

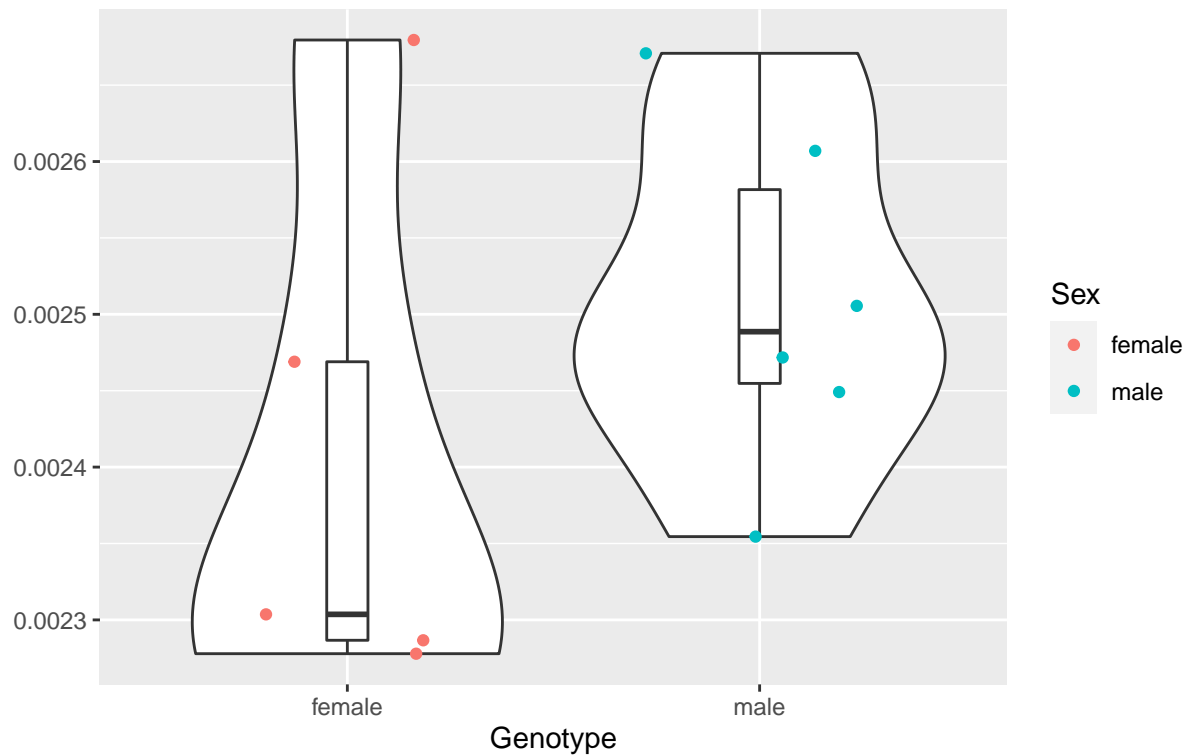
Right APOE2 Disaggregated by Sex

Anna MacFarlane

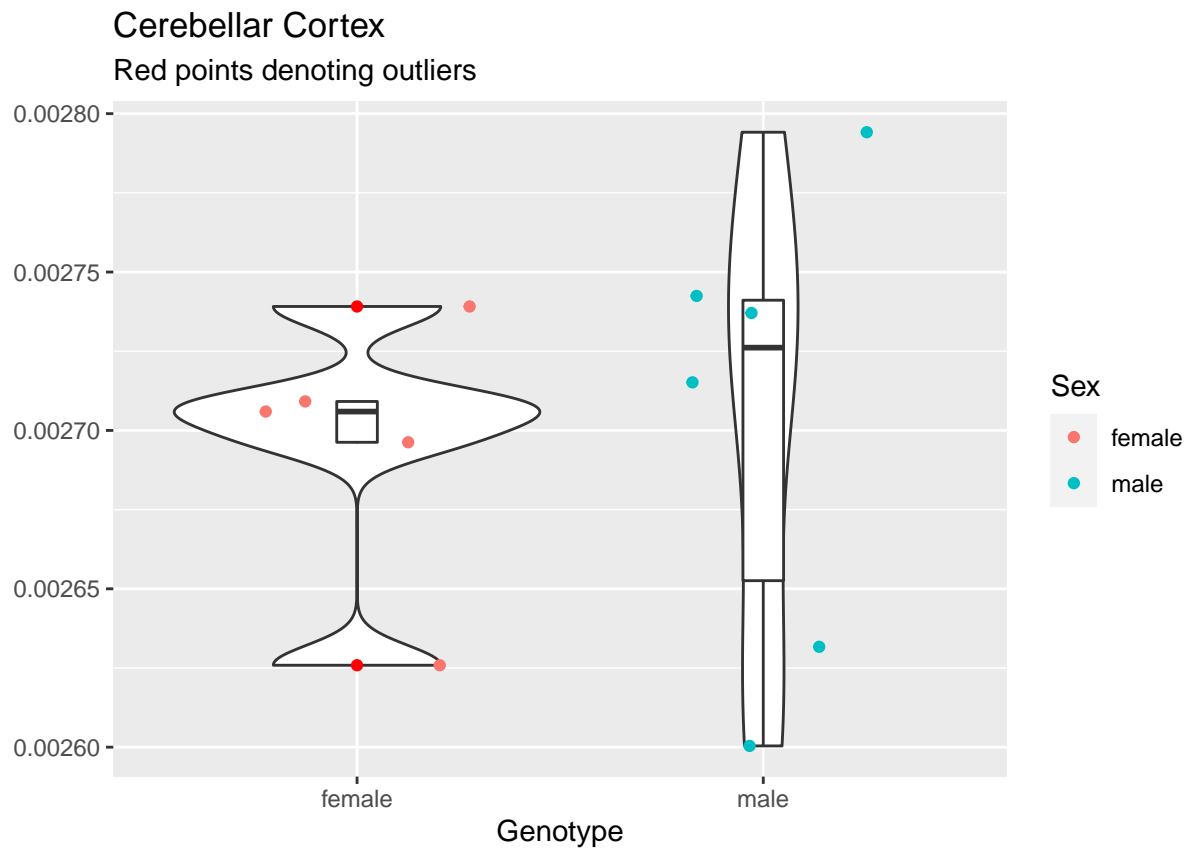
3/25/2021

Interpeduncular Nucleus

Red points denoting outliers

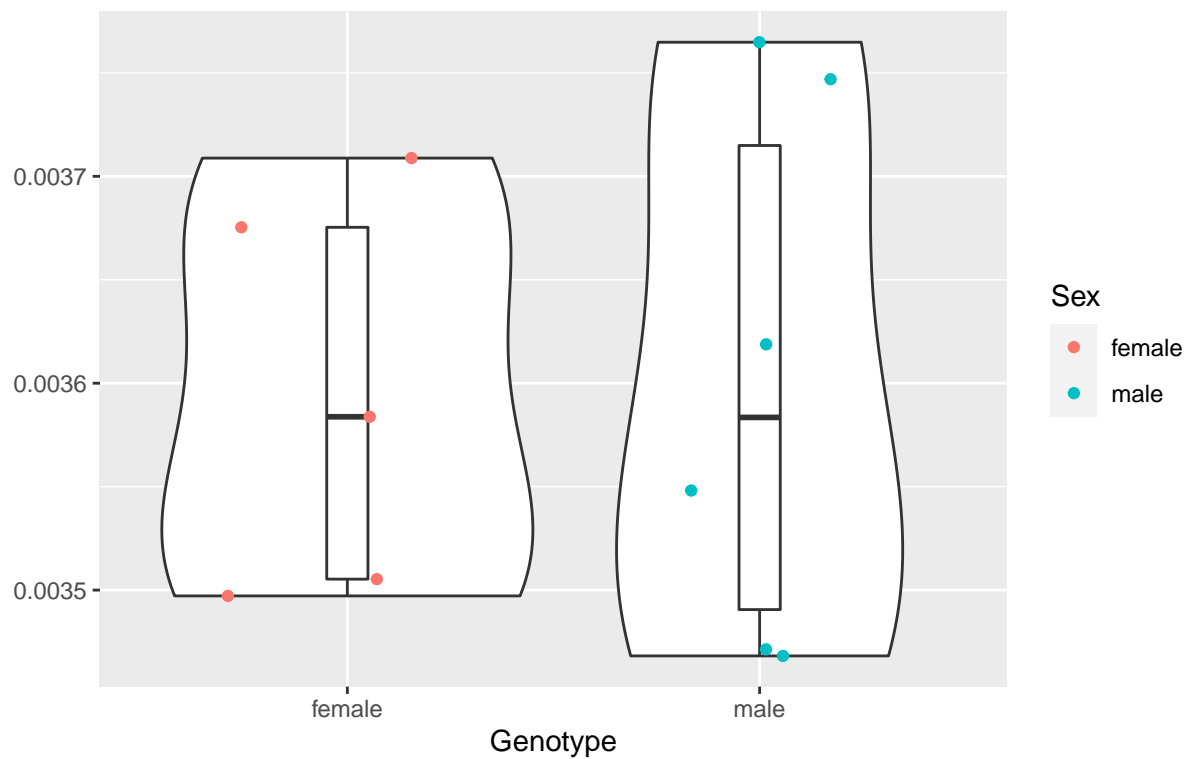


```
##          Df    Sum Sq  Mean Sq F value Pr(>F)
## Sex       1 3.091e-08 3.091e-08   1.507  0.251
## Residuals 9 1.845e-07 2.050e-08
```

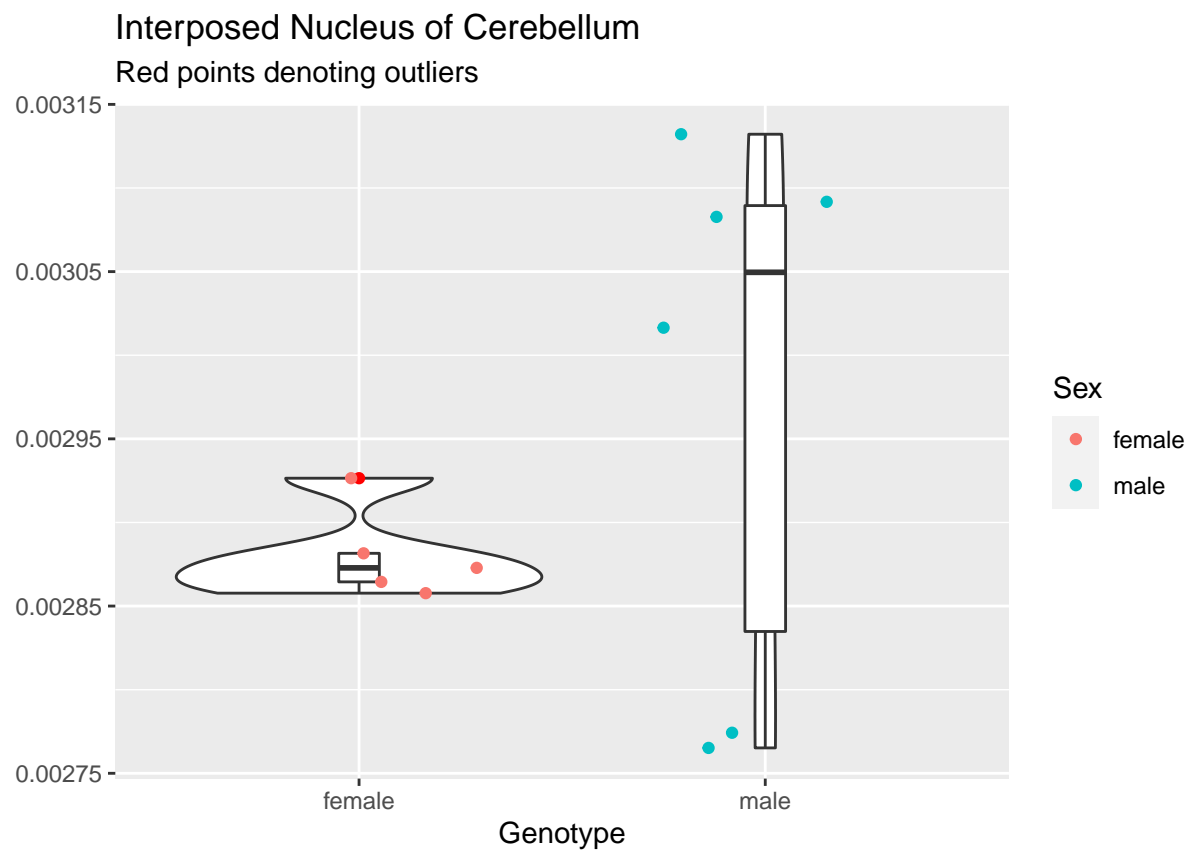


Dentate (Lateral) Nucleus of Cerebellum

Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.200e-10	2.190e-10	0.016	0.902
## Residuals	9	1.228e-07	1.365e-08		



Fastigial Medial Dorsolateral Nucleus of Cerebellum

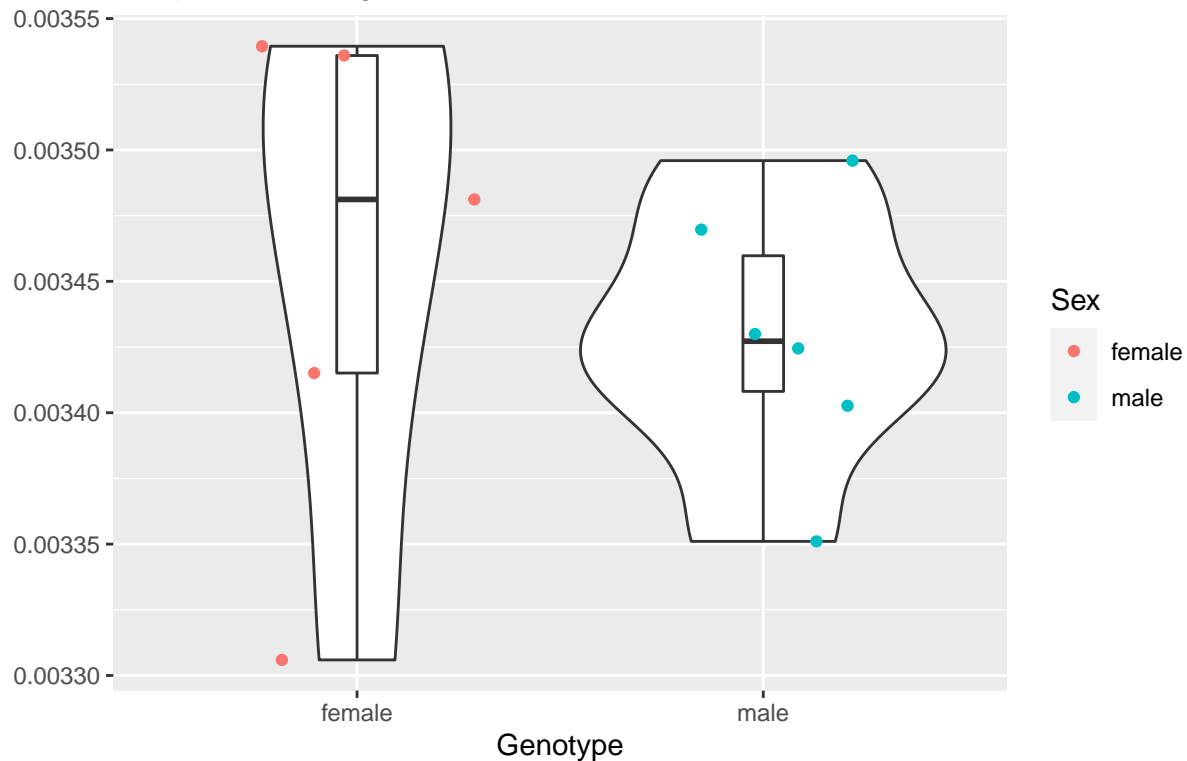
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.354e-08	2.354e-08	1.423	0.263
## Residuals	9	1.489e-07	1.654e-08		

Fastigial Medial Nucleus of Cerebellum

Red points denoting outliers



```
##           Df      Sum Sq   Mean Sq F value Pr(>F)
## Sex         1 1.920e-09 1.924e-09   0.339  0.575
## Residuals   9 5.111e-08 5.679e-09
```

```
#“{r VII, echo = FALSE} #ggplot(data = apoe2, aes(factor(Sex), VII)) + #geom_violin() +
#geom_boxplot(width = 0.1, outlier.color = “red”) + #geom_jitter(height = 0, width = 0.3) + #labs(x
= “Genotype”, #y = “”, #title = “Ventral Lateral Lemniscus Nucleus”, #subtitle = “Red points denoting
outliers”)
```

```
#res.aov <- aov(VII ~ Sex, data = apoe2) #summary(res.aov) #“
```

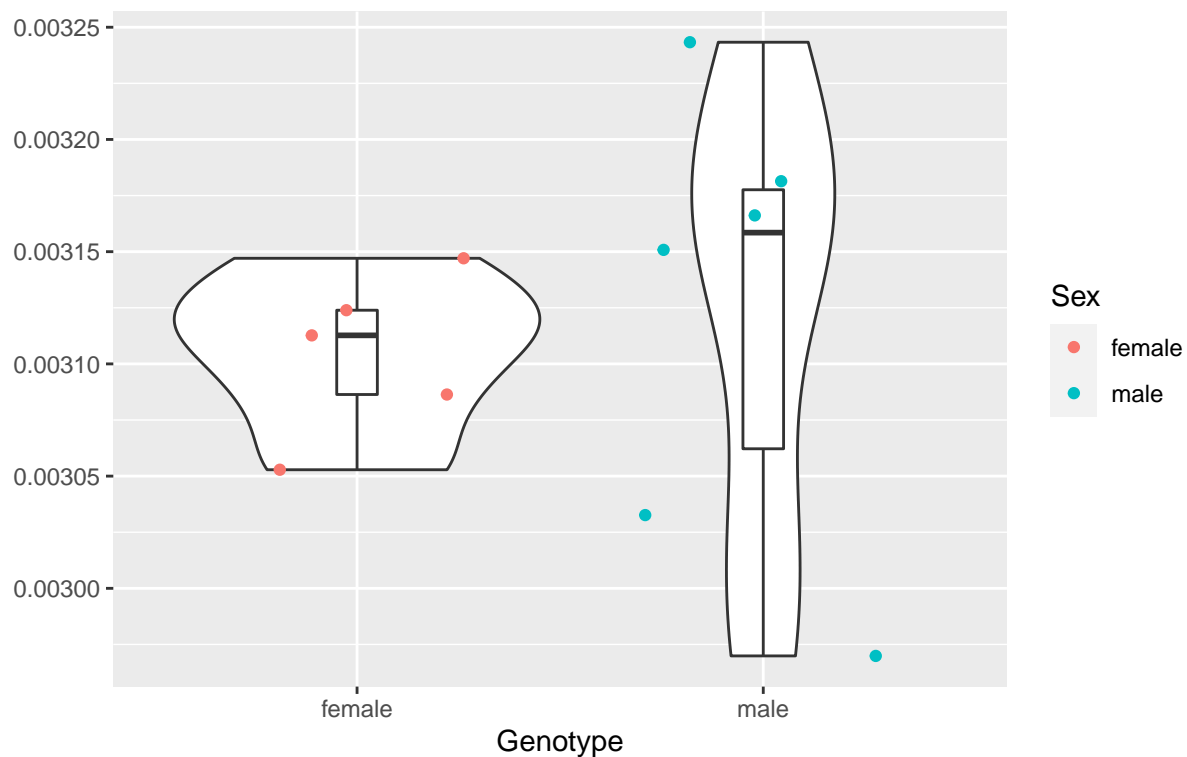
Parabrachial Nucleus

Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.510e-09	1.511e-09	0.107	0.751
## Residuals	9	1.267e-07	1.408e-08		

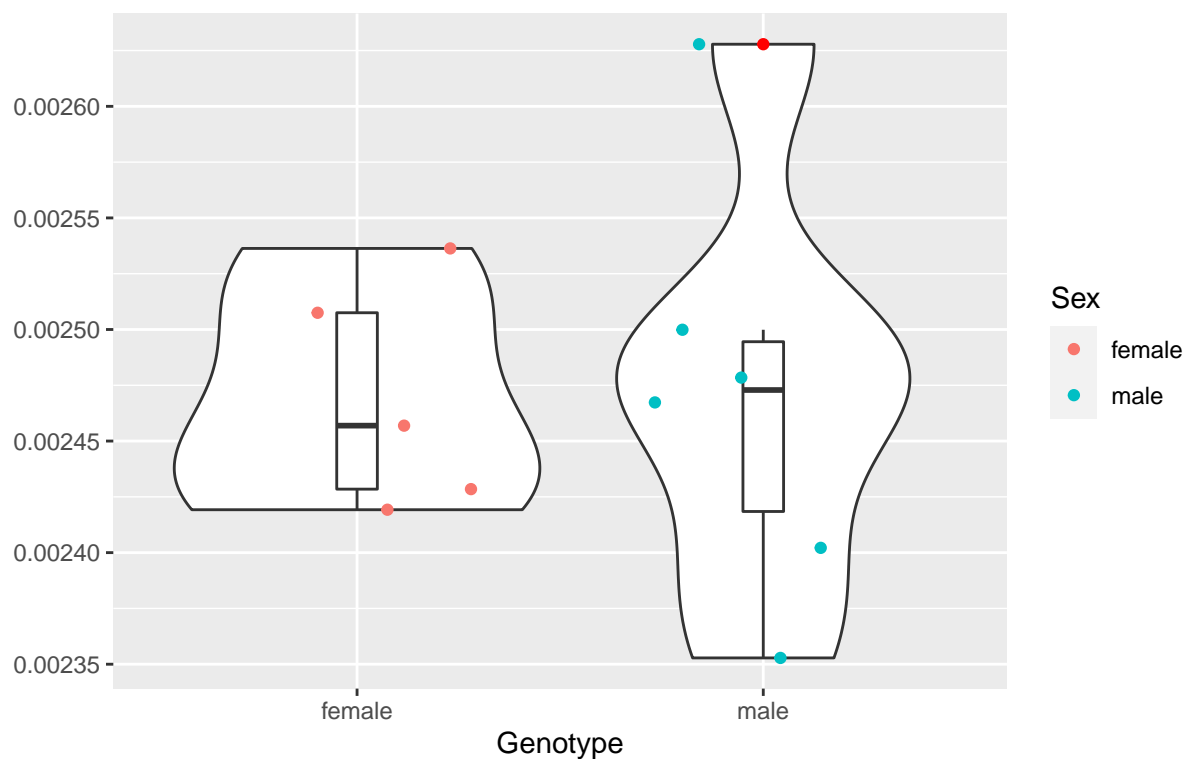
Parabrachial Medial Nucleus and Koelliker Fuse Nucleus Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.030e-09	1.033e-09	0.162	0.697
## Residuals	9	5.738e-08	6.376e-09		

Parvicellular Reticular Nucleus and Principal Sensory Trigeminal Nucleu

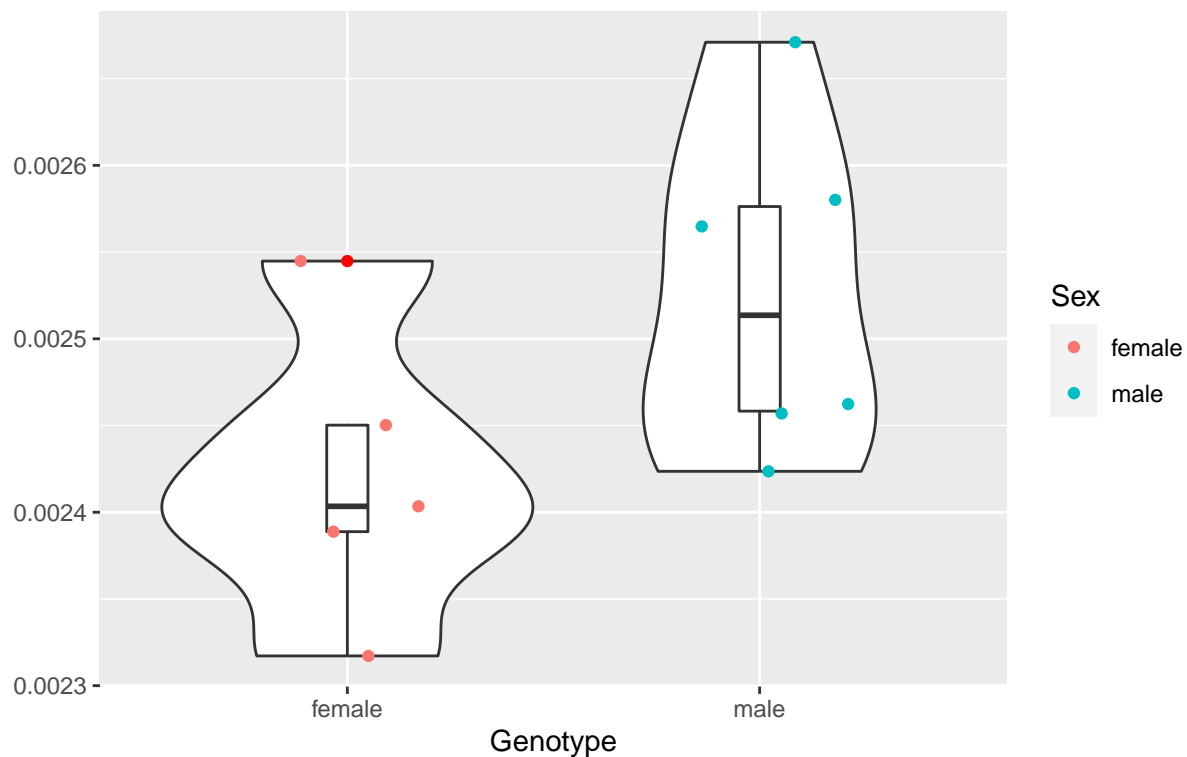
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.000e-11	8.000e-12	0.001	0.972
## Residuals	9	5.449e-08	6.054e-09		

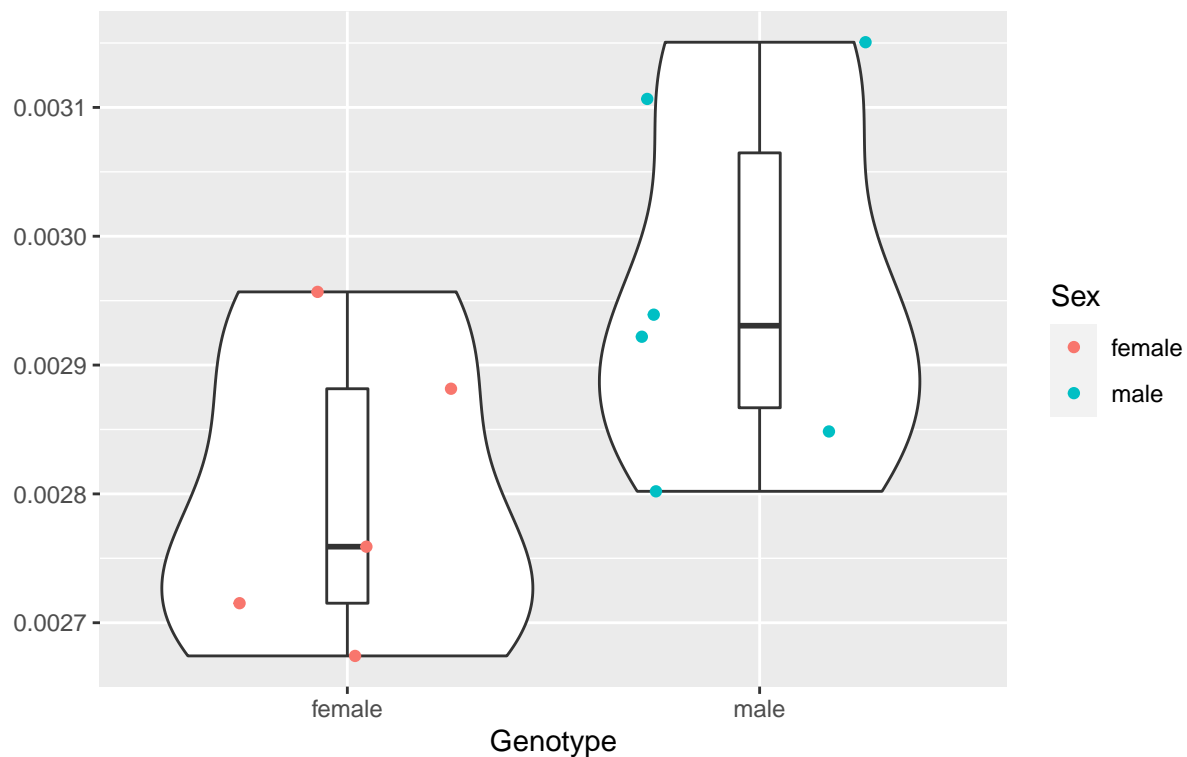
Central Gray

Red points denoting outliers



```
##           Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex         1 3.040e-08 3.040e-08   3.745  0.085 .
## Residuals   9 7.307e-08 8.119e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

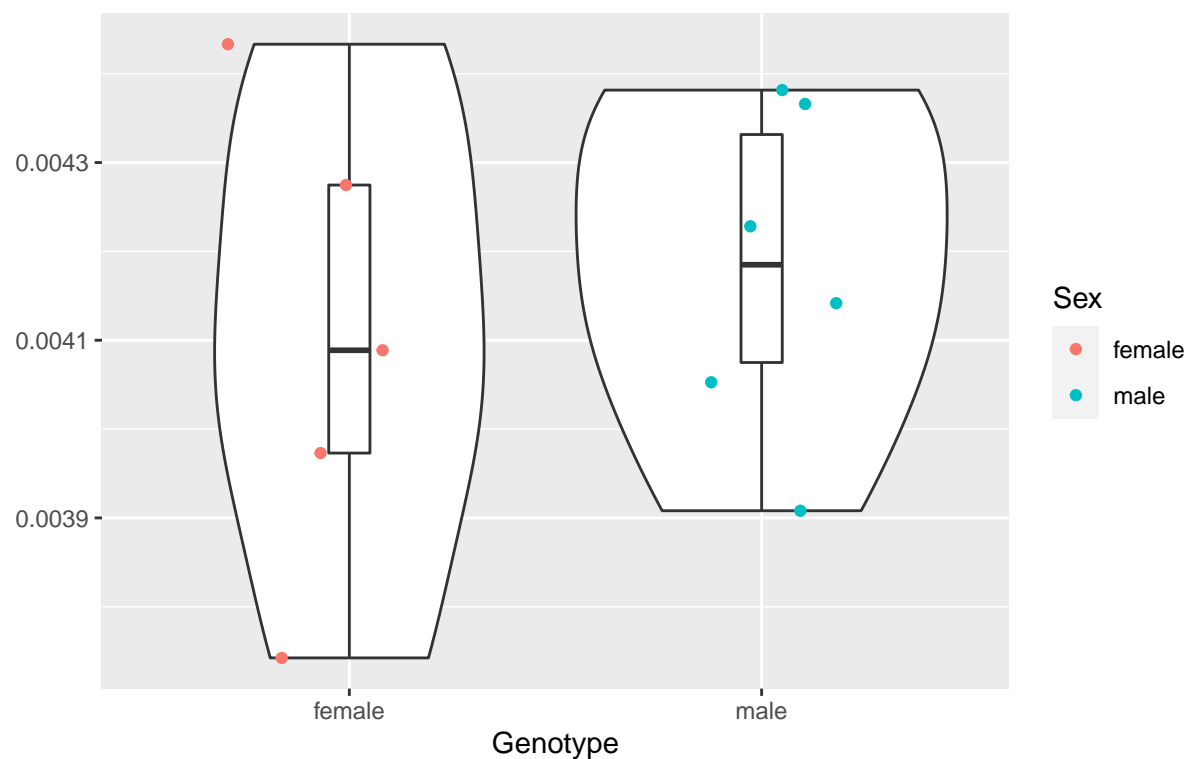
Pedunculotegmental Medial Paralemniscial and Supratrigeminal Nuclei Red points denoting outliers



```
##          Df    Sum Sq  Mean Sq F value Pr(>F)
## Sex          1 7.344e-08 7.344e-08   4.32 0.0674 .
## Residuals    9 1.530e-07 1.700e-08
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Motor Root of Trigeminal Nerve

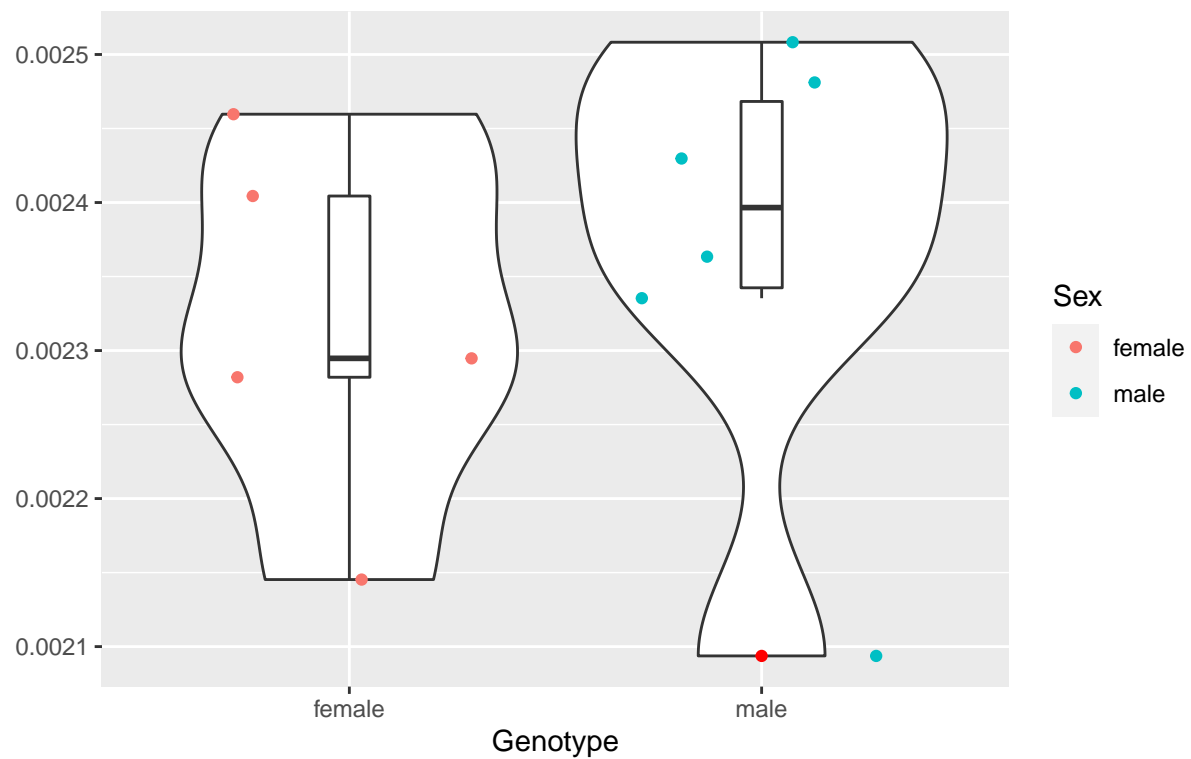
Red points denoting outliers



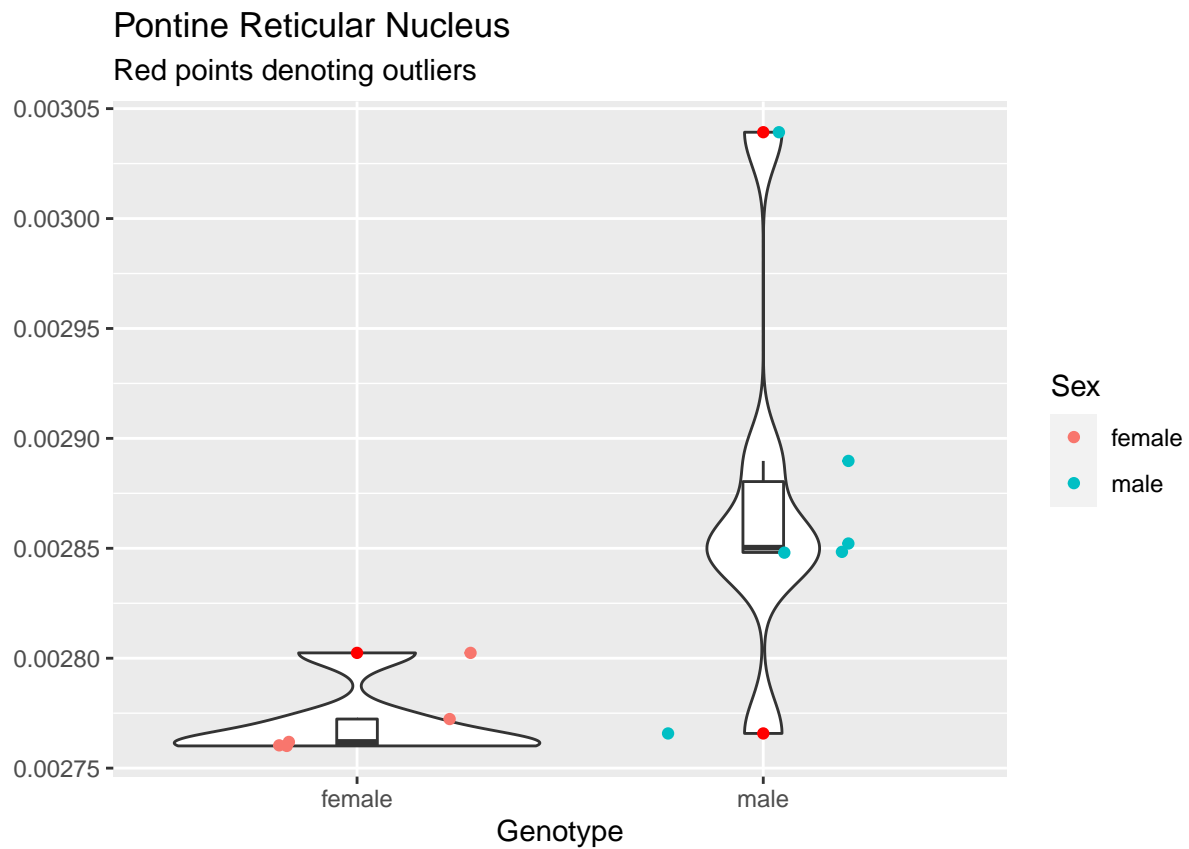
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.630e-08	1.630e-08	0.323	0.584
## Residuals	9	4.547e-07	5.052e-08		

Trigeminal Motor Nucleus

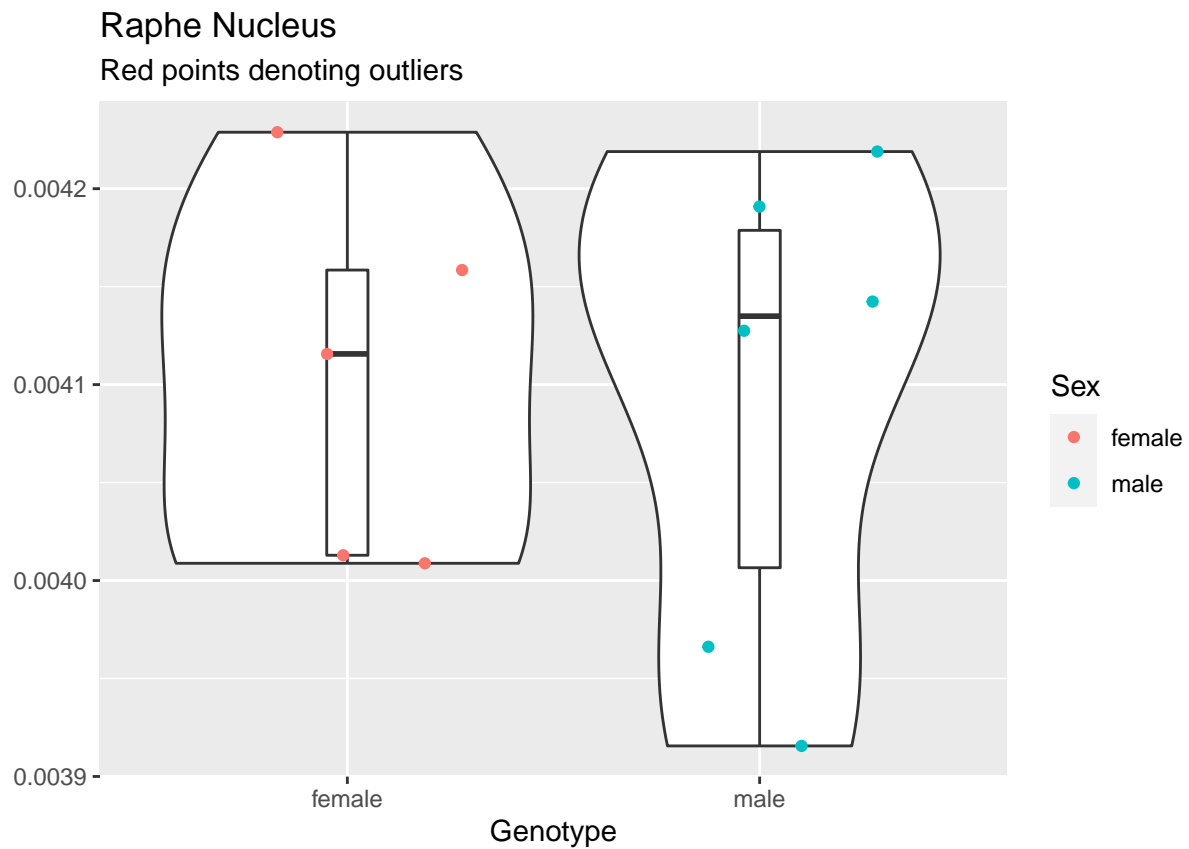
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	7.200e-09	7.204e-09	0.377	0.554
## Residuals	9	1.718e-07	1.909e-08		

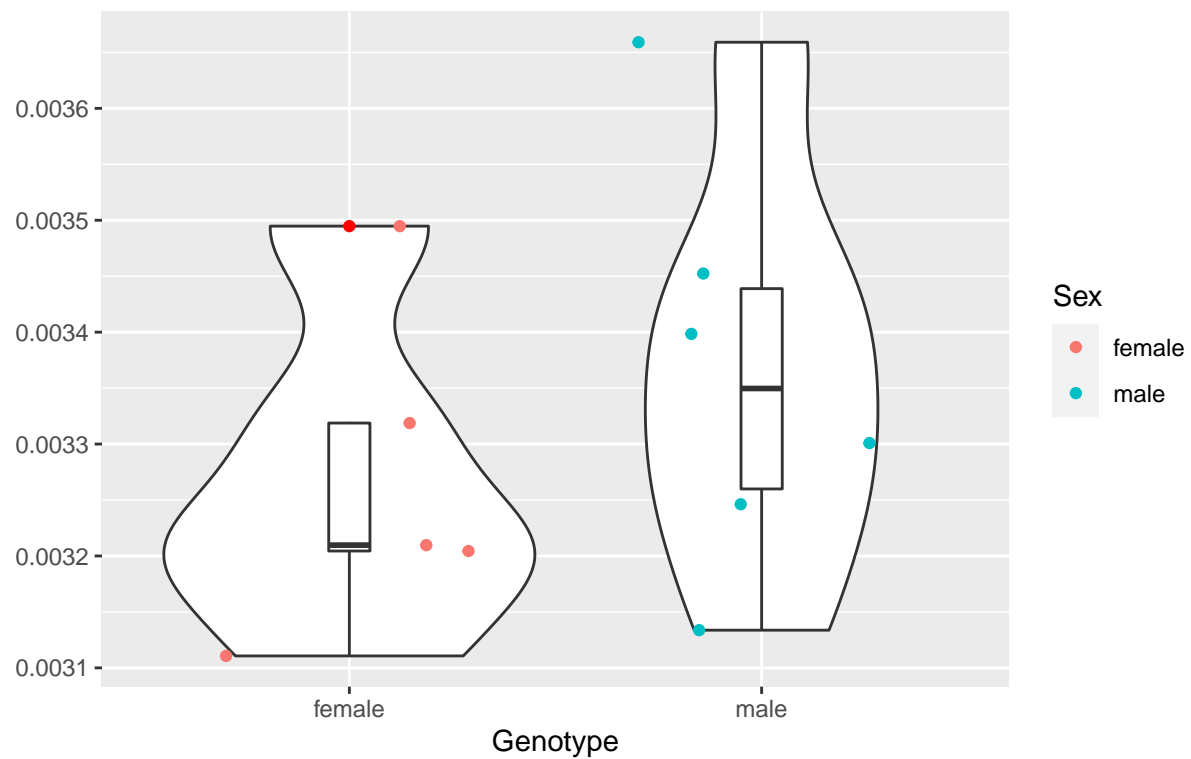


```
##          Df    Sum Sq  Mean Sq F value Pr(>F)
## Sex          1 2.863e-08 2.864e-08   6.08 0.0358 *
## Residuals    9 4.238e-08 4.709e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

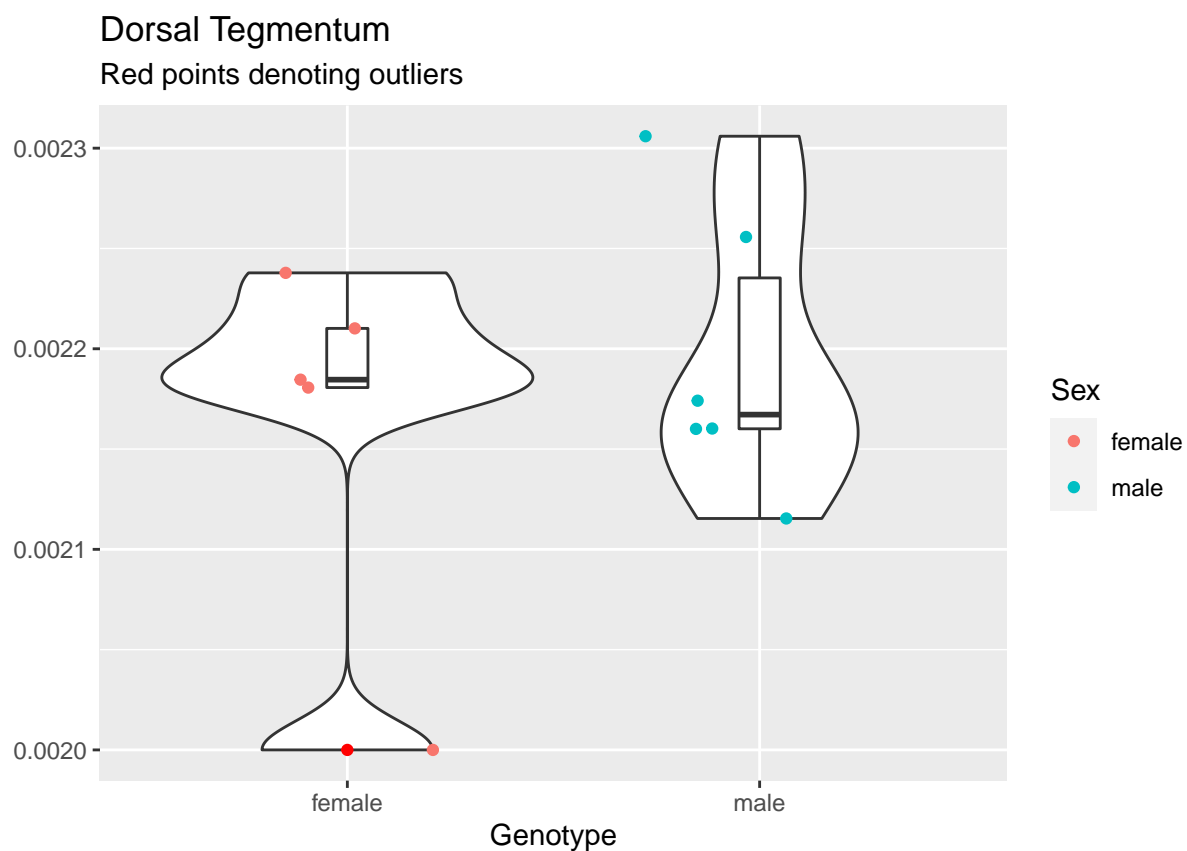


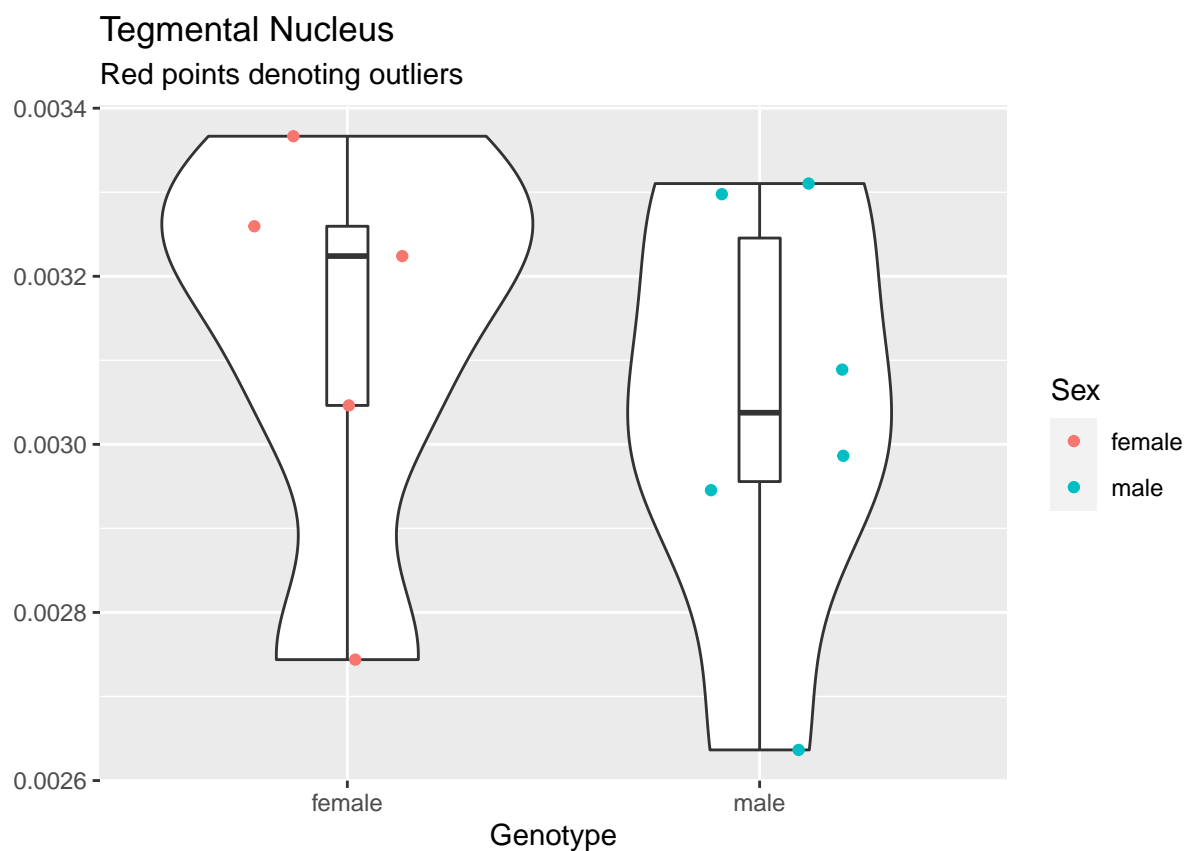
Trigeminal Sensory Nucleus

Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.591e-08	2.591e-08	0.921	0.362
## Residuals	9	2.532e-07	2.813e-08		

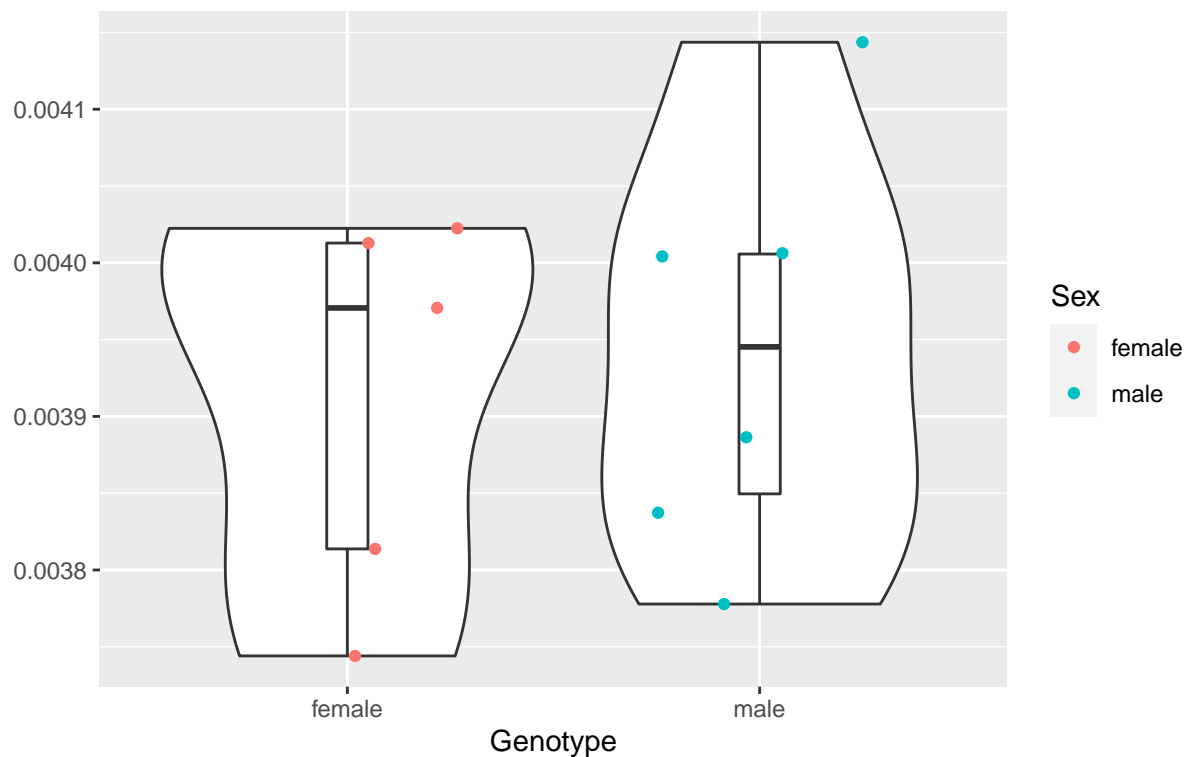




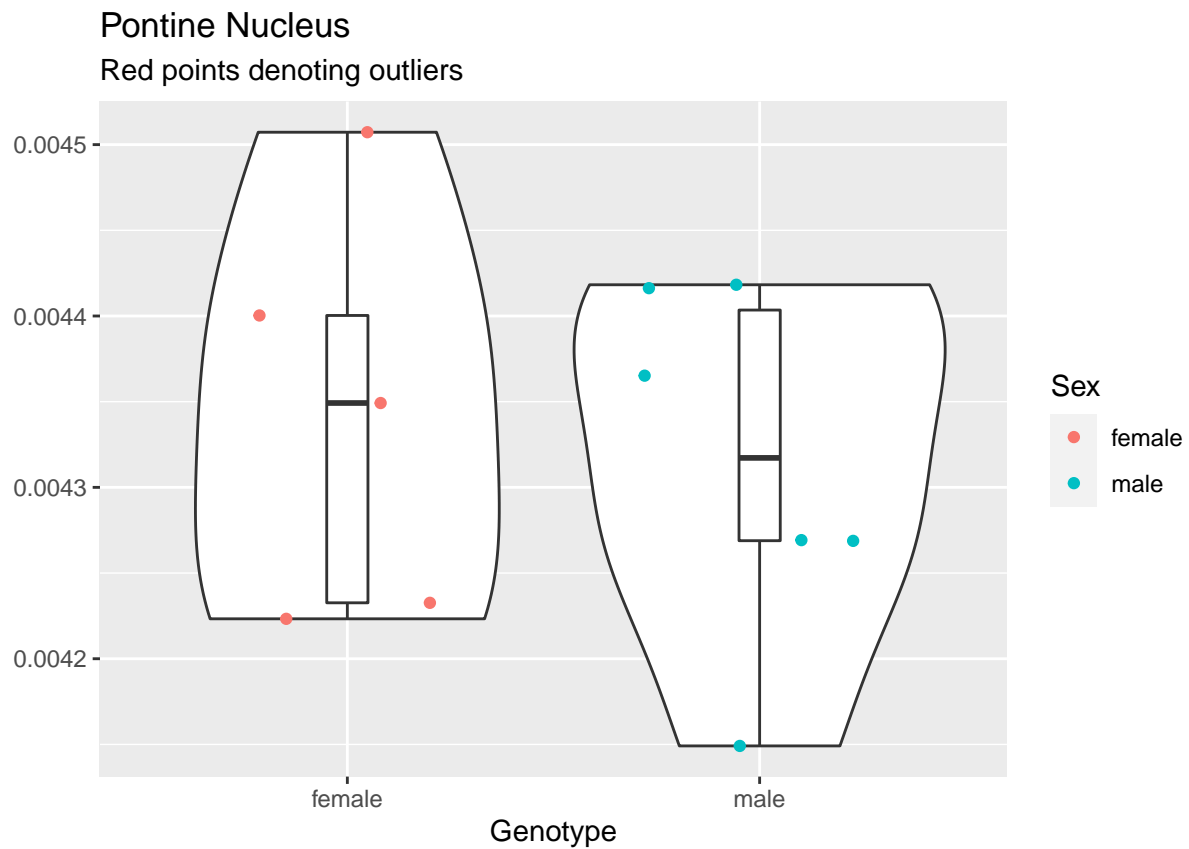
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.920e-08	1.918e-08	0.312	0.59
## Residuals	9	5.542e-07	6.157e-08		

Cochlear Nucleus

Red points denoting outliers



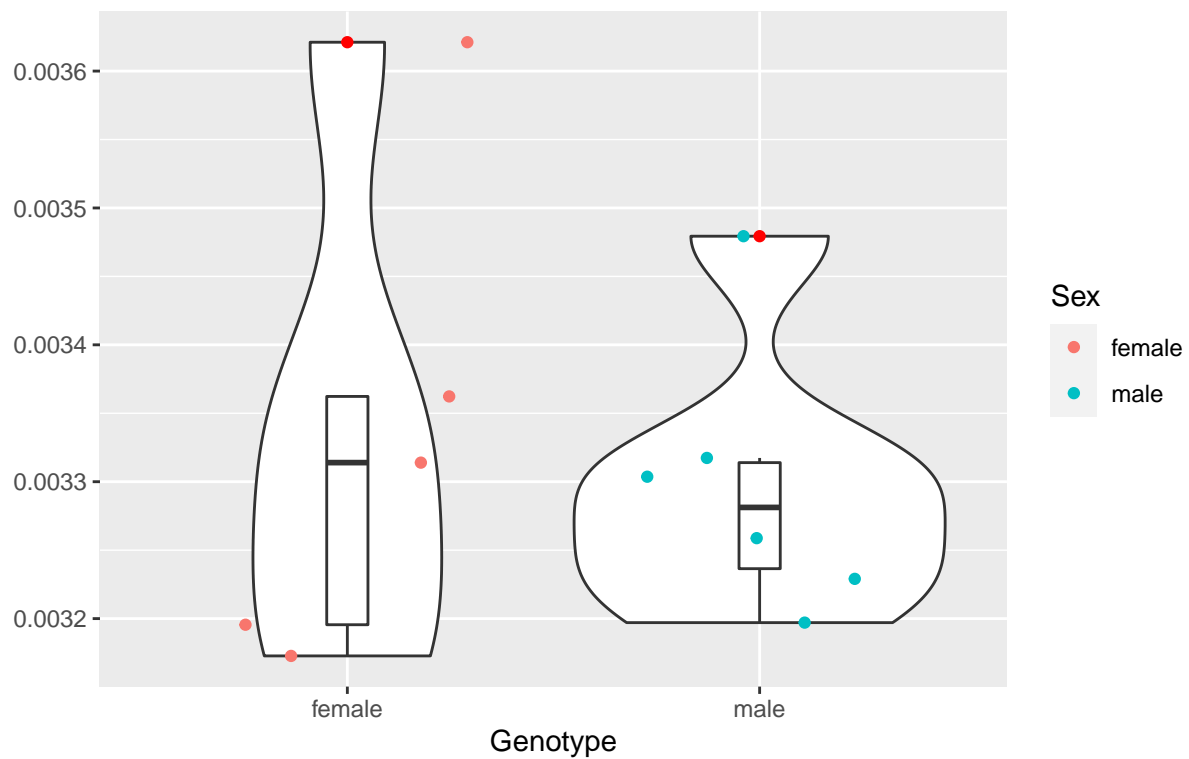
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.430e-09	2.430e-09	0.143	0.714
## Residuals	9	1.533e-07	1.704e-08		



