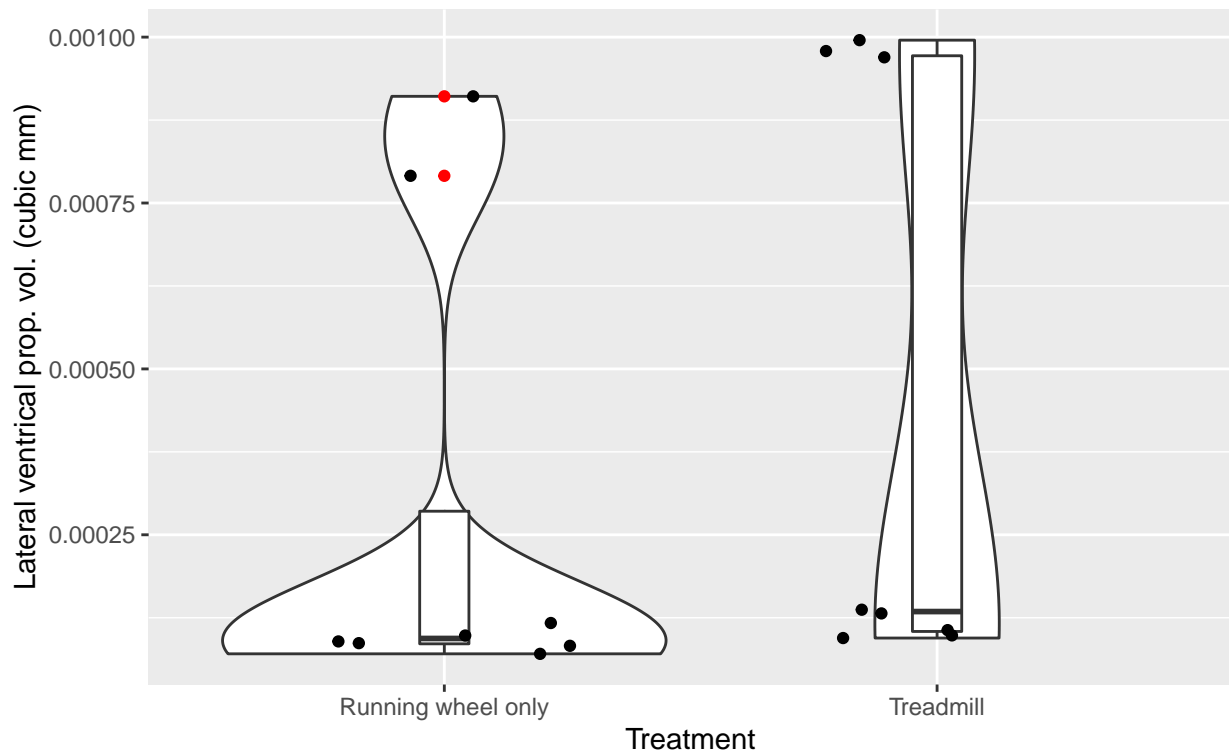




Distributions of cerebellum proportional volumes by genotype  
Red points denoting outliers



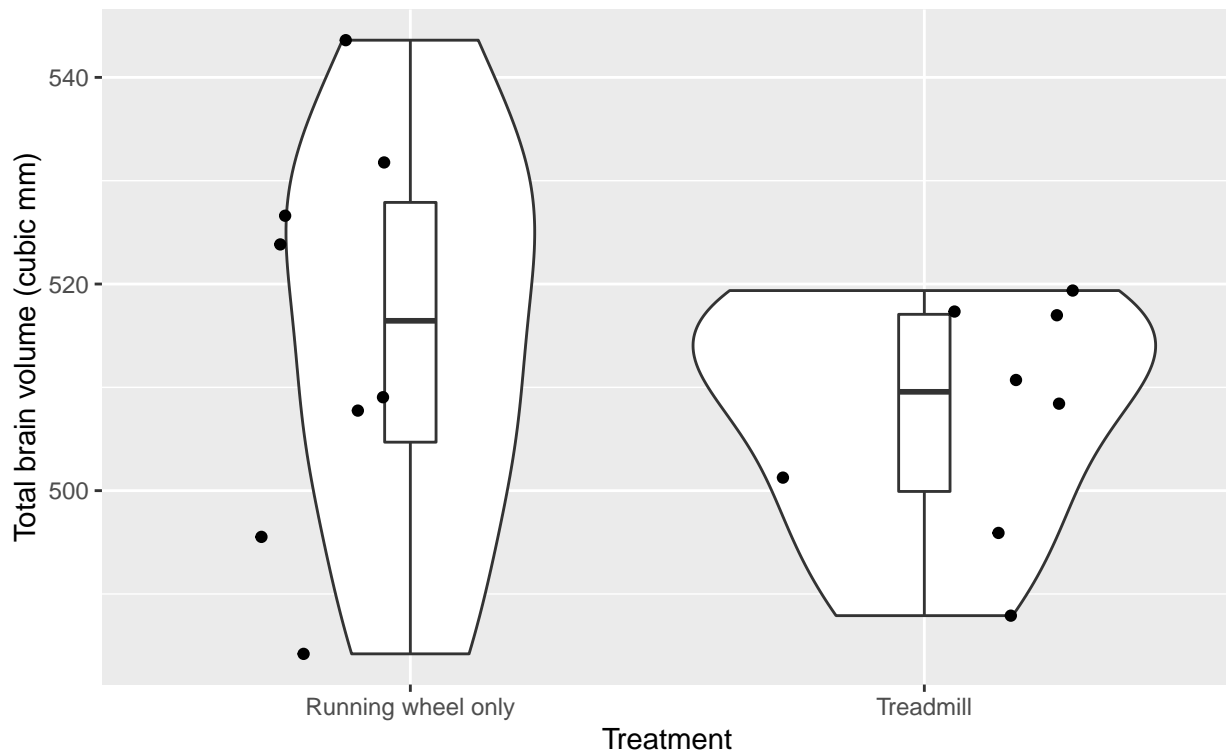
Distributions of lateral ventricular proportional volumes by treatment  