```
##          Df  Sum Sq  Mean Sq F value Pr(>F)
## Sex        1 2.15e-09 2.150e-09   0.173  0.687
## Residuals   9 1.12e-07 1.244e-08
```

Reticulotegmental Nucleus of Pons

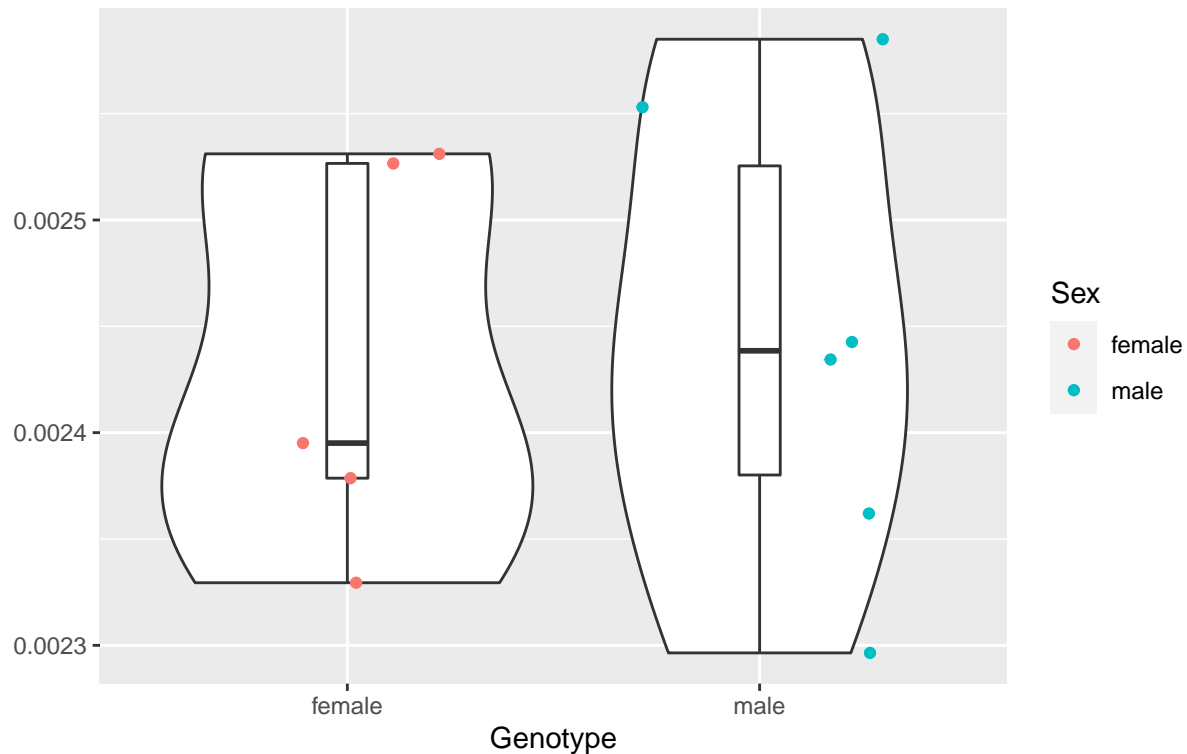
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	3.450e-09	3.452e-09	0.174	0.686
## Residuals	9	1.786e-07	1.984e-08		

Olivary Complex

Red points denoting outliers

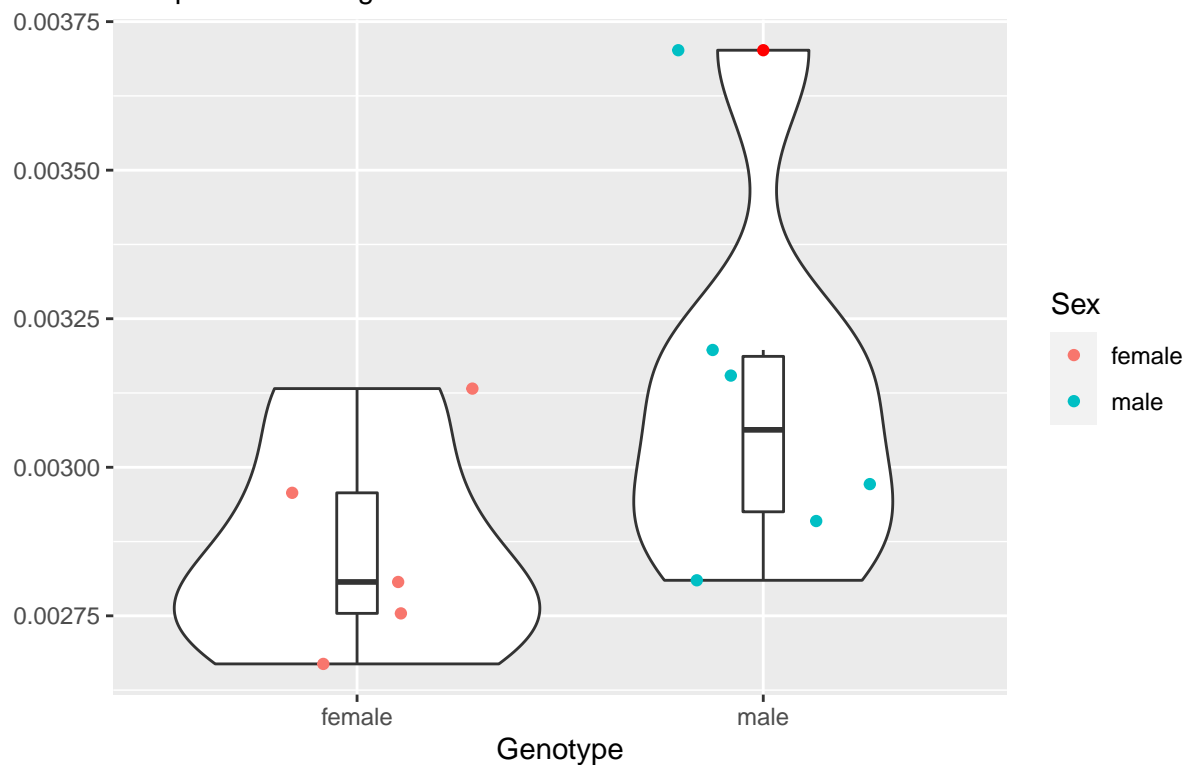


```
##           Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex         1 4.900e-10 4.910e-10   0.047  0.833
## Residuals   9 9.385e-08 1.043e-08
```

```
##“{r PnRt, echo = FALSE} #ggplot(data = apoe2, aes(factor(Sex), PnRt)) + # geom_violin() + #
geom_boxplot(width = 0.1, outlier.color = “red”) + # geom_jitter(height = 0, width = 0.3) + # labs(x =
“Genotype”, # y = “”, # title =”Pontine Reticular Nucleus“, # subtitle =”Red points denoting outliers”)
#res.aov <- aov(PnRt ~ Sex, data = apoe2) #summary(res.aov) #““
```

Spinal Trigeminal Nucleus

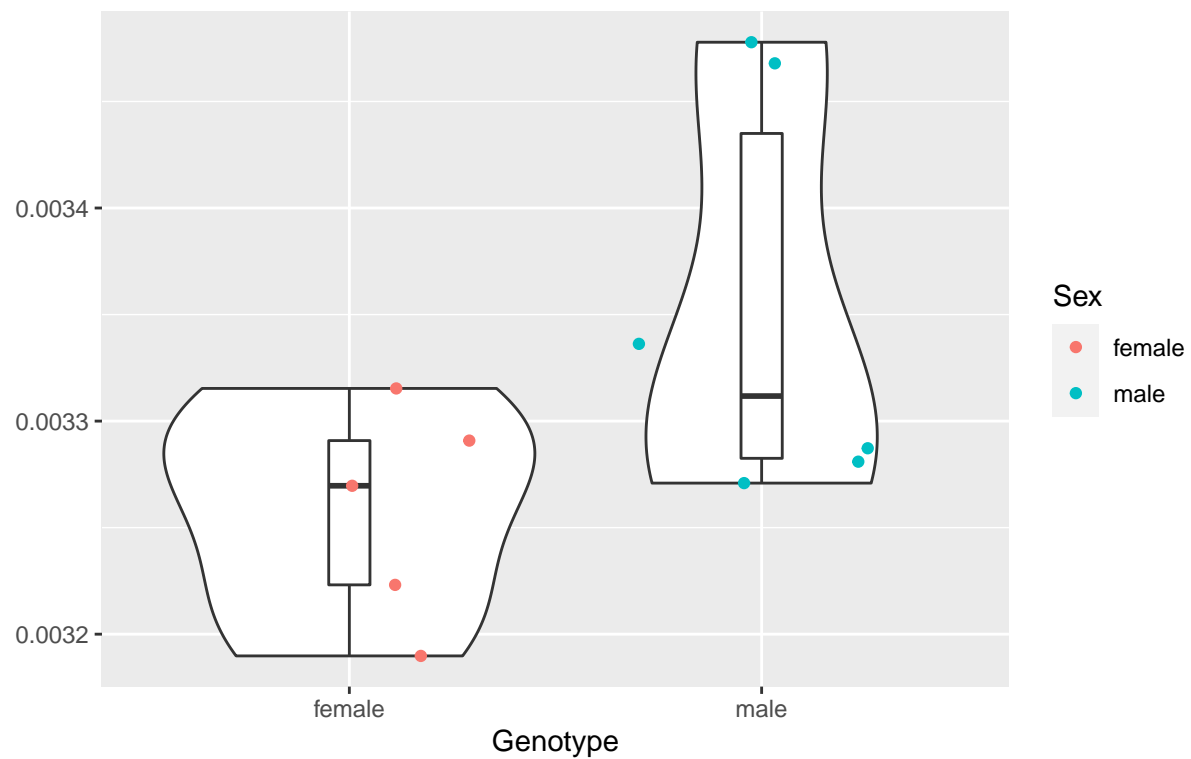
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.846e-07	1.846e-07	2.587	0.142
## Residuals	9	6.422e-07	7.136e-08		

Vestibular Nuclei

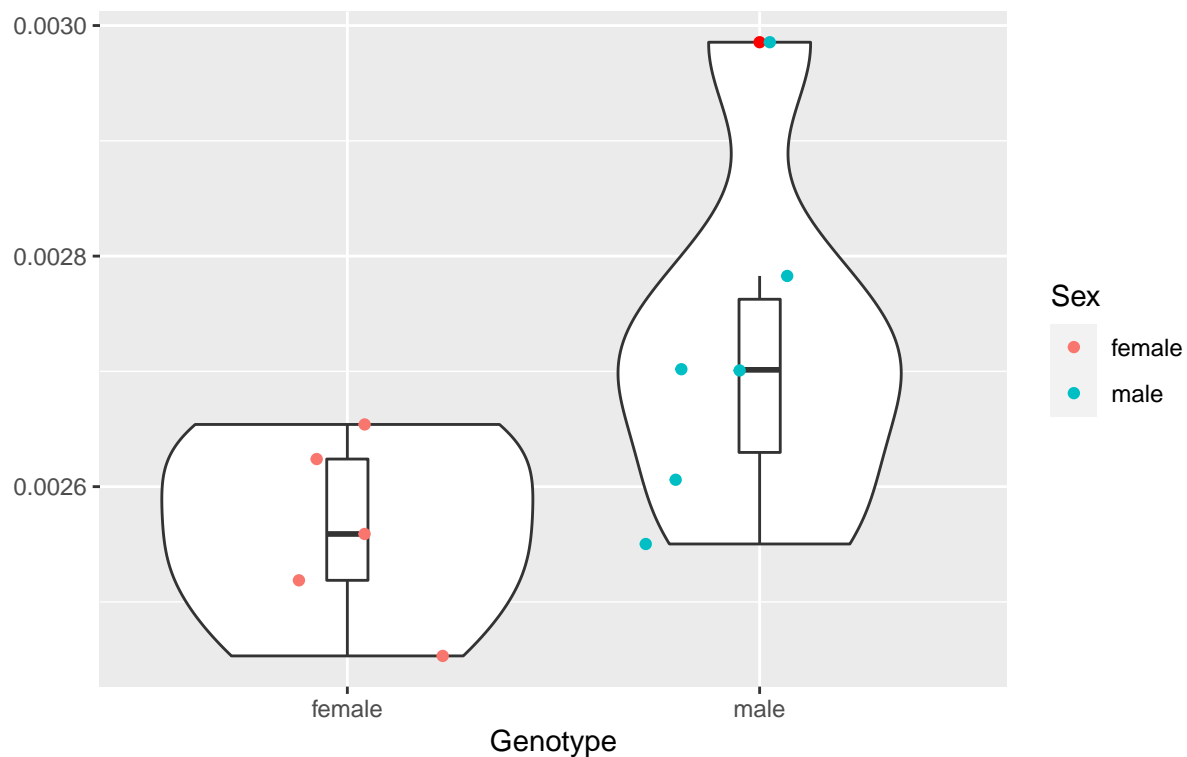
Red points denoting outliers



```
##          Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex          1 2.501e-08 2.501e-08   4.043 0.0753 .
## Residuals    9 5.568e-08 6.187e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```


Gigantocellular Reticular Nucleus

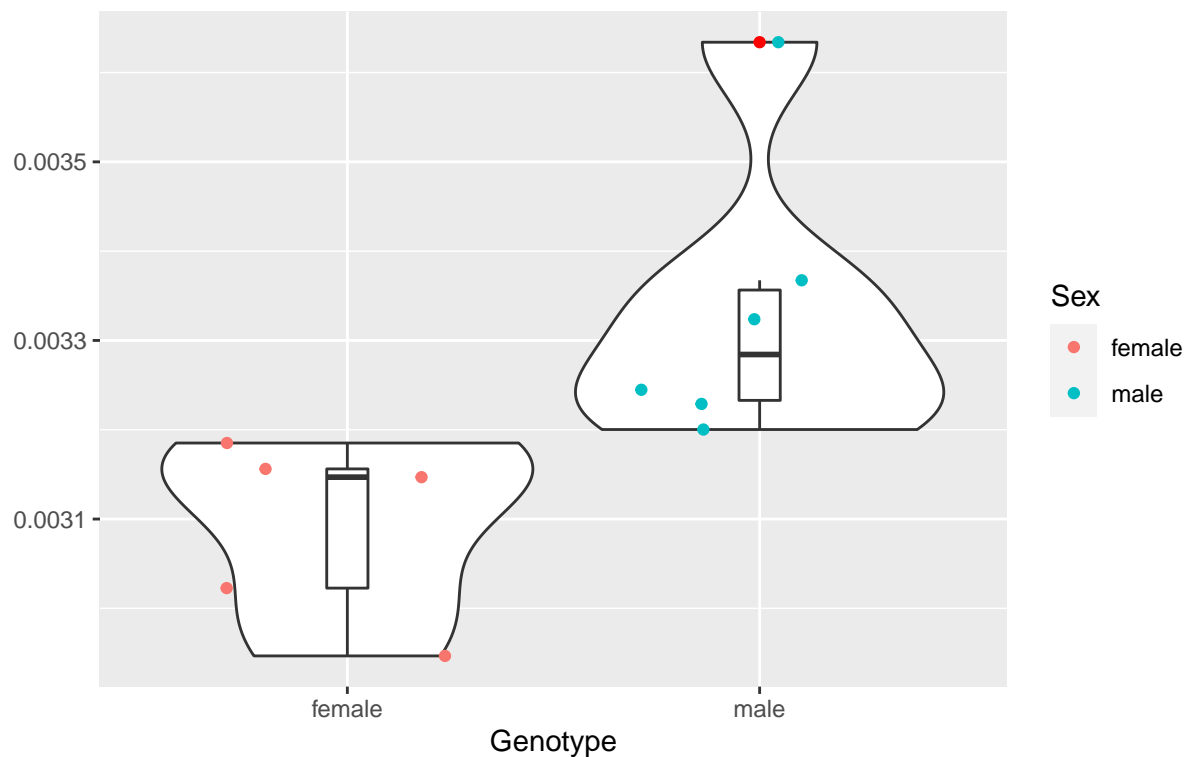
Red points denoting outliers



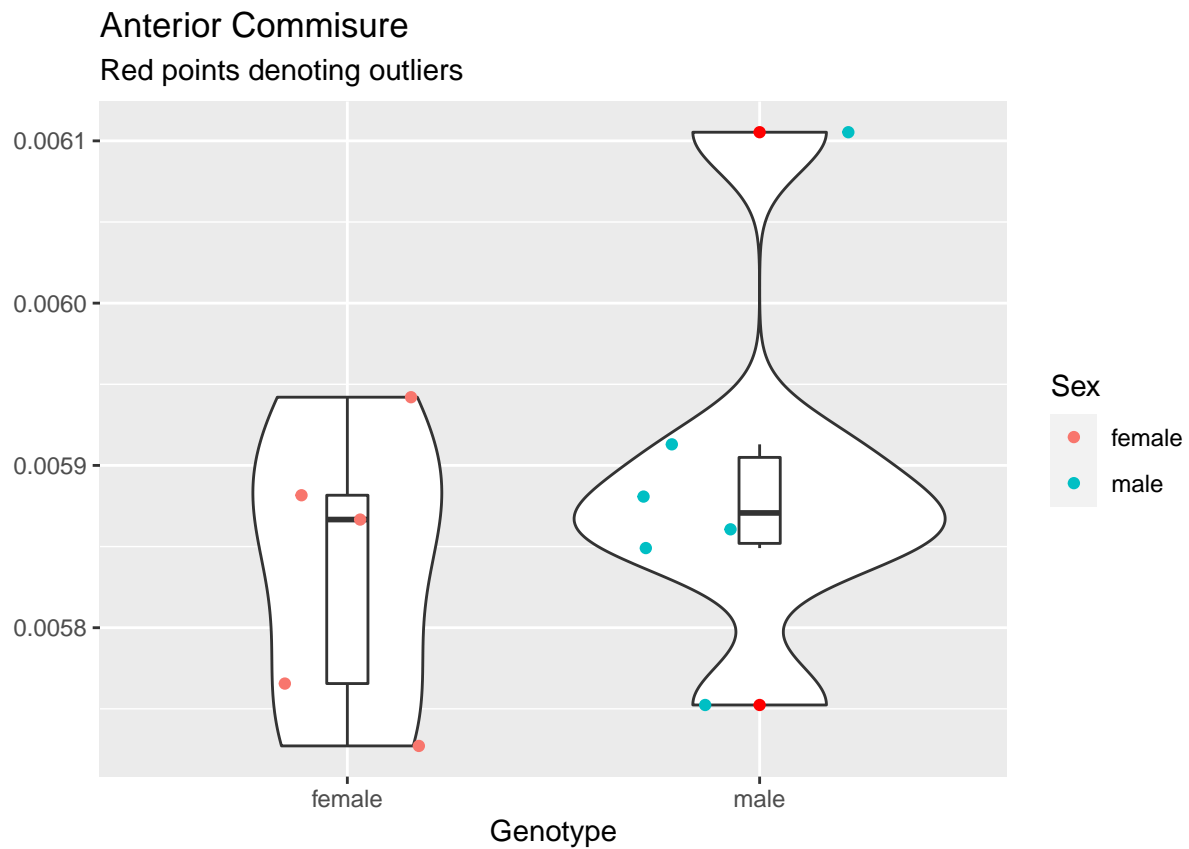
```
##           Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex         1 6.934e-08 6.934e-08   4.365 0.0663 .
## Residuals   9 1.430e-07 1.588e-08
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Cuneate Nucleus

Red points denoting outliers



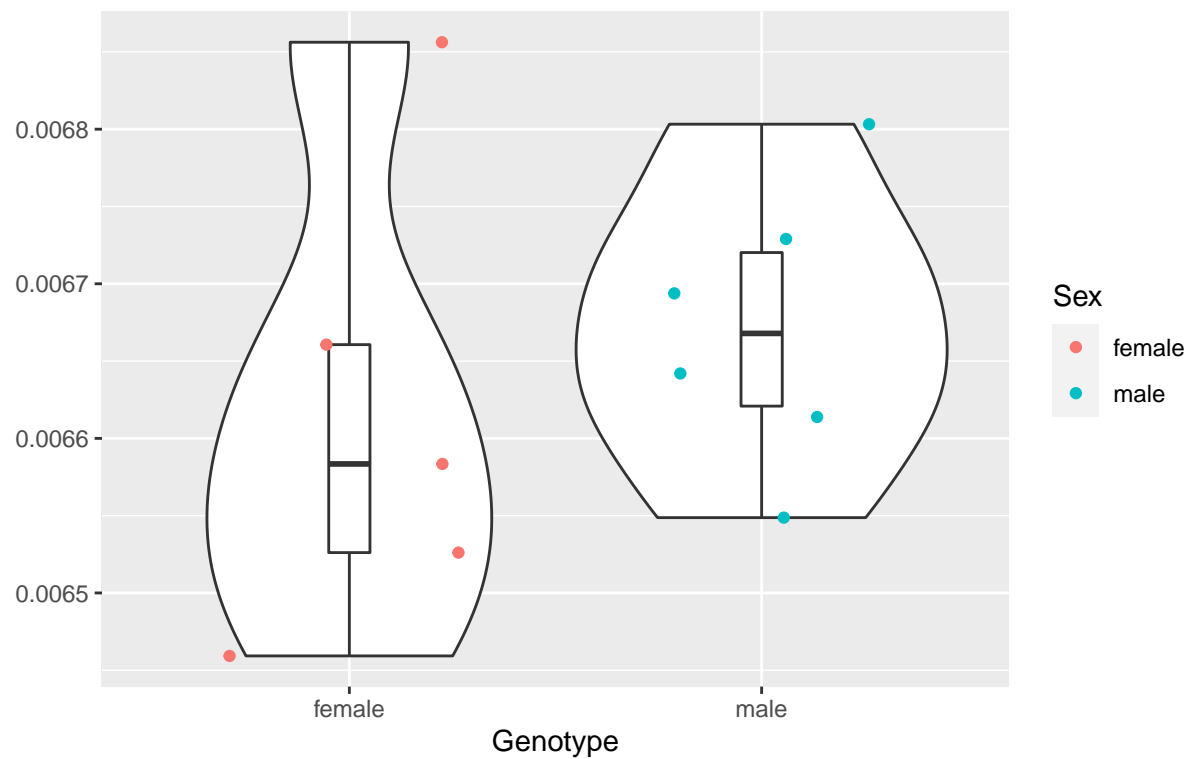
```
##           Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex           1 1.593e-07 1.593e-07   8.441 0.0174 *
## Residuals     9 1.698e-07 1.887e-08
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```



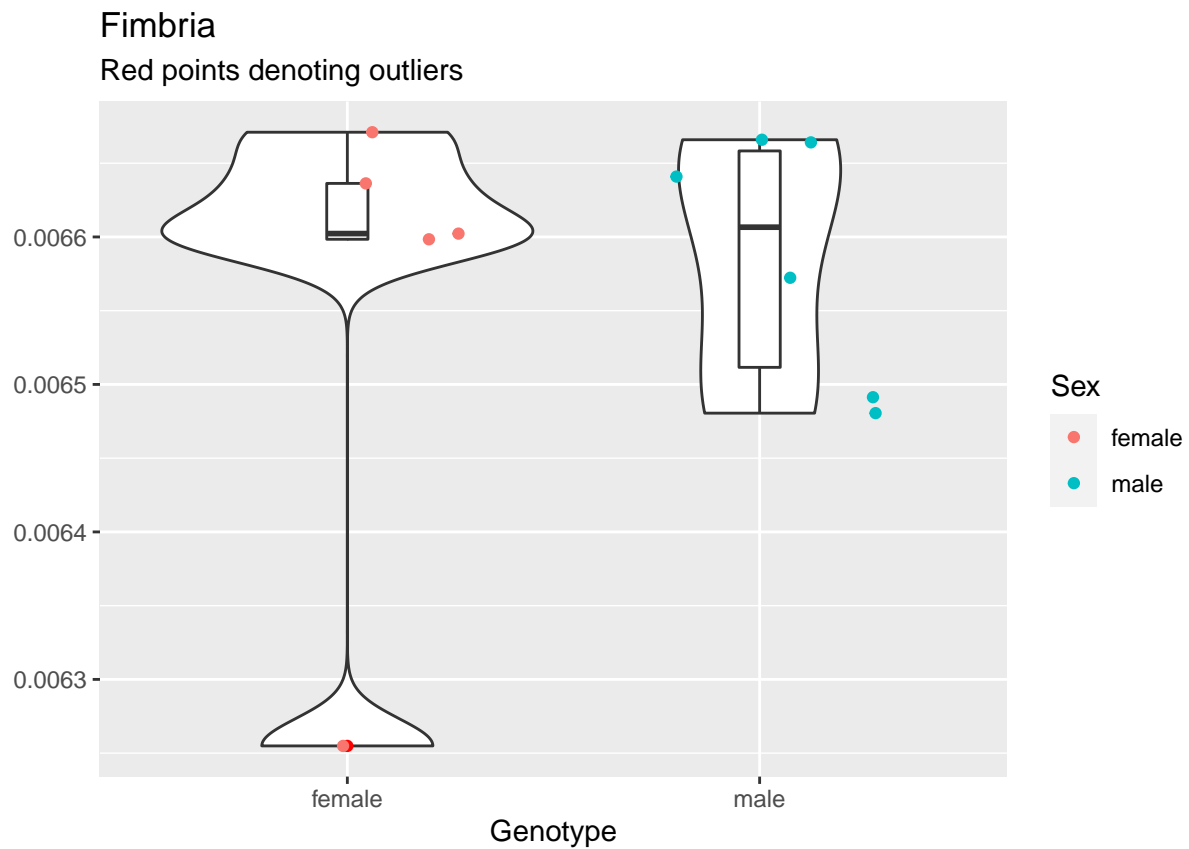
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	8.840e-09	8.837e-09	0.8	0.394
## Residuals	9	9.945e-08	1.105e-08		

Optic Tracts

Red points denoting outliers



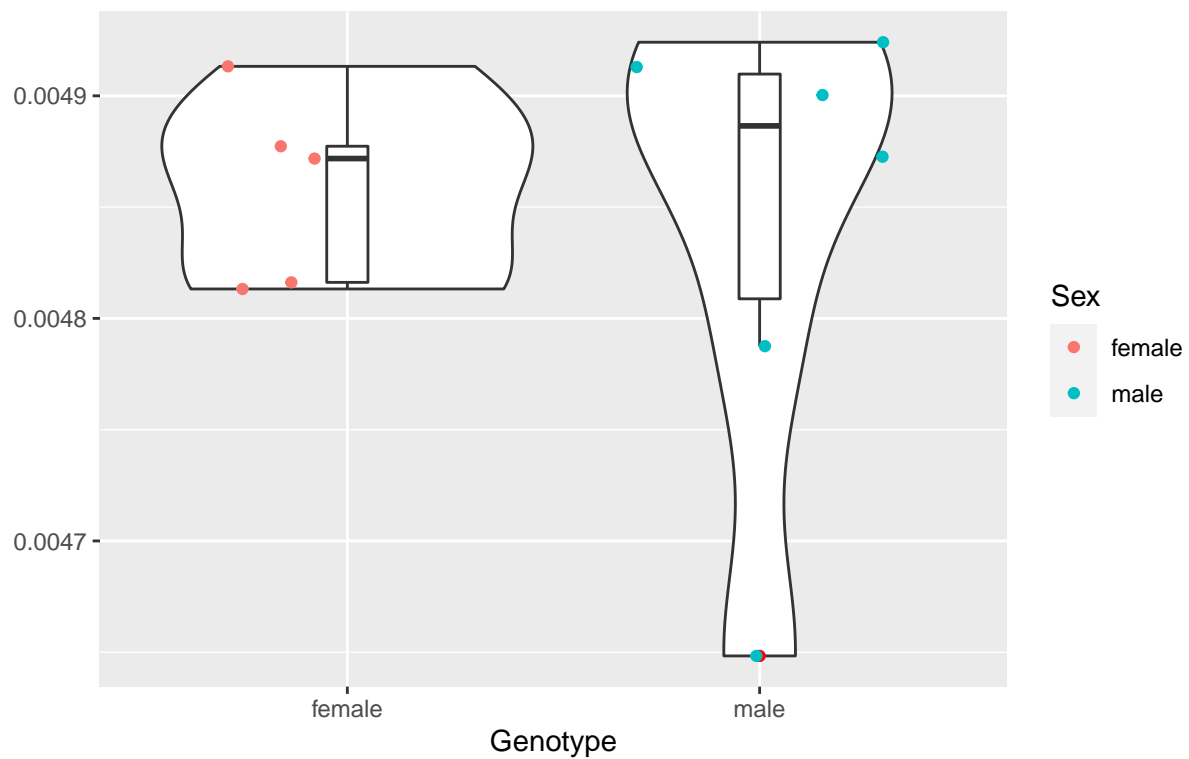
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	8.130e-09	8.134e-09	0.547	0.478
## Residuals	9	1.338e-07	1.487e-08		



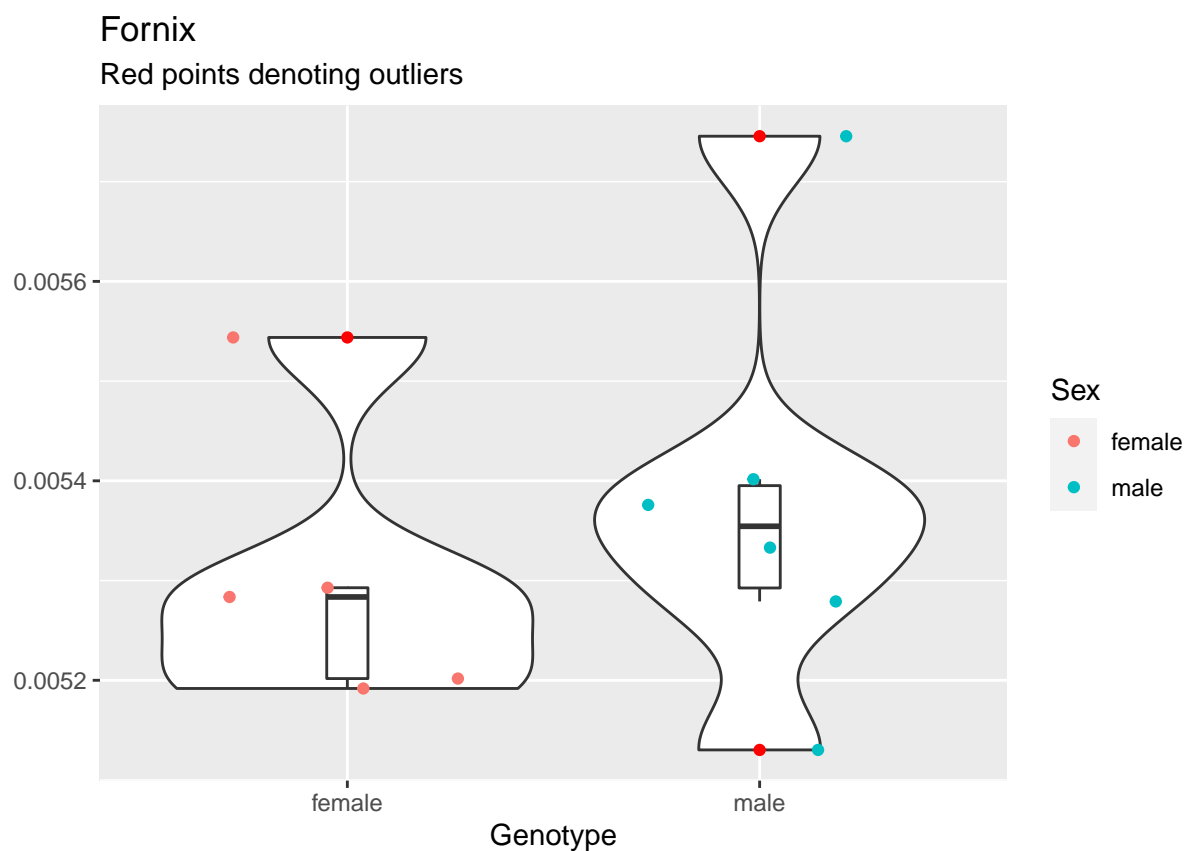
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	3.02e-09	3.020e-09	0.181	0.68
## Residuals	9	1.50e-07	1.667e-08		

Corpus Callosum

Red points denoting outliers



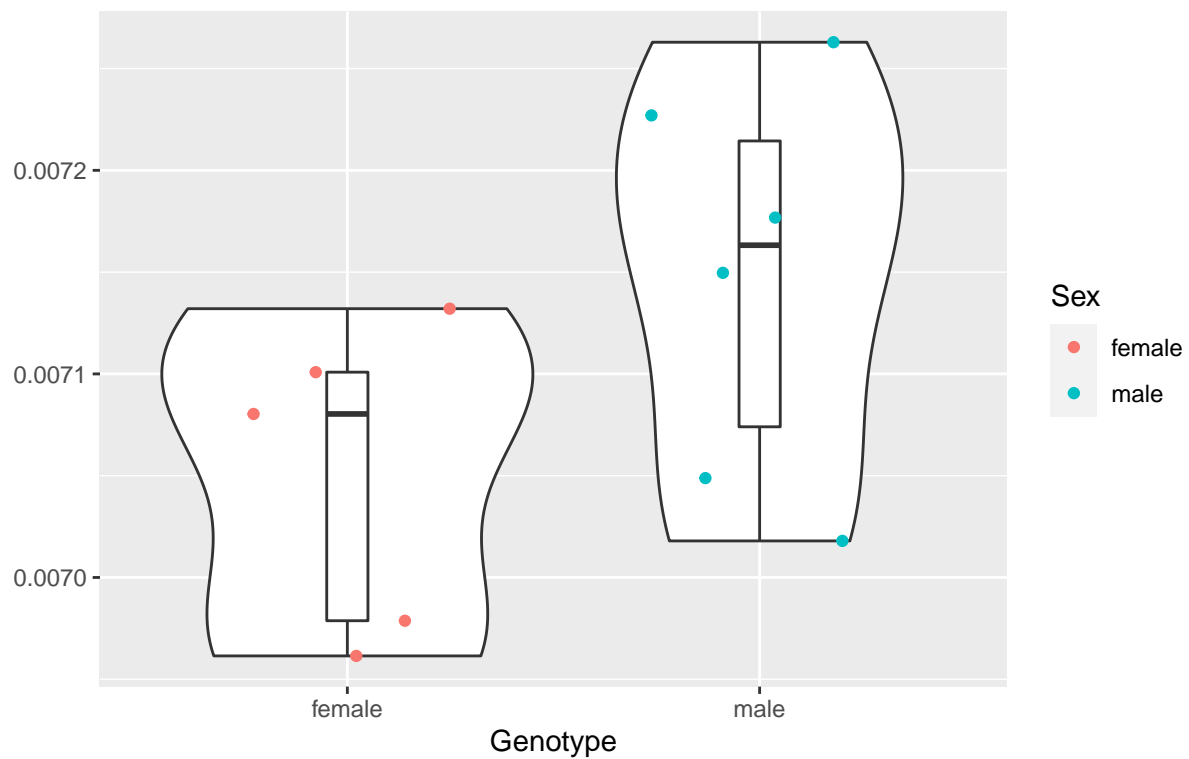
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	8.200e-10	8.240e-10	0.116	0.741
## Residuals	9	6.395e-08	7.105e-09		



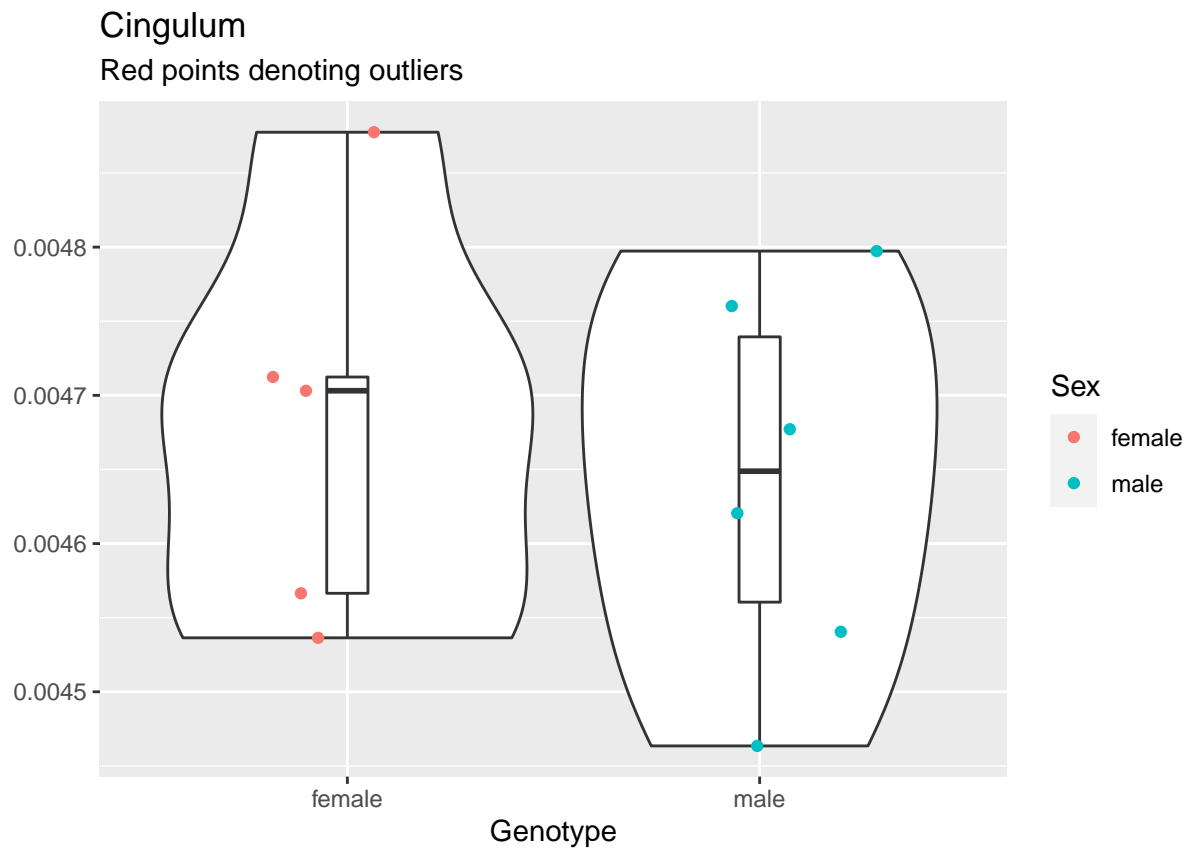
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.527e-08	1.527e-08	0.474	0.509
## Residuals	9	2.900e-07	3.222e-08		

Stria Terminalis

Red points denoting outliers



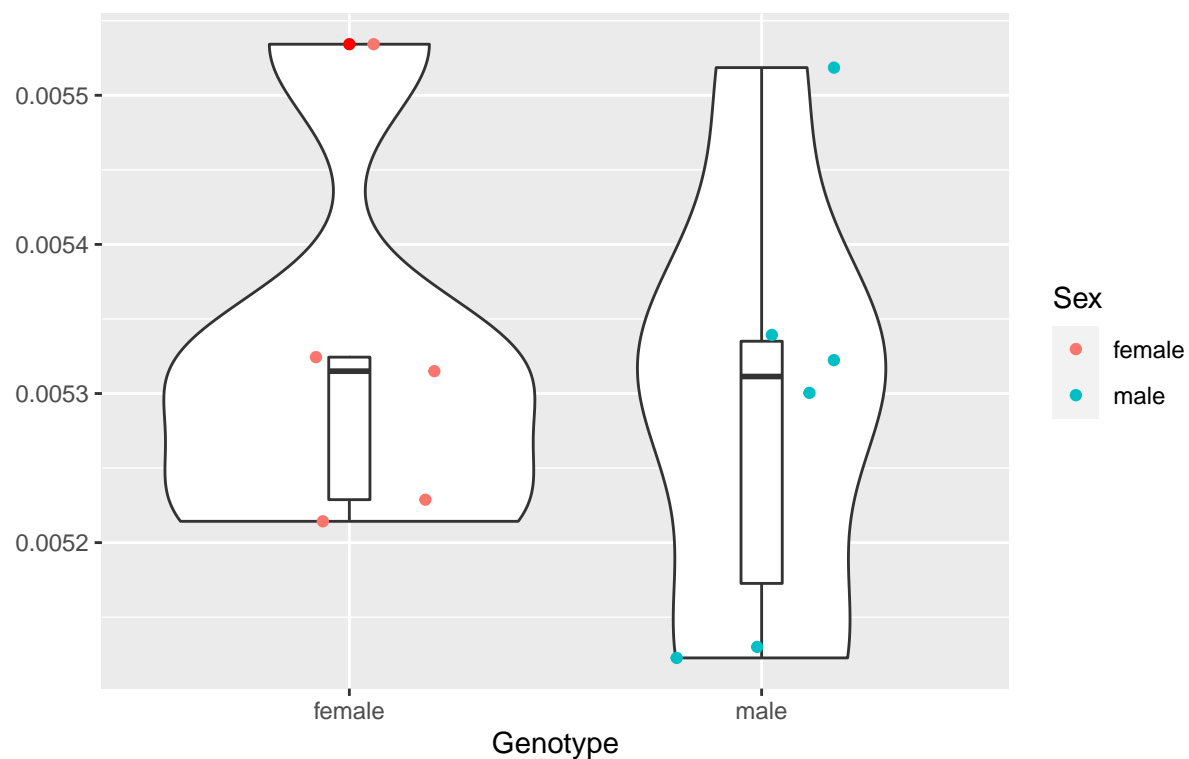
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.540e-08	2.540e-08	3.257	0.105
## Residuals	9	7.017e-08	7.797e-09		



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	3.530e-09	3.530e-09	0.204	0.663
## Residuals	9	1.561e-07	1.734e-08		

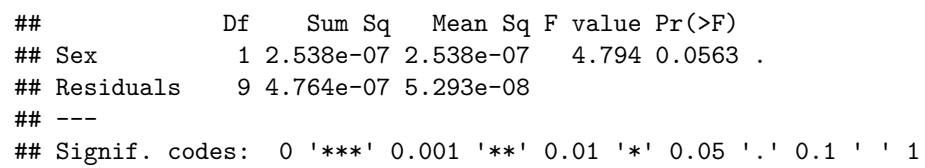
Lateral Olfactory Tract

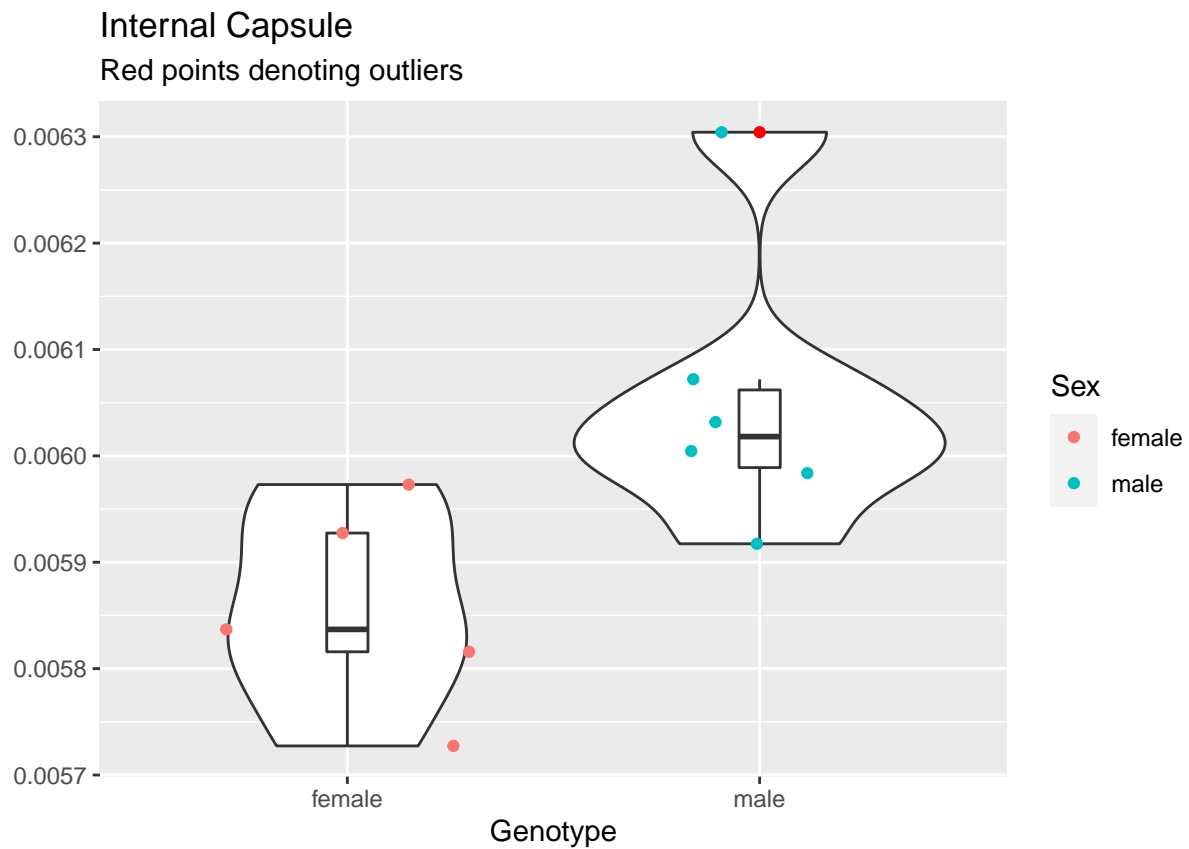
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	3.240e-09	3.236e-09	0.167	0.693
## Residuals	9	1.748e-07	1.942e-08		