In CVN genotype with red points denoting outliers



```
##
## 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

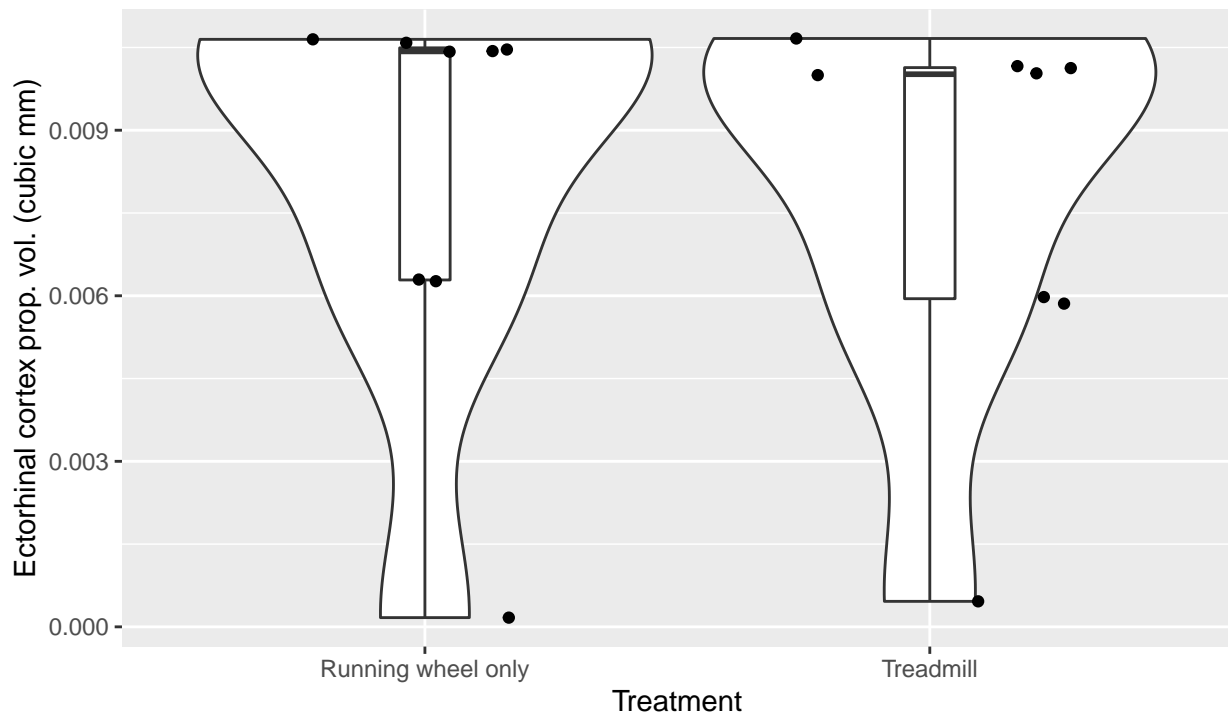
In CVN genotype with red points denoting outliers



```
##
## 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
```