Red points denoting outliers

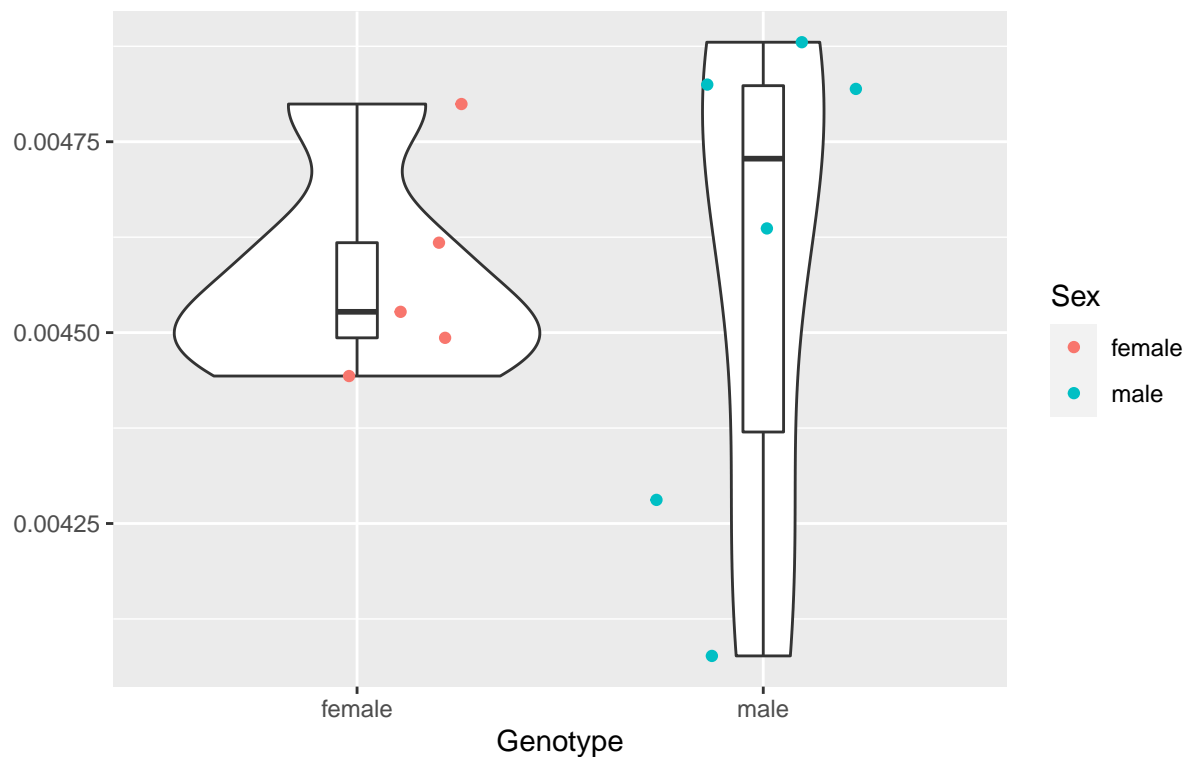




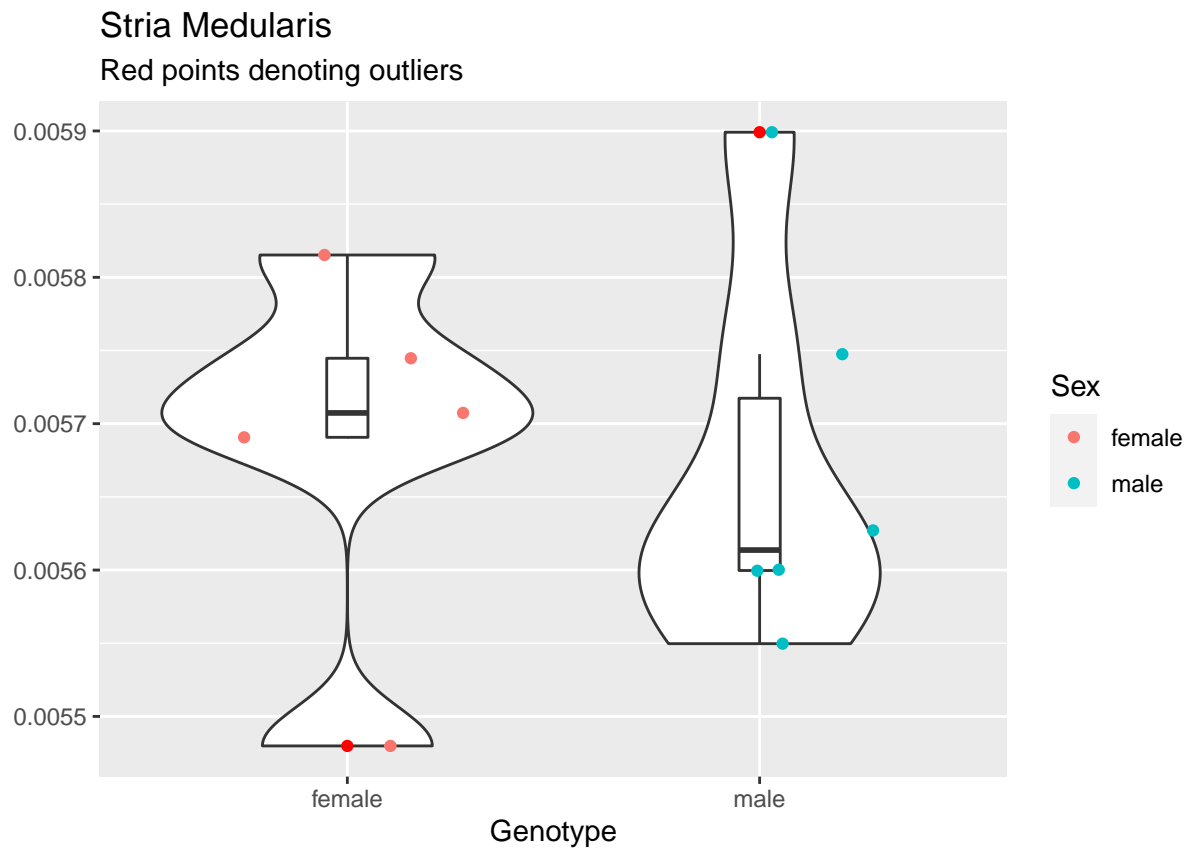
```
##           Df    Sum Sq  Mean Sq F value Pr(>F)
## Sex           1 1.050e-07 1.050e-07   7.45 0.0232 *
## Residuals     9 1.268e-07 1.409e-08
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Fasciculus Retroflexus

Red points denoting outliers

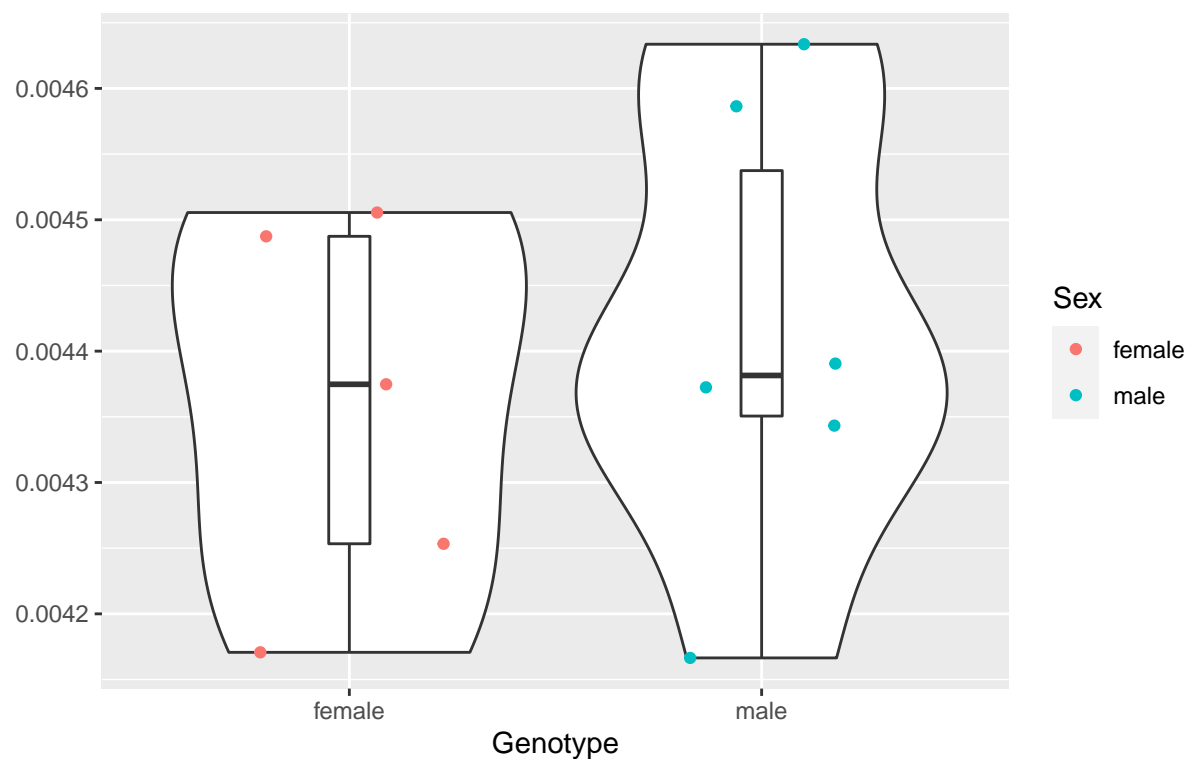


##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	3.000e-10	2.900e-10	0.004	0.95
## Residuals	9	6.316e-07	7.018e-08		



Mammillothalamic Tract

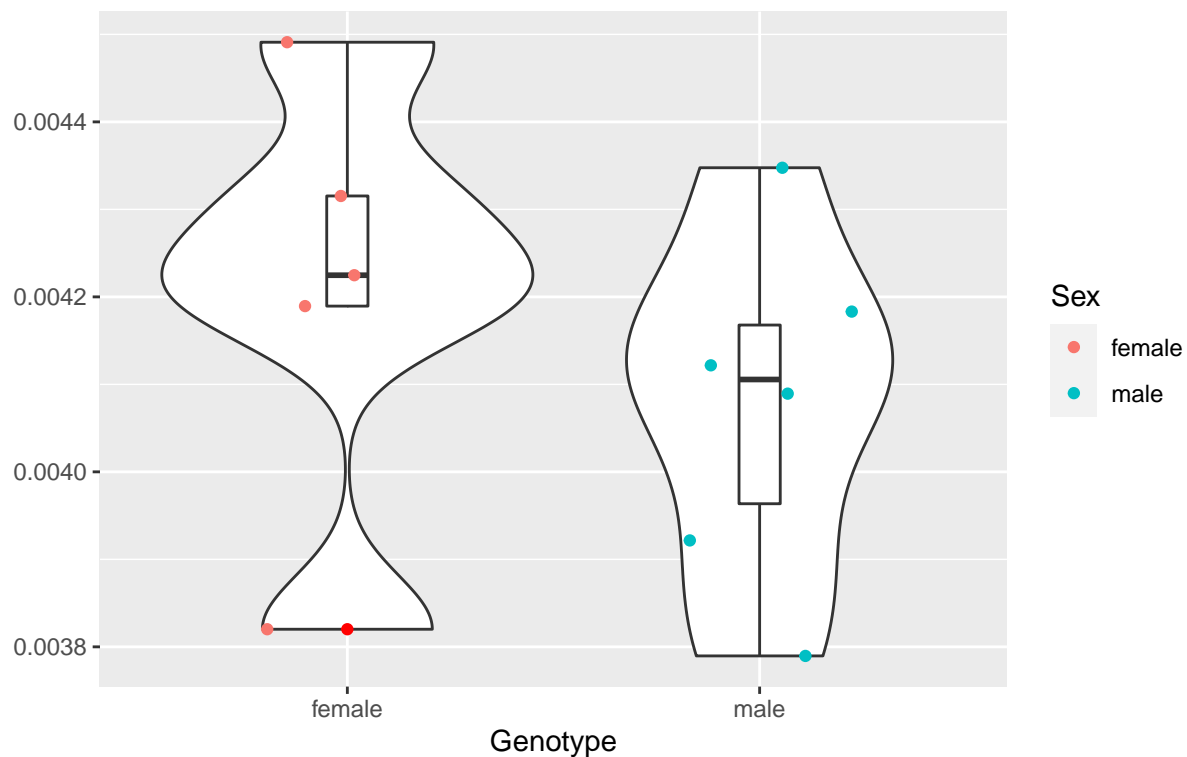
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	8.900e-09	8.896e-09	0.346	0.571
## Residuals	9	2.313e-07	2.570e-08		

Posterior Commissure

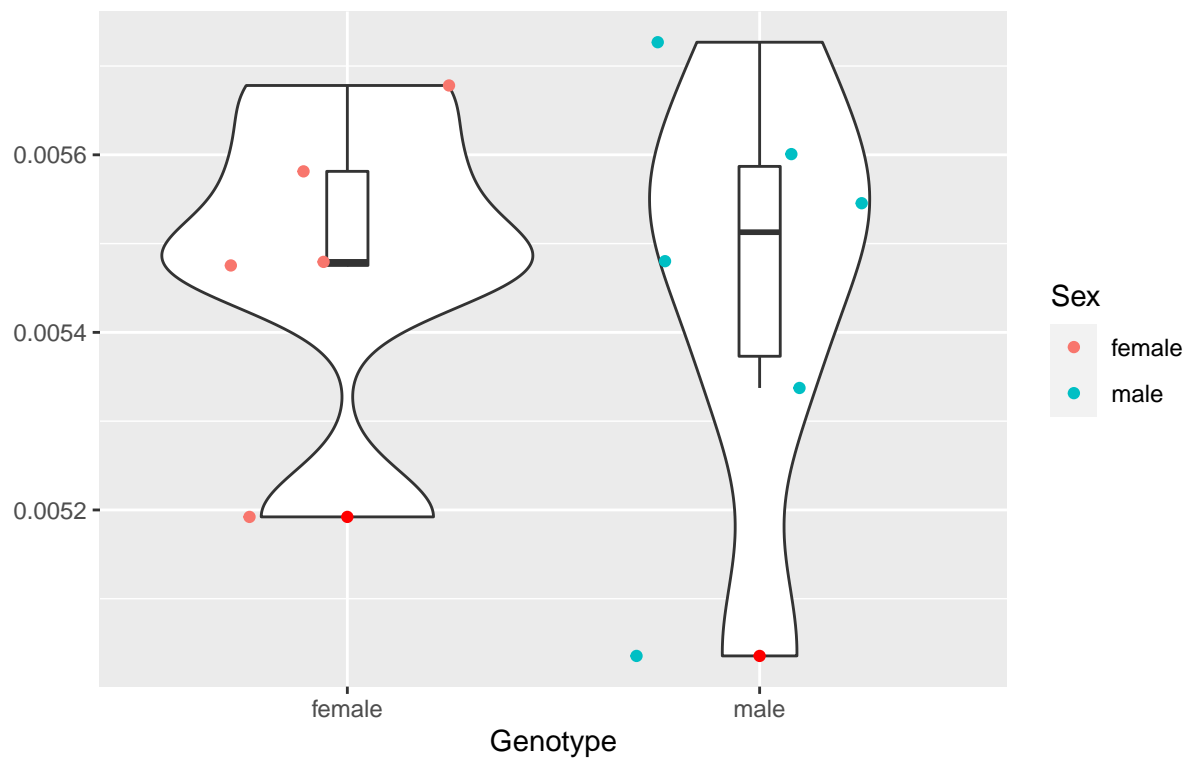
Red points denoting outliers



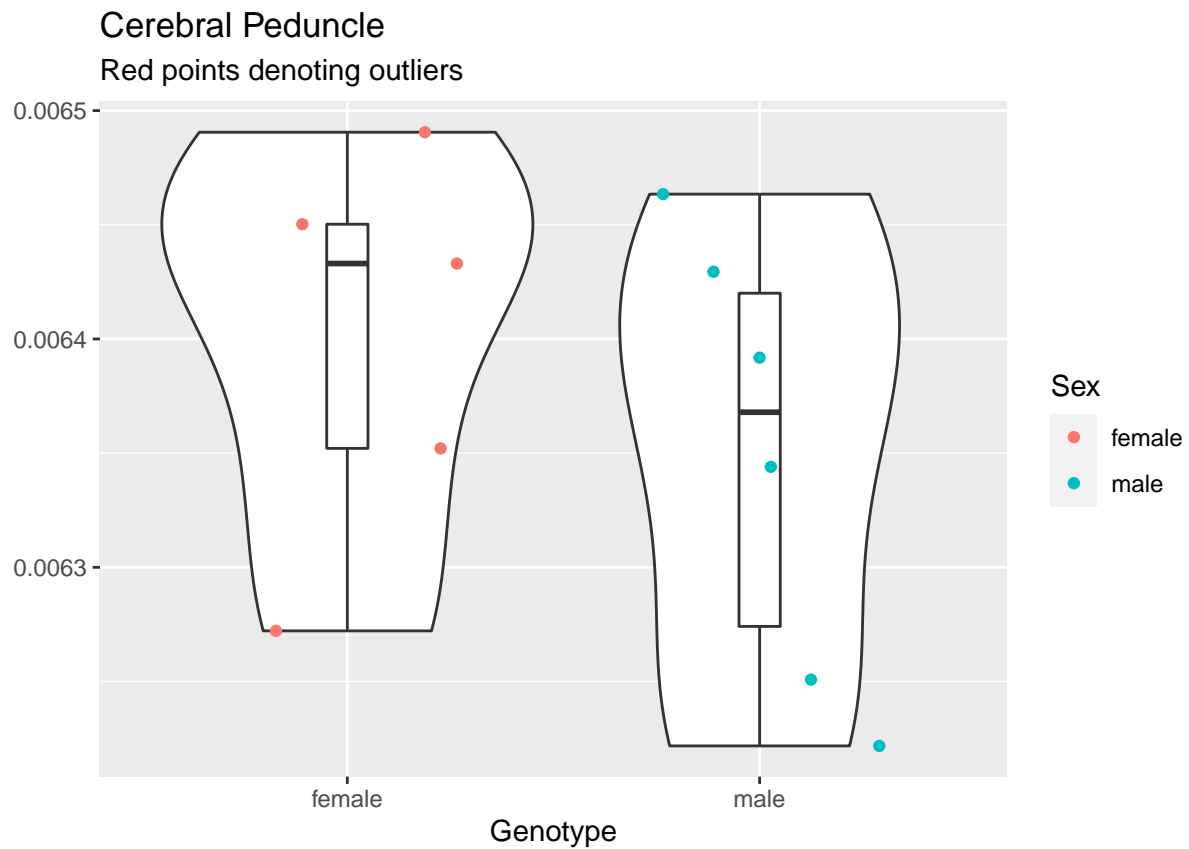
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	4.80e-08	4.796e-08	0.99	0.346
## Residuals	9	4.36e-07	4.845e-08		

Brachium of Superior Colliculus

Red points denoting outliers



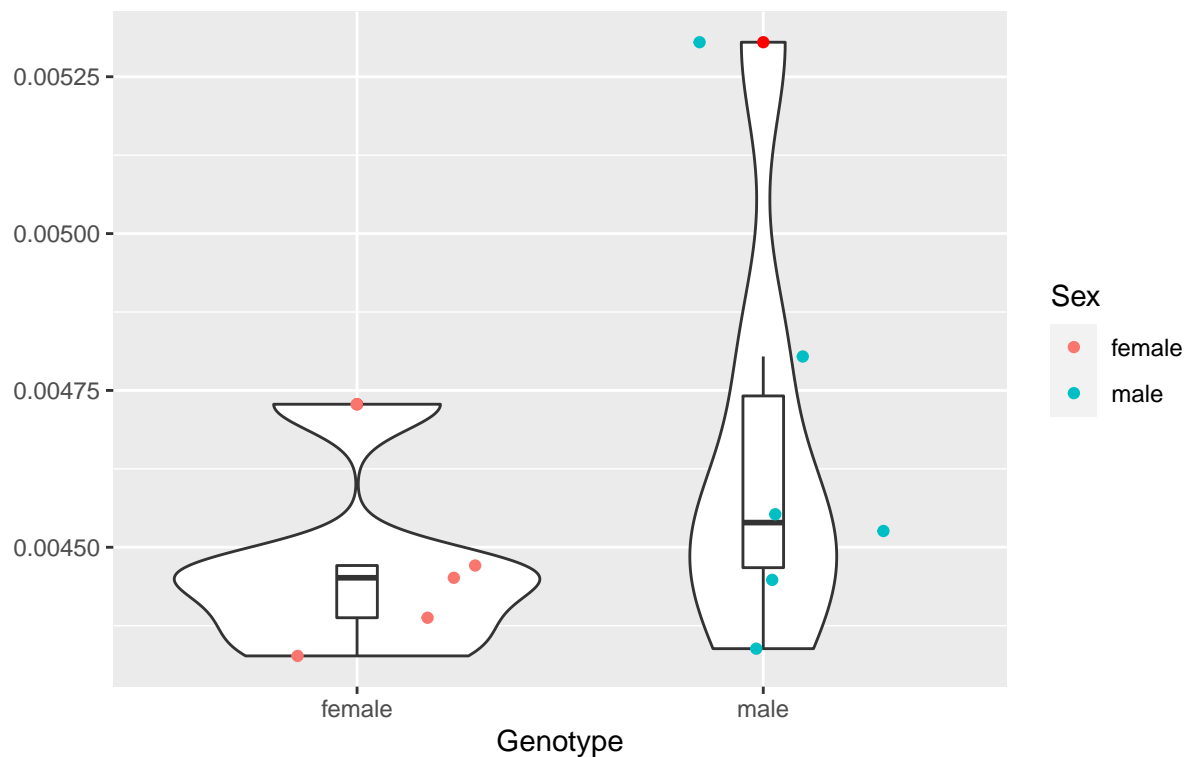
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.000e-09	1.980e-09	0.042	0.843
## Residuals	9	4.261e-07	4.734e-08		



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	6.660e-09	6.655e-09	0.771	0.403
## Residuals	9	7.769e-08	8.632e-09		

Lateral Lemniscus

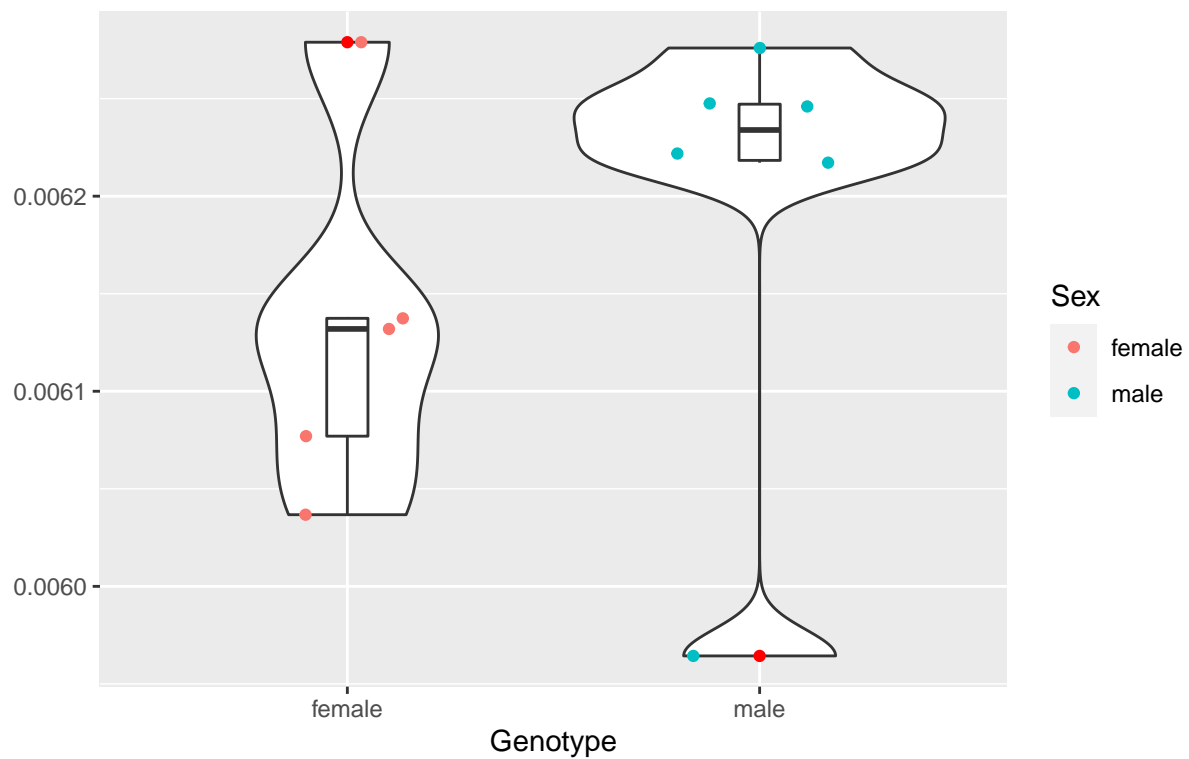
Red points denoting outliers



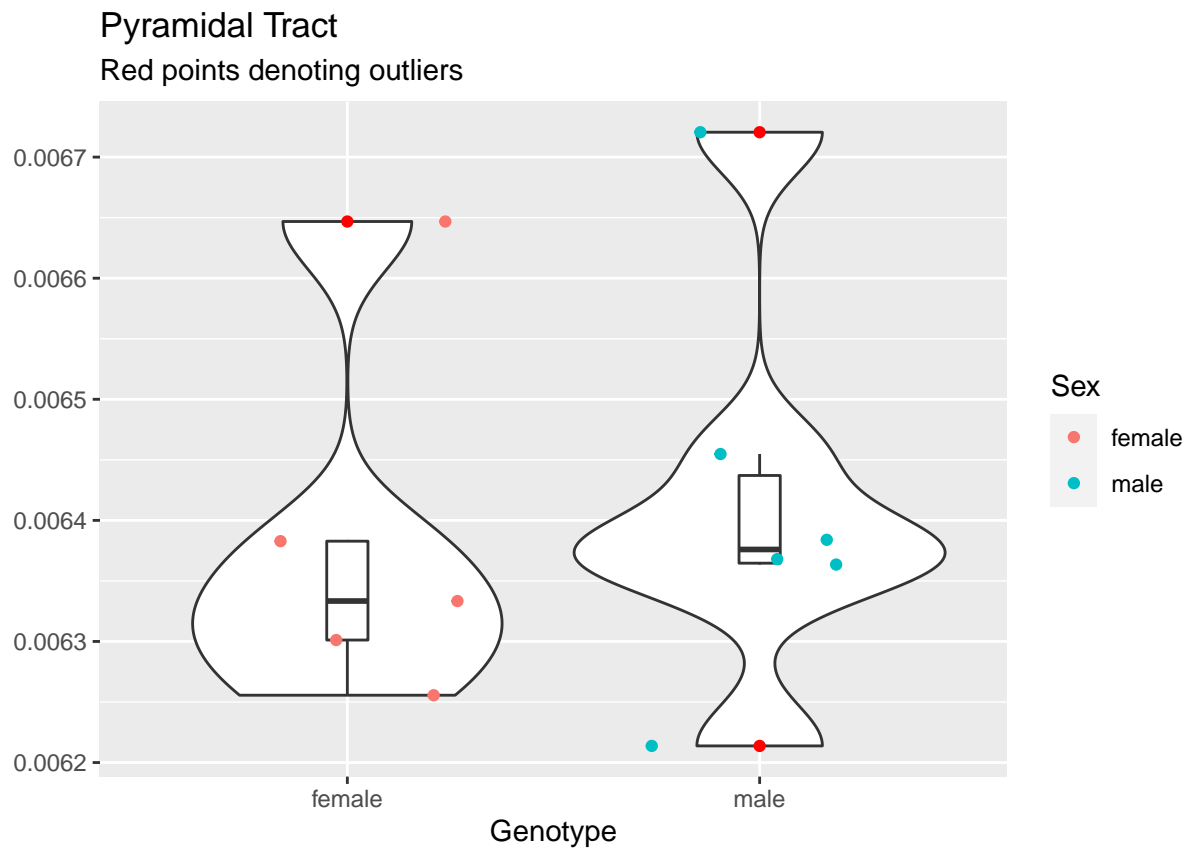
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	9.780e-08	9.783e-08	1.242	0.294
## Residuals	9	7.088e-07	7.875e-08		

Spinal Trigeminal Nerve

Red points denoting outliers

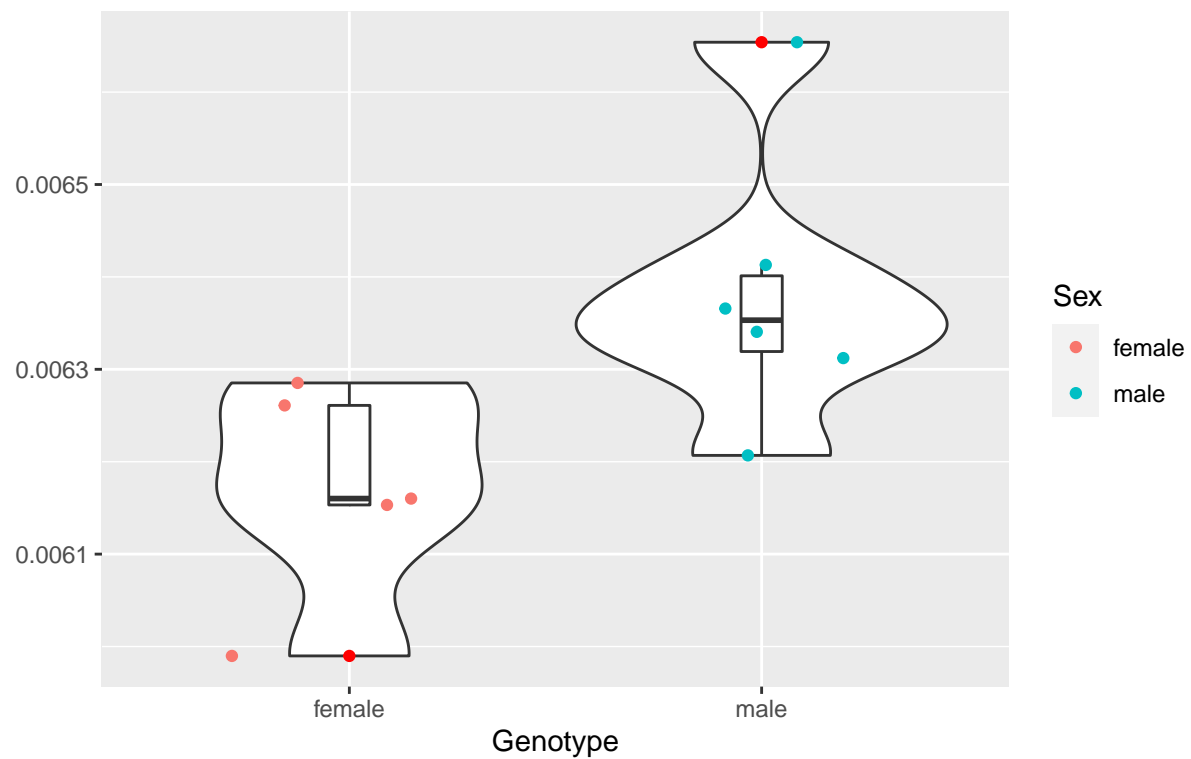


##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.085e-08	1.085e-08	0.976	0.349
## Residuals	9	1.000e-07	1.112e-08		



Vestibulocochlear Nerve

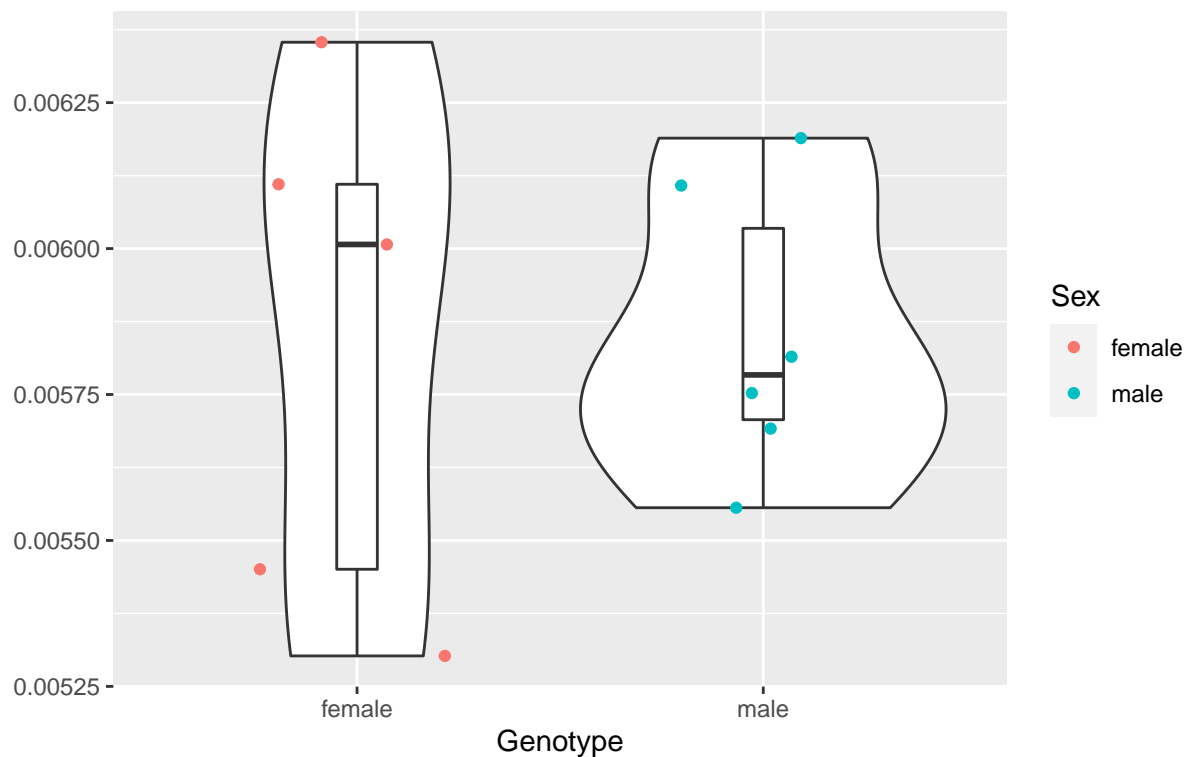
Red points denoting outliers



```
##          Df    Sum Sq  Mean Sq F value Pr(>F)
## Sex          1 1.228e-07 1.228e-07   6.621  0.03 *
## Residuals    9 1.669e-07 1.854e-08
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Facial Nerve

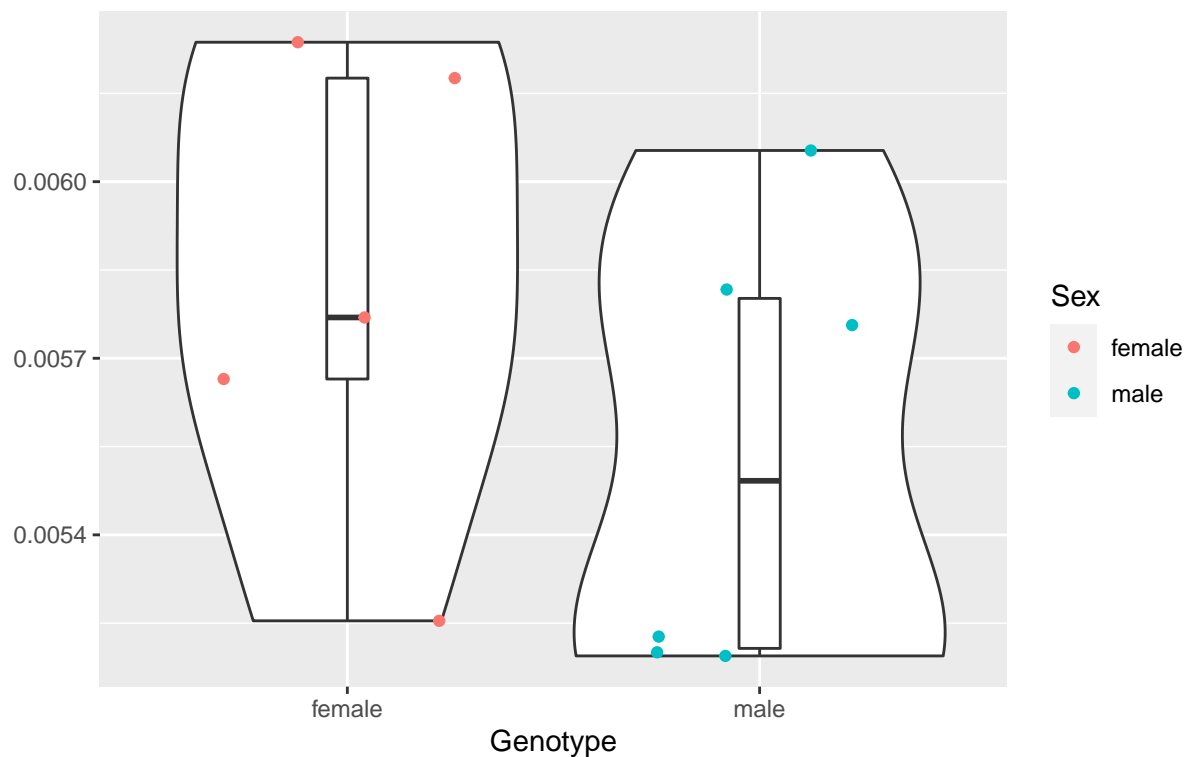
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.000e-10	1.400e-10	0.001	0.973
## Residuals	9	1.109e-06	1.232e-07		

Longitudinal Fasciculus of Pons

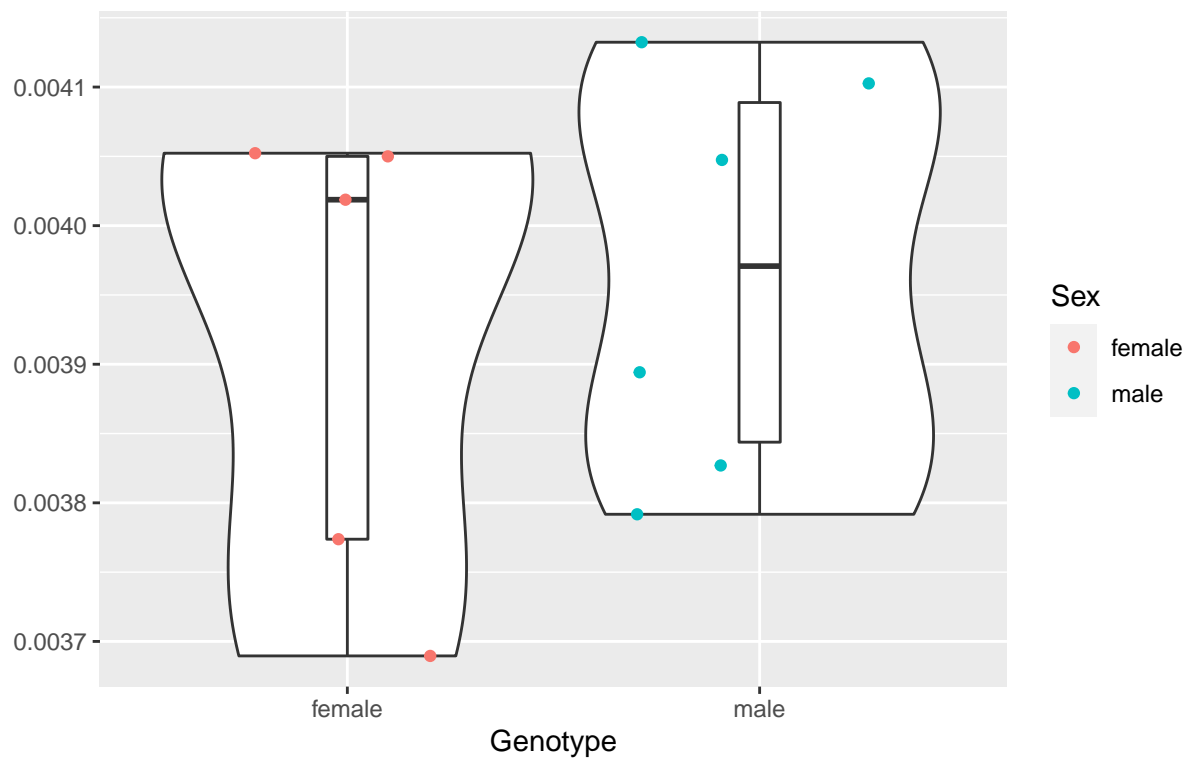
Red points denoting outliers



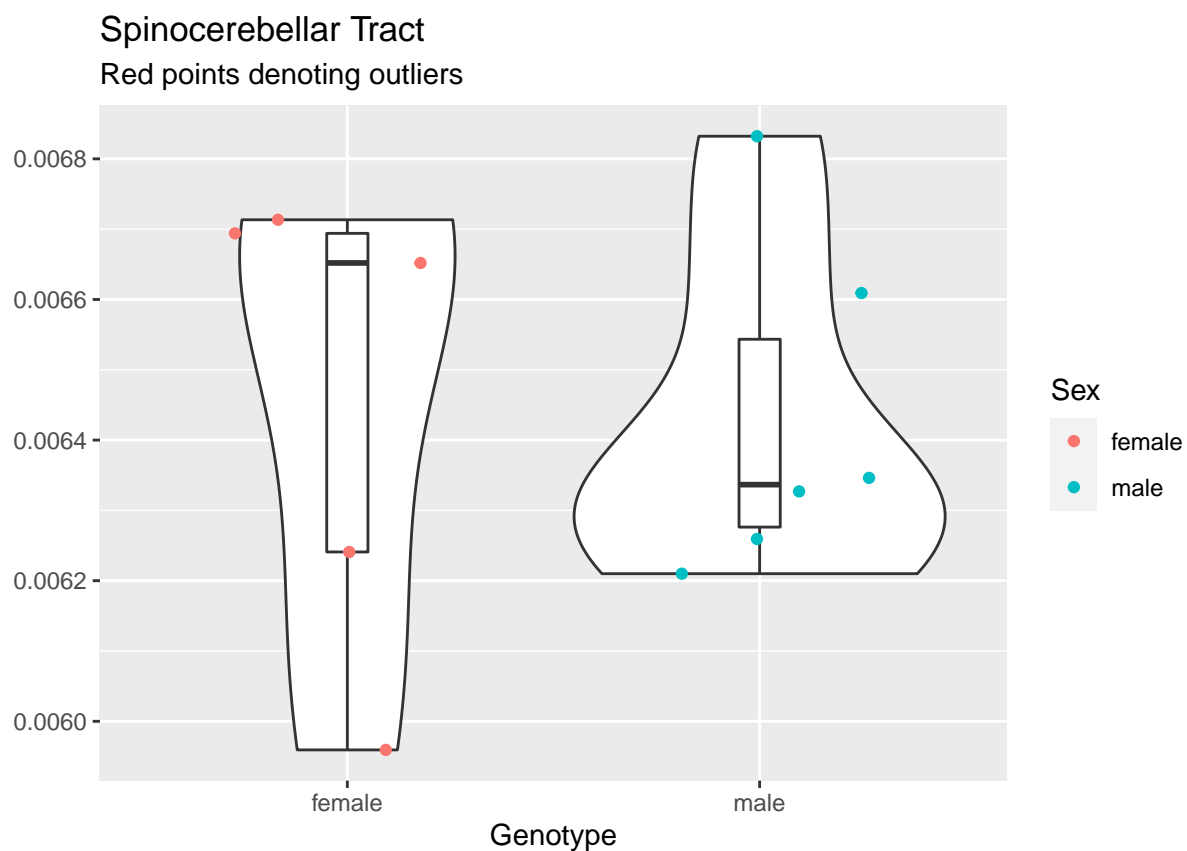
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.120e-07	2.120e-07	1.396	0.268
## Residuals	9	1.367e-06	1.519e-07		

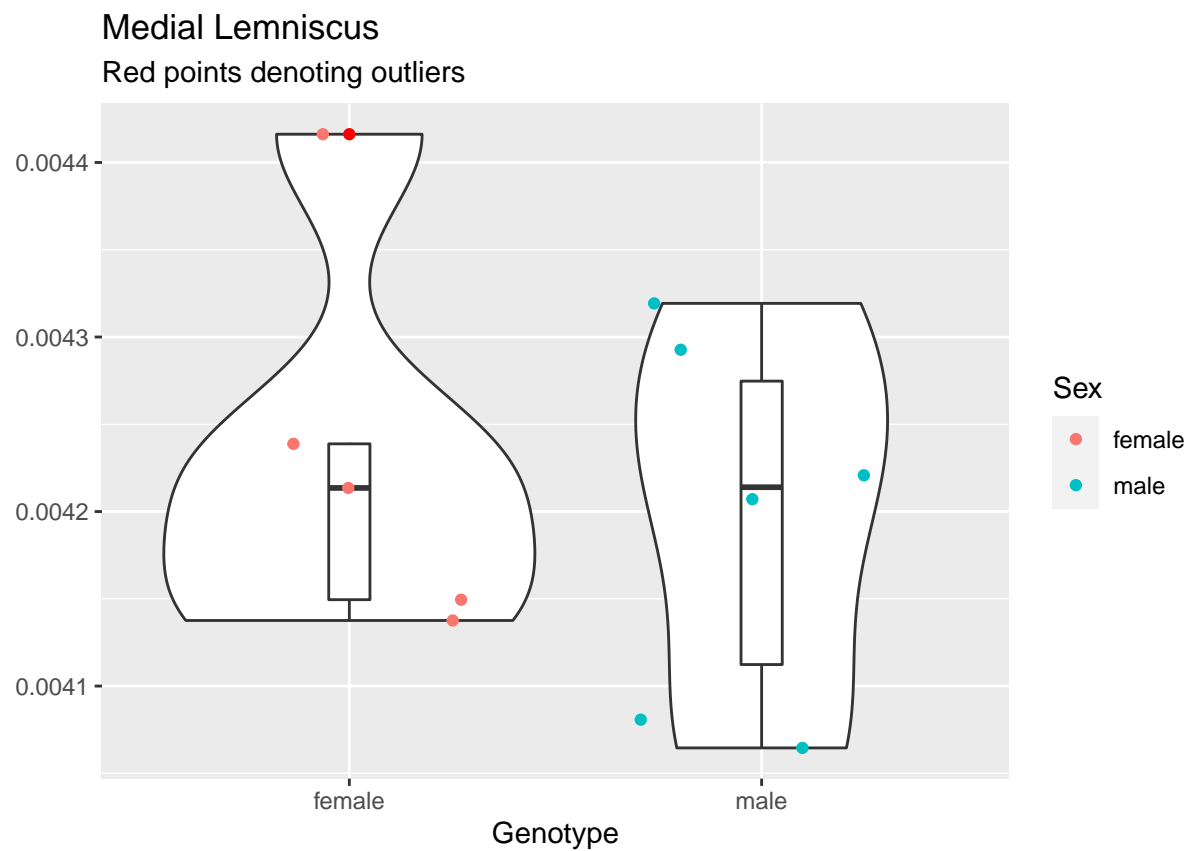
Medial Longitudinal Fasciculus and Tectospinal Tract

Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	6.560e-09	6.556e-09	0.261	0.622
## Residuals	9	2.264e-07	2.516e-08		

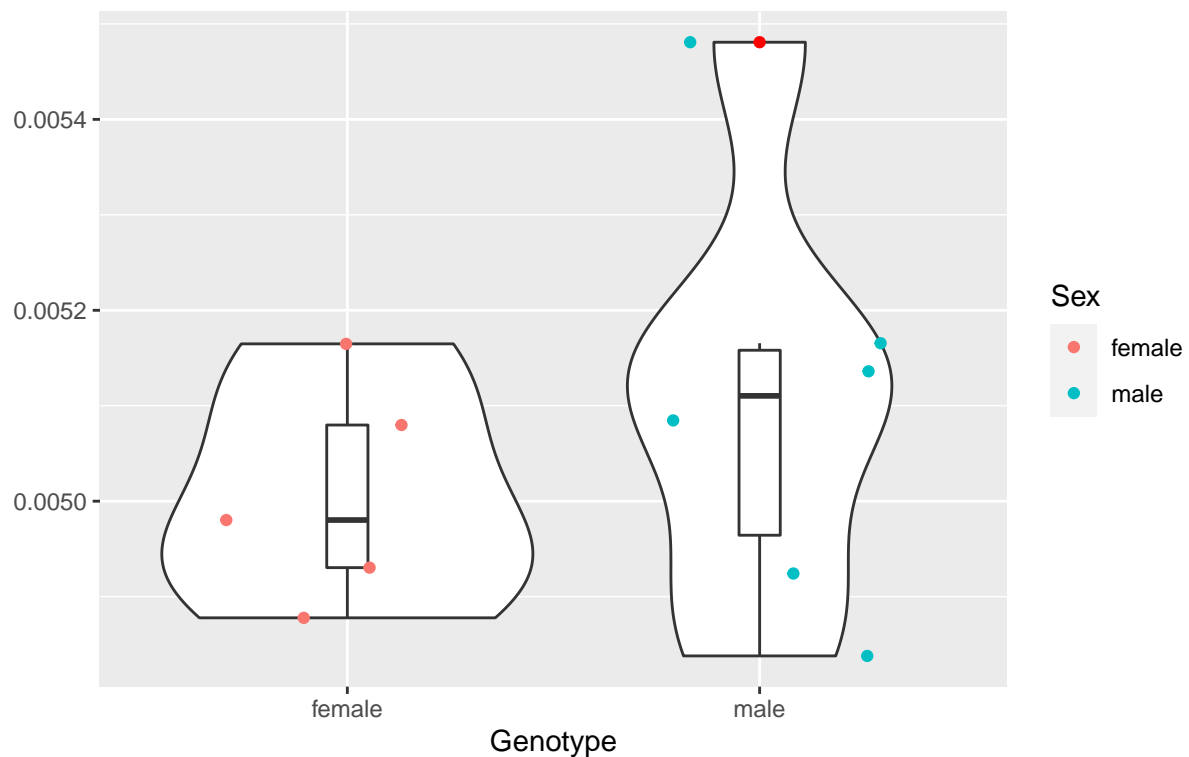




##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	3.080e-09	3.077e-09	0.262	0.621
## Residuals	9	1.058e-07	1.176e-08		

Ventral Spinocerebellar Tract

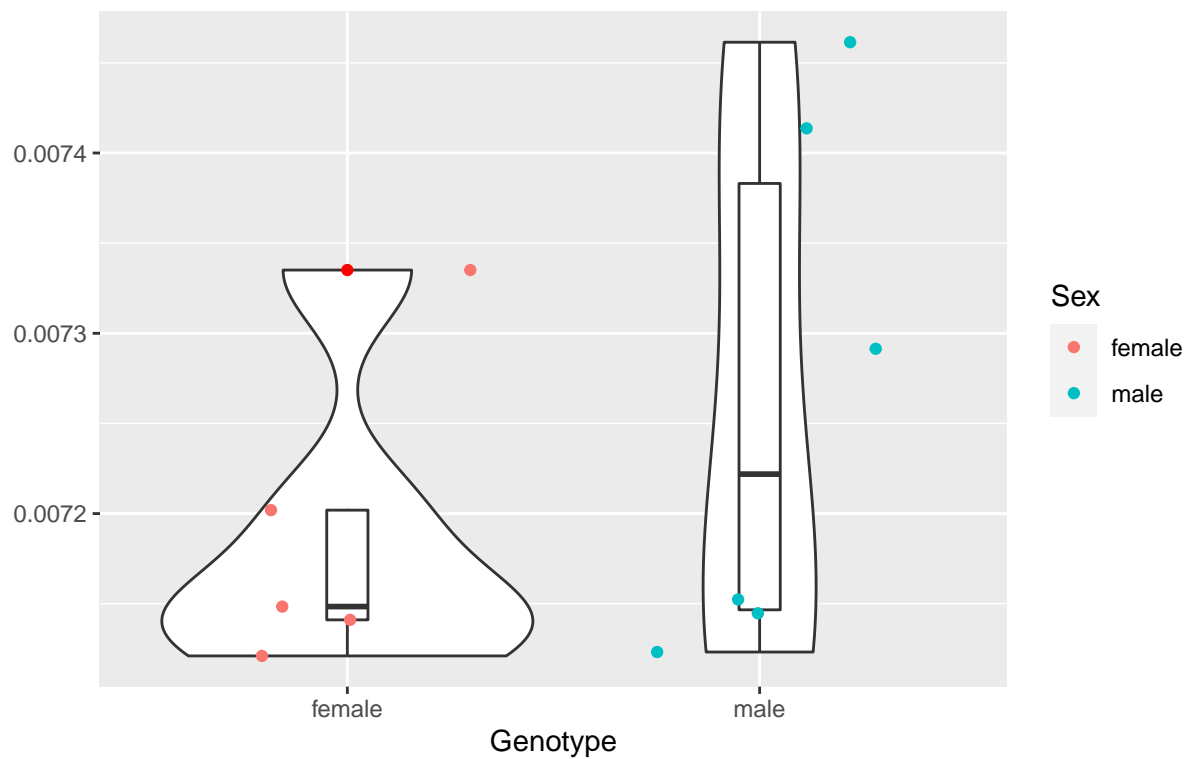
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.633e-08	2.633e-08	0.78	0.4
## Residuals	9	3.038e-07	3.376e-08		

Middle Cerebellar Peduncle

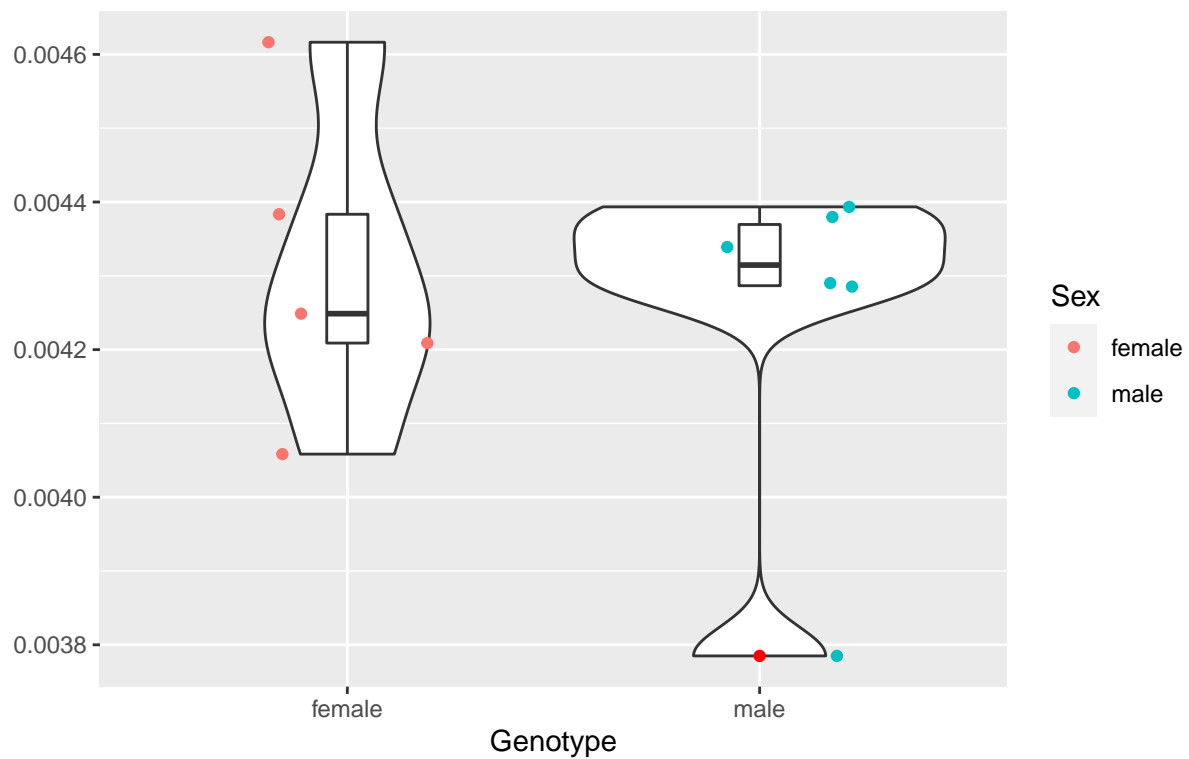
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.533e-08	1.533e-08	0.994	0.345
## Residuals	9	1.387e-07	1.541e-08		

Superior Cerebellar Peduncle

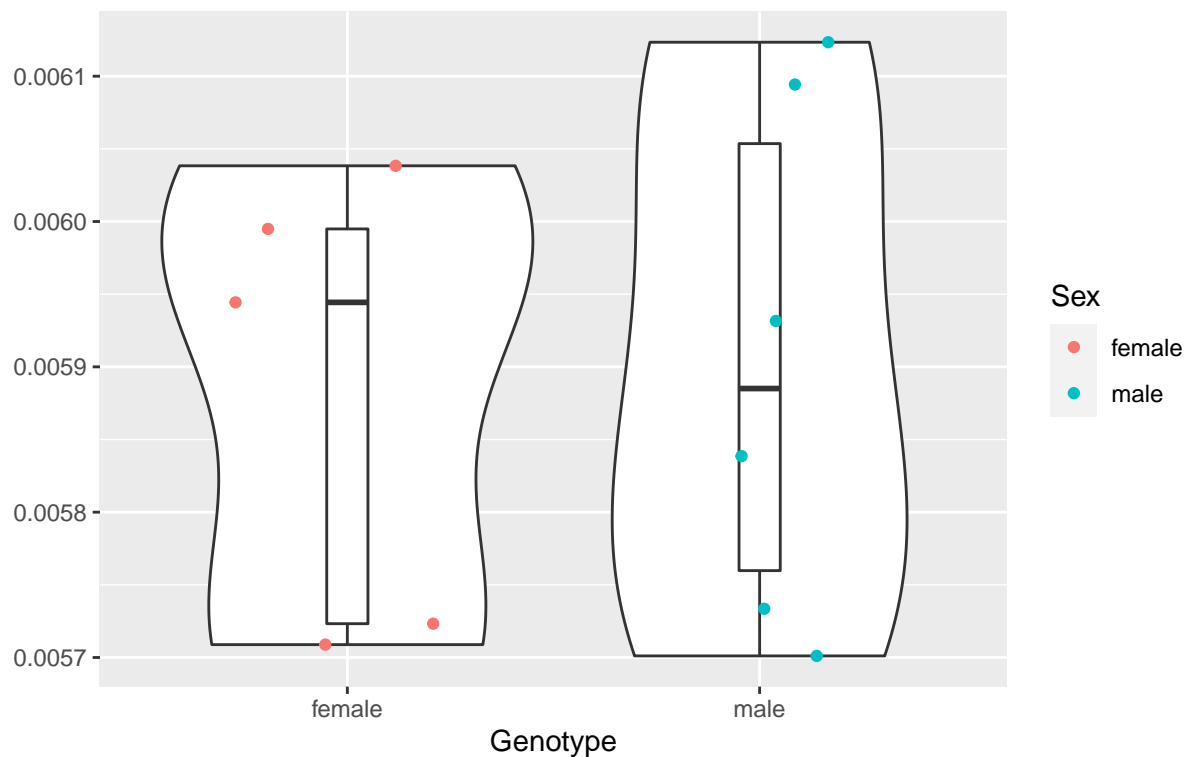
Red points denoting outliers



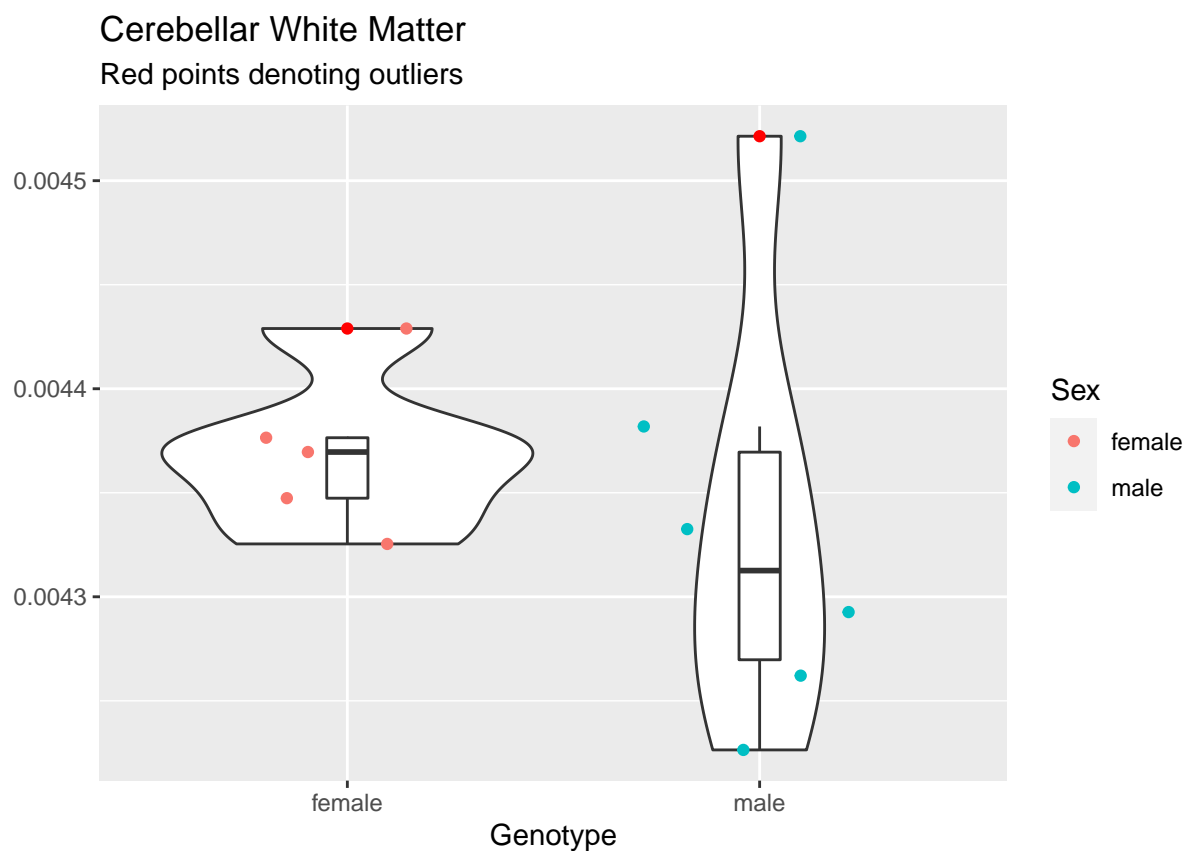
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	9.100e-09	9.090e-09	0.186	0.677
## Residuals	9	4.407e-07	4.897e-08		

Inferior Cerebellar Peduncle

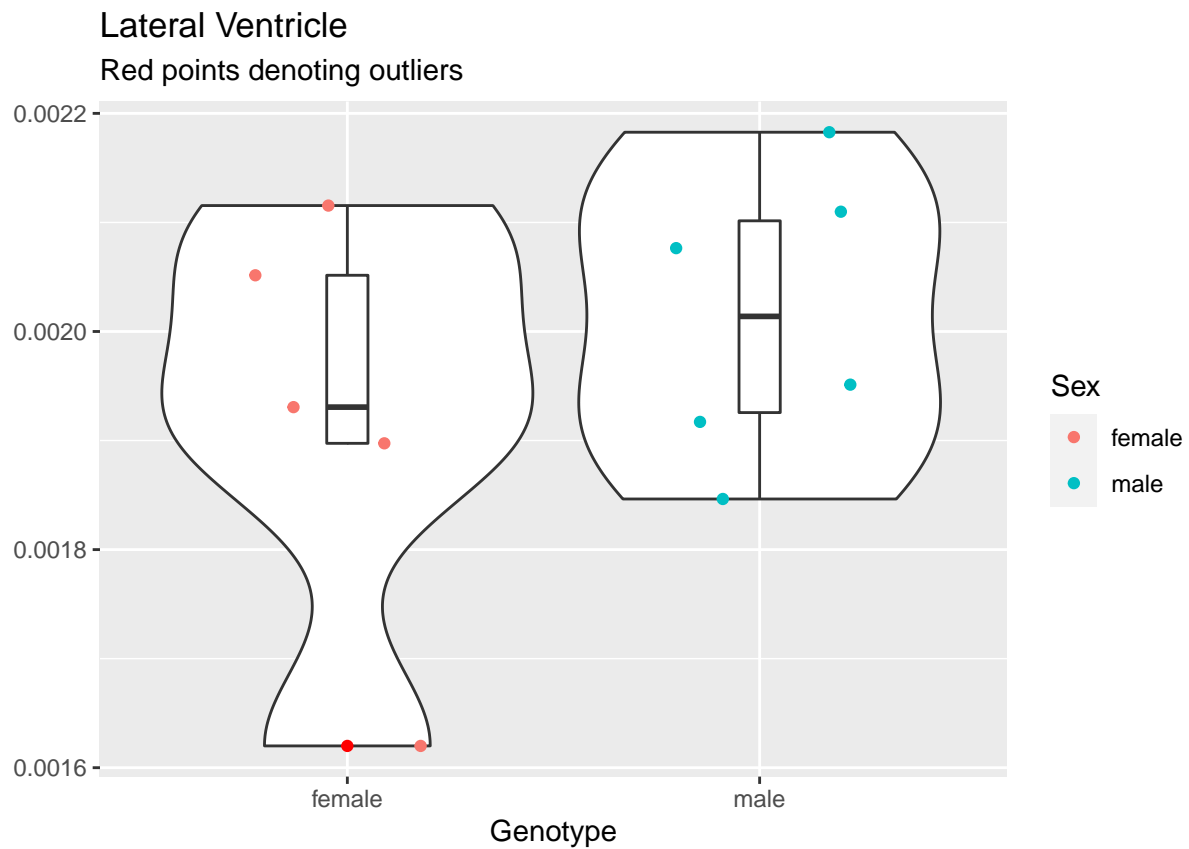
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.300e-09	1.295e-09	0.046	0.836
## Residuals	9	2.558e-07	2.843e-08		