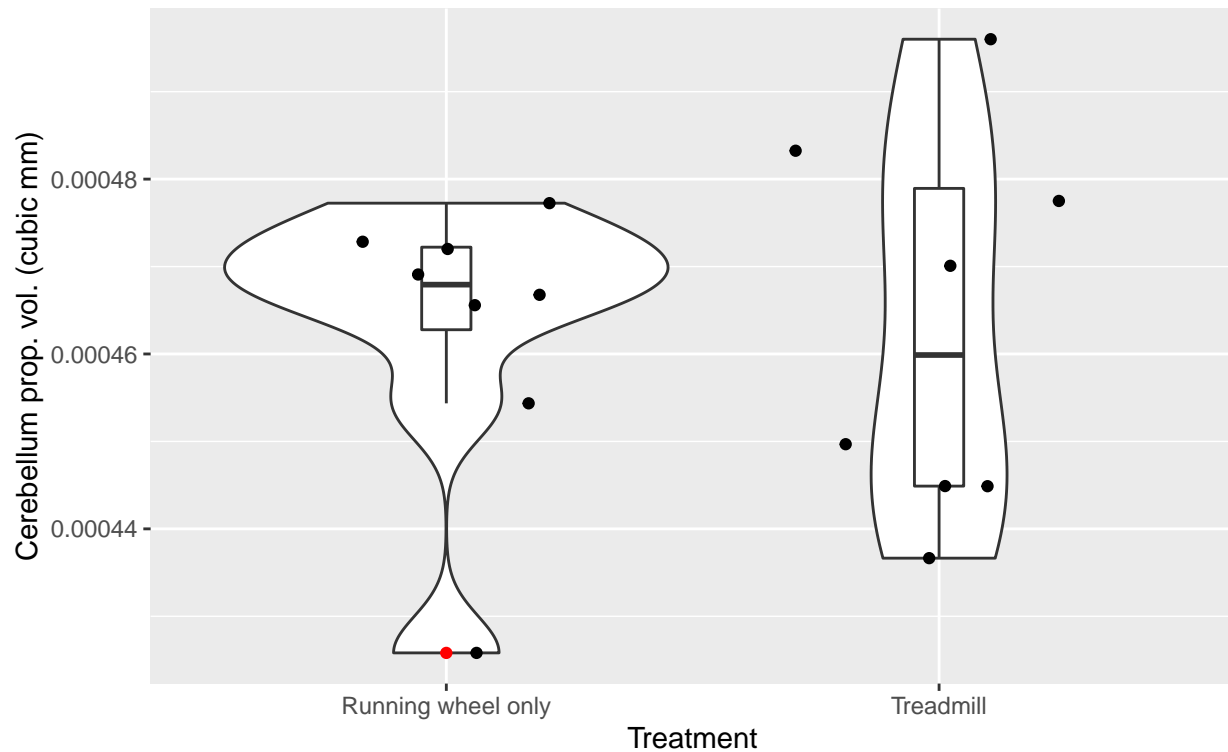
# Distributions of ectorhinal cortex proportional volumes by treatment

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 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
```