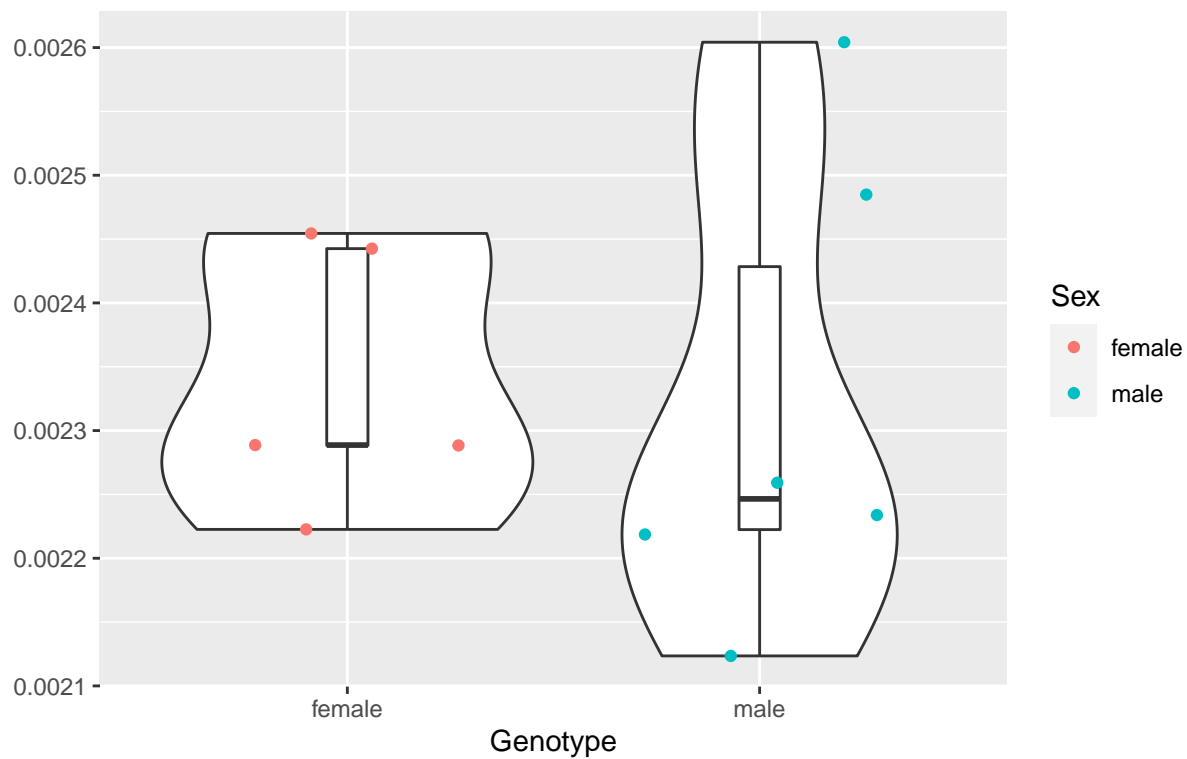


##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	3.040e-09	3.044e-09	0.443	0.522
## Residuals	9	6.185e-08	6.873e-09		



Cingulate Cortex Area 25

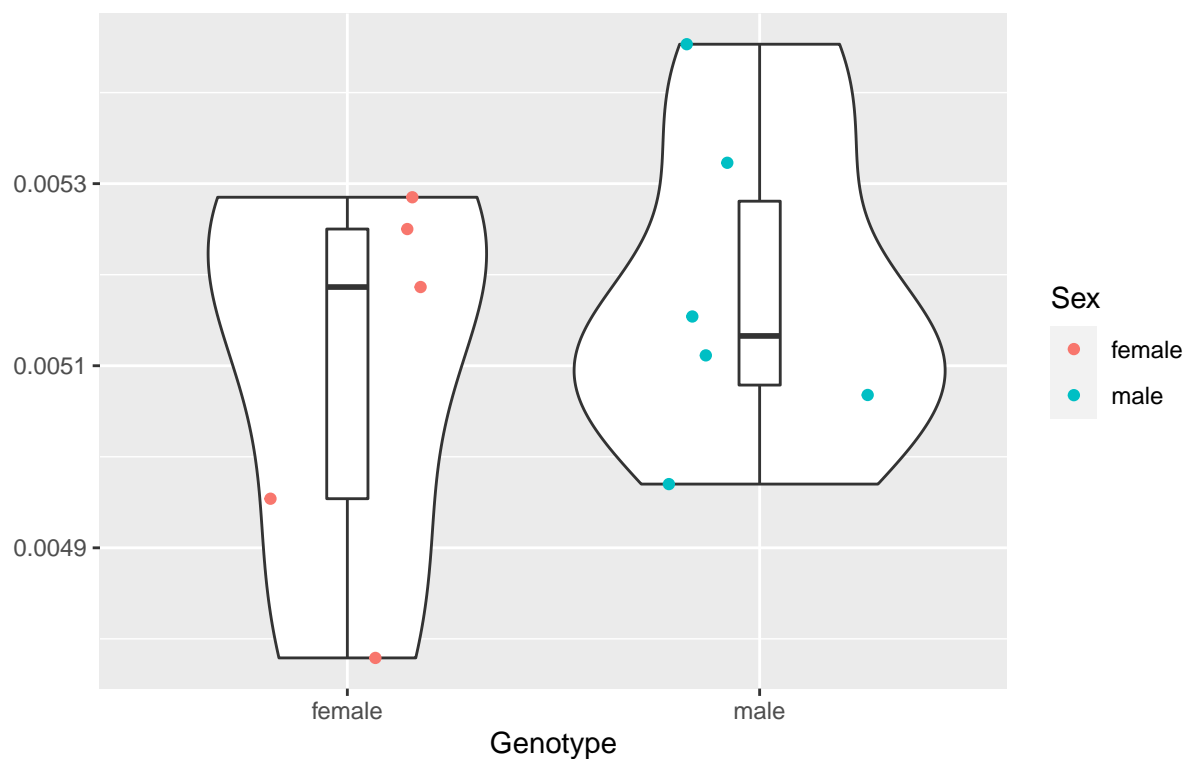
Red points denoting outliers



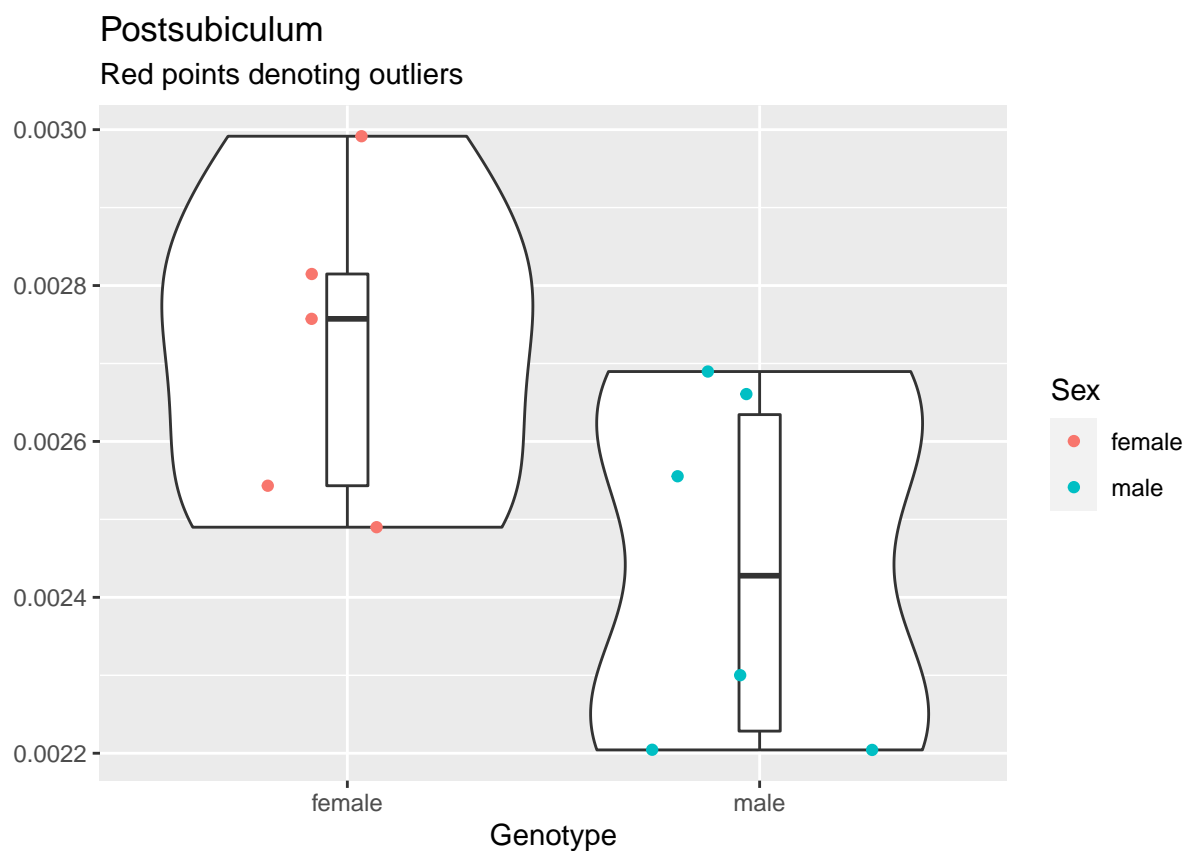
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	9.500e-10	9.460e-10	0.04	0.845
## Residuals	9	2.106e-07	2.341e-08		

Dorsal Acoustic Stria

Red points denoting outliers



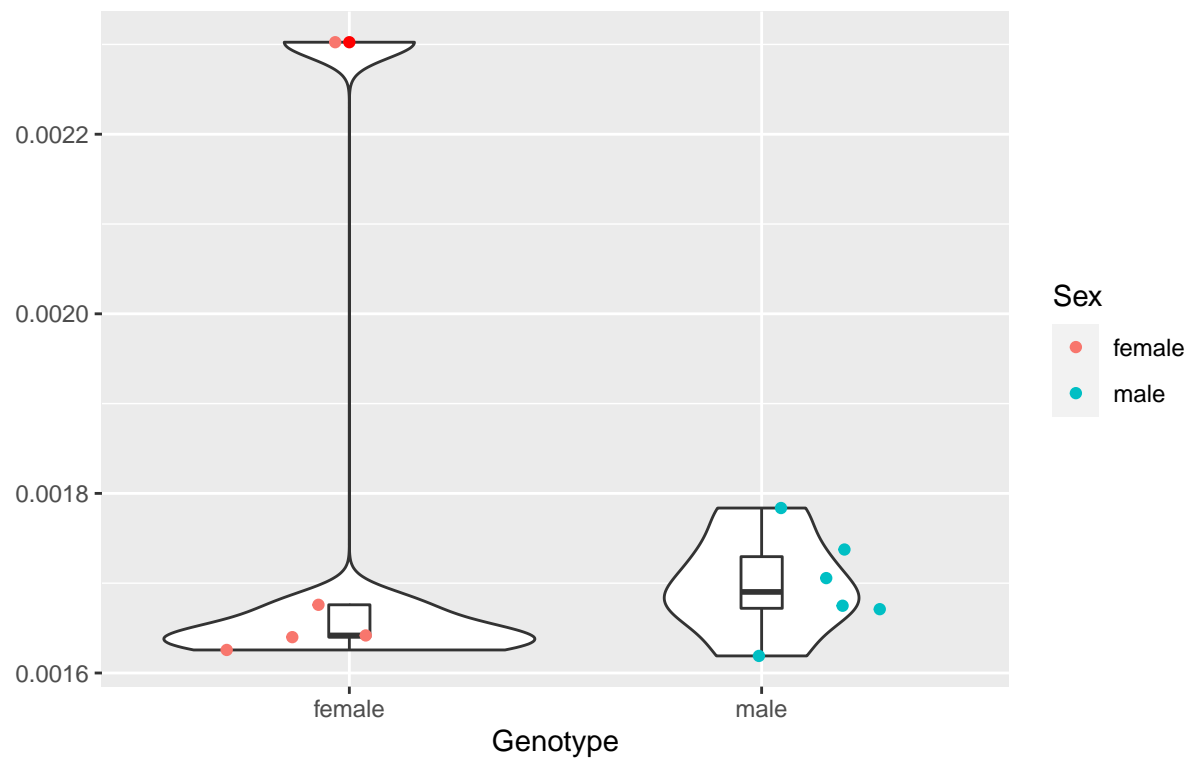
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.160e-08	2.159e-08	0.563	0.472
## Residuals	9	3.453e-07	3.837e-08		



```
##          Df    Sum Sq  Mean Sq F value Pr(>F)
## Sex          1 2.193e-07 2.193e-07   4.665 0.0591 .
## Residuals    9 4.232e-07 4.702e-08
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Ventricular System 4th Ventricle

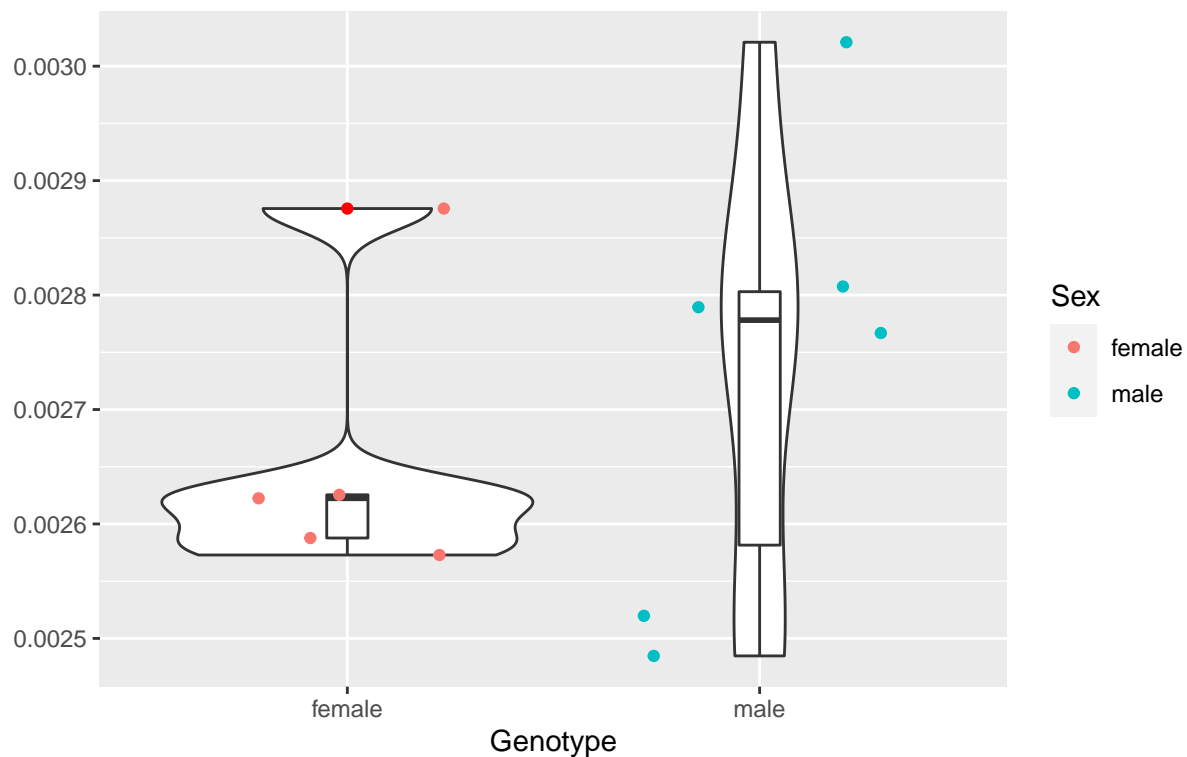
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.680e-08	1.681e-08	0.417	0.534
## Residuals	9	3.627e-07	4.030e-08		

Microcellular Tegmental Nucleus

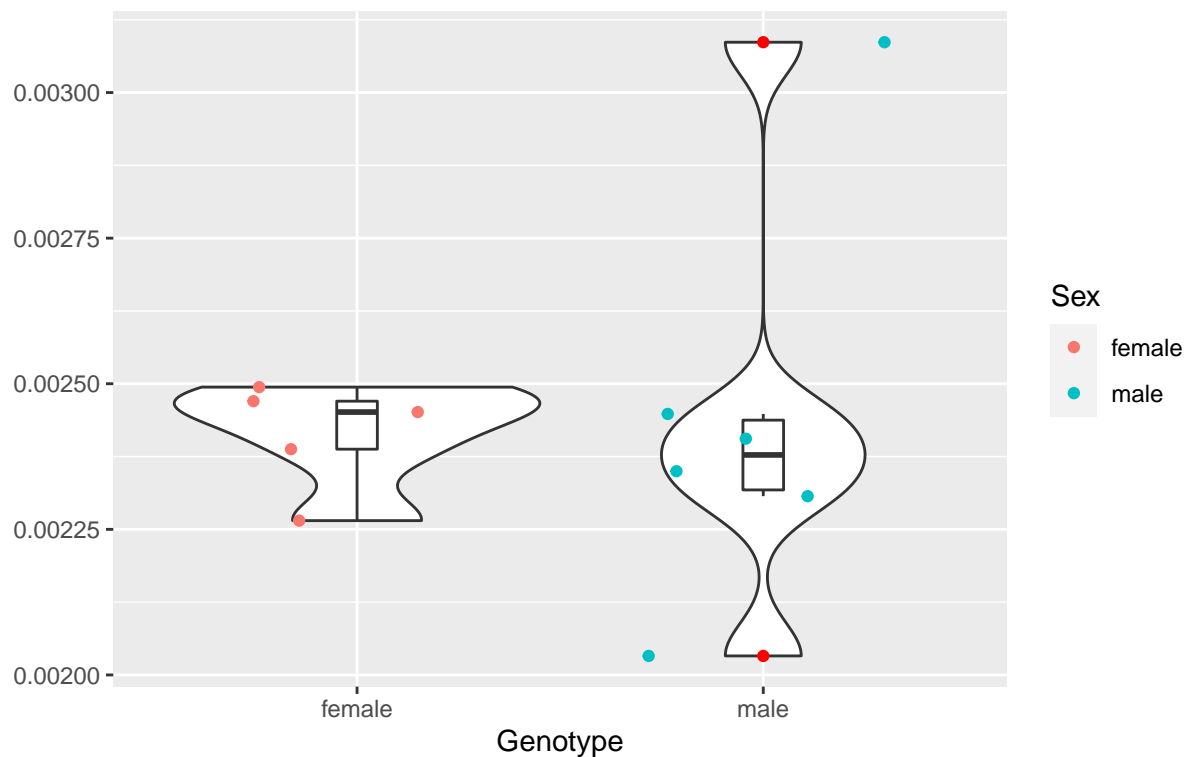
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.521e-08	1.522e-08	0.523	0.488
## Residuals	9	2.618e-07	2.908e-08		

Pretectal Nucleus

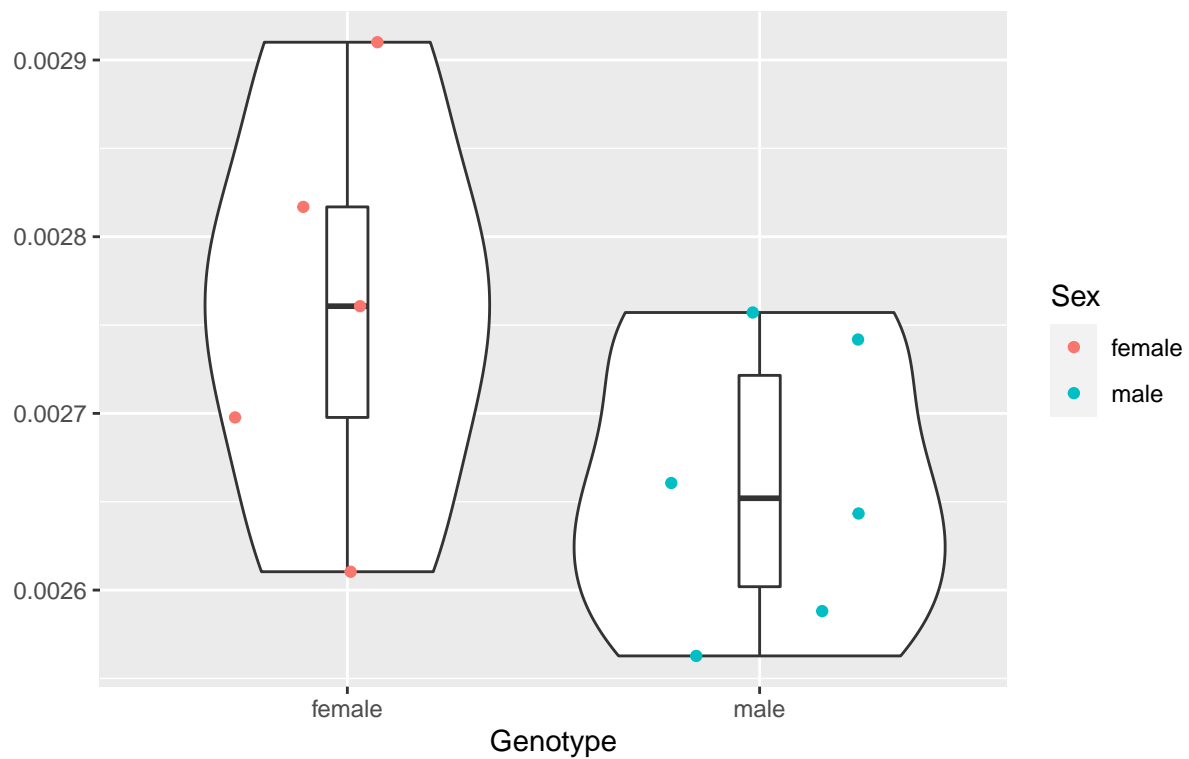
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.700e-09	1.650e-09	0.023	0.883
## Residuals	9	6.447e-07	7.164e-08		

Latero Dorsal Thalamic Nucleus Ventro Lateral

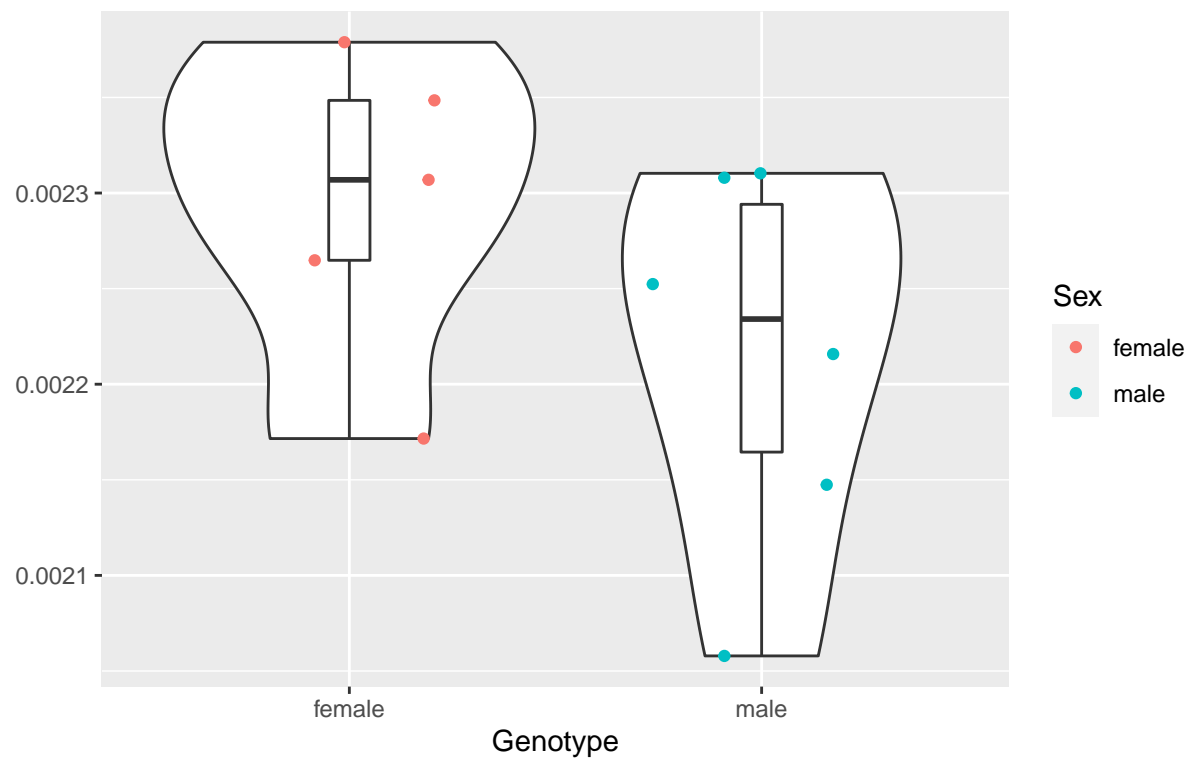
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.738e-08	2.738e-08	2.967	0.119
## Residuals	9	8.304e-08	9.227e-09		

Latero Posterior Nuclei of Thalamus

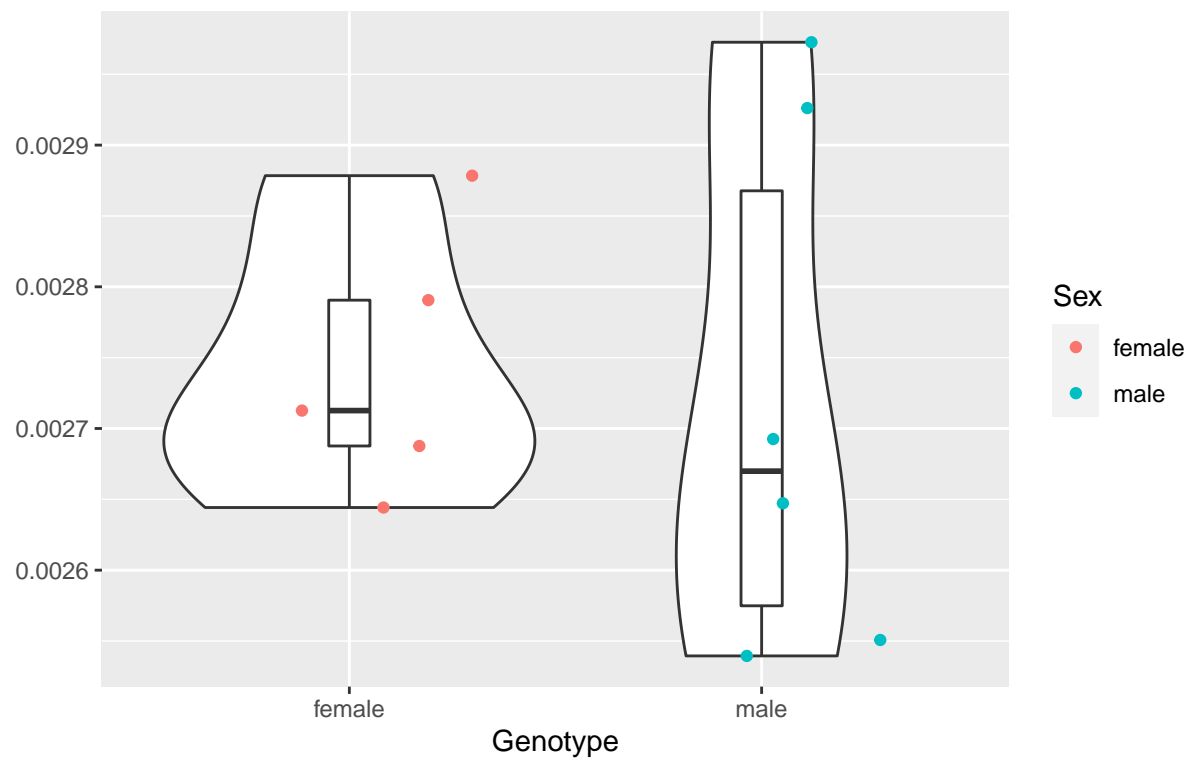
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.695e-08	1.695e-08	2.046	0.186
## Residuals	9	7.456e-08	8.285e-09		

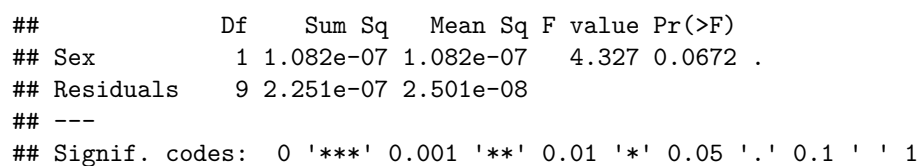
Anterior Thalamic Nuclei

Red points denoting outliers



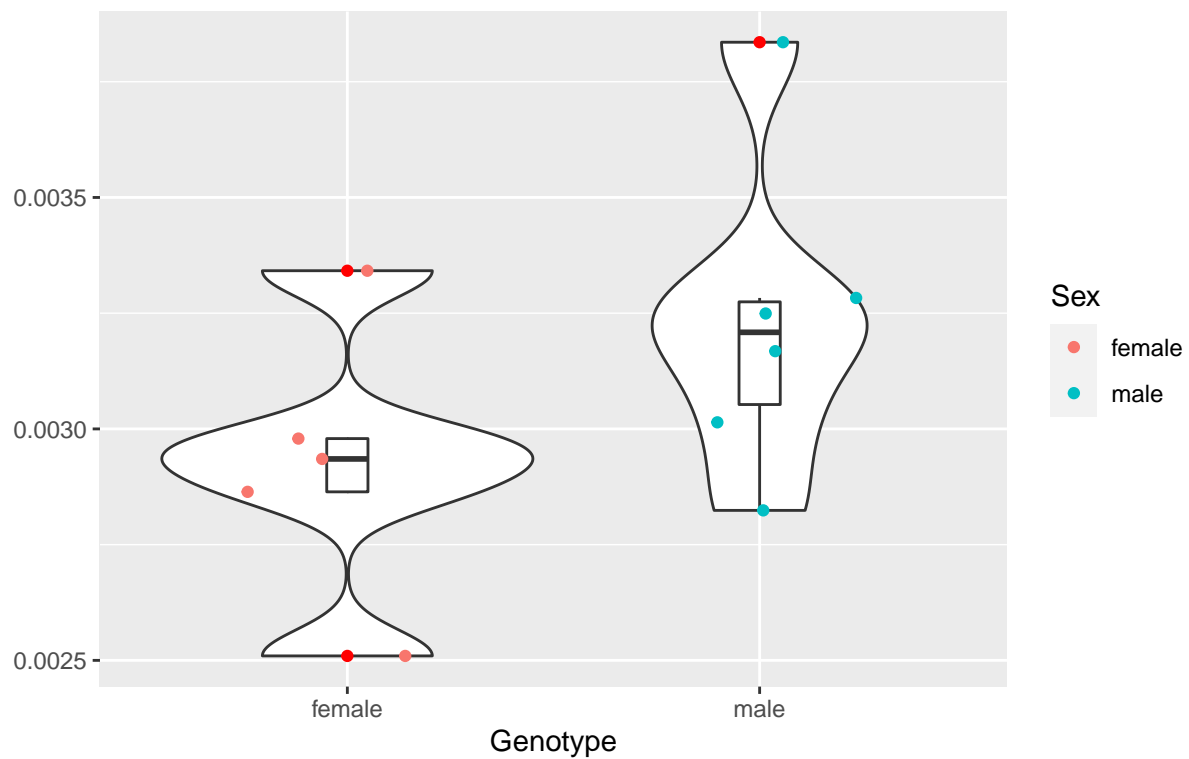
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.230e-09	1.231e-09	0.053	0.823
## Residuals	9	2.079e-07	2.310e-08		

Red points denoting outliers

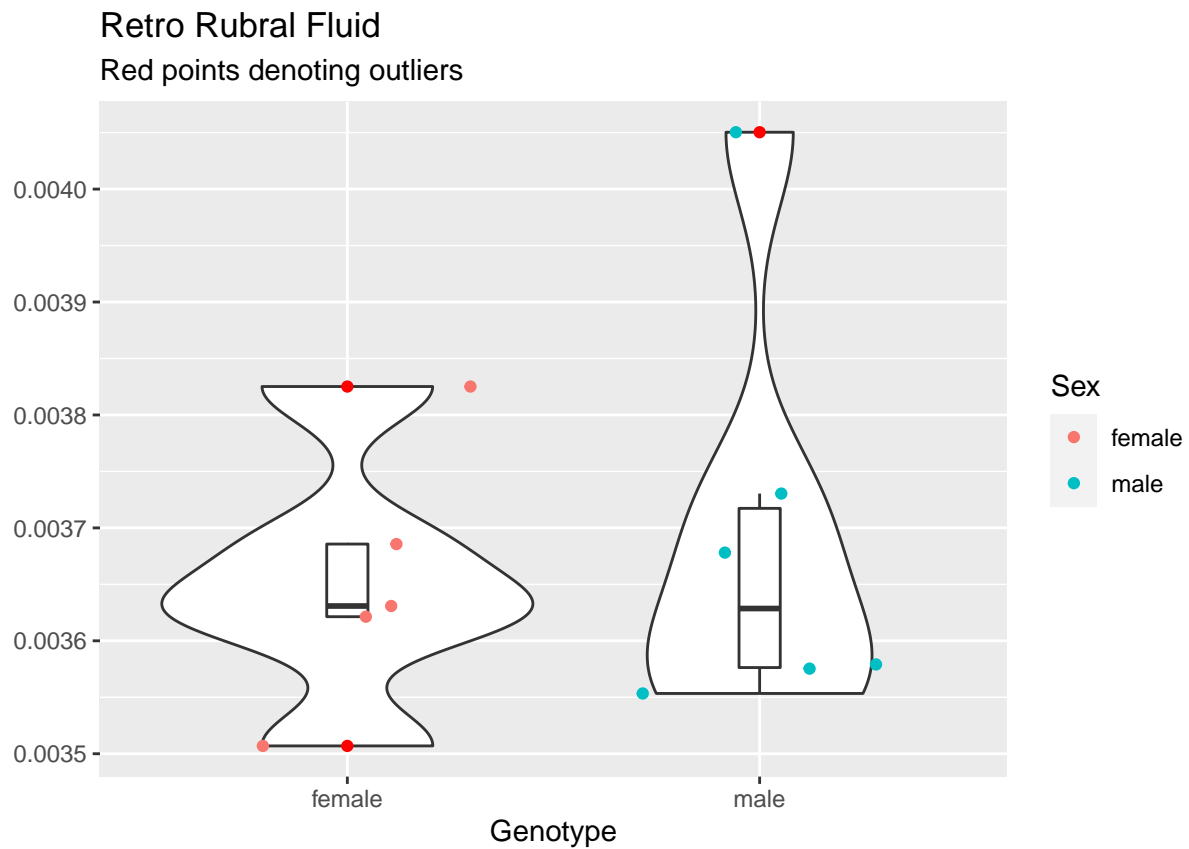


Pararubral Nucleus

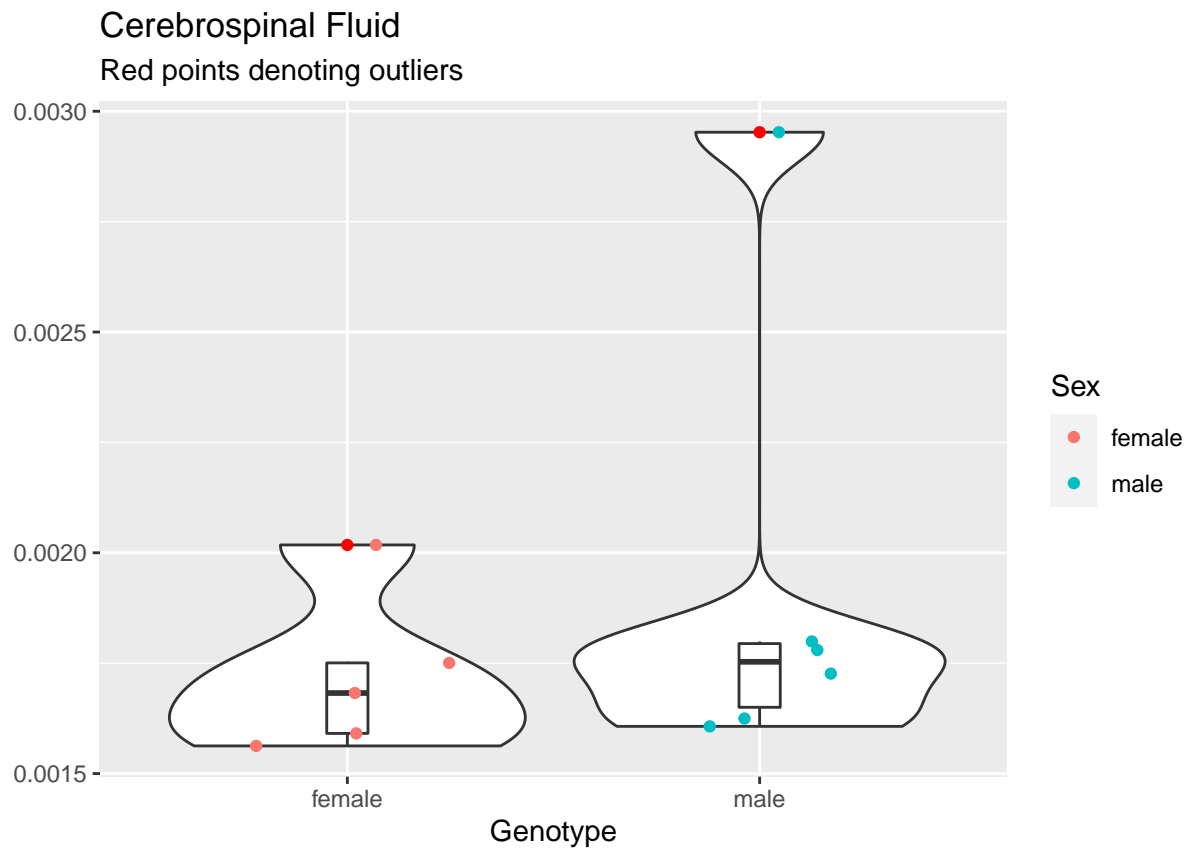
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.504e-07	2.504e-07	2.404	0.155
## Residuals	9	9.378e-07	1.042e-07		

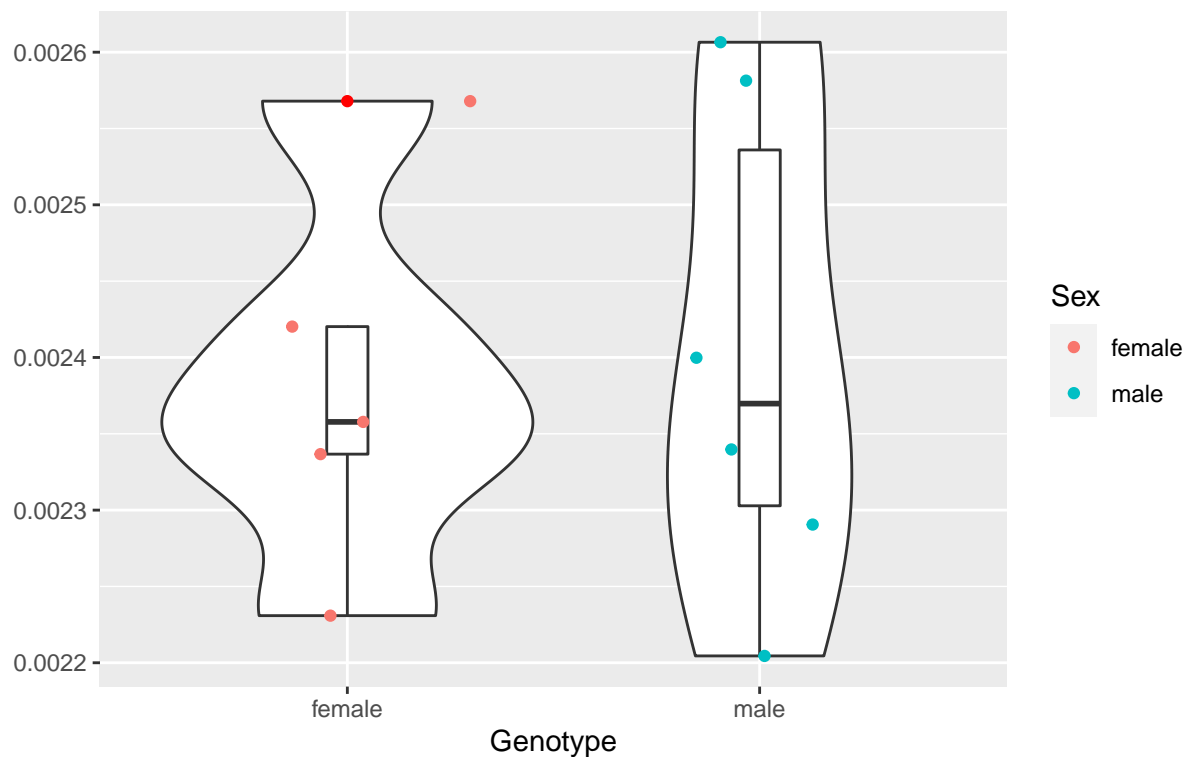


##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	4.460e-09	4.465e-09	0.175	0.685
## Residuals	9	2.292e-07	2.546e-08		



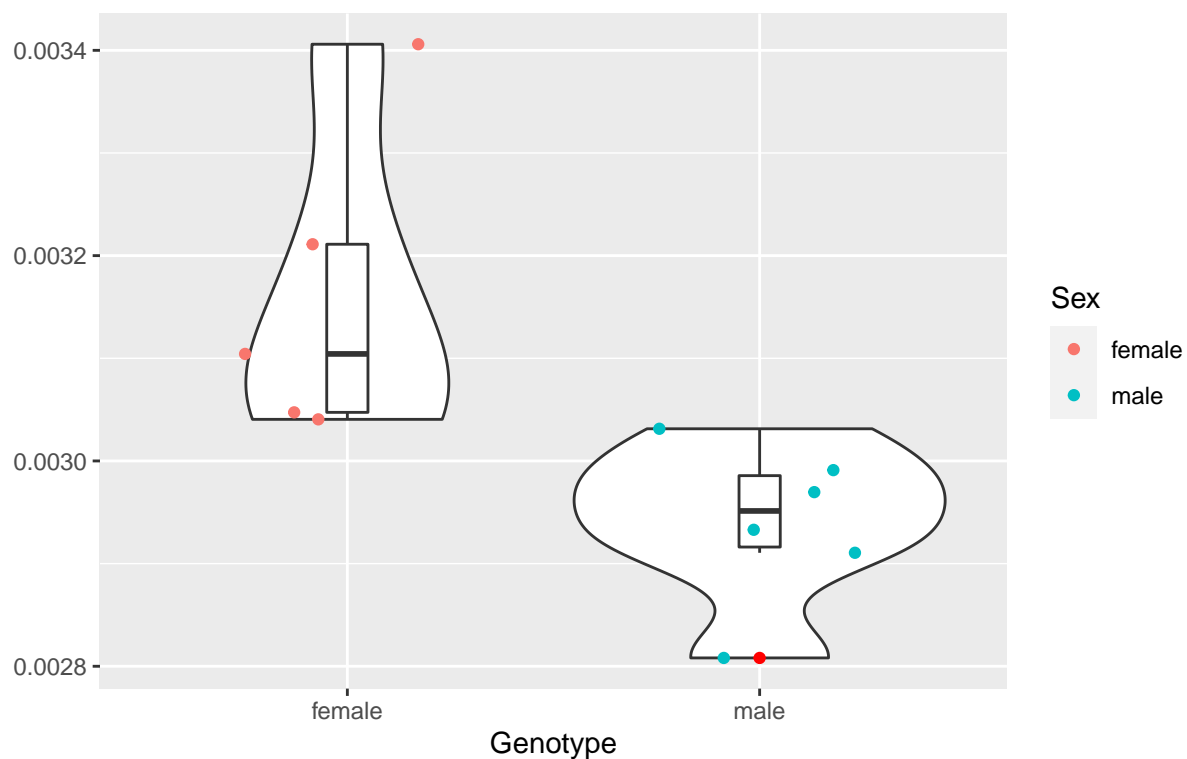
Intermediate Reticular Nucleus

Red points denoting outliers

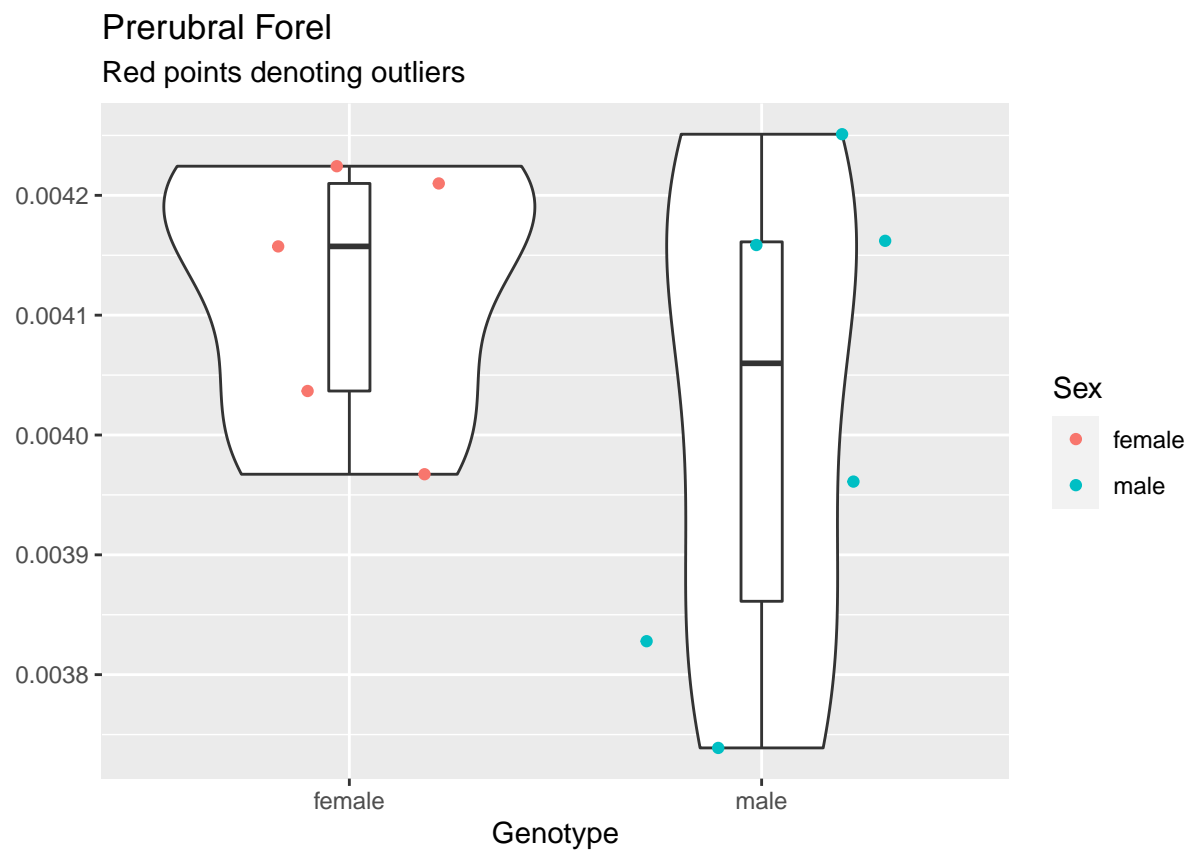


##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.210e-09	1.208e-09	0.057	0.817
## Residuals	9	1.908e-07	2.120e-08		

Posterior Dorsal Paraventricular Medial Parvicellular Posterior Lateral Hy
Red points denoting outliers

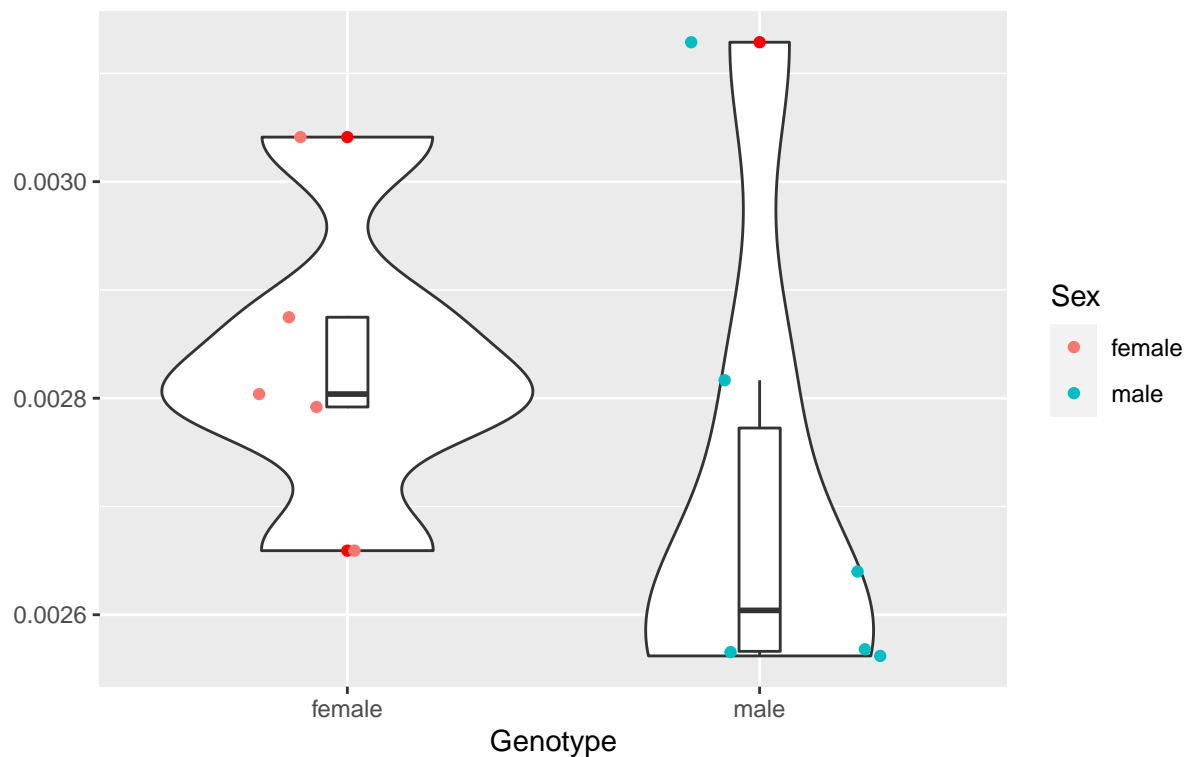


```
##          Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex          1 1.335e-07 1.335e-07   9.743 0.0123 *
## Residuals    9 1.233e-07 1.370e-08
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

PVG of Hypothalamus

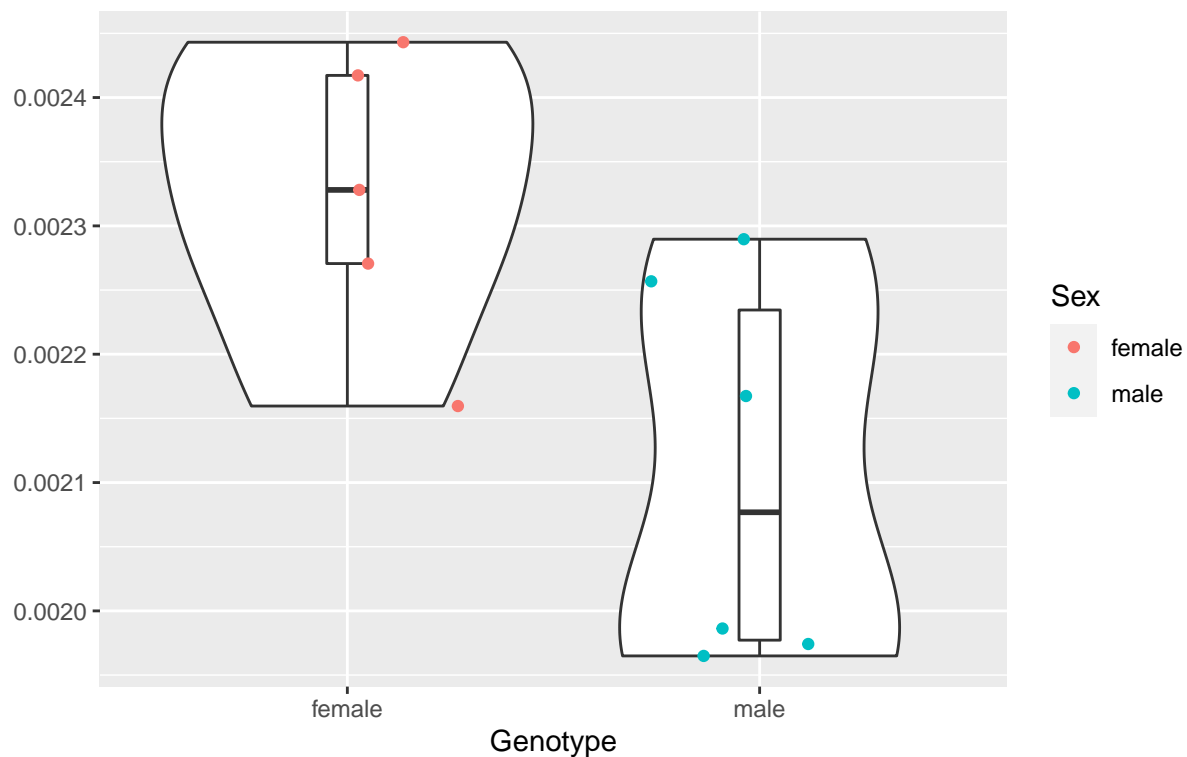
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	3.970e-08	3.971e-08	1.076	0.327
## Residuals	9	3.323e-07	3.692e-08		

Basal Lateral Amygdala

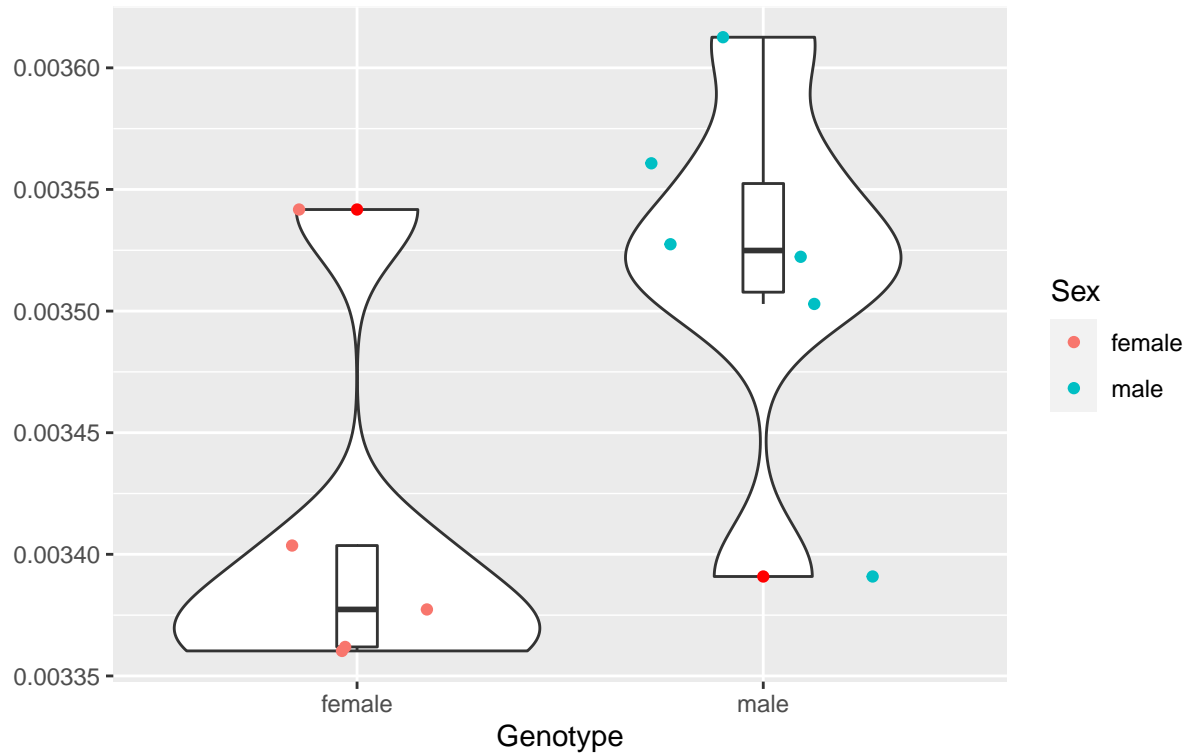
Red points denoting outliers



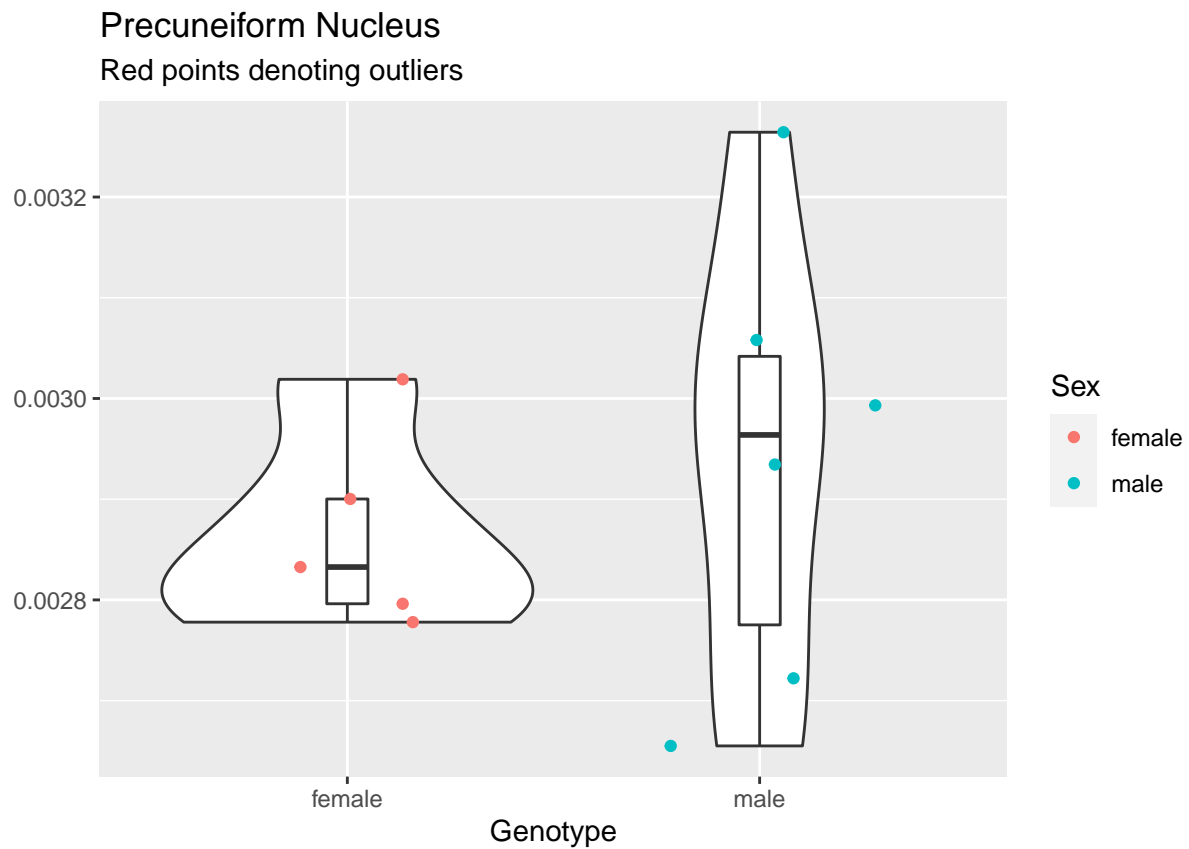
```
##           Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex           1 1.287e-07 1.287e-07   7.034 0.0264 *
## Residuals     9 1.646e-07 1.829e-08
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Brain Stem Rest

Red points denoting outliers



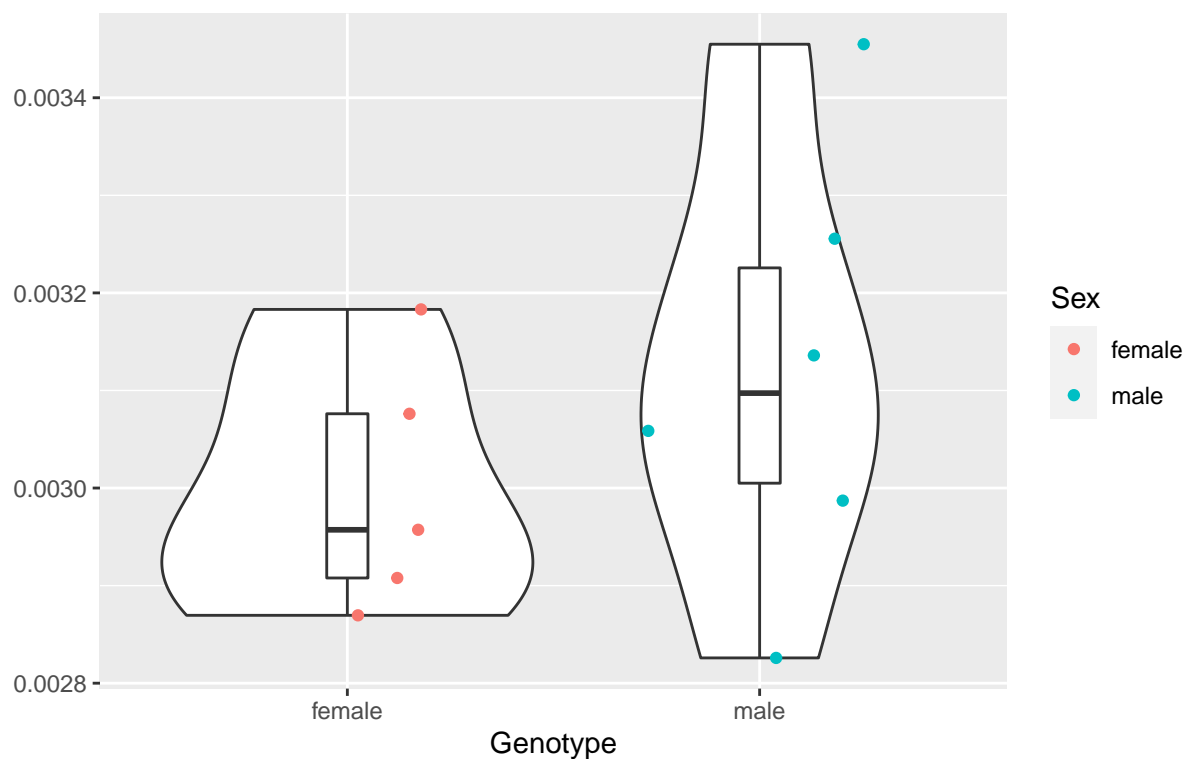
```
##           Df  Sum Sq Mean Sq F value Pr(>F)
## Sex           1 3.33e-08 3.33e-08   5.936 0.0376 *
## Residuals     9 5.05e-08 5.61e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.440e-08	1.44e-08	0.449	0.52
## Residuals	9	2.889e-07	3.21e-08		

Cuneiform Nucleus

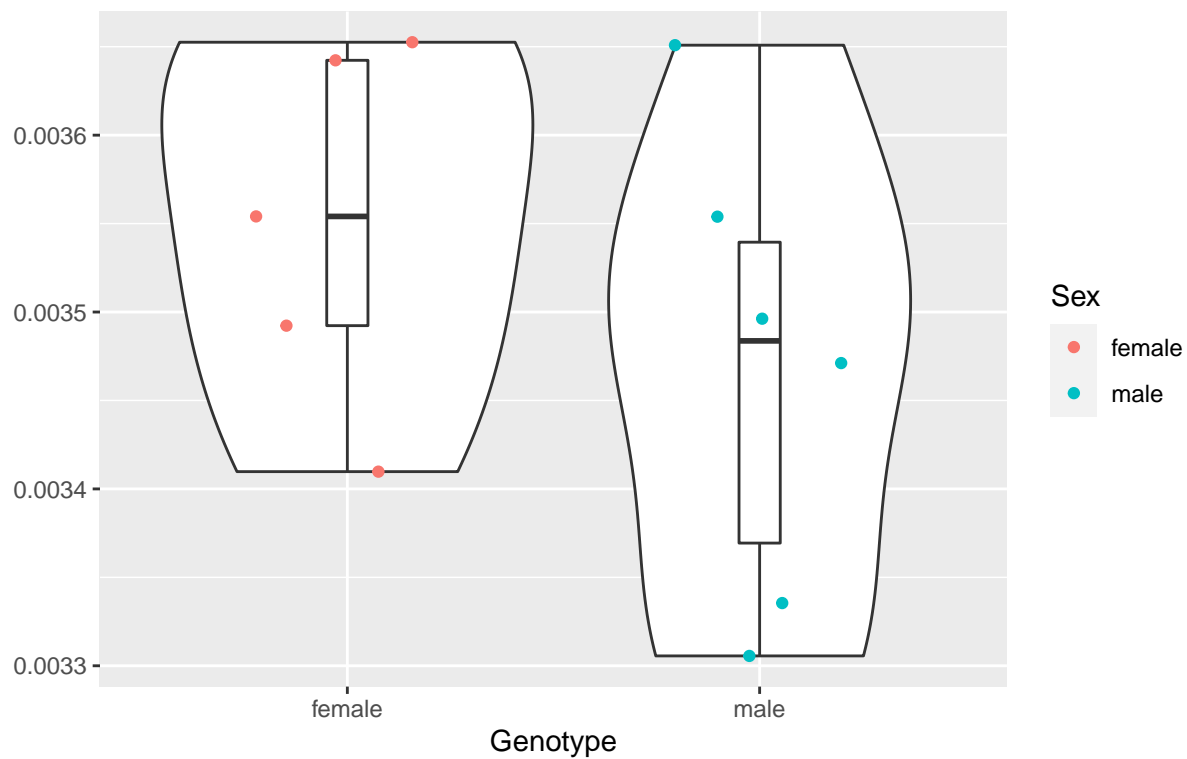
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	3.986e-08	3.986e-08	1.175	0.307
## Residuals	9	3.053e-07	3.393e-08		

Midbrain Linear Nucleus

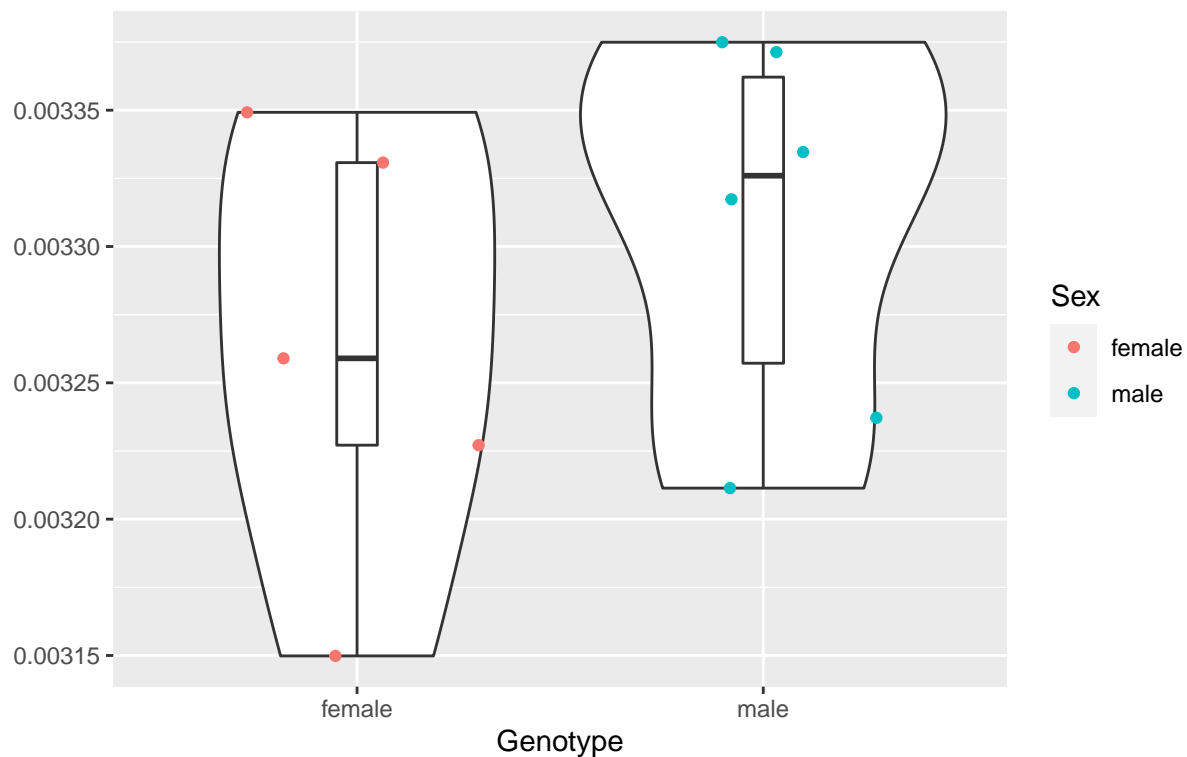
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.804e-08	1.804e-08	1.272	0.289
## Residuals	9	1.276e-07	1.418e-08		

Midbrain Reticular Nucleus

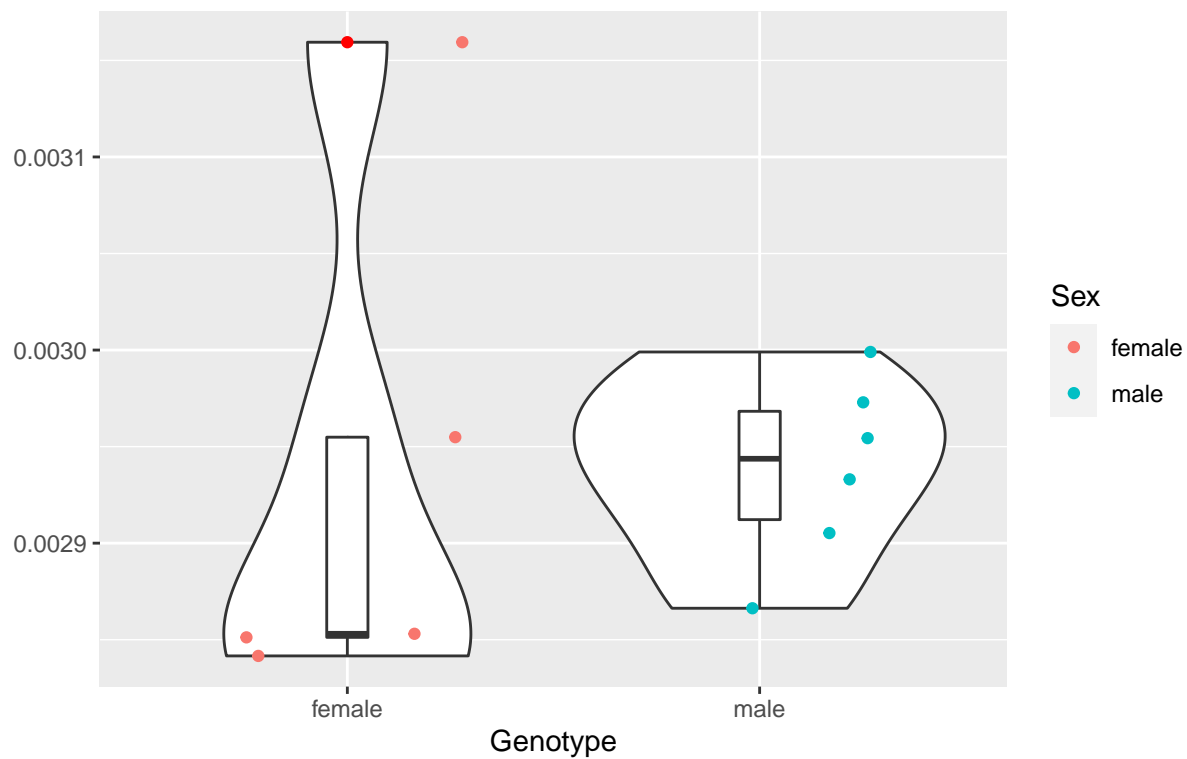
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	5.420e-09	5.425e-09	0.981	0.348
## Residuals	9	4.977e-08	5.530e-09		

Red Nucleus Parvicellular

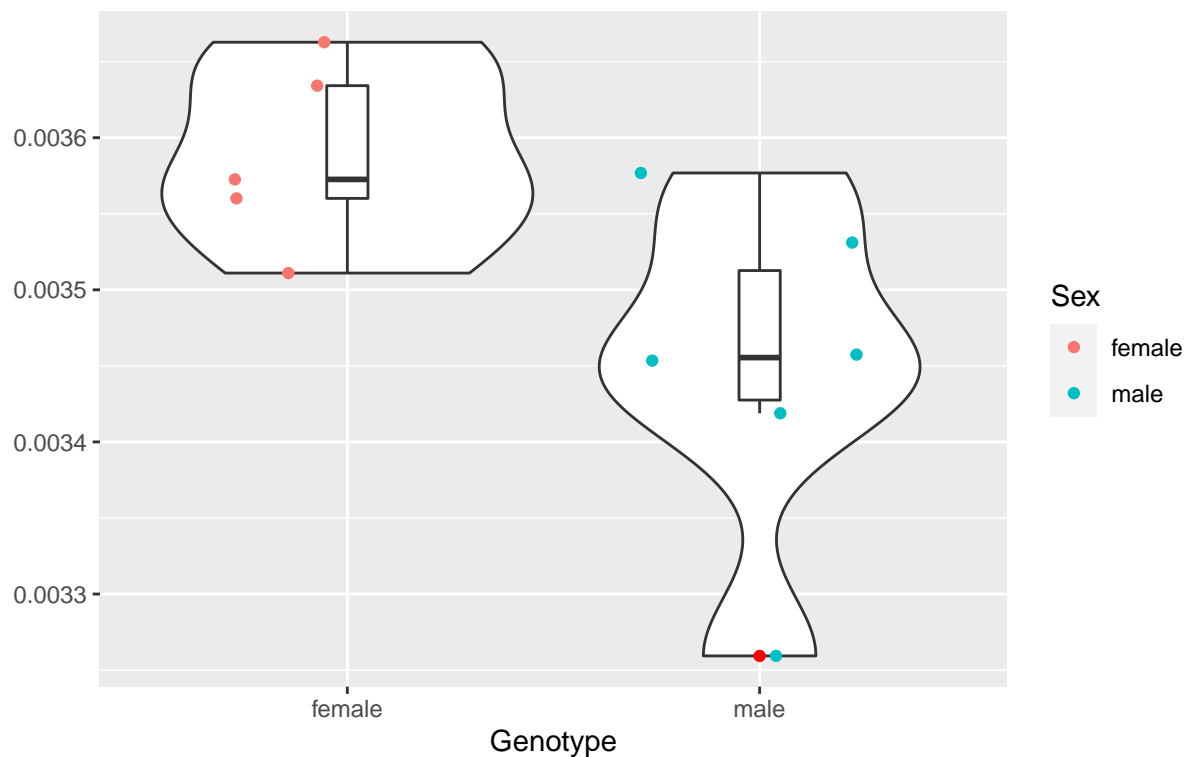
Red points denoting outliers



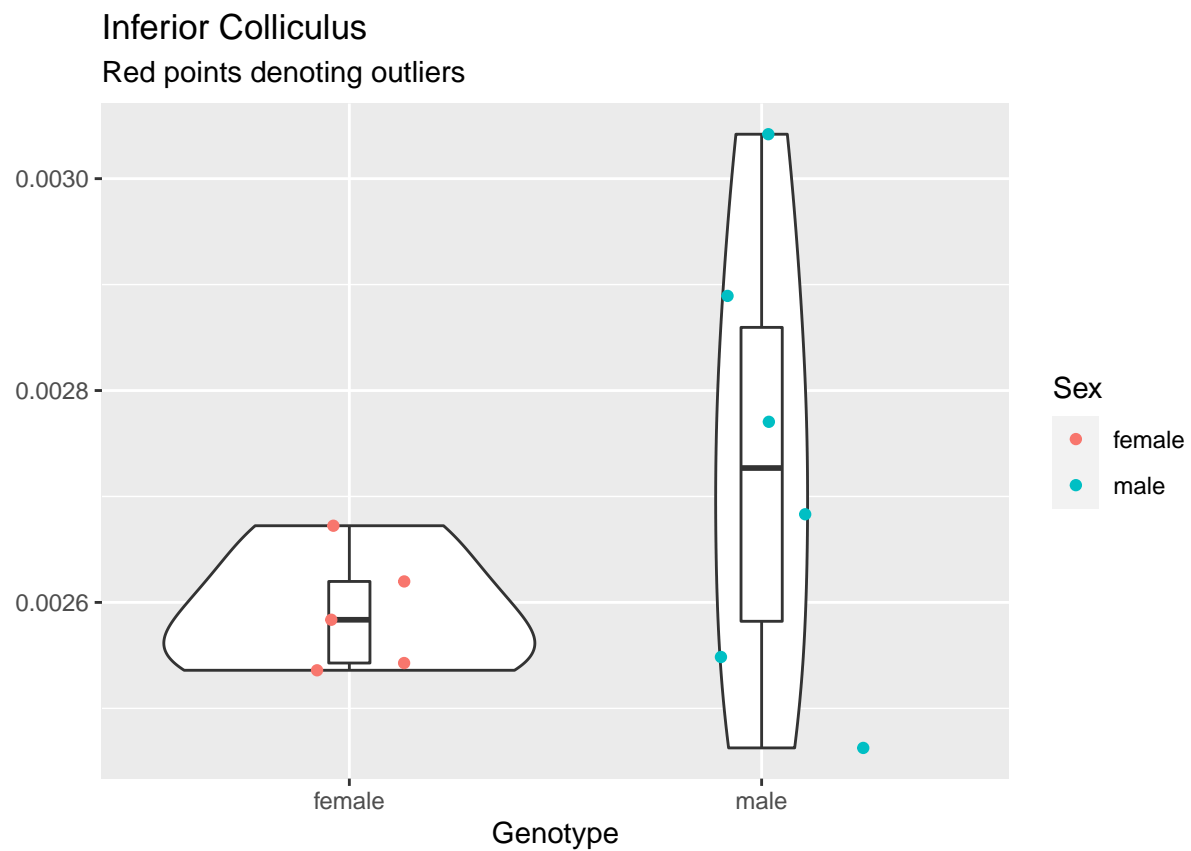
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.100e-10	1.130e-10	0.012	0.915
## Residuals	9	8.462e-08	9.403e-09		

Substantia Nigra

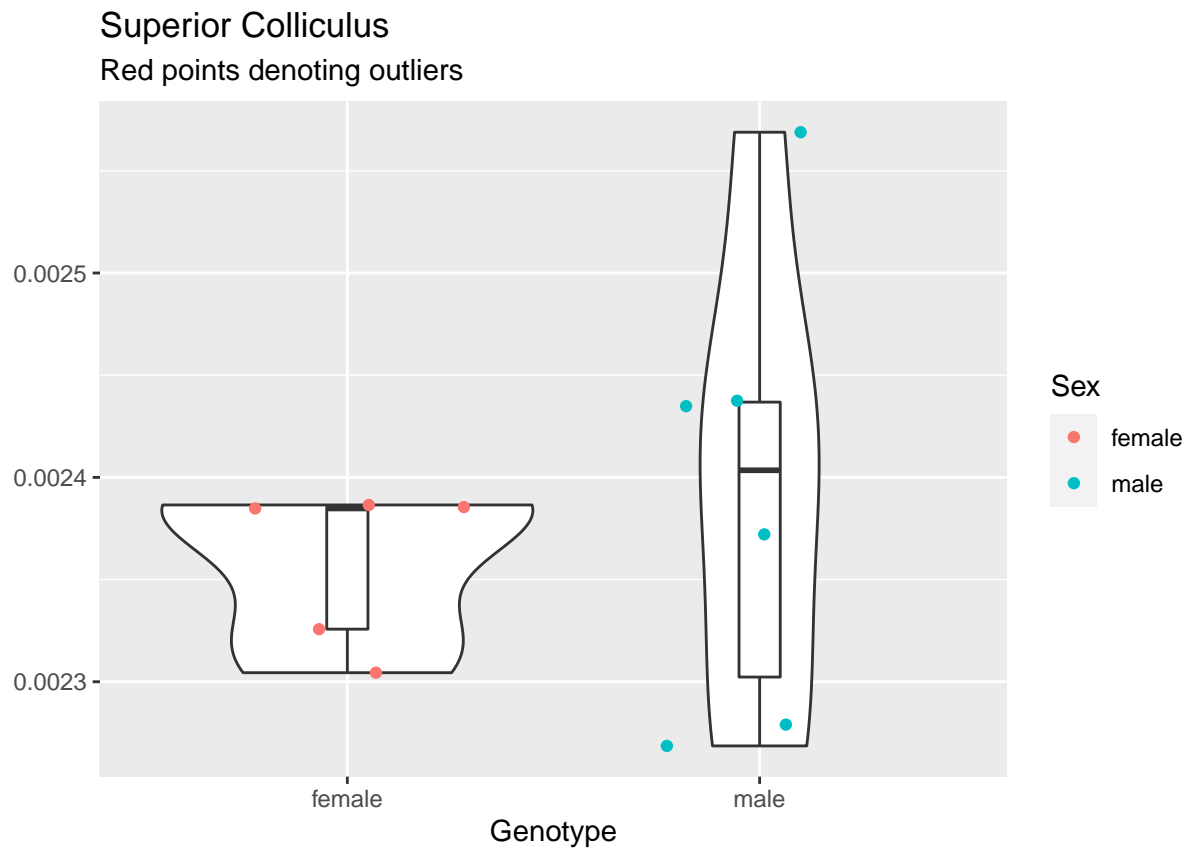
Red points denoting outliers



```
##           Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex         1 5.241e-08 5.241e-08   6.314 0.0332 *
## Residuals   9 7.471e-08 8.300e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```



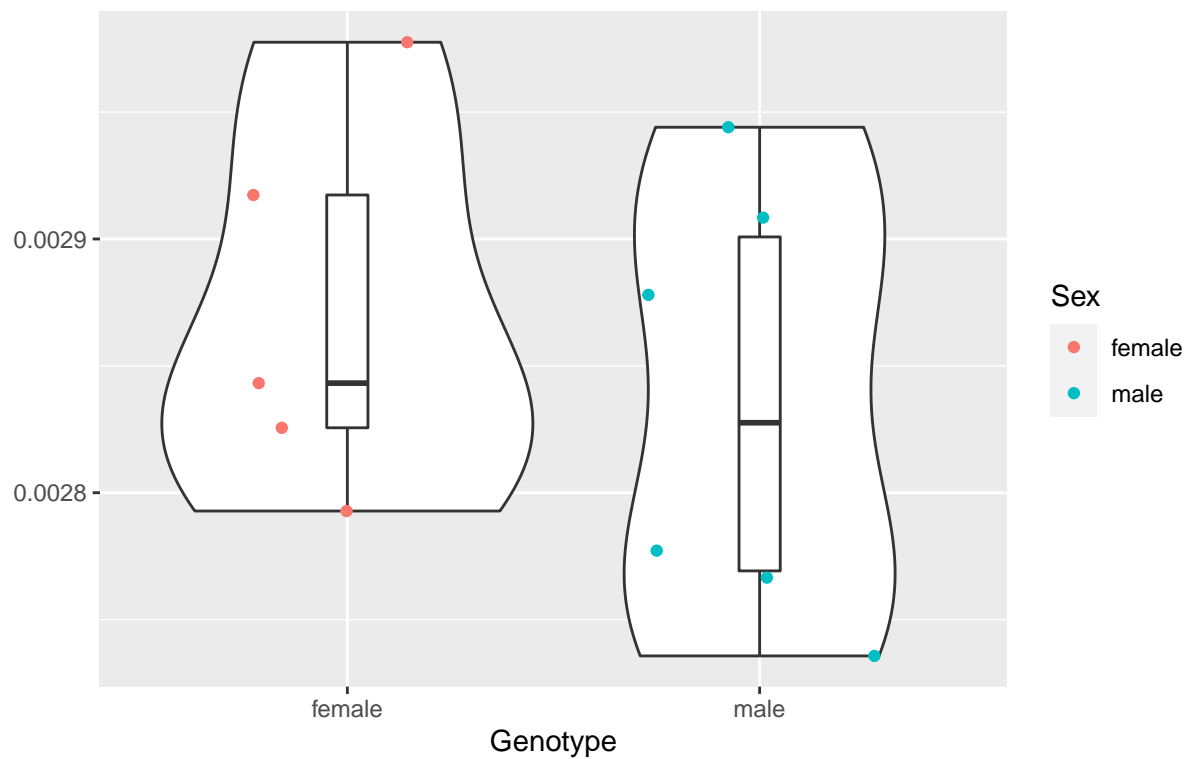
```
##          Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex        1 5.482e-08 5.482e-08   2.024  0.189
## Residuals  9 2.438e-07 2.708e-08
```



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	3.560e-09	3.556e-09	0.459	0.515
## Residuals	9	6.976e-08	7.751e-09		

Deep Mesencephalic Nuclei

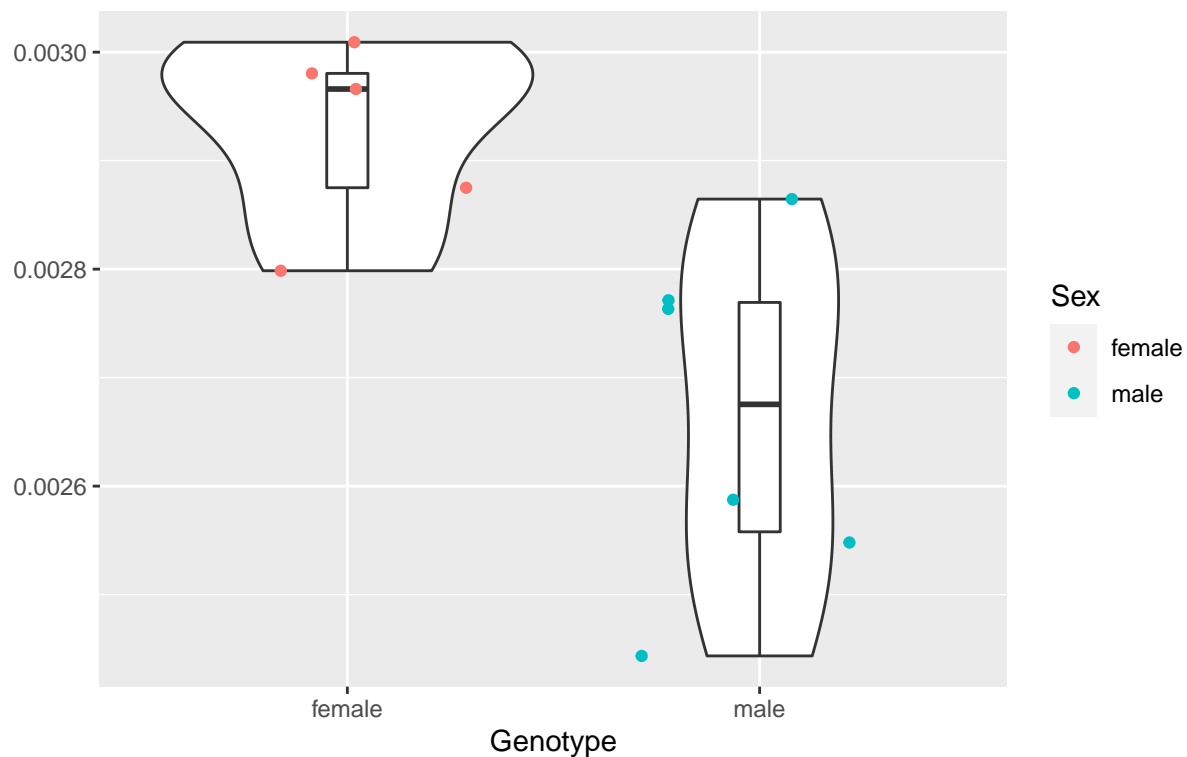
Red points denoting outliers



```
##          Df    Sum Sq  Mean Sq F value Pr(>F)
## Sex       1 3.600e-09 3.597e-09   0.544  0.479
## Residuals 9 5.947e-08 6.608e-09
```

Subbrachial Nucleus and Peripeduncular Nucleus

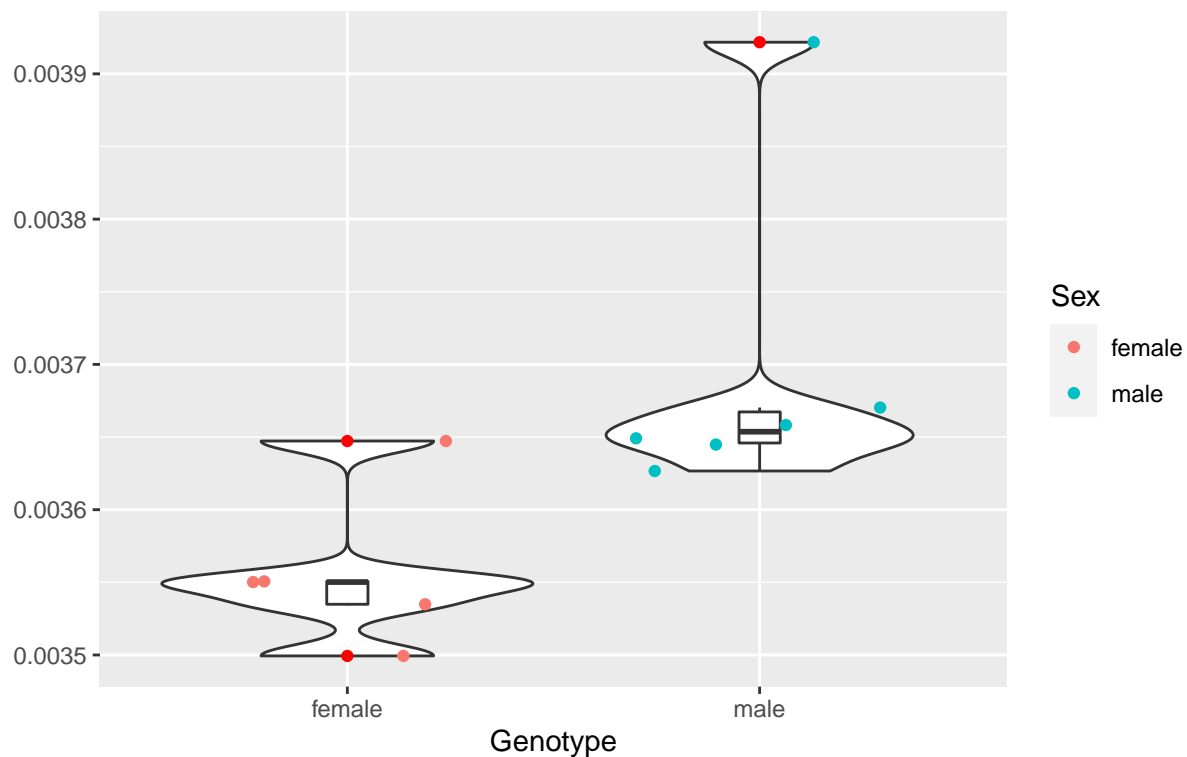
Red points denoting outliers



```
##          Df    Sum Sq  Mean Sq F value Pr(>F)
## Sex          1 1.883e-07 1.883e-07   10.6 0.0099 **
## Residuals    9 1.599e-07 1.776e-08
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Reticular Nucleus of Thalamus

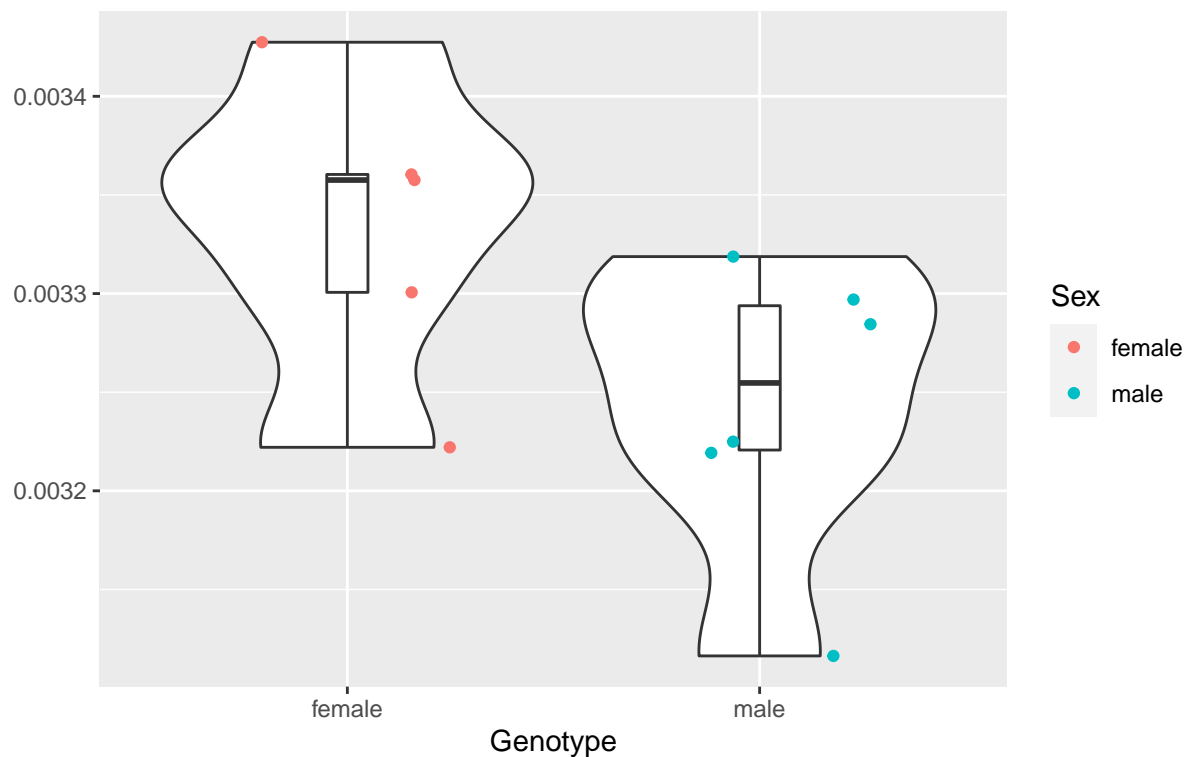
Red points denoting outliers



```
##          Df    Sum Sq  Mean Sq F value Pr(>F)
## Sex          1 5.246e-08 5.246e-08   6.318 0.0331 *
## Residuals    9 7.473e-08 8.300e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Zona Incerta

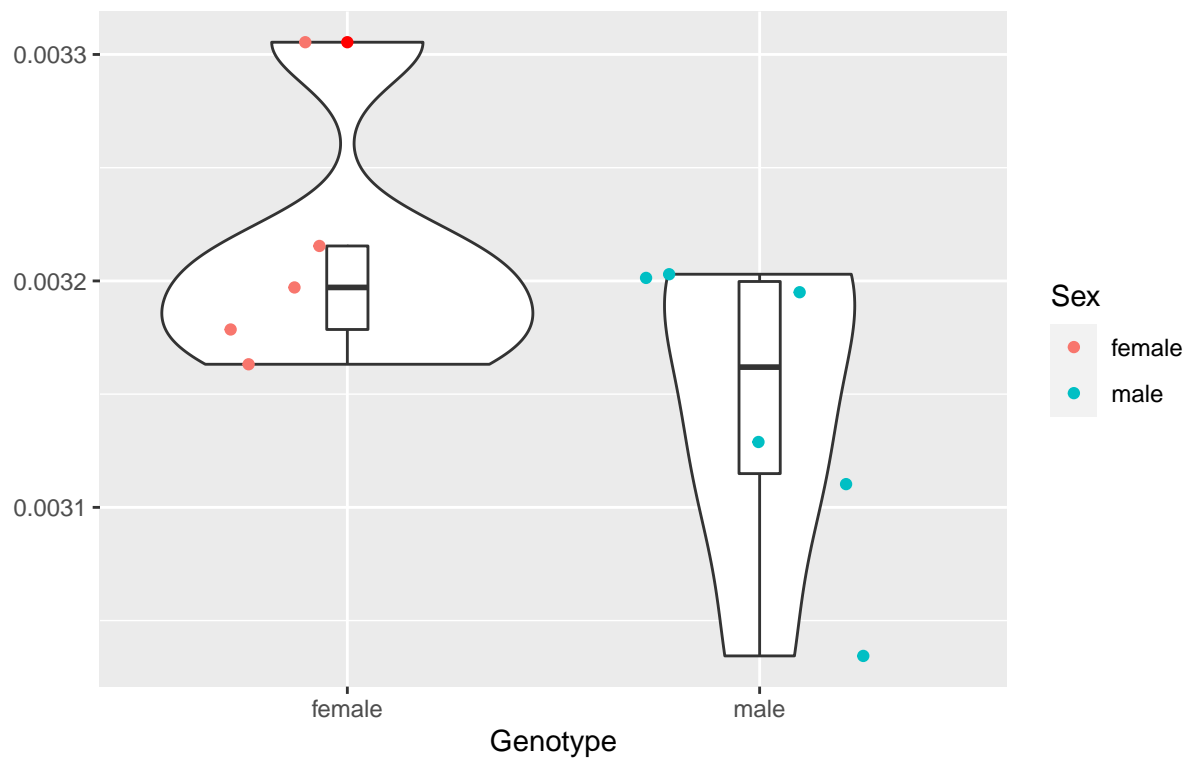
Red points denoting outliers



```
##           Df    Sum Sq  Mean Sq F value Pr(>F)
## Sex          1 2.217e-08 2.217e-08   3.918 0.0791 .
## Residuals    9 5.092e-08 5.658e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```


Lateral Geniculate Nucleus

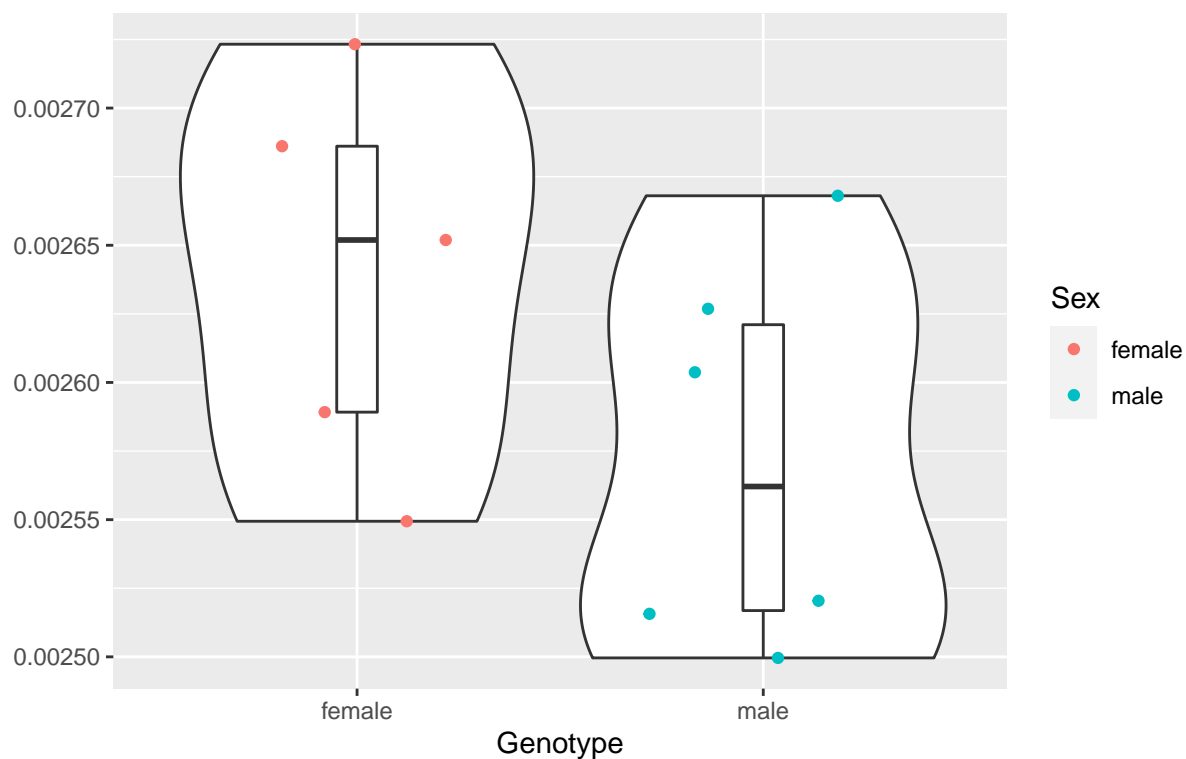
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.205e-08	1.205e-08	3.082	0.113
## Residuals	9	3.519e-08	3.910e-09		

Medial Geniculate Nucleus

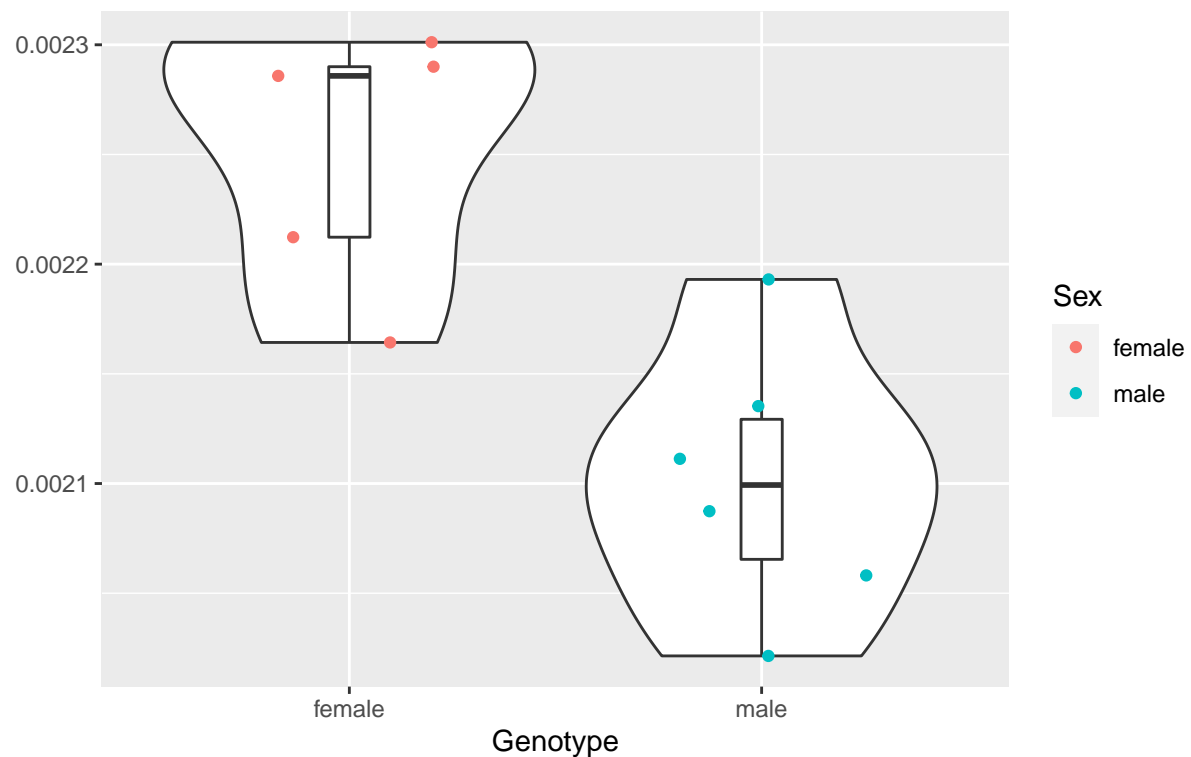
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.246e-08	1.246e-08	2.532	0.146
## Residuals	9	4.429e-08	4.922e-09		

Latero Dorsal Nucleus of Thalamus

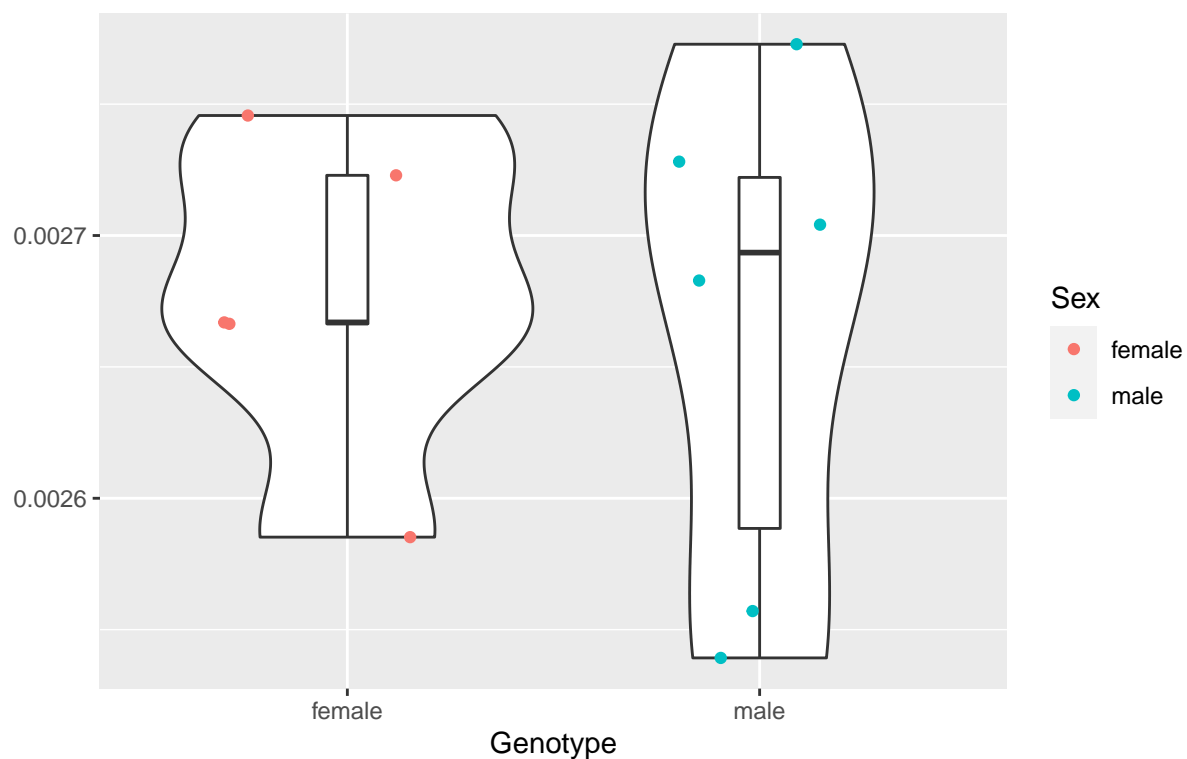
Red points denoting outliers



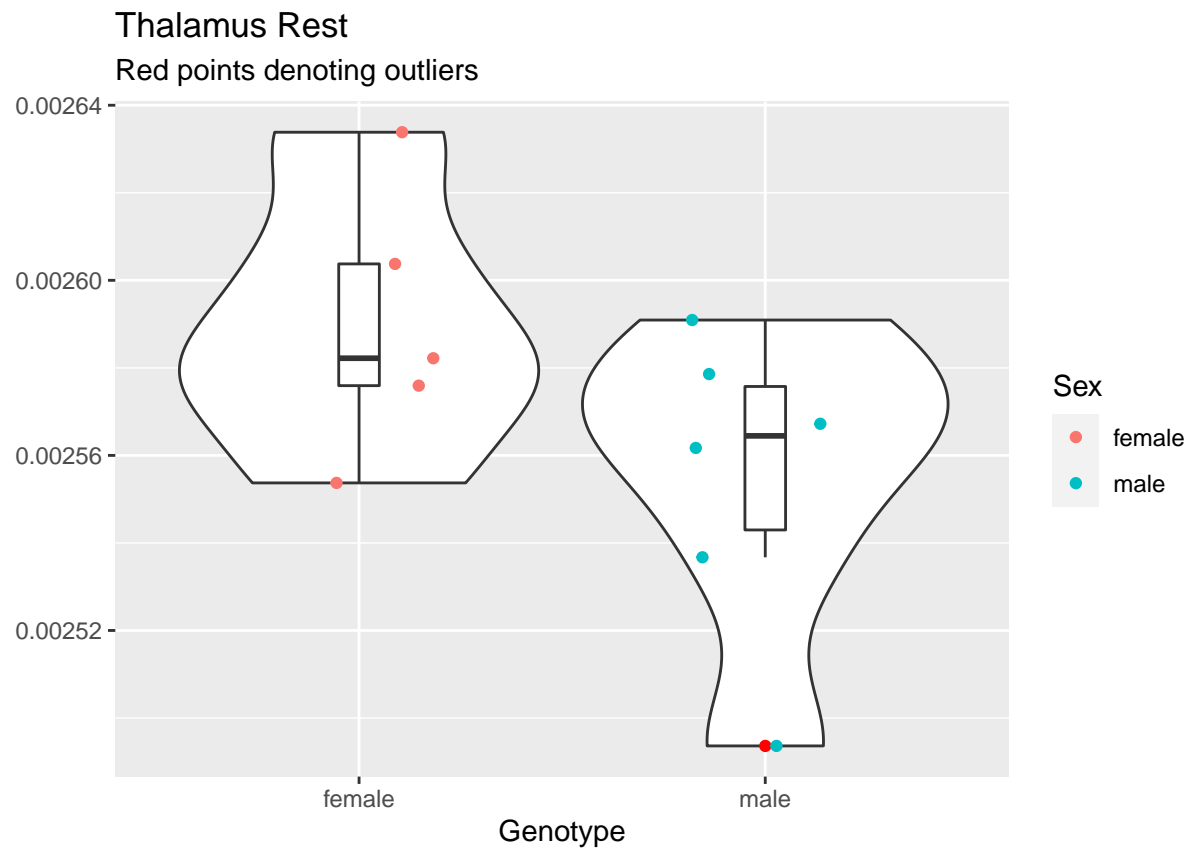
```
##           Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex         1 6.107e-08 6.107e-08   16.98 0.0026 **
## Residuals   9 3.238e-08 3.600e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Ventral Thalamic Nuclei

Red points denoting outliers

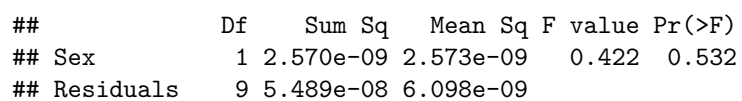


##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	4.900e-10	4.890e-10	0.073	0.793
## Residuals	9	6.037e-08	6.708e-09		



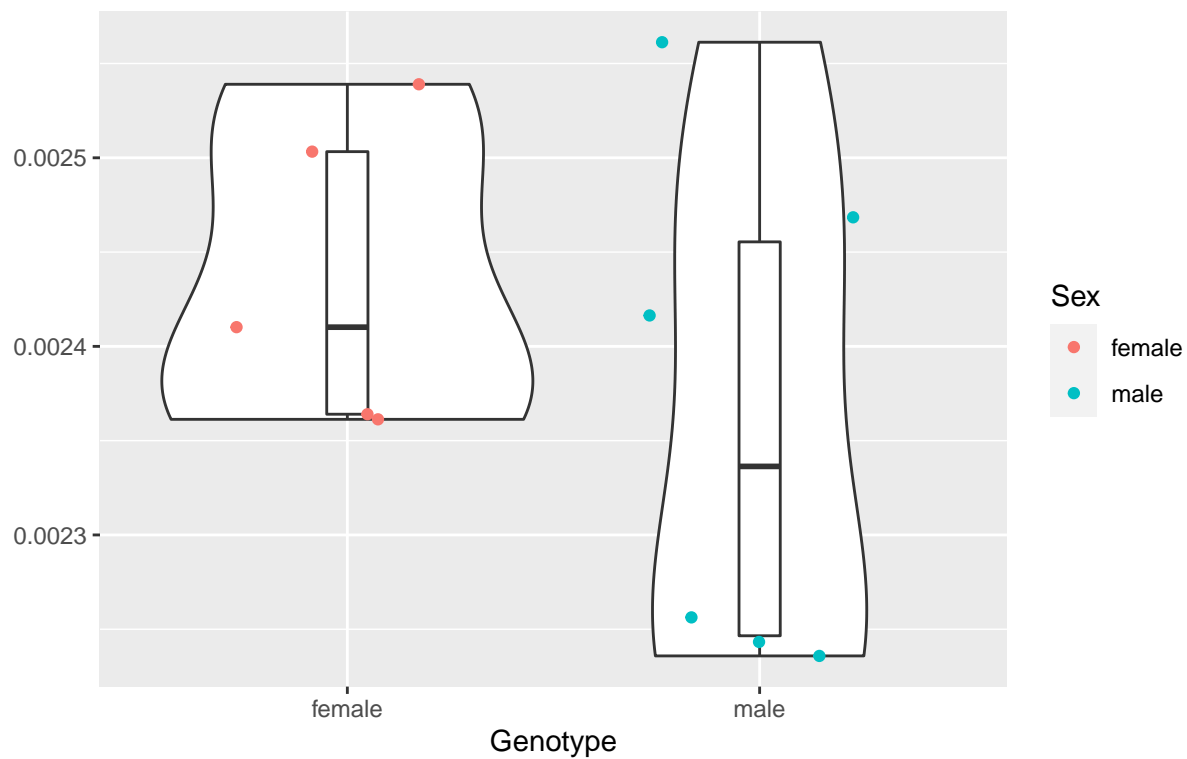
```
##          Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex       1 3.358e-09 3.358e-09   3.075  0.113
## Residuals 9 9.829e-09 1.092e-09
```

Red points denoting outliers



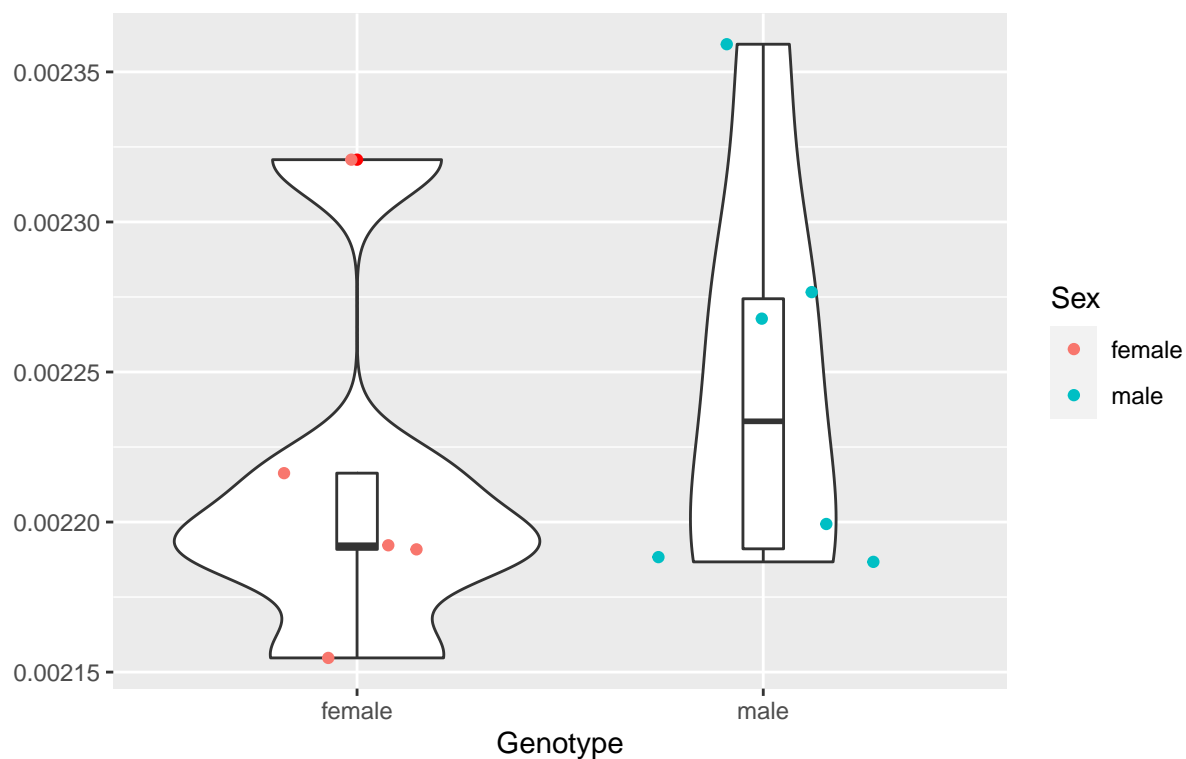
Anterior Pretectal Nucleus

Red points denoting outliers

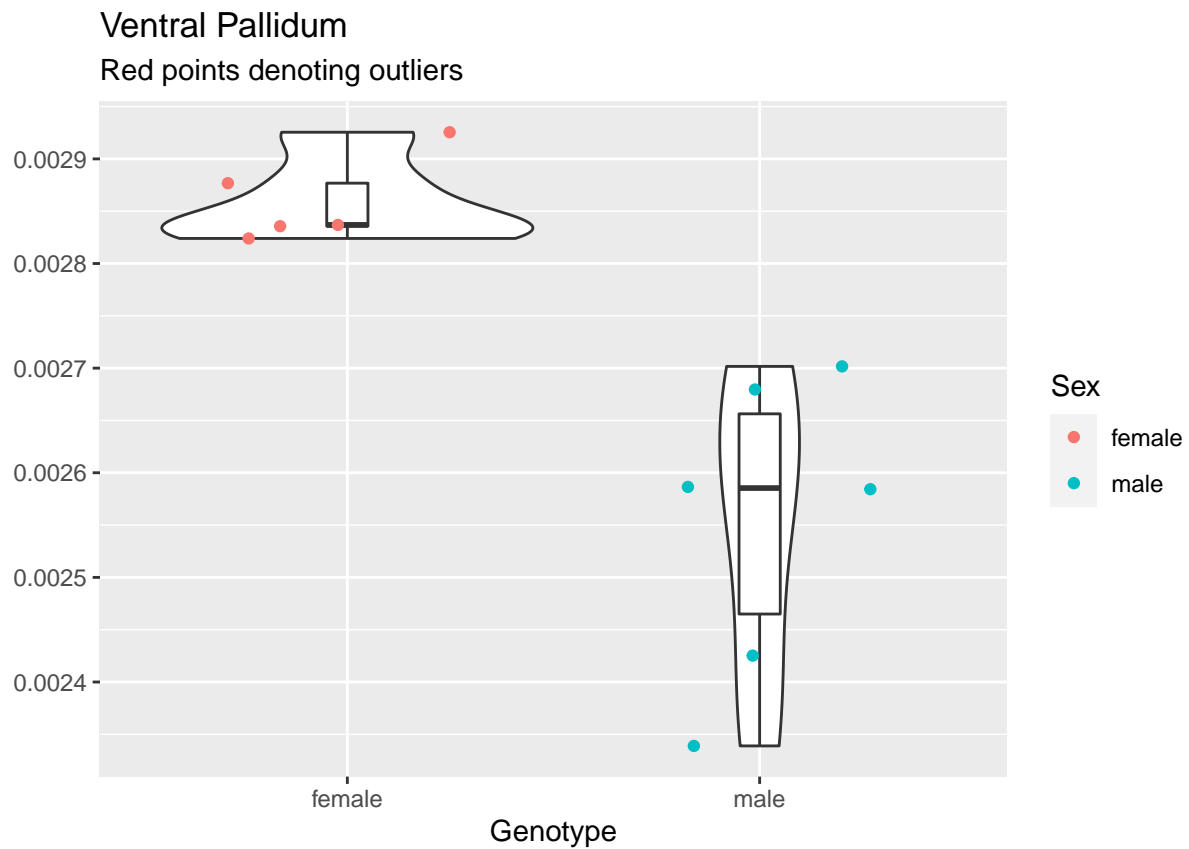


##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.412e-08	1.412e-08	1.044	0.334
## Residuals	9	1.217e-07	1.353e-08		

Periaqueductal Grey Red points denoting outliers



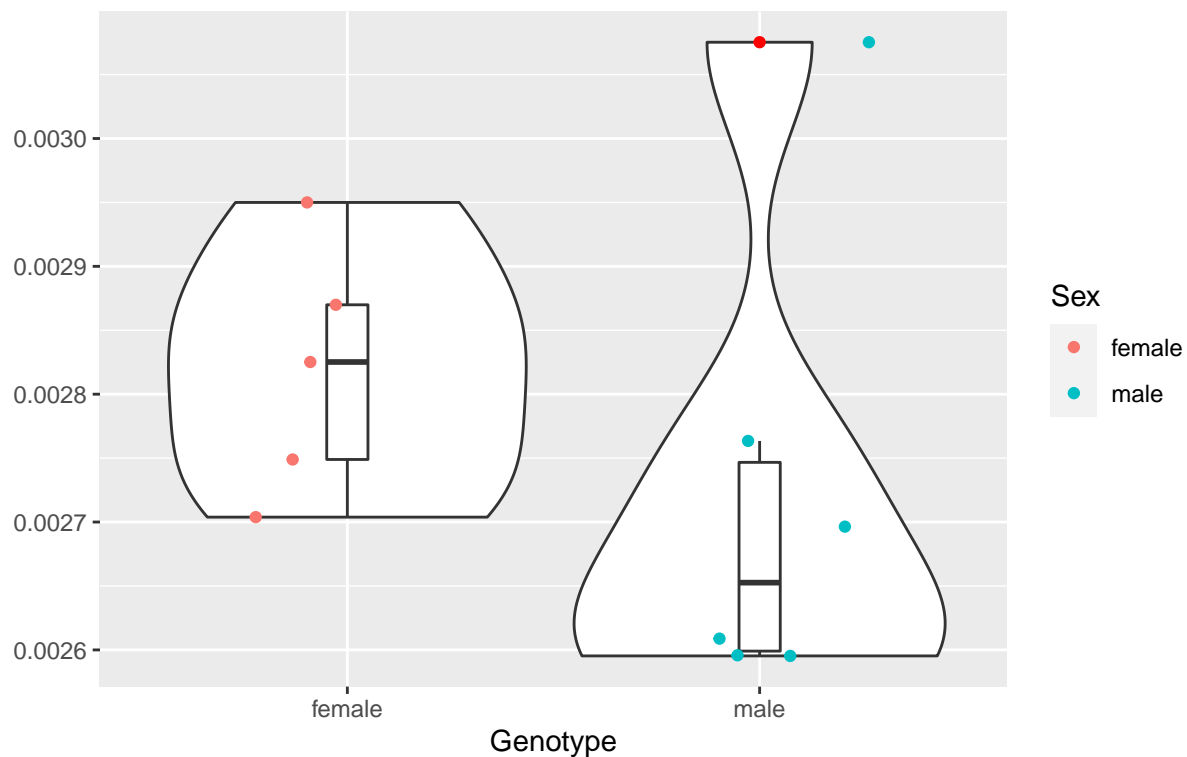
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.680e-09	2.681e-09	0.616	0.453
## Residuals	9	3.917e-08	4.352e-09		



```
##          Df    Sum Sq   Mean Sq F value    Pr(>F)
## Sex          1 2.571e-07 2.571e-07    21.16 0.00129 **
## Residuals    9 1.094e-07 1.215e-08
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Bed Nucleus of the Stria Terminalis

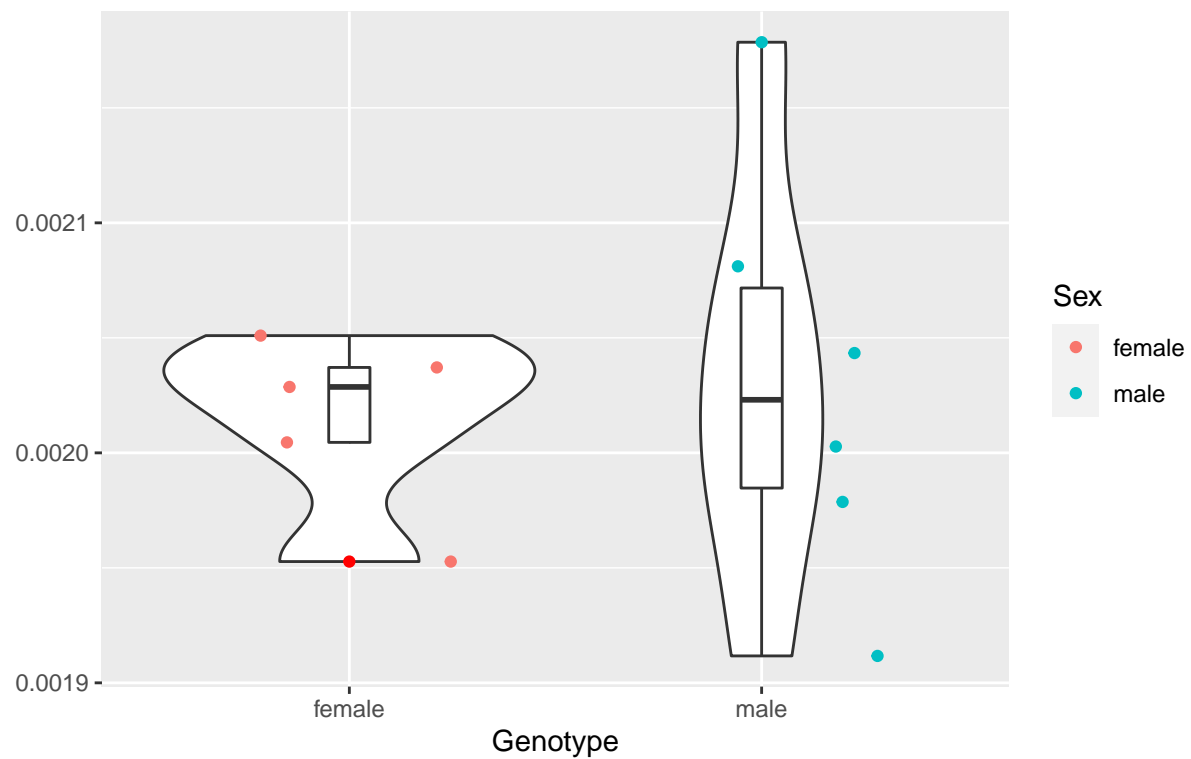
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.570e-08	2.570e-08	1.102	0.321
## Residuals	9	2.099e-07	2.333e-08		

Acumbens

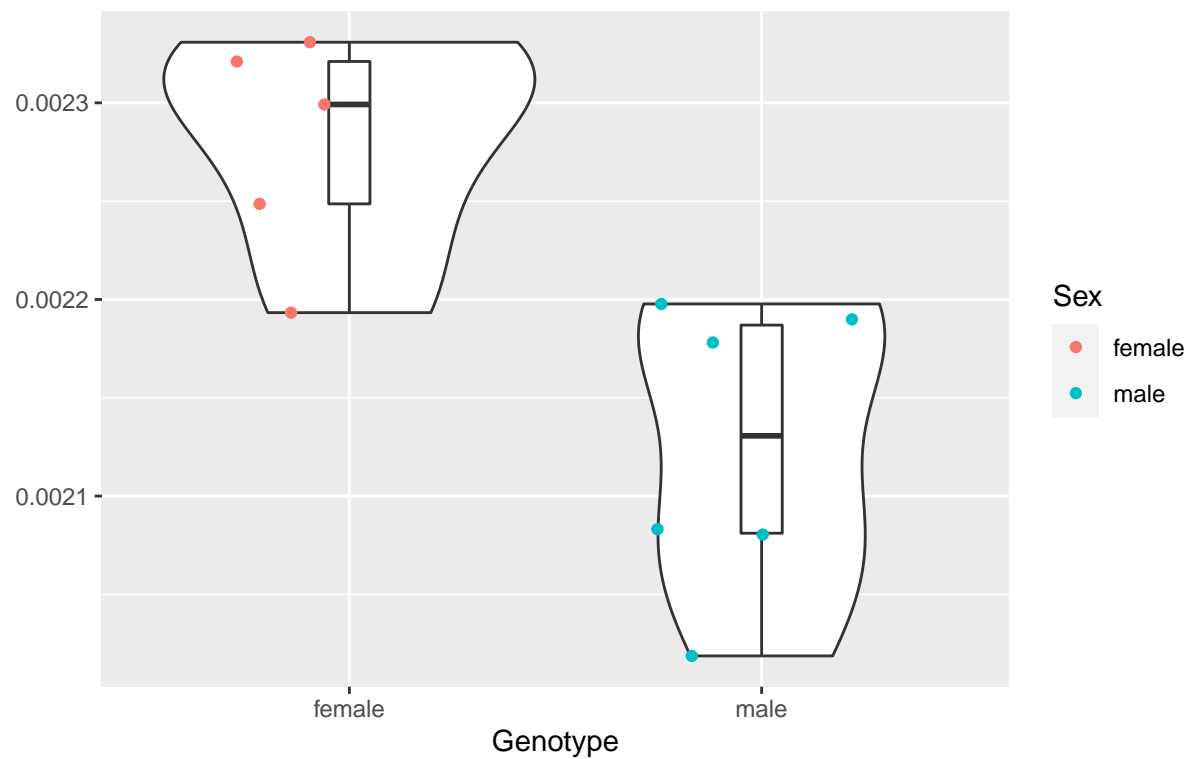
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	8.700e-10	8.720e-10	0.163	0.696
## Residuals	9	4.814e-08	5.349e-09		

Amygdala

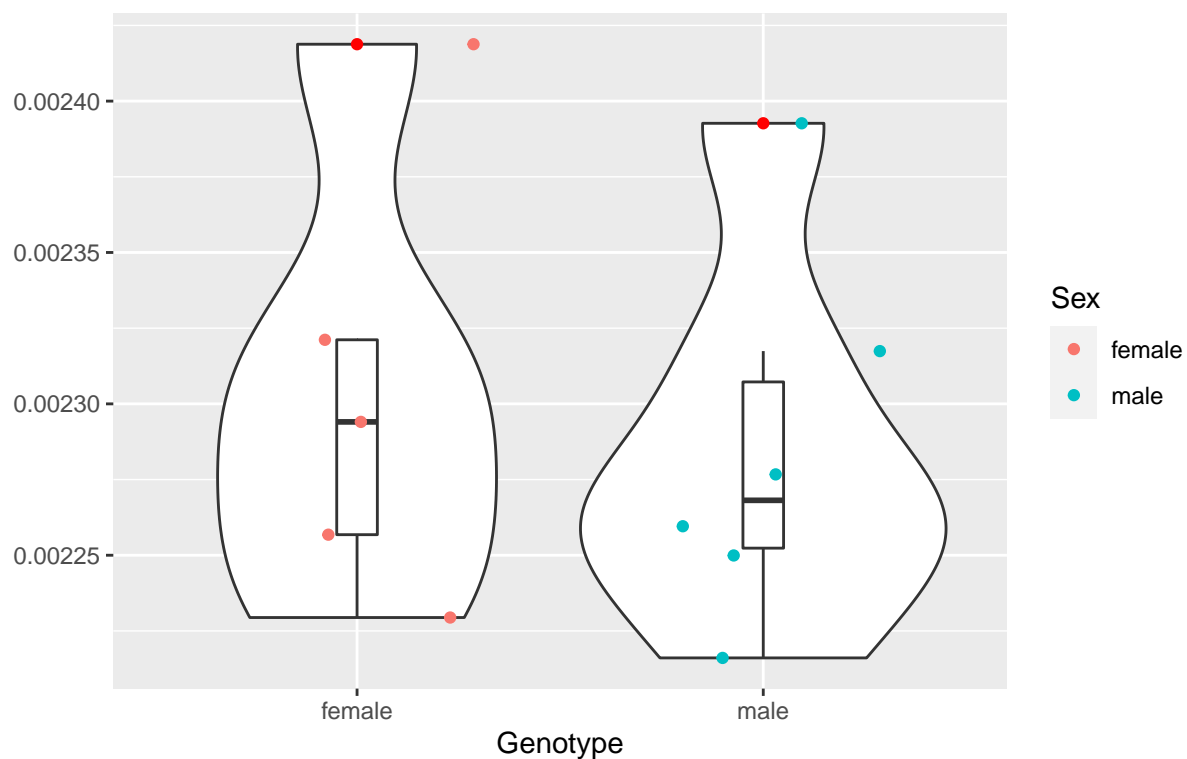
Red points denoting outliers



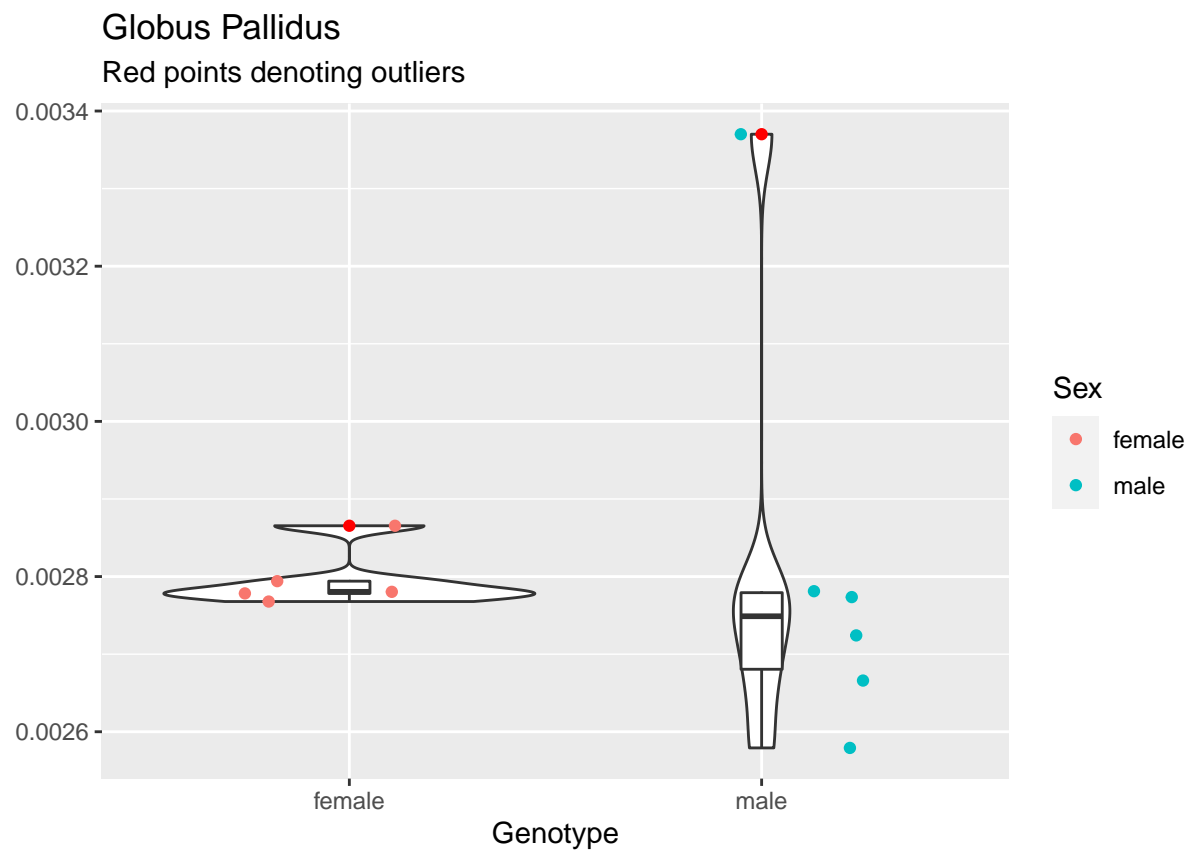
```
##           Df    Sum Sq   Mean Sq F value    Pr(>F)
## Sex          1 6.462e-08 6.462e-08   14.37 0.00428 **
## Residuals    9 4.048e-08 4.500e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

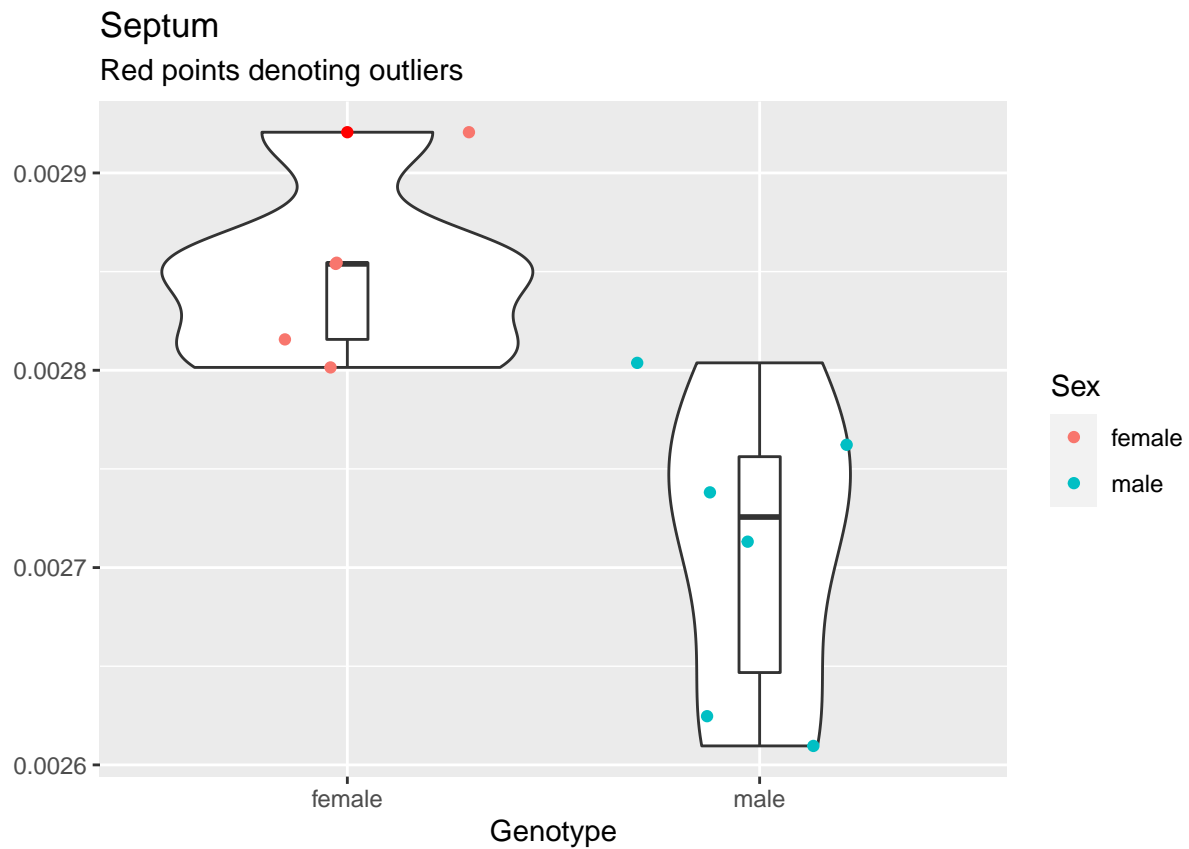
Striatum

Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	9.50e-10	9.480e-10	0.21	0.658
## Residuals	9	4.07e-08	4.522e-09		

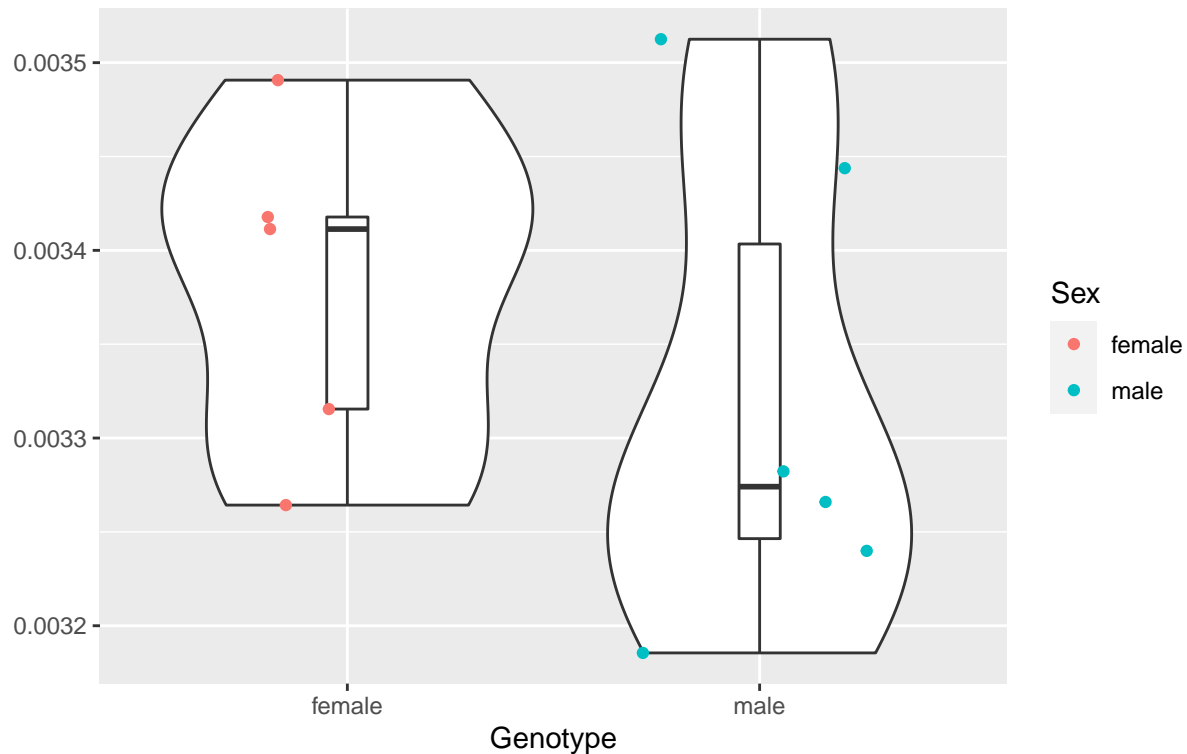




```
##          Df    Sum Sq   Mean Sq F value    Pr(>F)
## Sex          1 5.394e-08 5.394e-08    12.7 0.00608 **
## Residuals    9 3.822e-08 4.250e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Subthalamic Nucleus

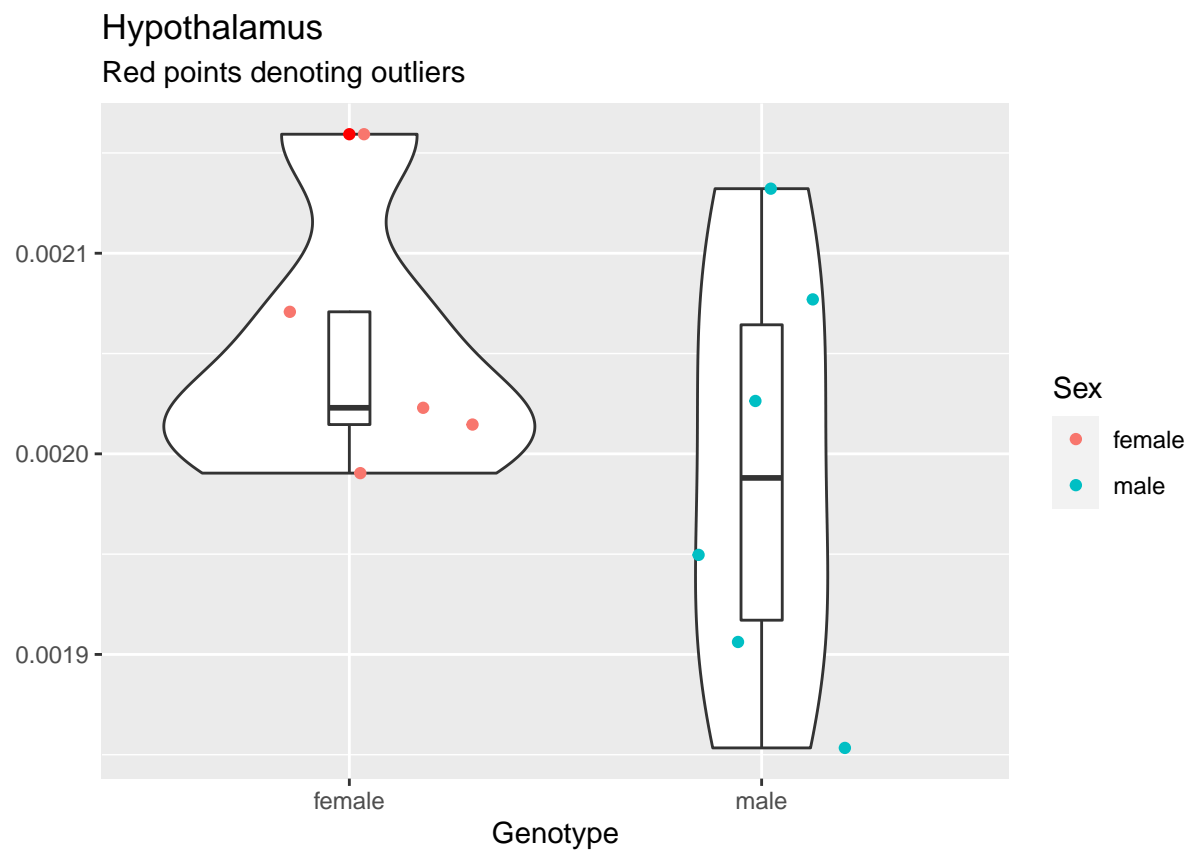
Red points denoting outliers



```
##           Df      Sum Sq   Mean Sq F value Pr(>F)
## Sex         1 9.260e-09 9.256e-09   0.734  0.414
## Residuals   9 1.134e-07 1.260e-08
```

```
#“{r Preopt, echo = FALSE} #ggplot(data = apoe2, aes(factor(Sex), Preopt)) + #geom_violin() +
#geom_boxplot(width = 0.1, outlier.color = “red”) + #geom_jitter(height = 0, width = 0.3, aes(color
= Sex)) + #labs(x = “Genotype”, #y = “”, #title =”Preoptic Telencephalon“, #subtitle =”Red points
denoting outliers”)
```

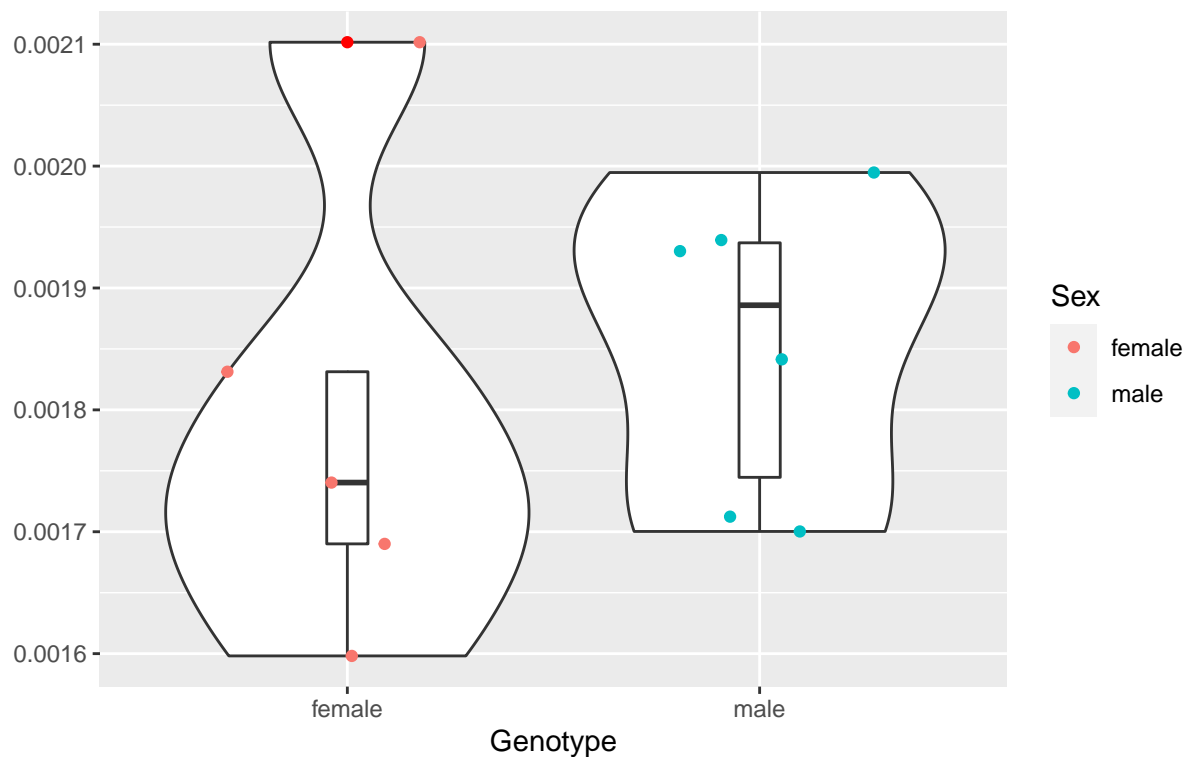
```
#res.aov <- aov(Preopt ~ Sex, data = apoe2) #summary(res.aov) #“
```

##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.009e-08	1.009e-08	1.221	0.298
## Residuals	9	7.433e-08	8.258e-09		

Amygdalopiriform Transition Area

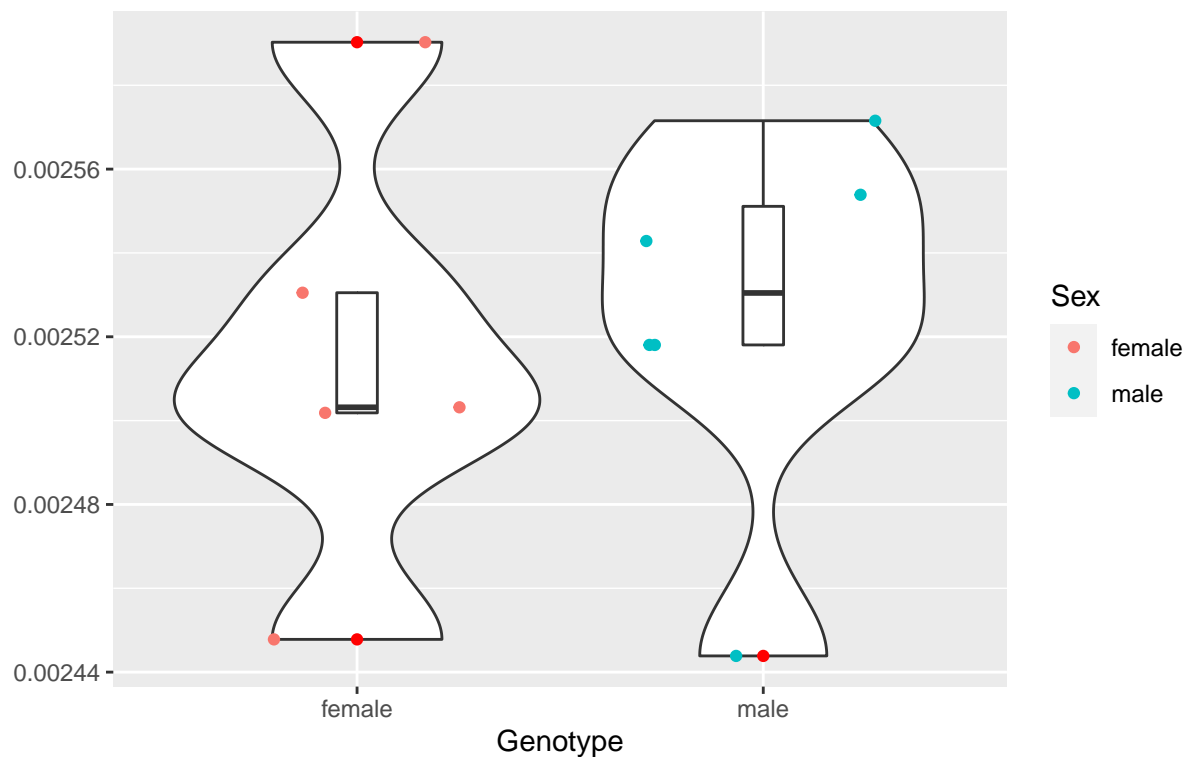
Red points denoting outliers



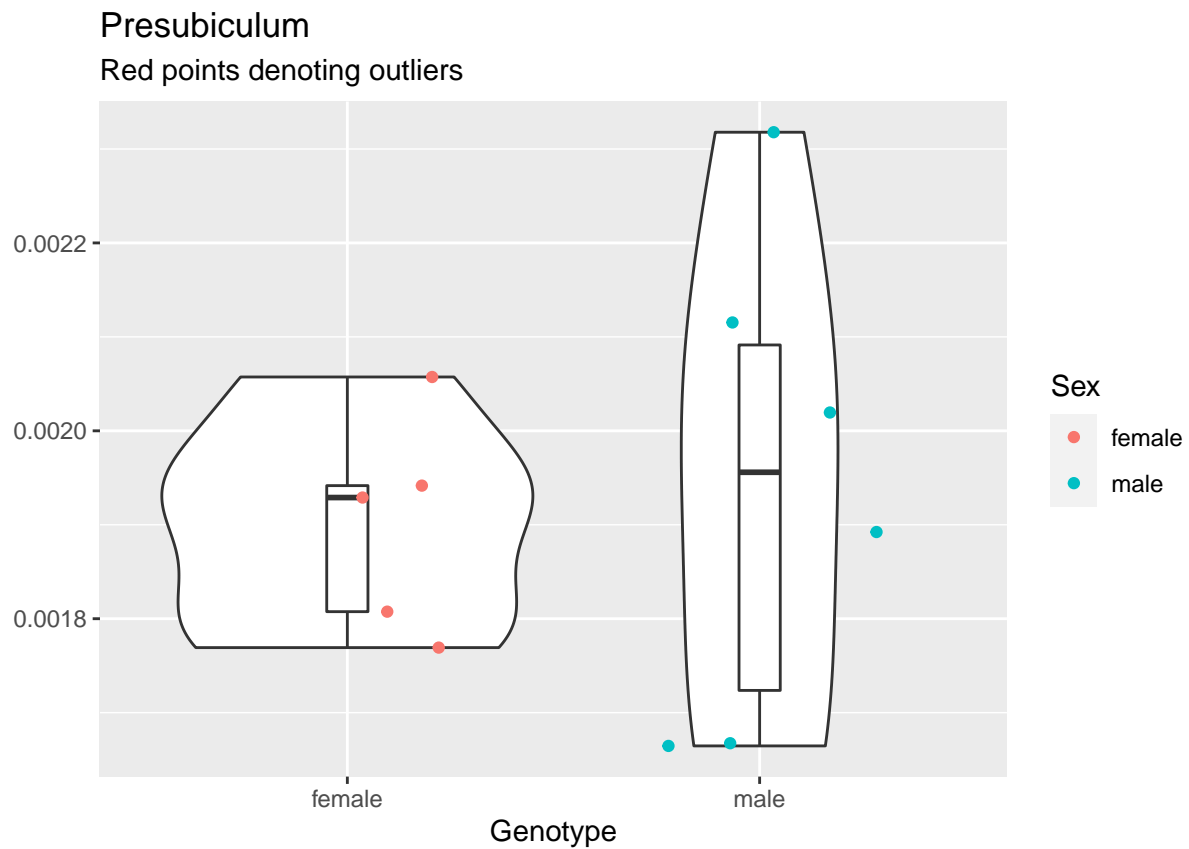
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.007e-08	1.007e-08	0.403	0.541
## Residuals	9	2.248e-07	2.498e-08		

Periform Cortex

Red points denoting outliers



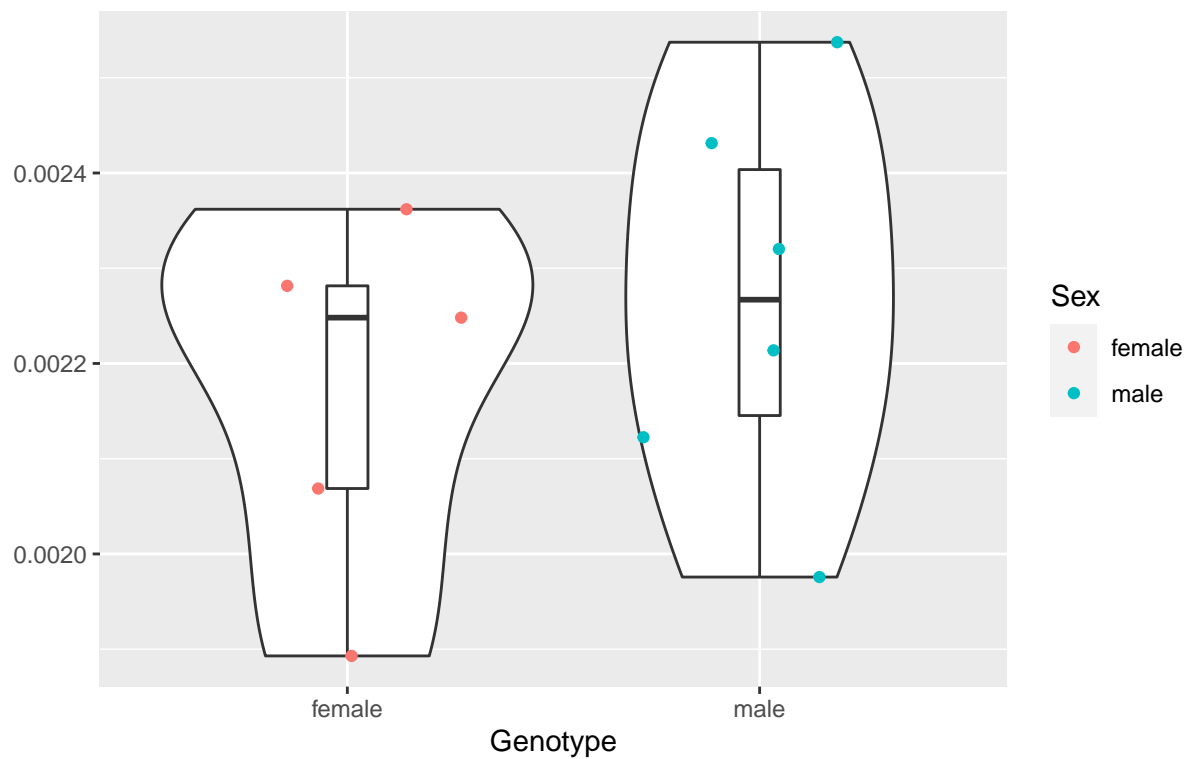
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.720e-10	2.718e-10	0.118	0.739
## Residuals	9	2.073e-08	2.303e-09		



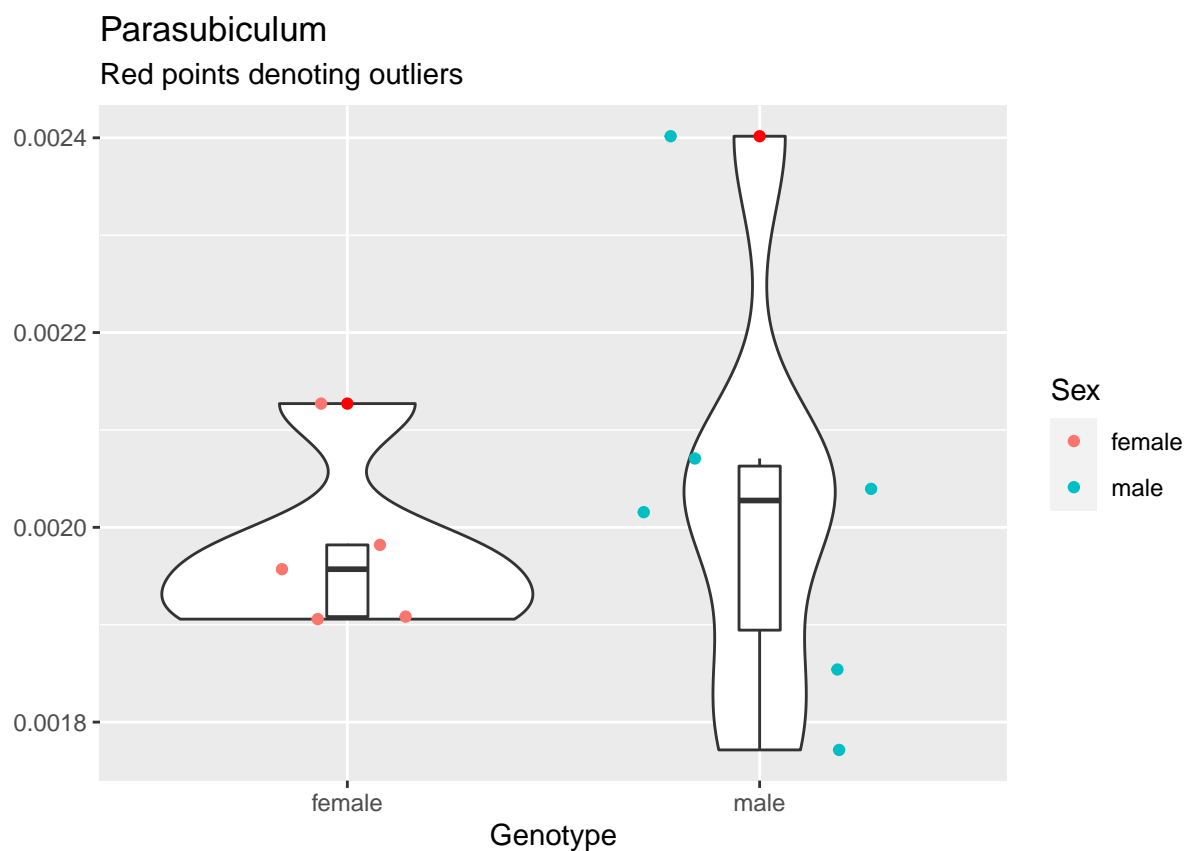
```
##          Df Sum Sq Mean Sq F value Pr(>F)
## Sex      1 5.60e-09 5.580e-09    0.13  0.726
## Residuals 9 3.85e-07 4.278e-08
```

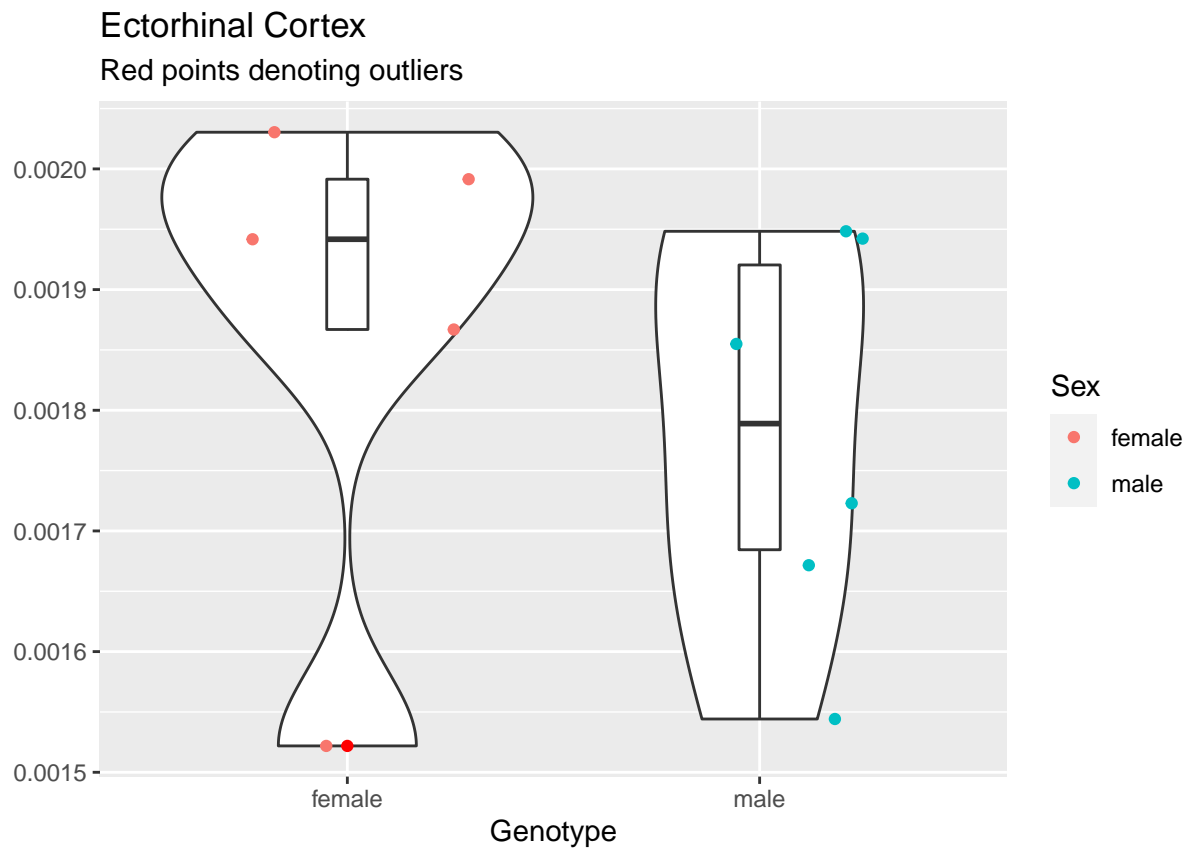
Perirhinal Cortex

Red points denoting outliers



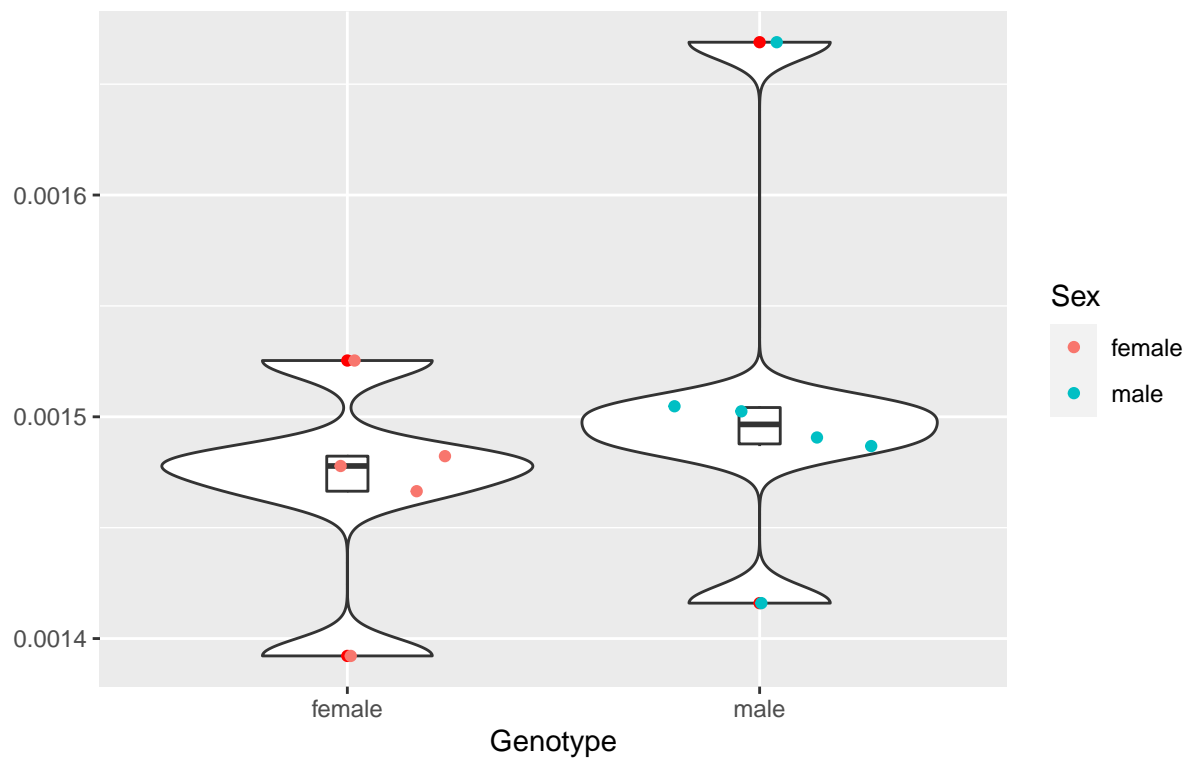
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.520e-08	2.523e-08	0.642	0.444
## Residuals	9	3.539e-07	3.932e-08		





Dorsal Tenia Tecta

Red points denoting outliers



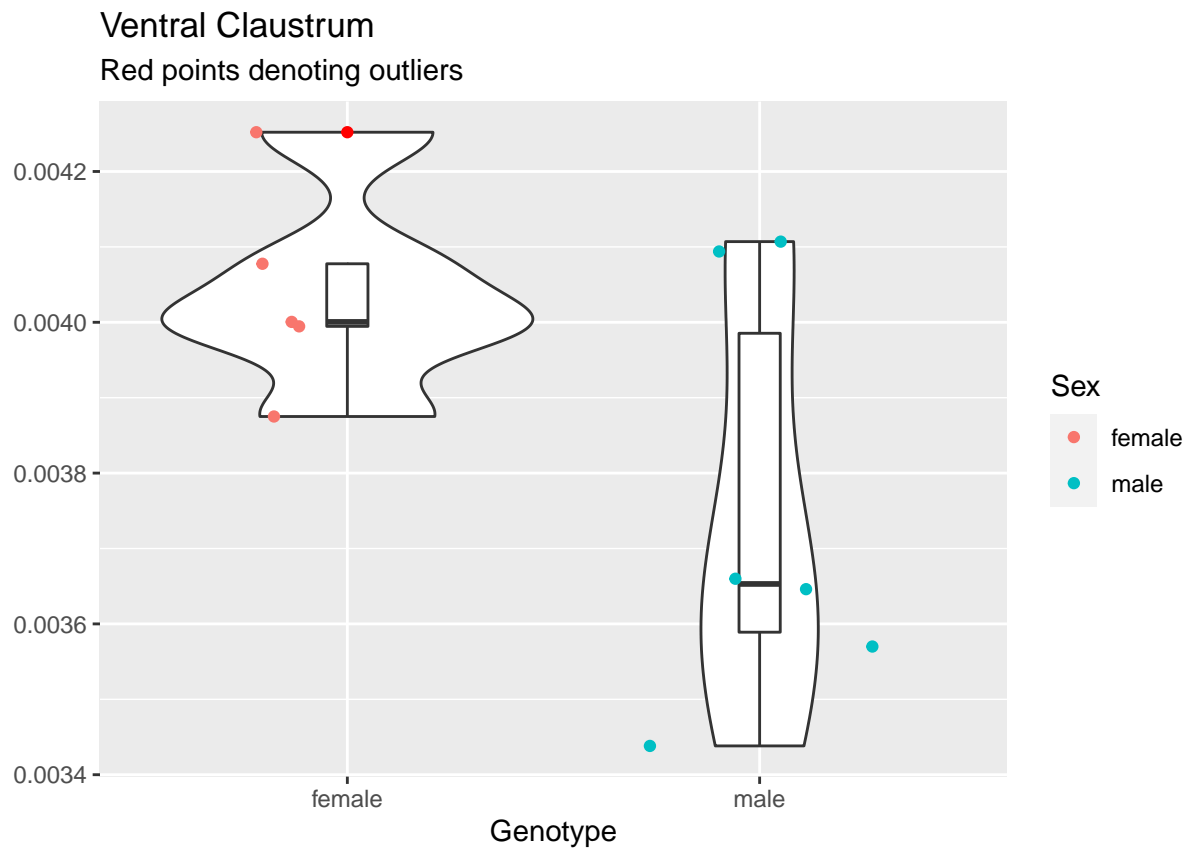
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	4.990e-09	4.992e-09	1.012	0.341
## Residuals	9	4.442e-08	4.935e-09		

Hippocampus

Red points denoting outliers



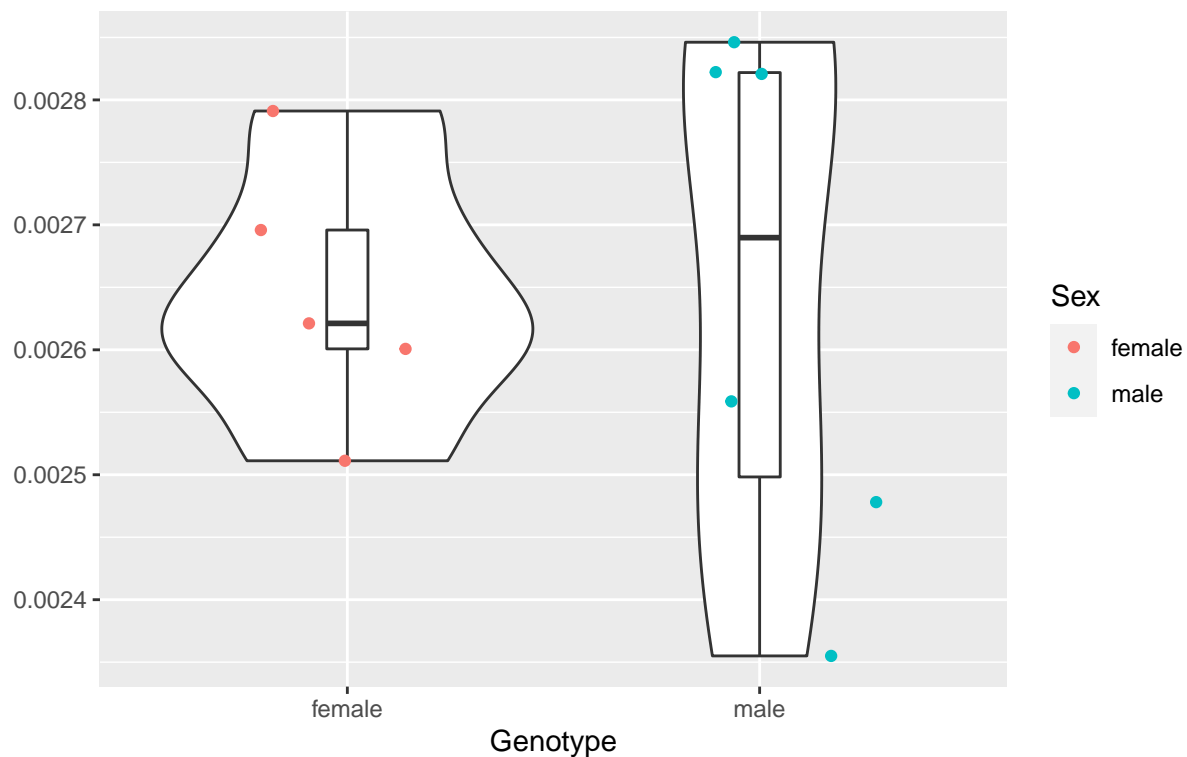
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.171e-09	1.171e-09	0.451	0.519
## Residuals	9	2.336e-08	2.596e-09		



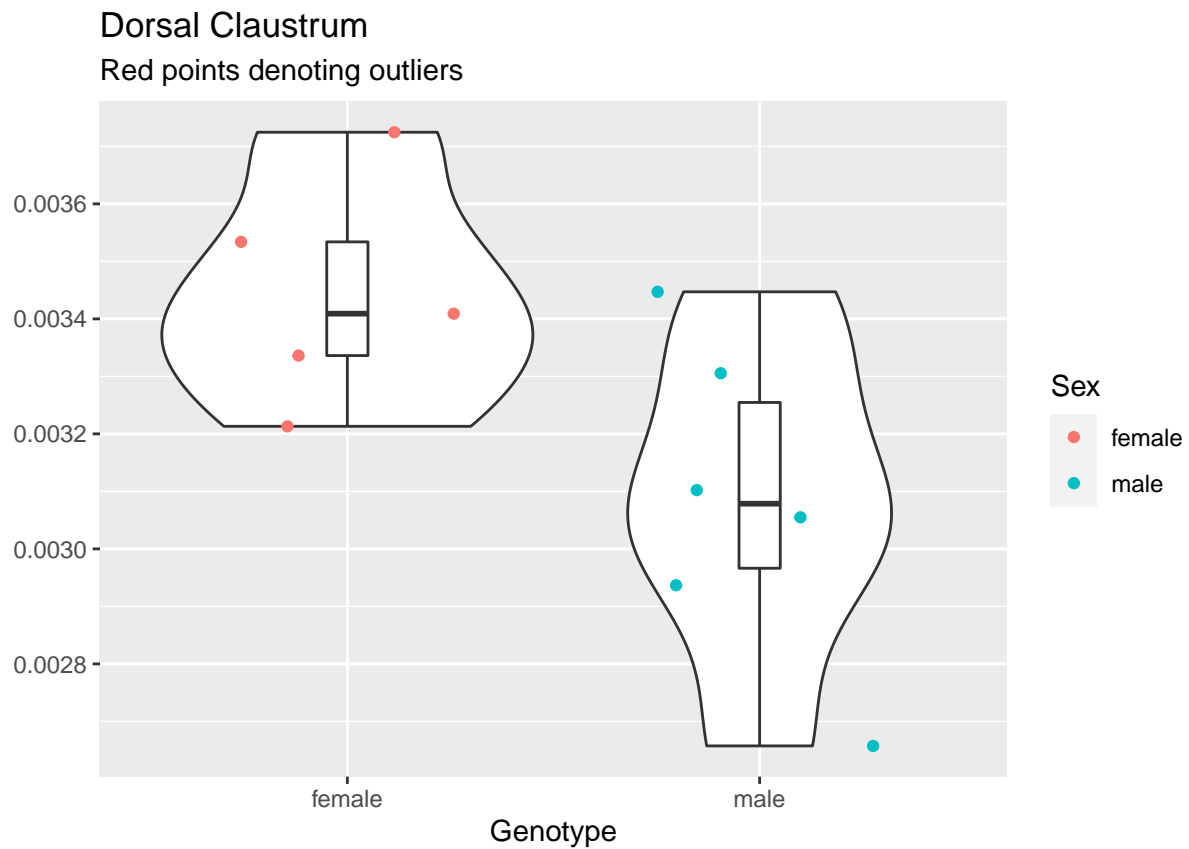
```
##          Df    Sum Sq  Mean Sq F value Pr(>F)
## Sex          1 2.254e-07 2.254e-07   4.302 0.0679 .
## Residuals    9 4.715e-07 5.238e-08
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Posterolateral Cortical Amygdaloid Area

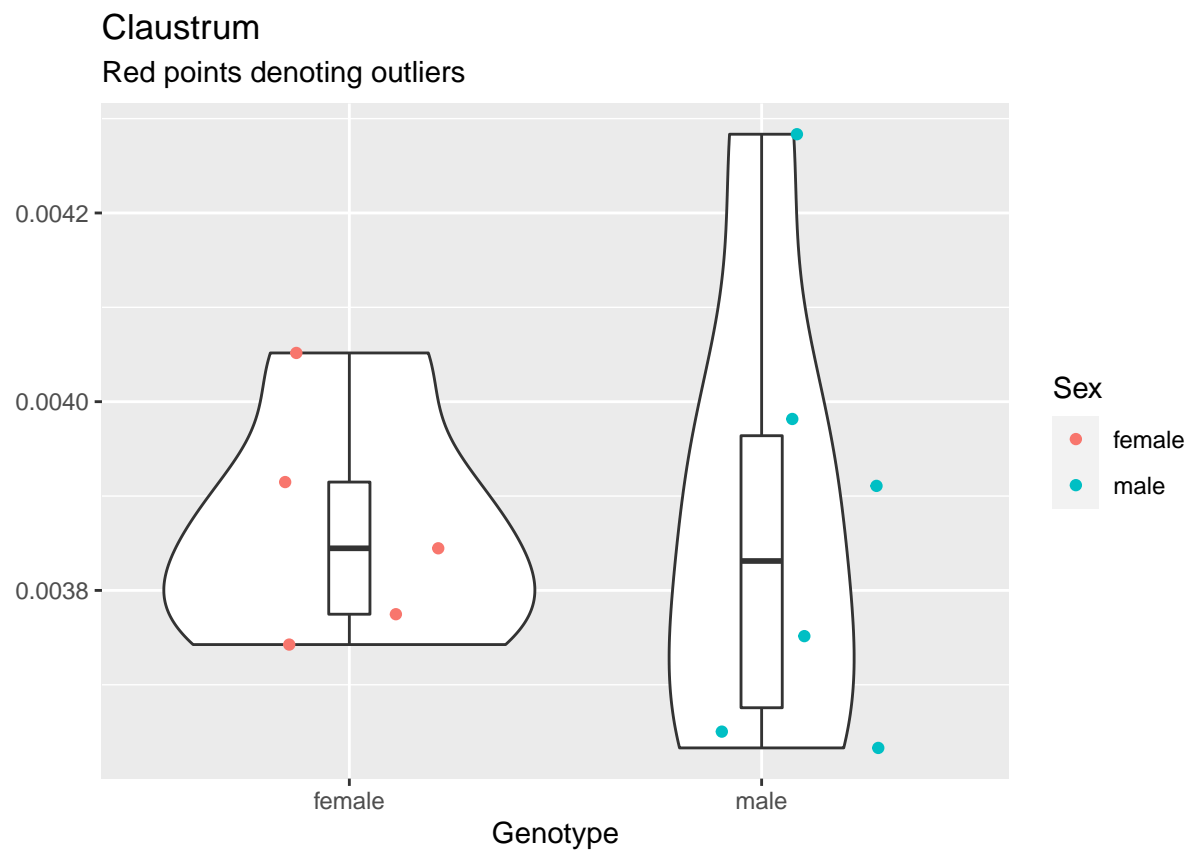
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.000e-11	2.200e-11	0.001	0.979
## Residuals	9	2.666e-07	2.962e-08		



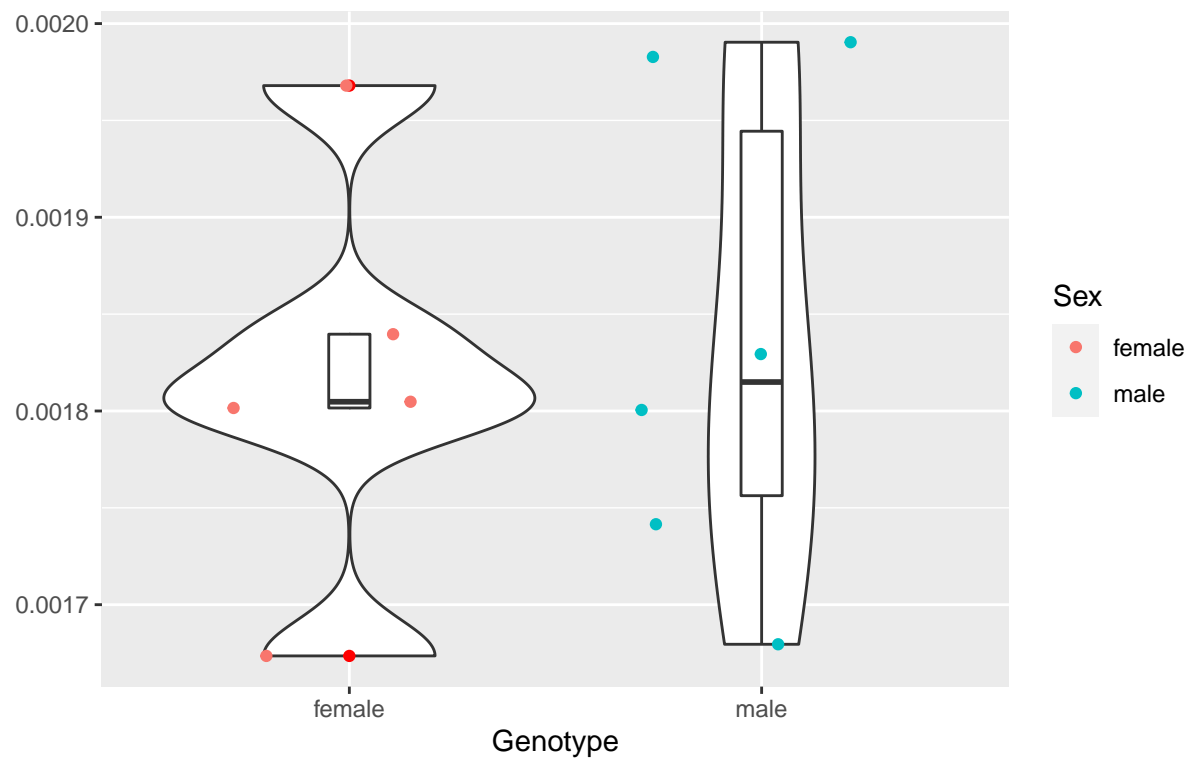
```
##          Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex          1 3.523e-07 3.523e-07   5.886 0.0382 *
## Residuals    9 5.387e-07 5.990e-08
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	0.000e+00	2.000e-11	0.001	0.982
## Residuals	9	3.644e-07	4.049e-08		

Ventral Intermediate Entorhinal Cortex

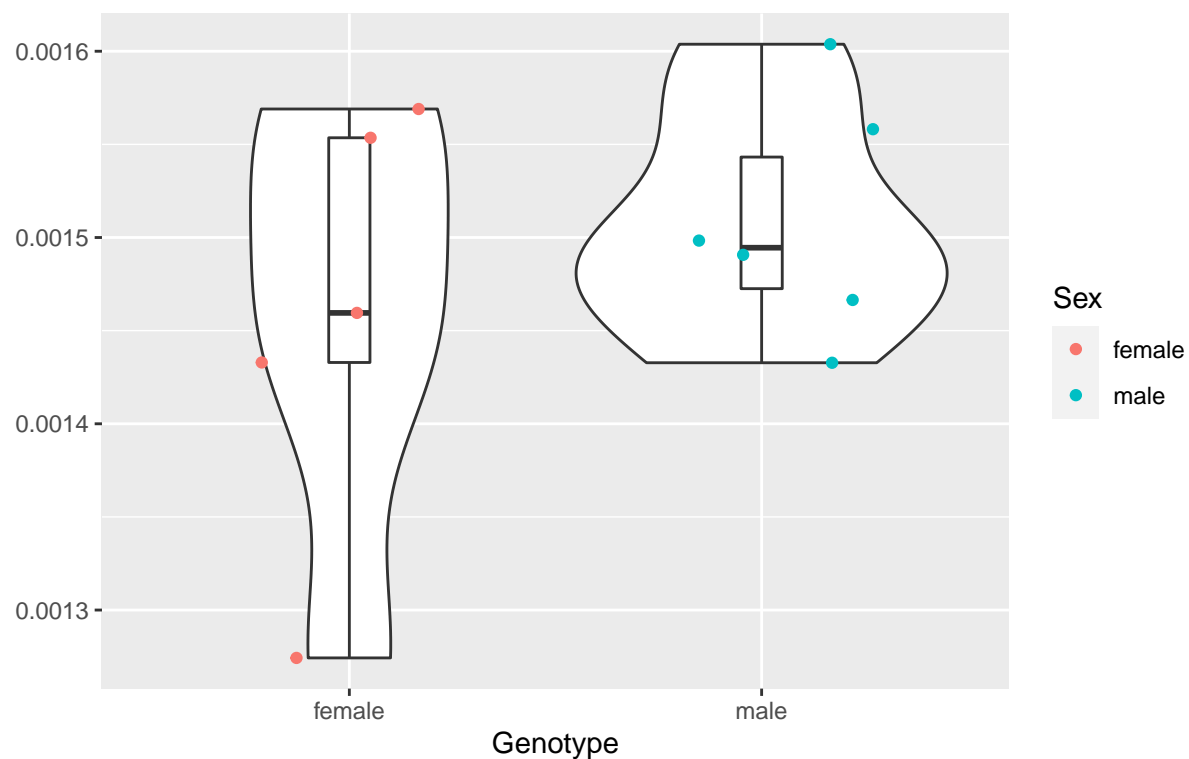
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.080e-09	1.076e-09	0.078	0.787
## Residuals	9	1.243e-07	1.382e-08		

Left Caudomedial Entorhinal Cortex

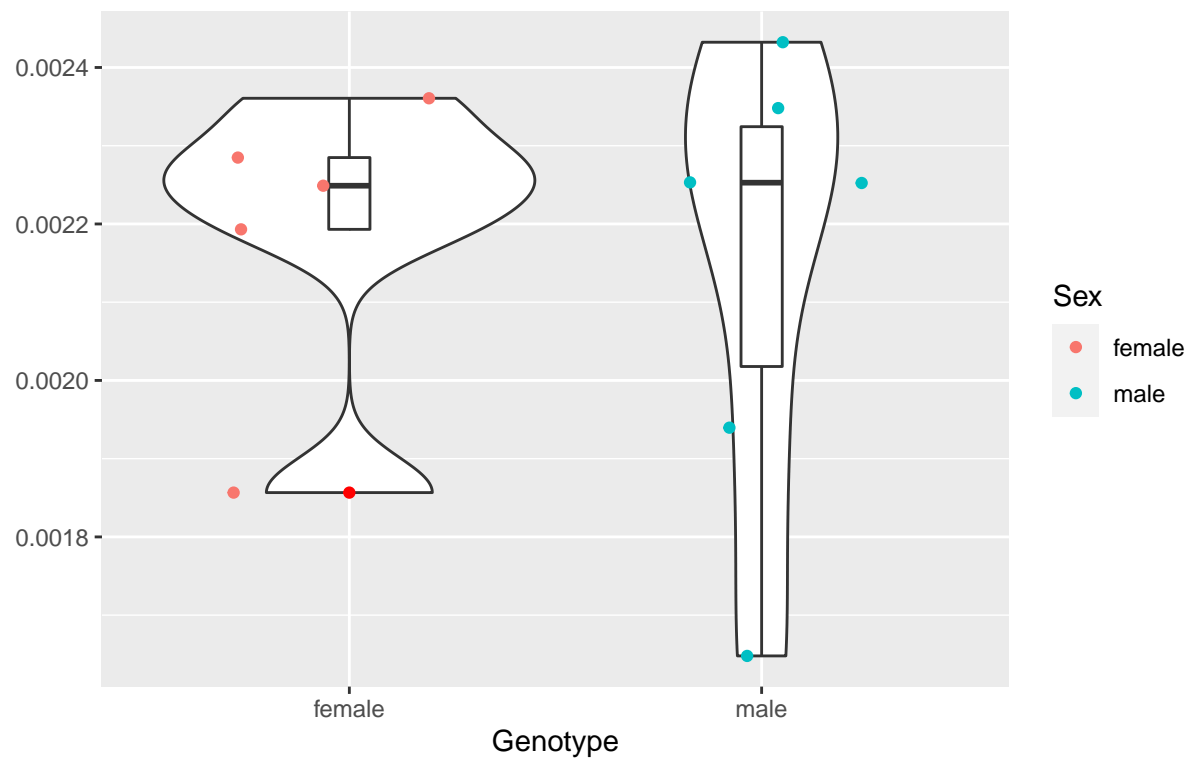
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	6.960e-09	6.960e-09	0.832	0.385
## Residuals	9	7.529e-08	8.365e-09		

Left Dorsolateral Entorhinal Cortex

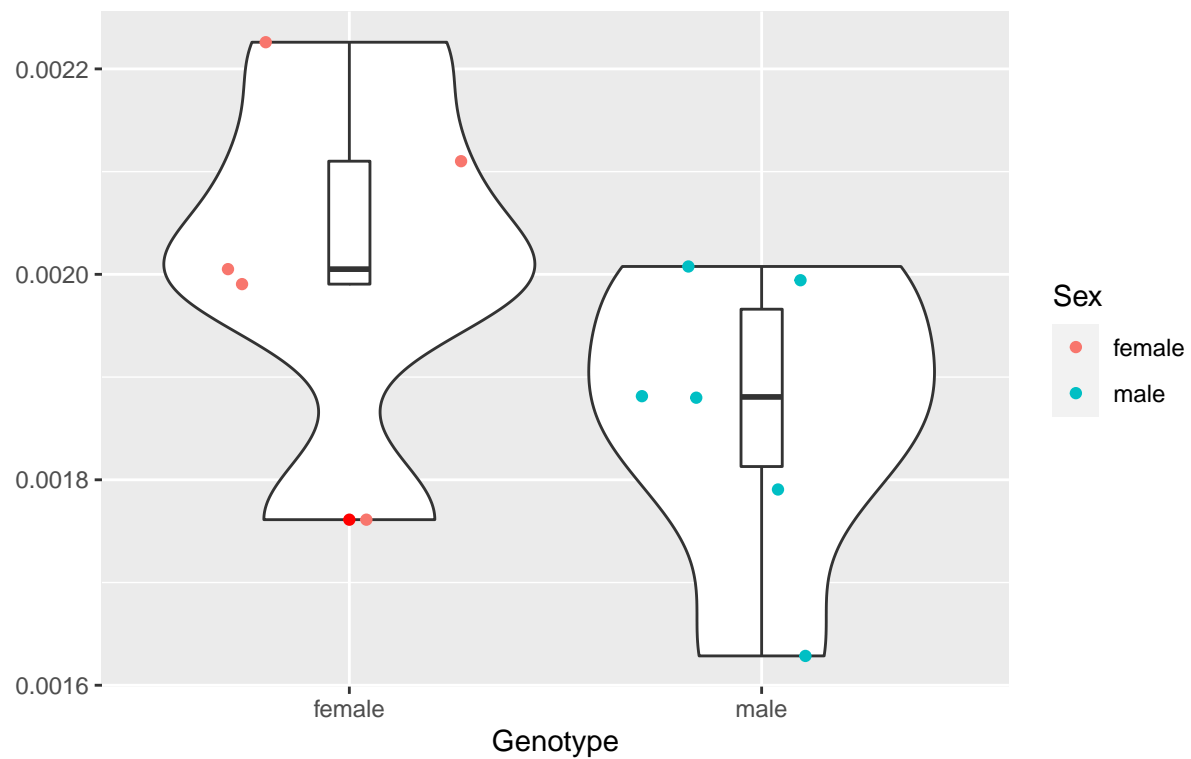
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	5.10e-09	5.100e-09	0.078	0.786
## Residuals	9	5.89e-07	6.545e-08		

Left Dorsal Intermediate Entorhinal Cortex

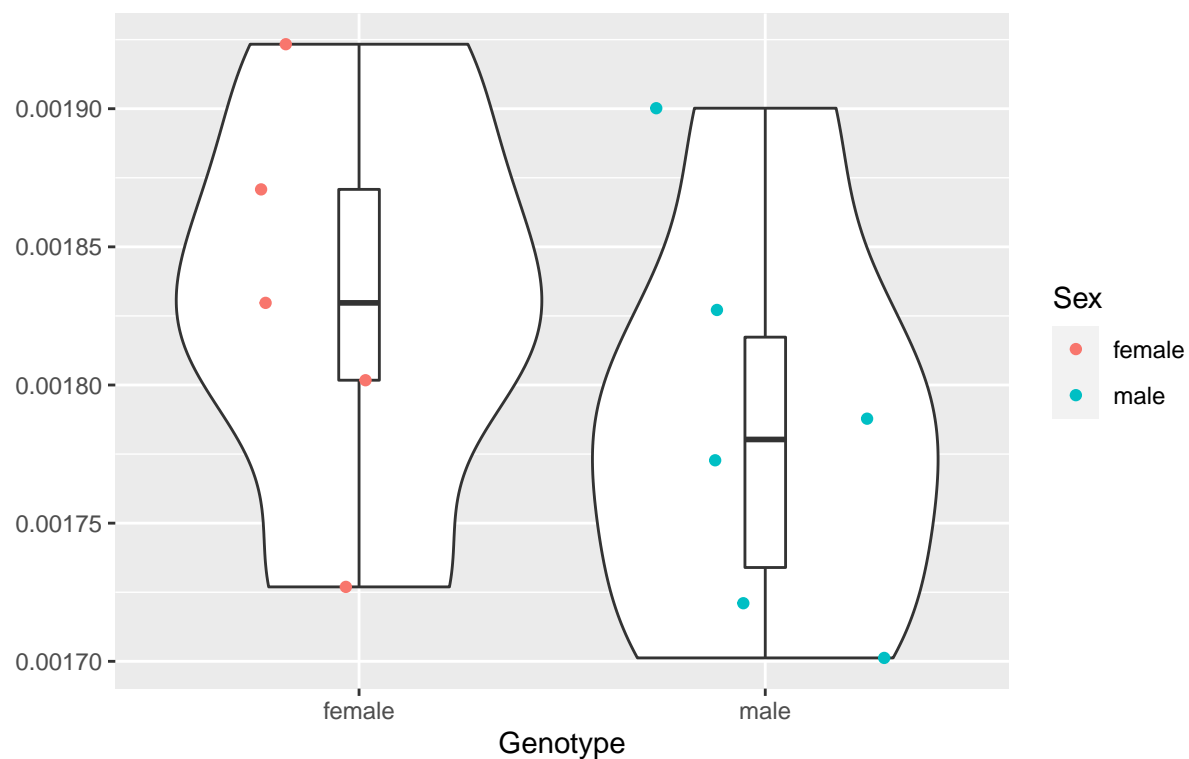
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	6.539e-08	6.539e-08	2.704	0.134
## Residuals	9	2.176e-07	2.418e-08		

Left Caudomedial Entorhinal Cortex

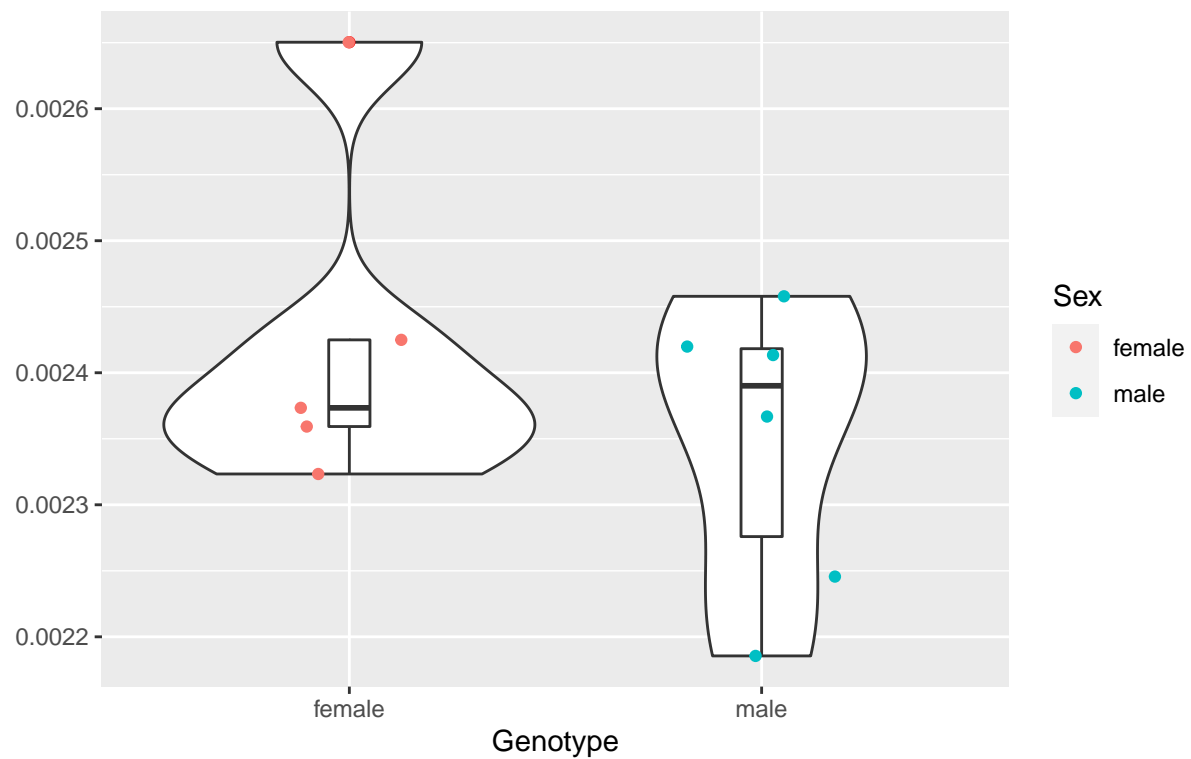
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	5.640e-09	5.641e-09	1.055	0.331
## Residuals	9	4.811e-08	5.345e-09		

Left Ventral Orbital Cortex

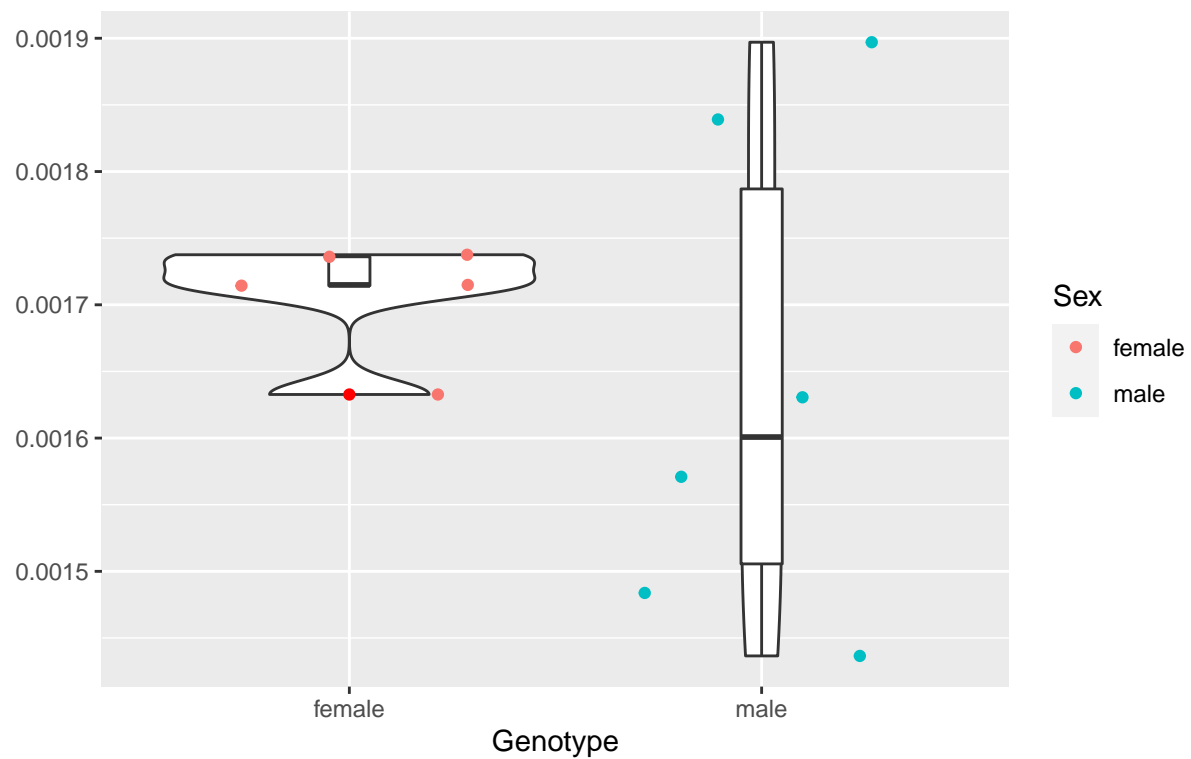
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.662e-08	1.662e-08	1.179	0.306
## Residuals	9	1.268e-07	1.409e-08		

Left Secondary Visual Cortex Mediomedial Area

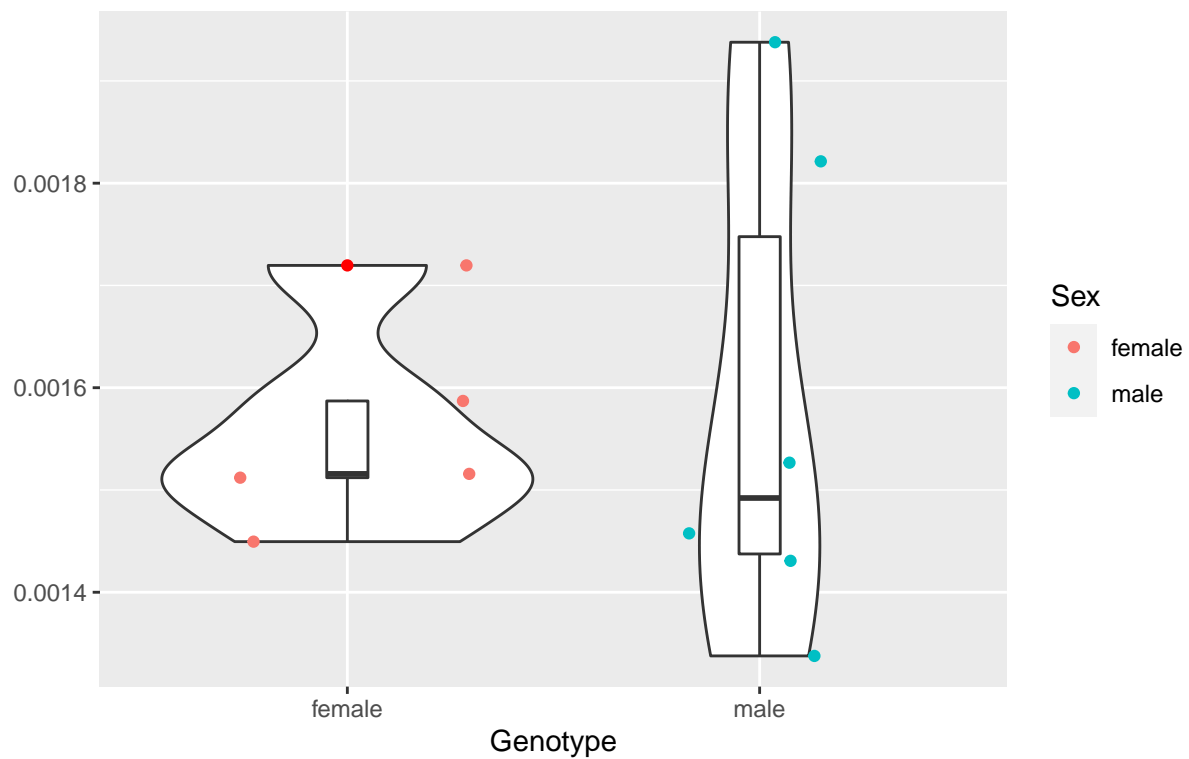
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.121e-08	1.121e-08	0.549	0.478
## Residuals	9	1.837e-07	2.041e-08		

Left Secondary Visual Cortex Mediolateral Area

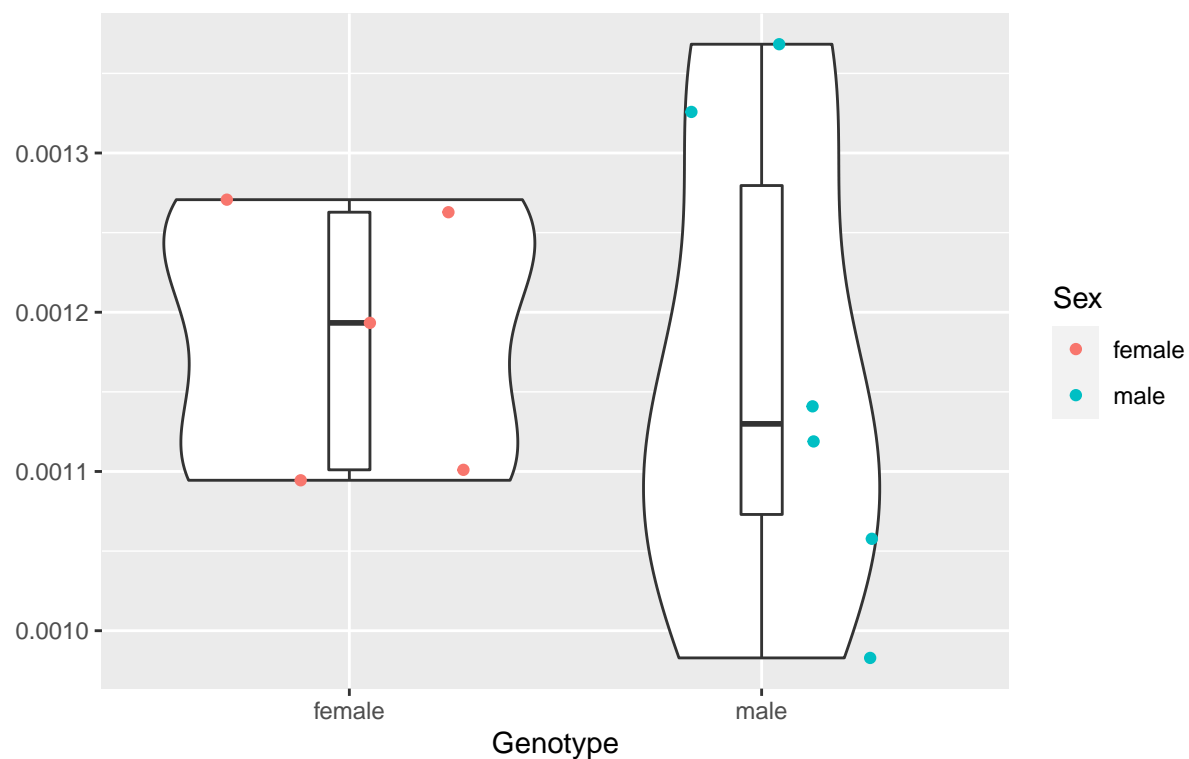
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.200e-09	2.22e-09	0.061	0.81
## Residuals	9	3.276e-07	3.64e-08		

Left Secondary Visual Cortex Lateral Area

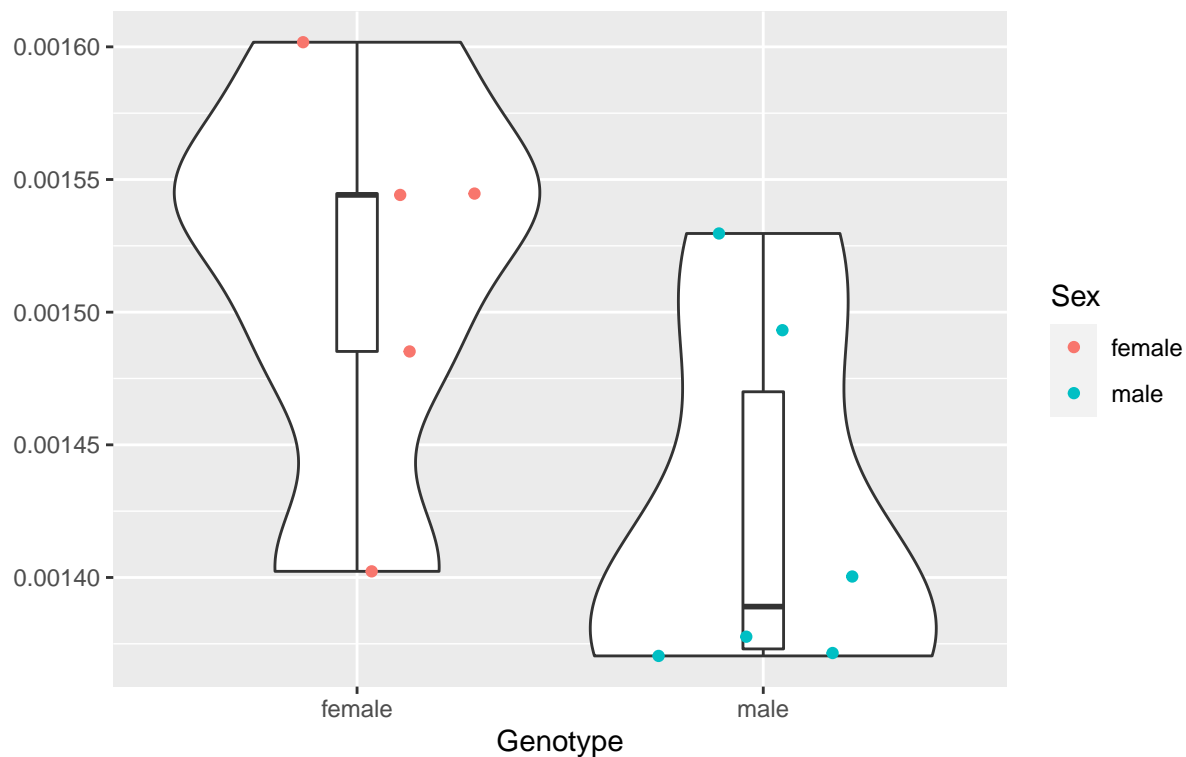
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	9.600e-10	9.550e-10	0.06	0.812
## Residuals	9	1.433e-07	1.592e-08		

Left Primary Visual Cortex Monocular Area

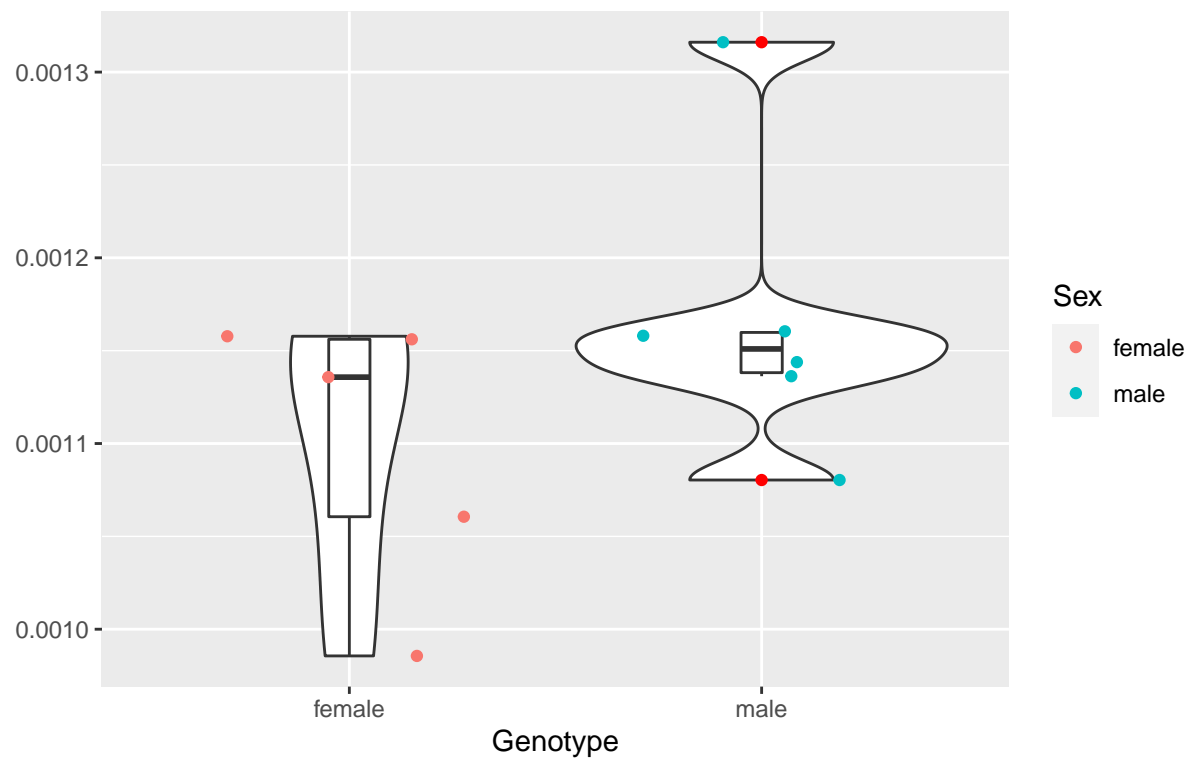
Red points denoting outliers



```
##          Df    Sum Sq  Mean Sq F value Pr(>F)
## Sex          1 2.299e-08 2.299e-08   4.391 0.0656 .
## Residuals    9 4.712e-08 5.236e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Left Primary Visual Cortex Binocular Area

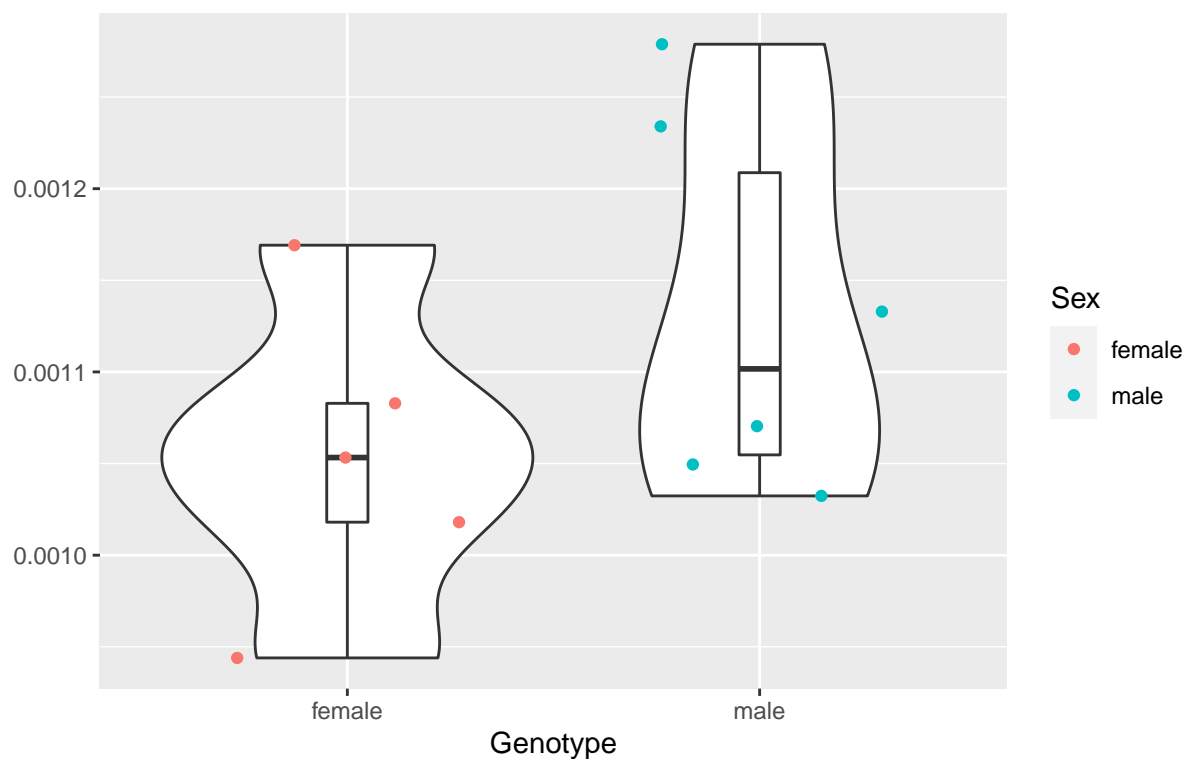
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.212e-08	1.212e-08	2.03	0.188
## Residuals	9	5.373e-08	5.970e-09		

Left Primary Visual Cortex

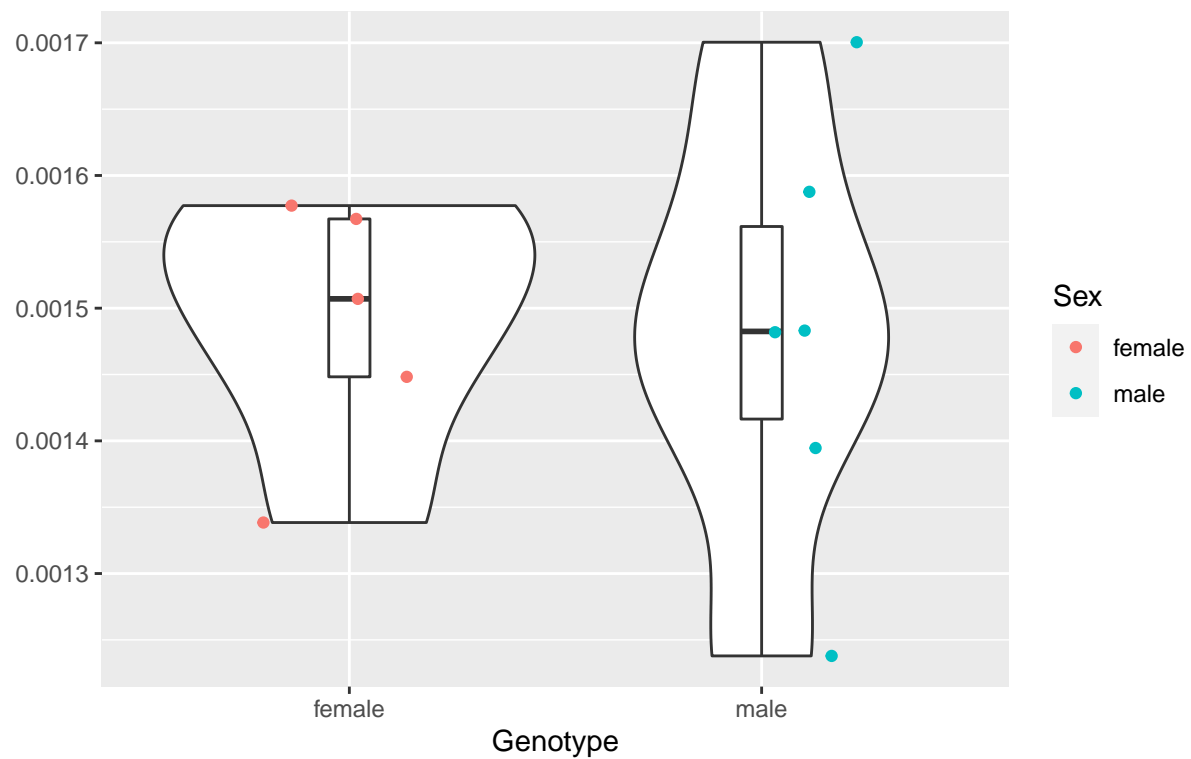
Red points denoting outliers



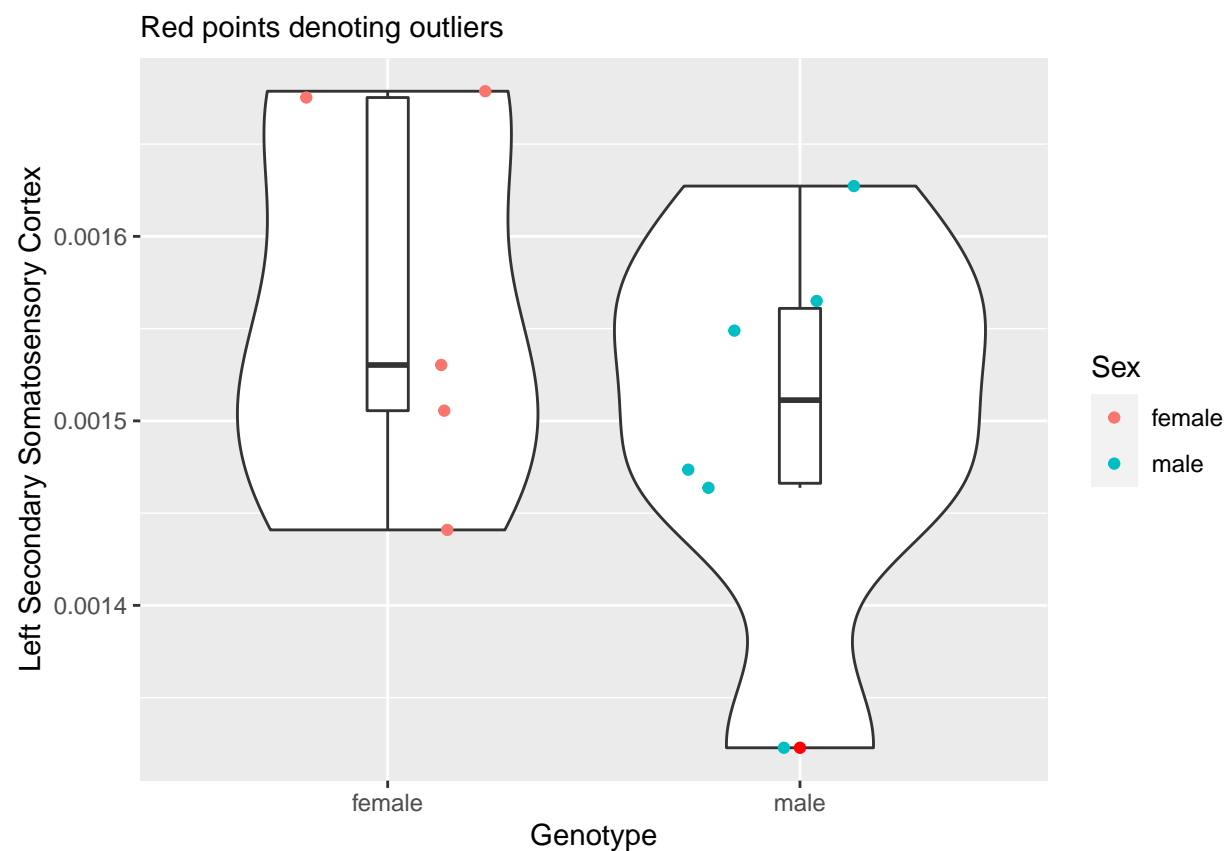
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.726e-08	1.726e-08	1.942	0.197
## Residuals	9	7.999e-08	8.888e-09		

Left Temporal Association Cortex

Red points denoting outliers



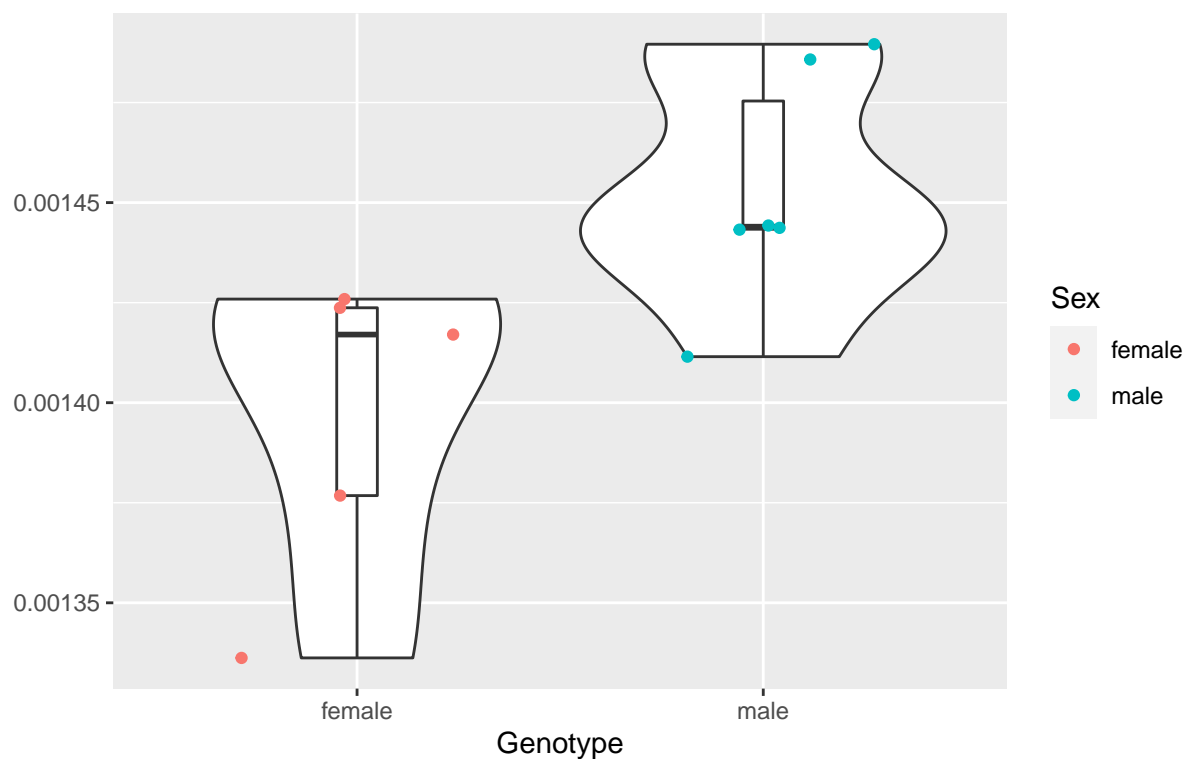
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.200e-10	1.23e-10	0.007	0.936
## Residuals	9	1.647e-07	1.83e-08		



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.185e-08	1.185e-08	1.051	0.332
## Residuals	9	1.014e-07	1.127e-08		

Left Primary Somatosensory Cortex Upper Lip Region

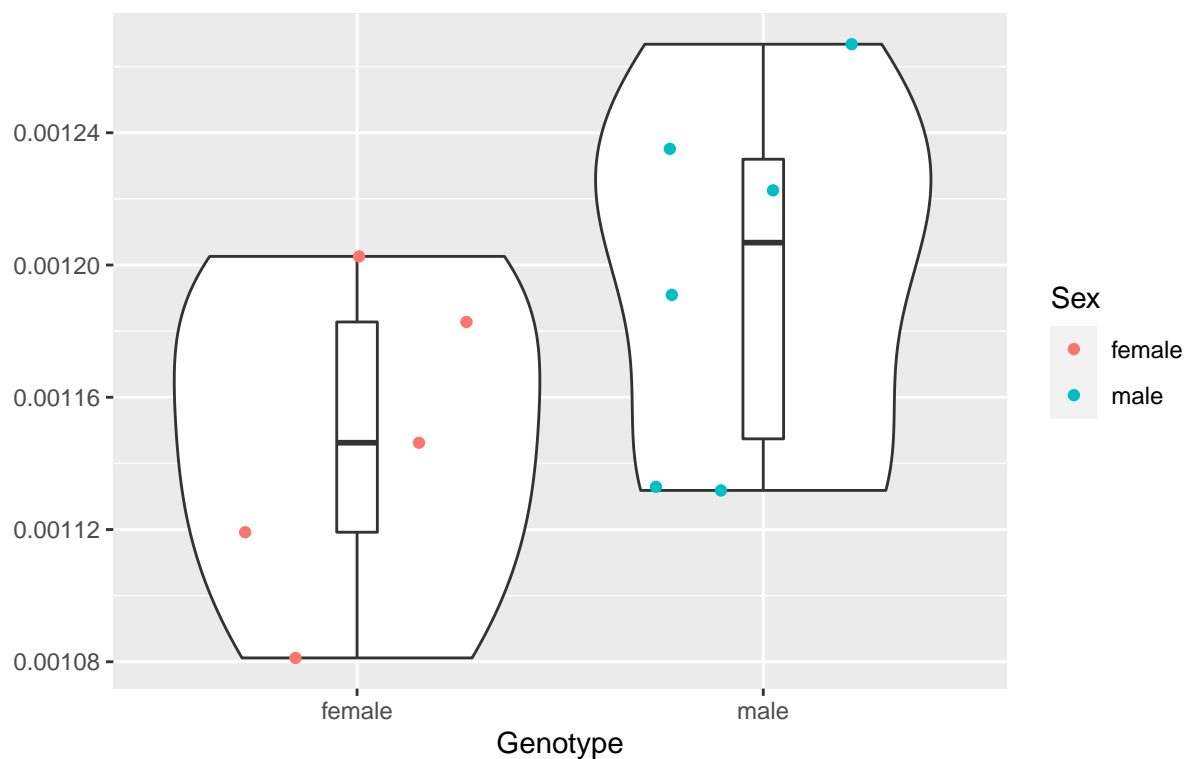
Red points denoting outliers



```
##          Df    Sum Sq  Mean Sq F value Pr(>F)
## Sex          1 8.884e-09 8.884e-09   7.662 0.0218 *
## Residuals    9 1.044e-08 1.160e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Left Primary Somatosensory Cortex Trunk Region

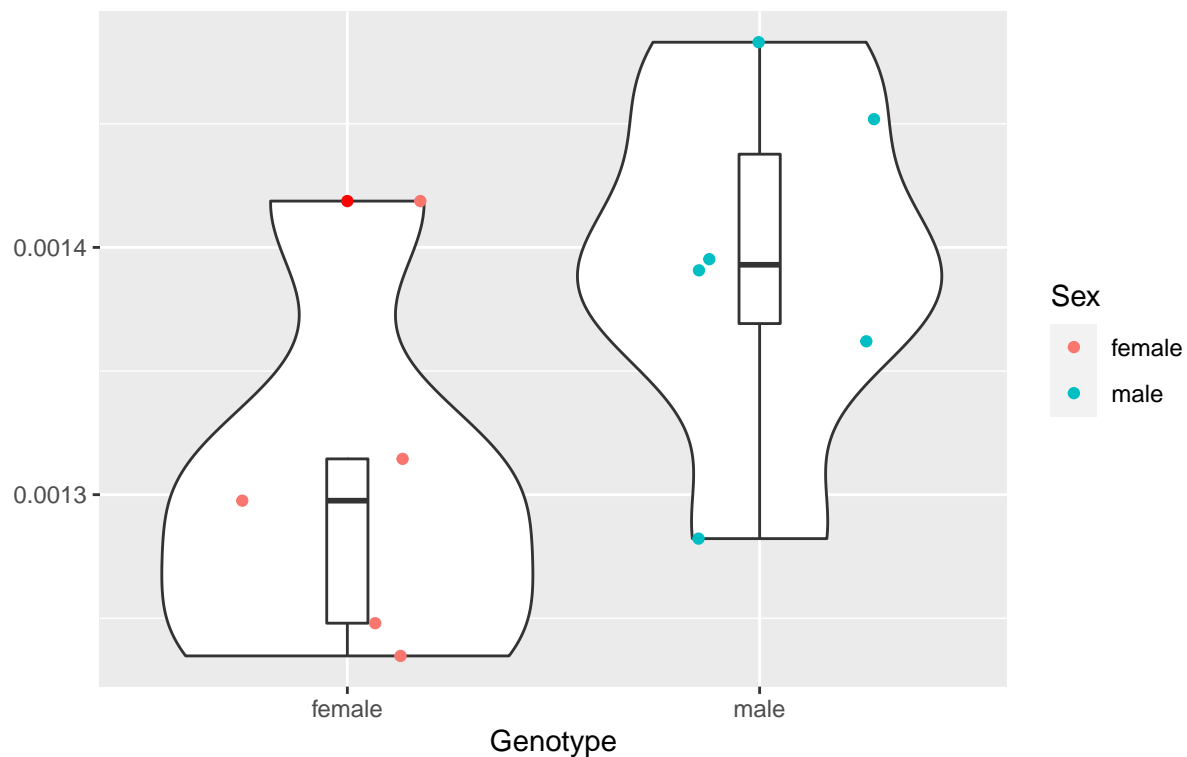
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	6.901e-09	6.901e-09	2.5	0.148
## Residuals	9	2.485e-08	2.761e-09		

Left Primary Somatosensory Cortex Shoulder Region

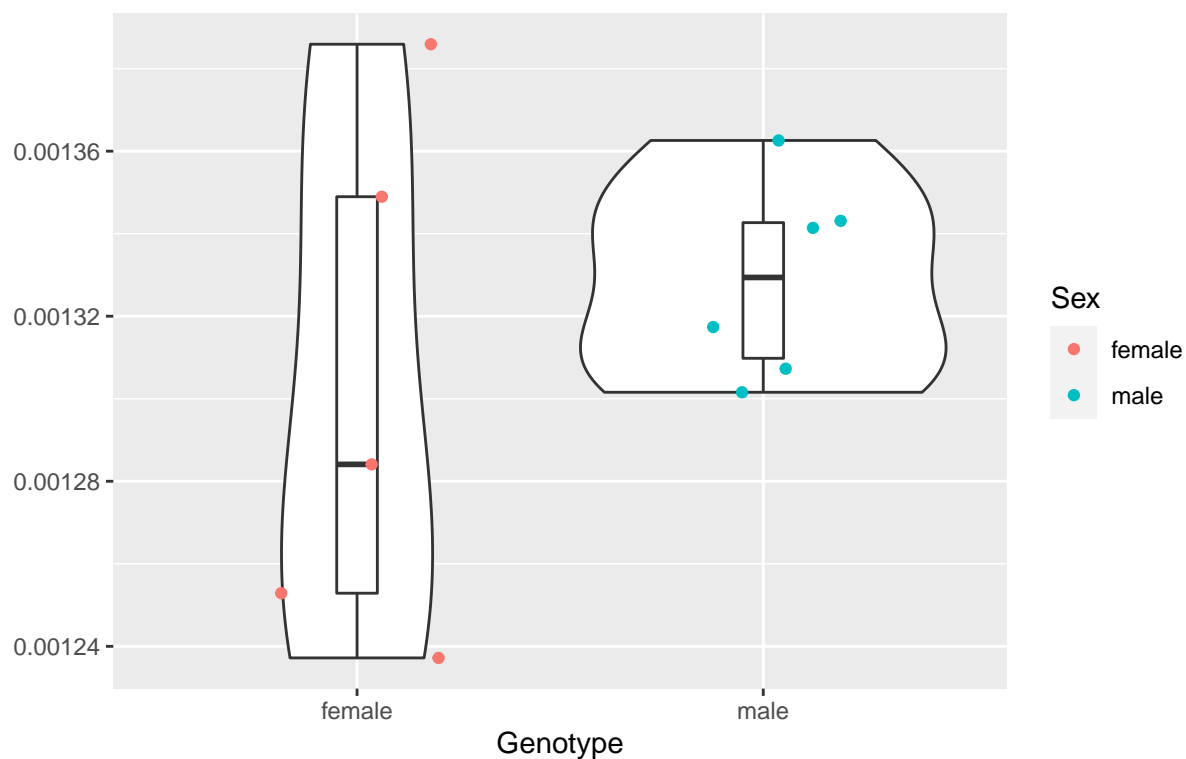
Red points denoting outliers



```
##           Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex           1 2.281e-08 2.281e-08   4.461 0.0639 .
## Residuals     9 4.603e-08 5.114e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Left Primary Somatosensory Cortex Jaw Region

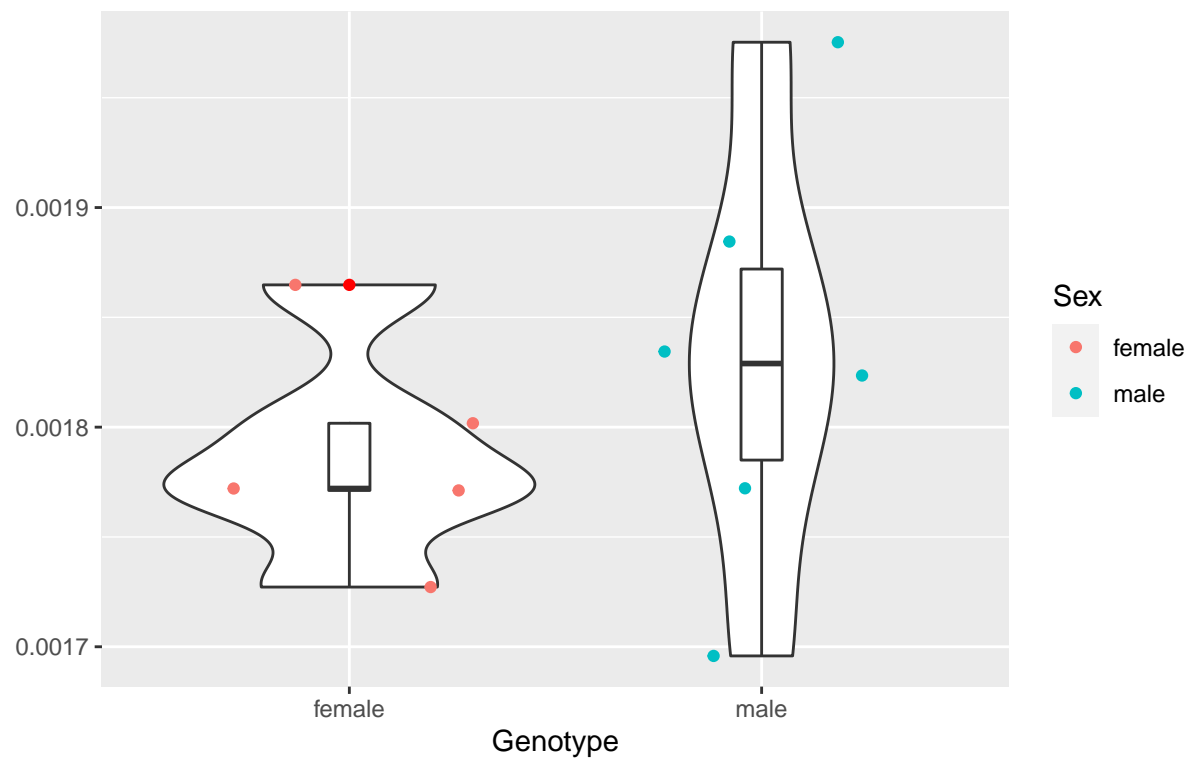
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.000e-09	2.000e-09	0.946	0.356
## Residuals	9	1.902e-08	2.114e-09		

Left Primary Somatosensory Cortex Hindlimb Region

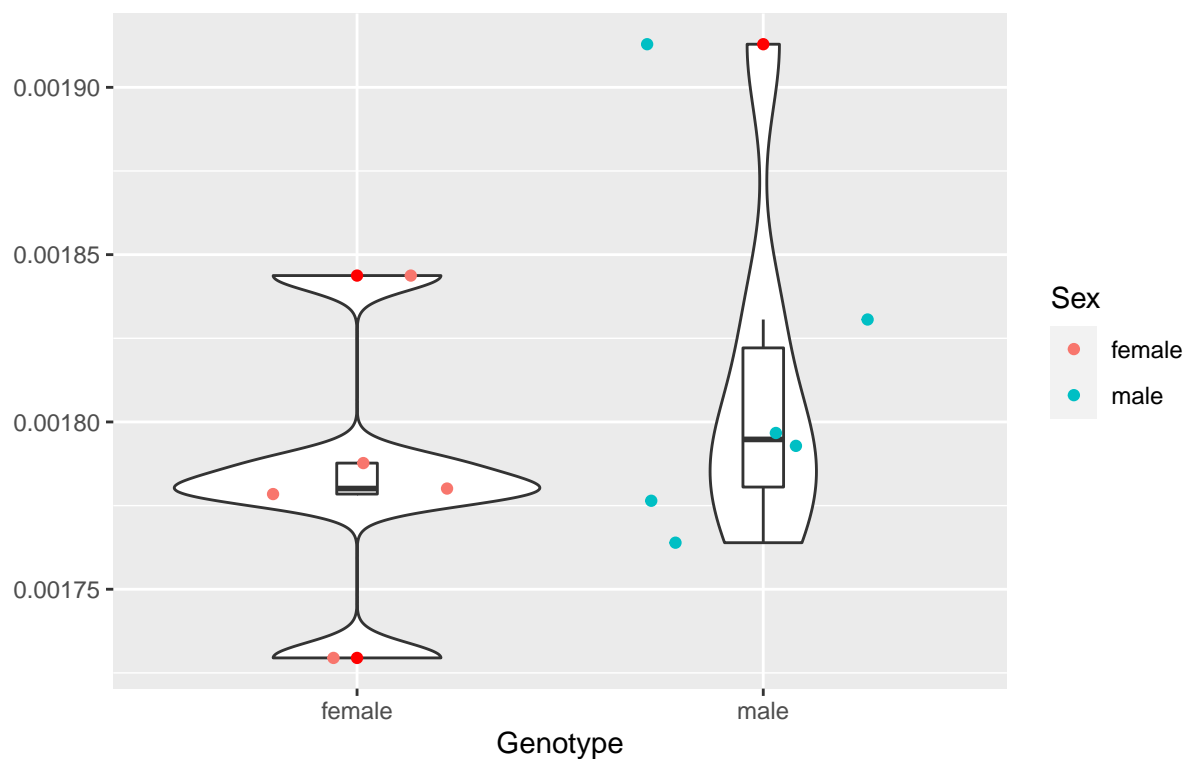
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	5.170e-09	5.173e-09	0.835	0.385
## Residuals	9	5.579e-08	6.199e-09		

Left Primary Somatosensory Cortex Forelimb Region

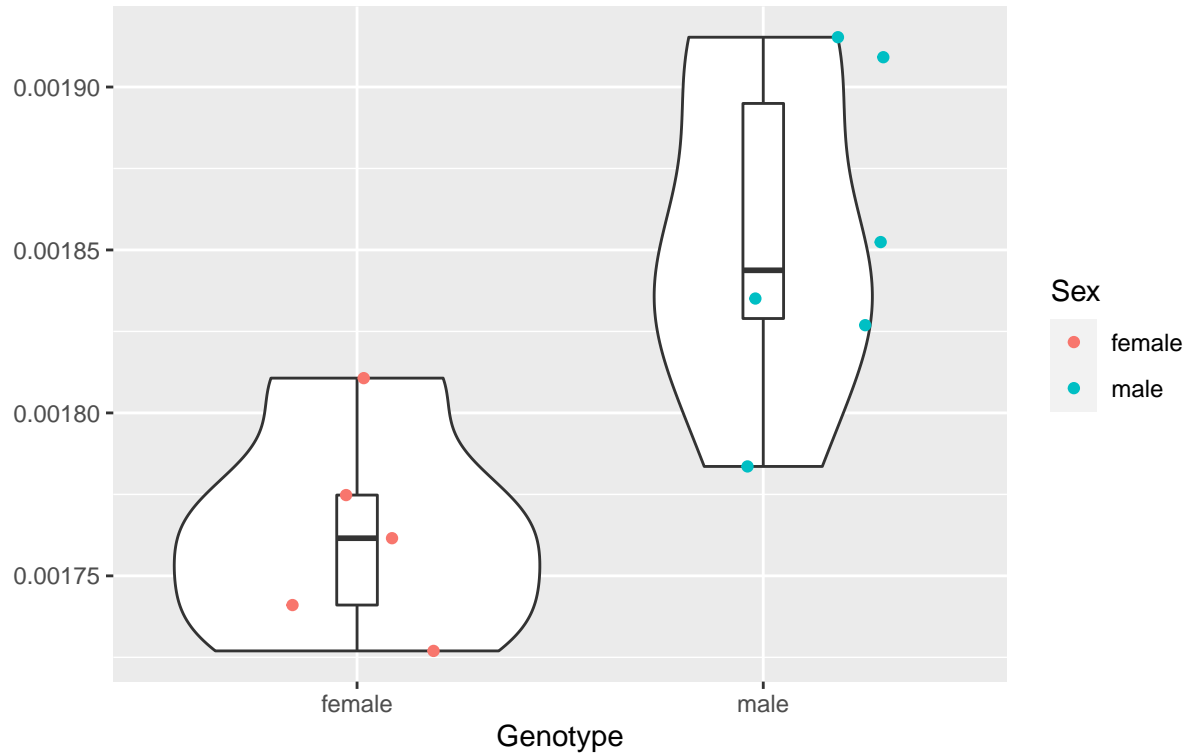
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.191e-09	2.191e-09	0.926	0.361
## Residuals	9	2.130e-08	2.367e-09		

Left Primary Somatosensory Cortex Dysgranular Zone

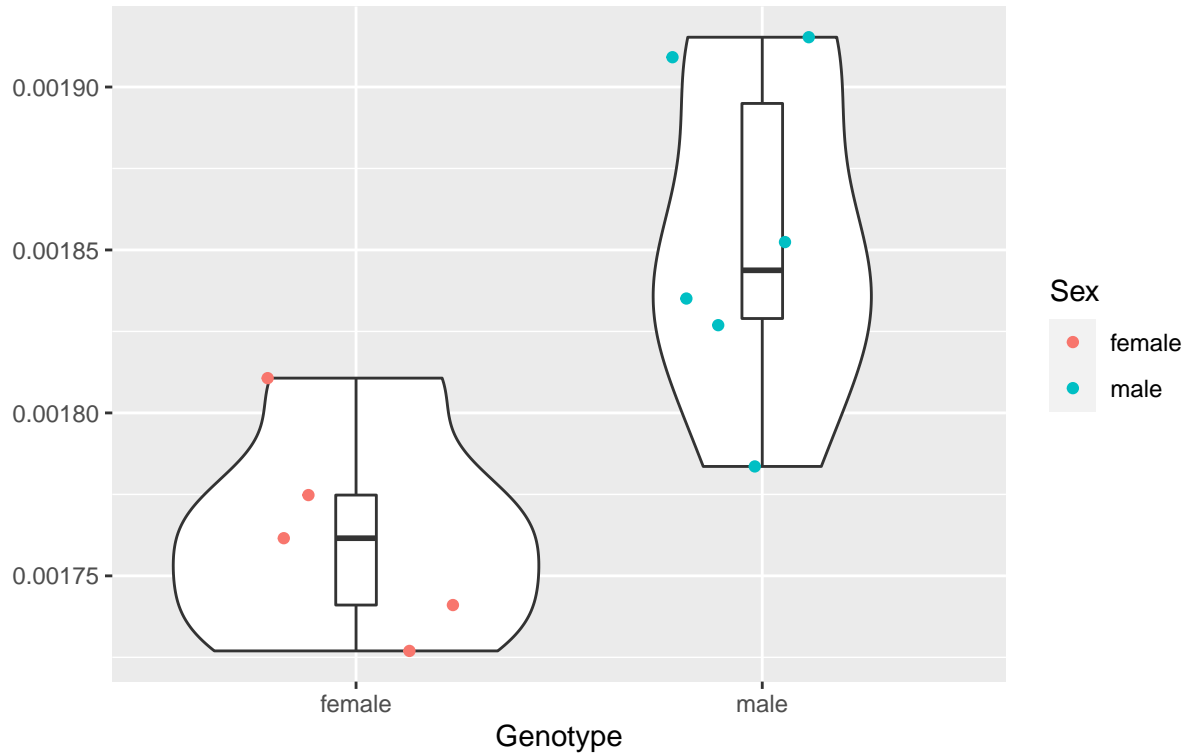
Red points denoting outliers



```
##           Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex           1 1.087e-08 1.087e-08   3.421 0.0974 .
## Residuals     9 2.860e-08 3.178e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Left Primary Somatosensory Cortex Barrel Field

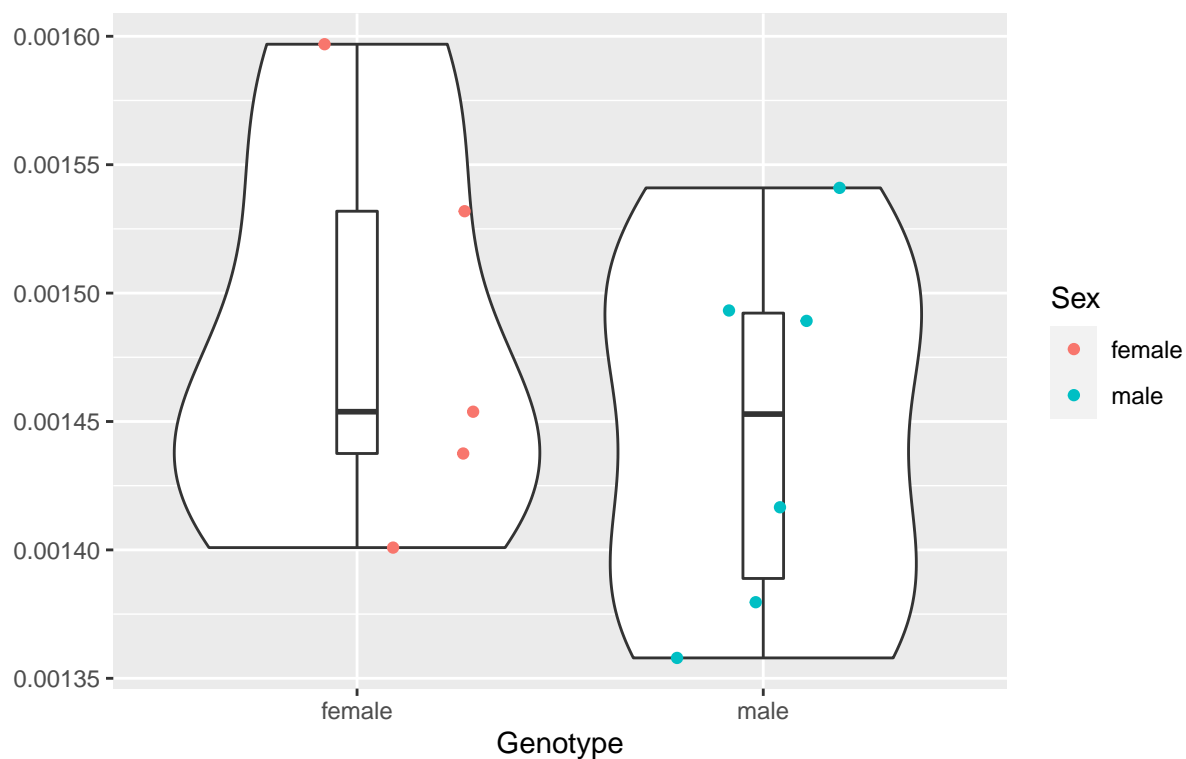
Red points denoting outliers



```
##           Df    Sum Sq  Mean Sq F value    Pr(>F)
## Sex           1 2.245e-08 2.245e-08   11.86 0.00735 **
## Residuals     9 1.704e-08 1.894e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Left Primary Somatosensory Cortex

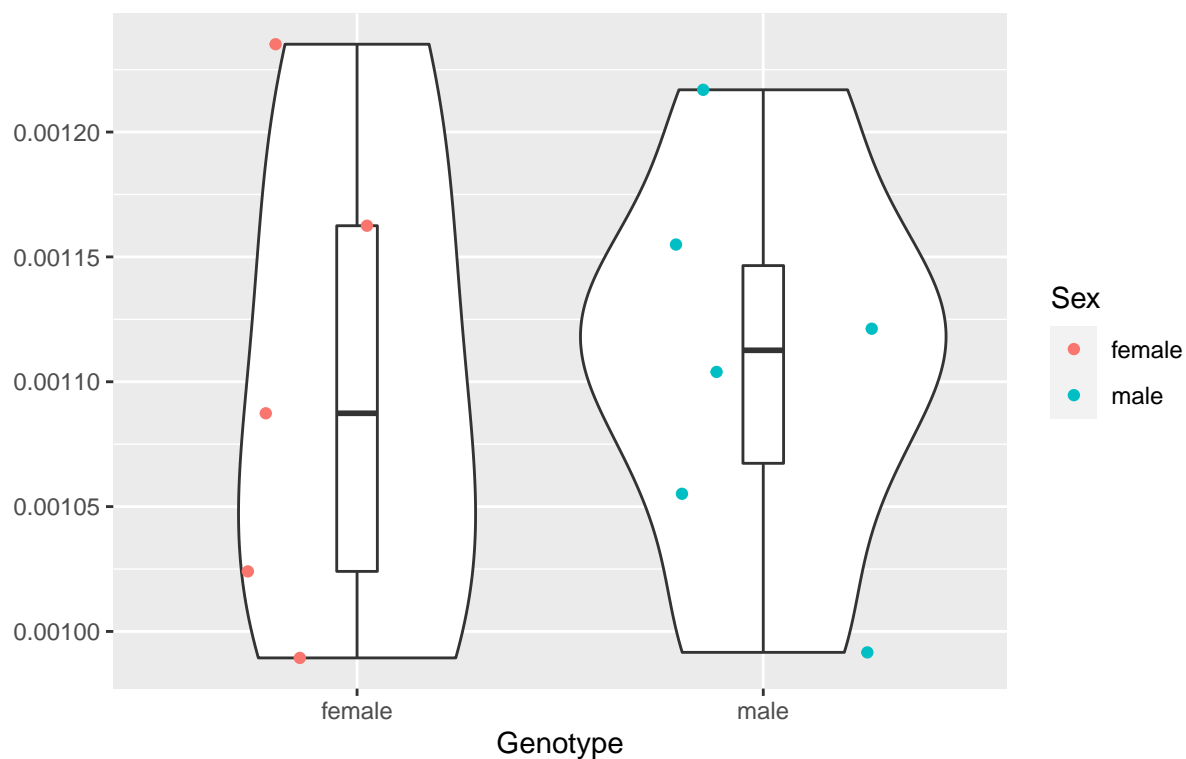
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	3.930e-09	3.927e-09	0.691	0.427
## Residuals	9	5.115e-08	5.684e-09		

Left Parietal Cortex Posterial Area Rostral Part

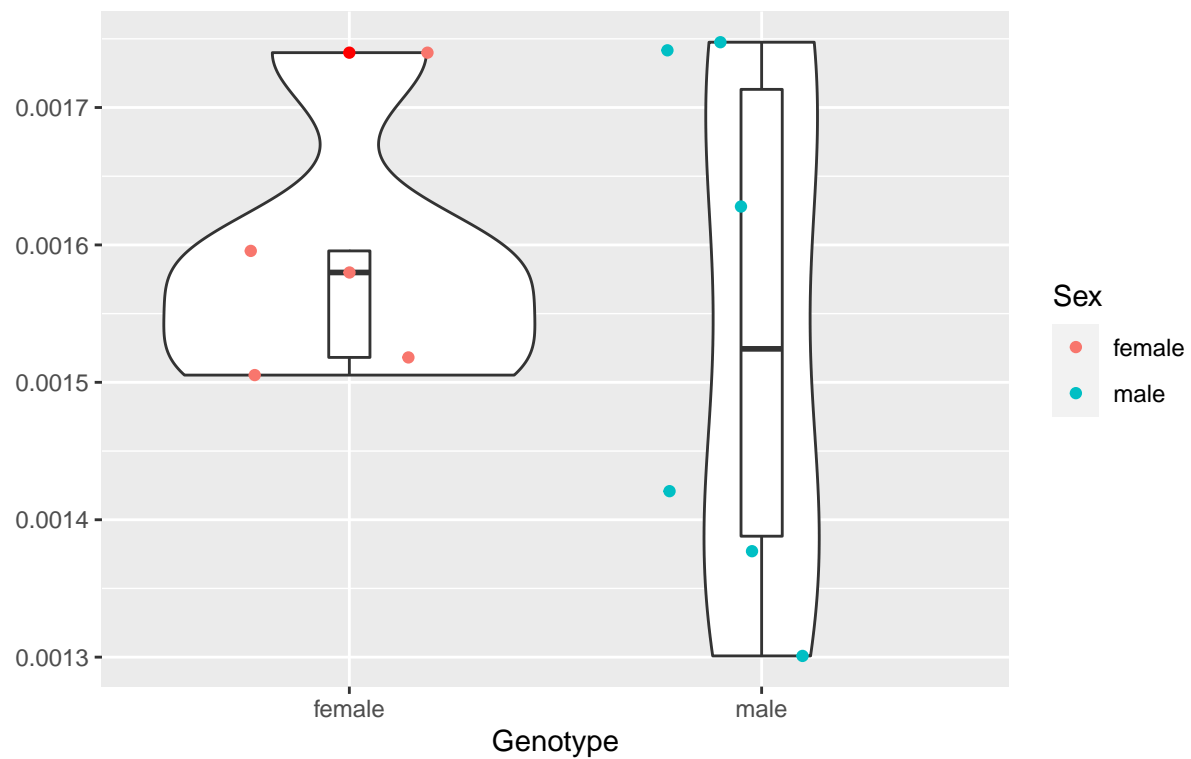
Red points denoting outliers



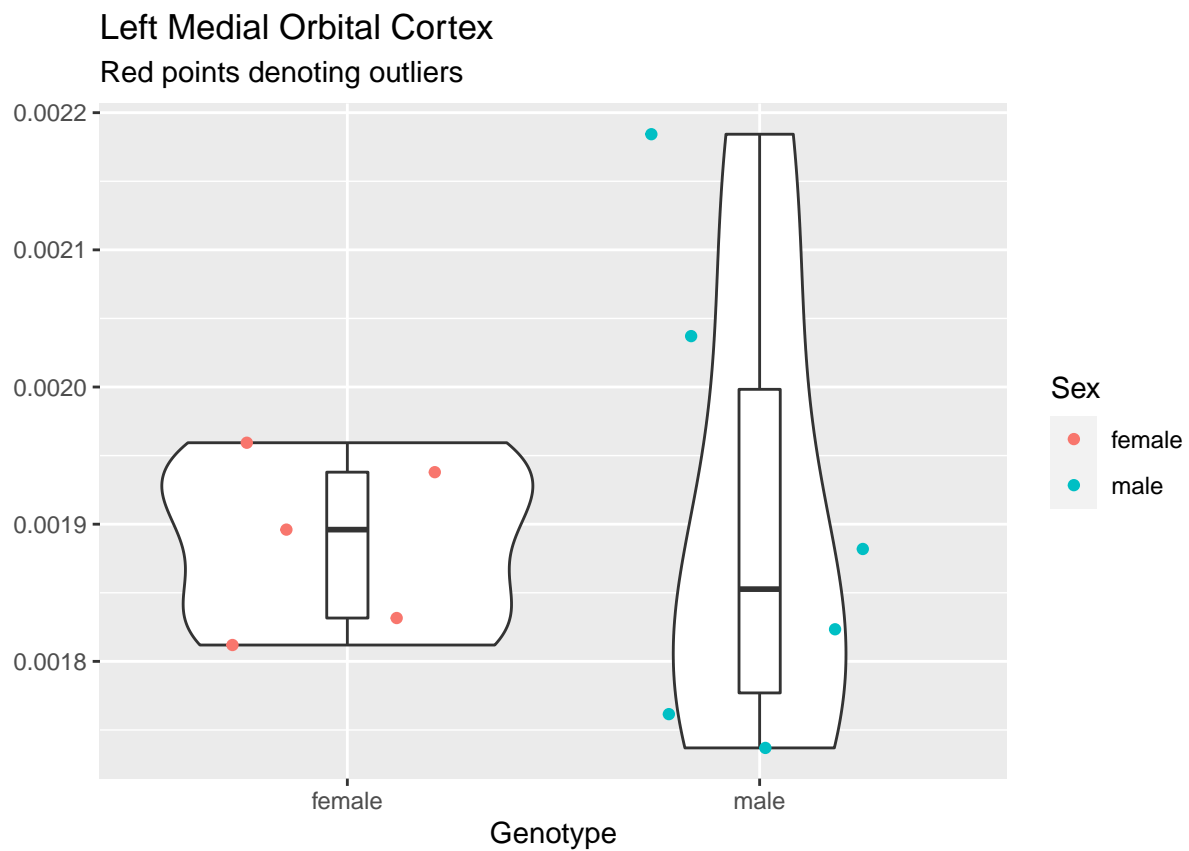
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.600e-10	1.580e-10	0.02	0.891
## Residuals	9	7.093e-08	7.881e-09		

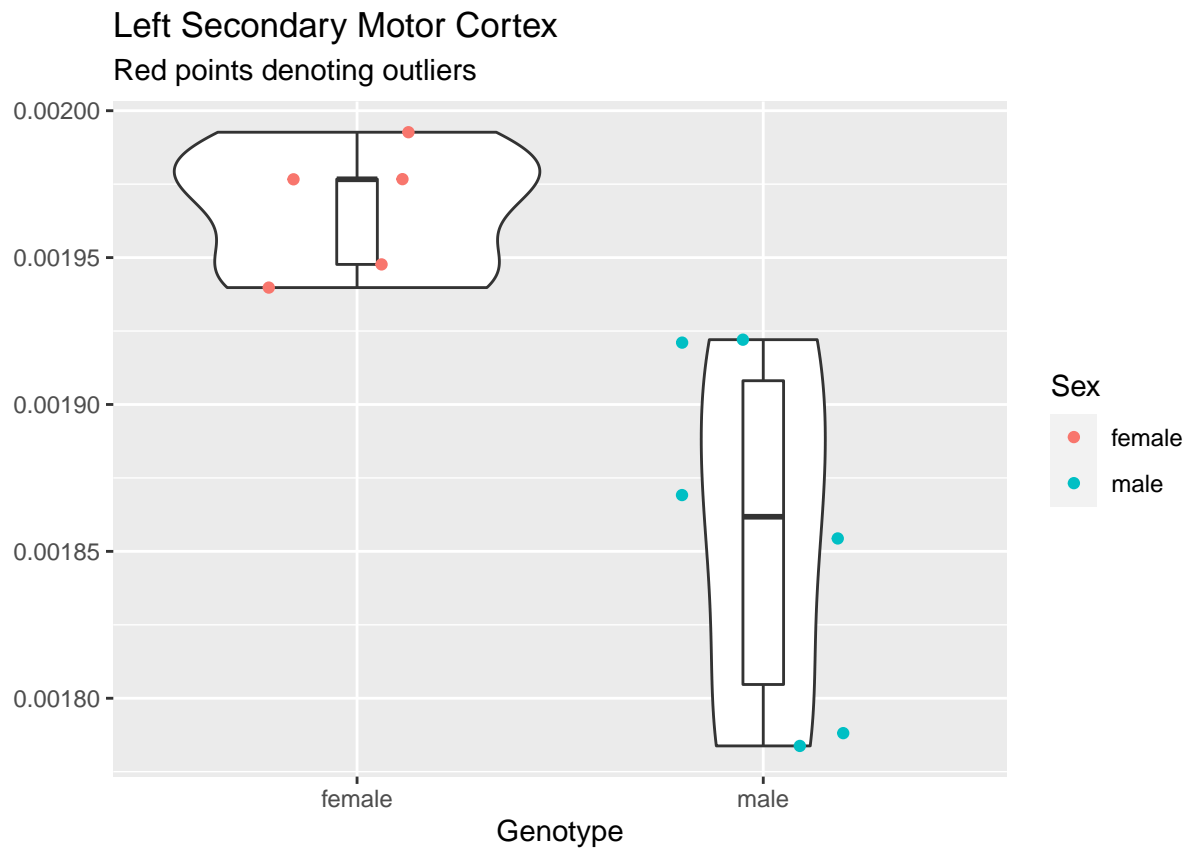
Left Medial Parietal Association Cortex

Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	7.320e-09	7.319e-09	0.294	0.601
## Residuals	9	2.242e-07	2.491e-08		

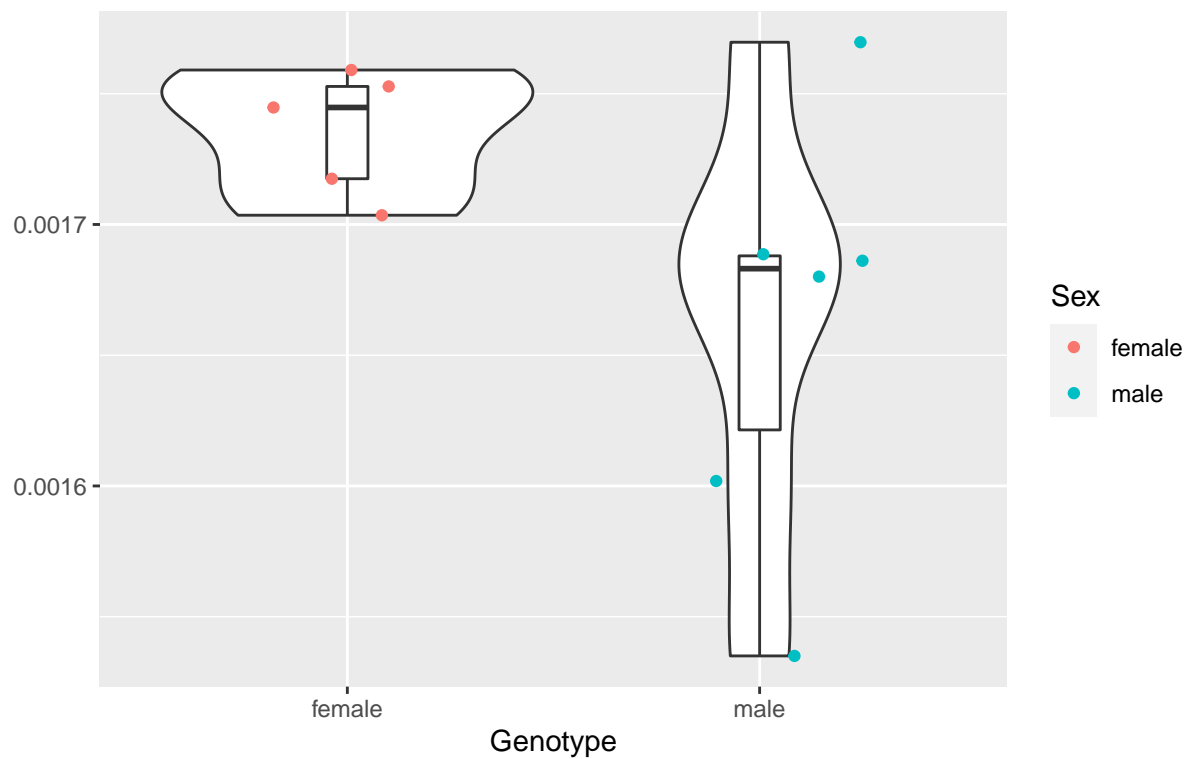




```
##          Df    Sum Sq  Mean Sq F value    Pr(>F)
## Sex           1 3.317e-08 3.317e-08   14.52 0.00415 **
## Residuals     9 2.056e-08 2.280e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```


Left Primary Motor Cortex

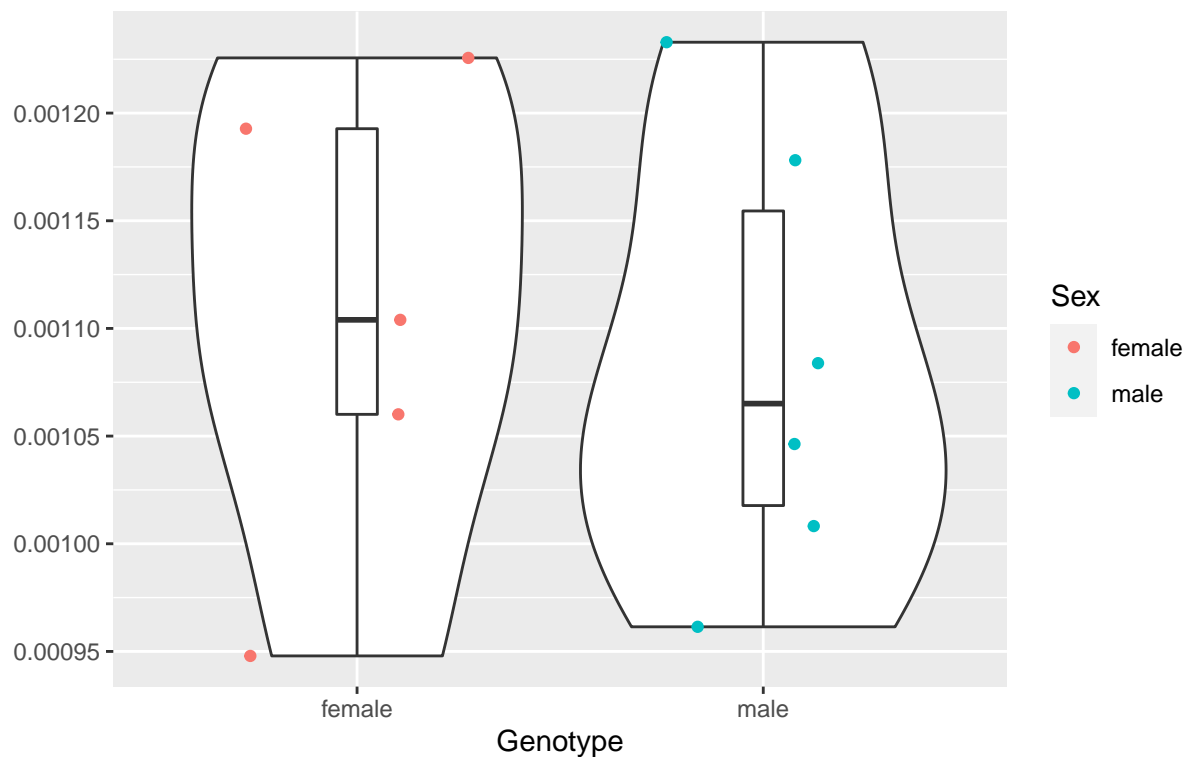
Red points denoting outliers



```
##          Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex          1 1.546e-08 1.546e-08    3.95 0.0781 .
## Residuals    9 3.523e-08 3.914e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Left Lateral Parietal Association Cortex

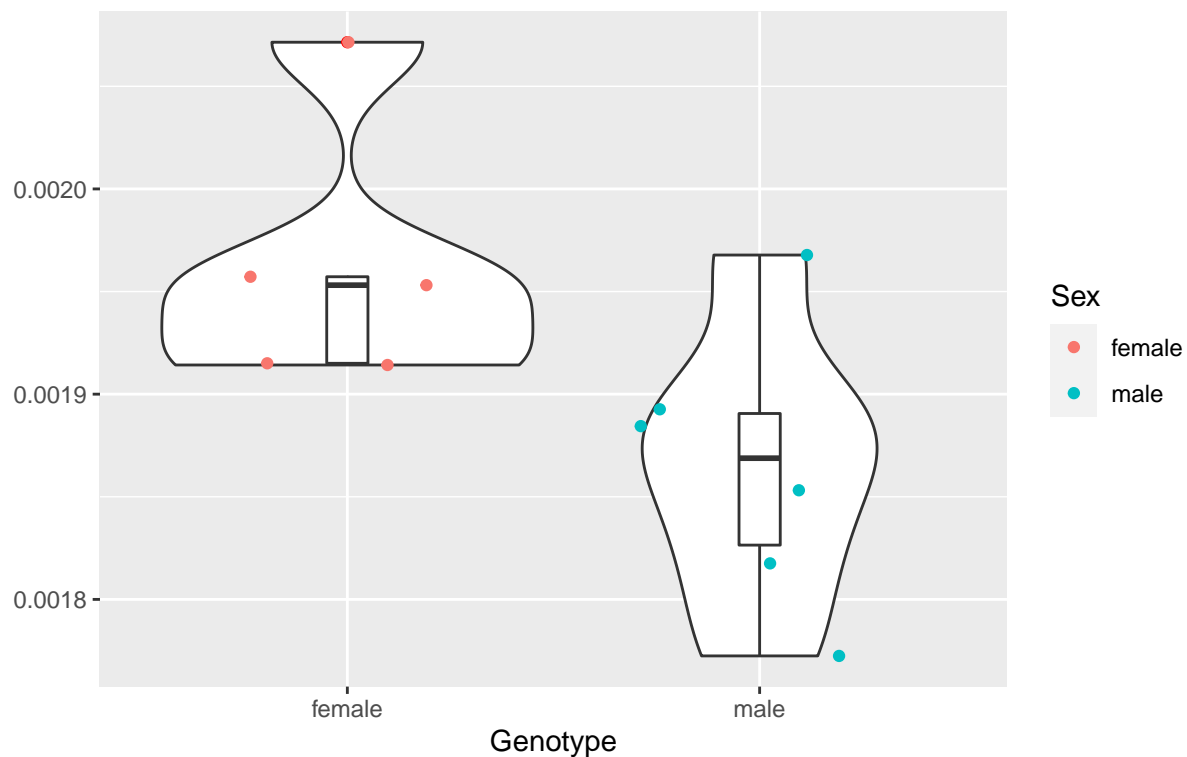
Red points denoting outliers



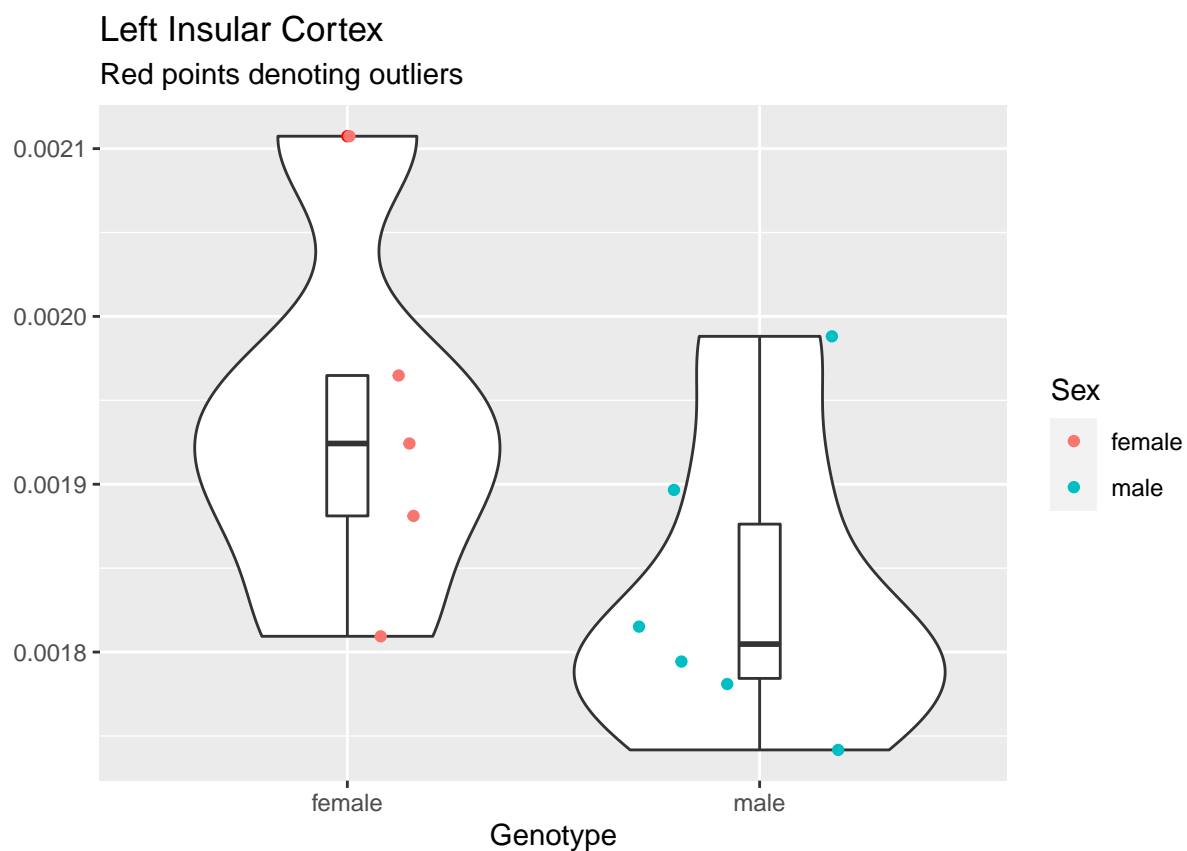
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.190e-09	1.195e-09	0.105	0.753
## Residuals	9	1.021e-07	1.135e-08		

Left Lateral Orbital Cortex

Red points denoting outliers



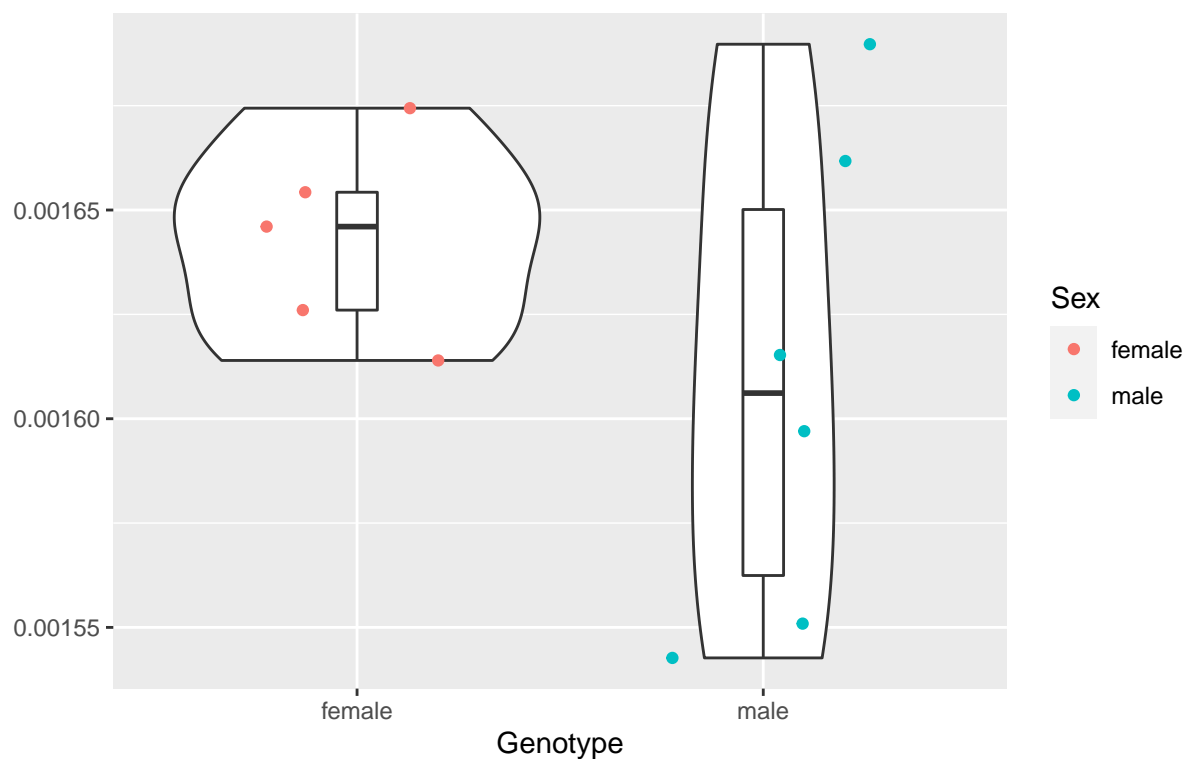
```
##           Df      Sum Sq   Mean Sq F value Pr(>F)
## Sex          1 2.594e-08 2.594e-08   5.951 0.0374 *
## Residuals    9 3.923e-08 4.359e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	2.796e-08	2.796e-08	2.787	0.129
## Residuals	9	9.029e-08	1.003e-08		

Left Frontal Association Cortex

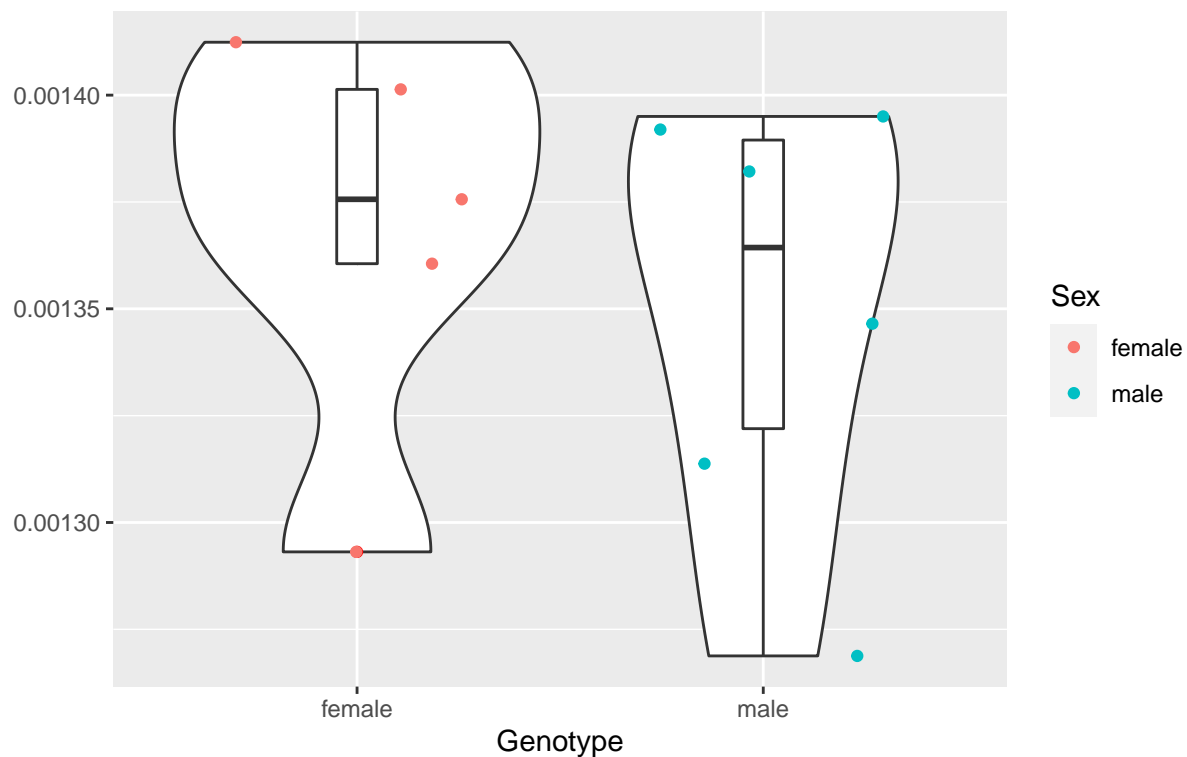
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	3.038e-09	3.038e-09	1.402	0.267
## Residuals	9	1.950e-08	2.167e-09		

Left Frontal Cortex Area 3

Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	9.760e-10	9.755e-10	0.408	0.539
## Residuals	9	2.154e-08	2.394e-09		

Left Dorsolateral Orbital Cortex

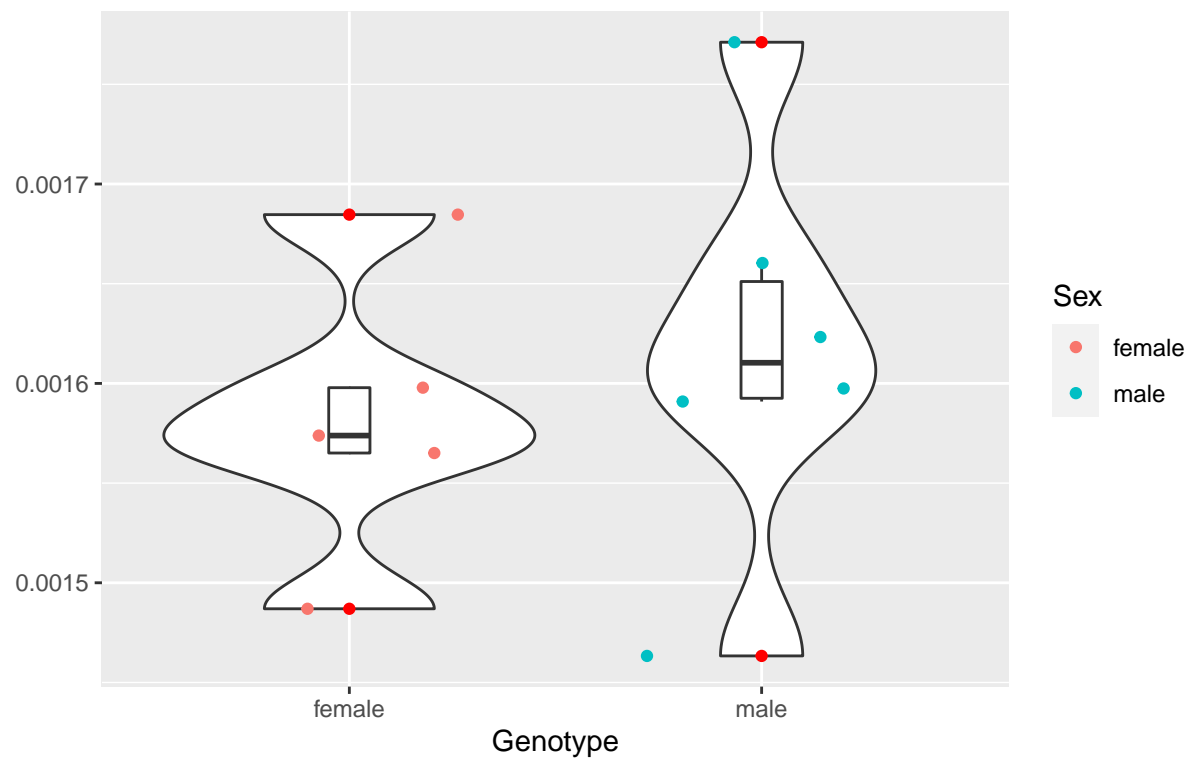
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	5.000e-11	4.900e-11	0.007	0.936
## Residuals	9	6.448e-08	7.164e-09		

Left Secondary Auditory Cortex Ventral Part

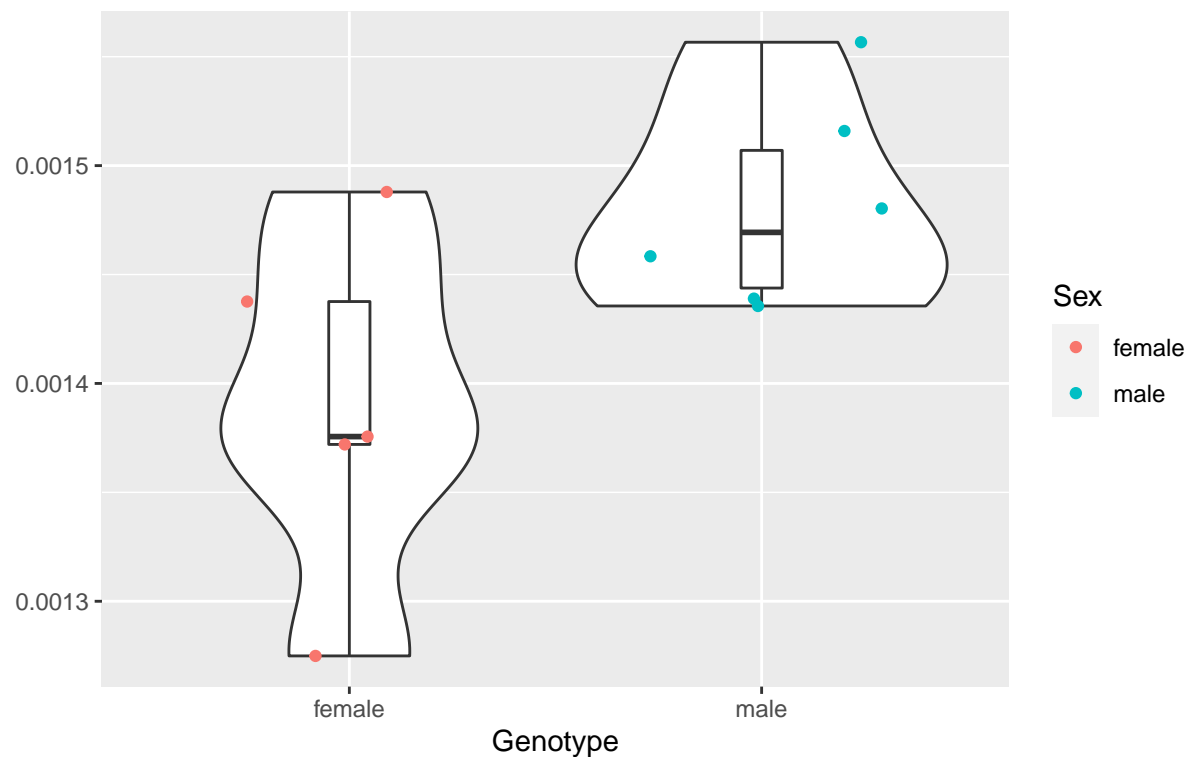
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	3.550e-09	3.548e-09	0.453	0.518
## Residuals	9	7.053e-08	7.836e-09		

Left Secondary Auditory Cortex Dorsal Part

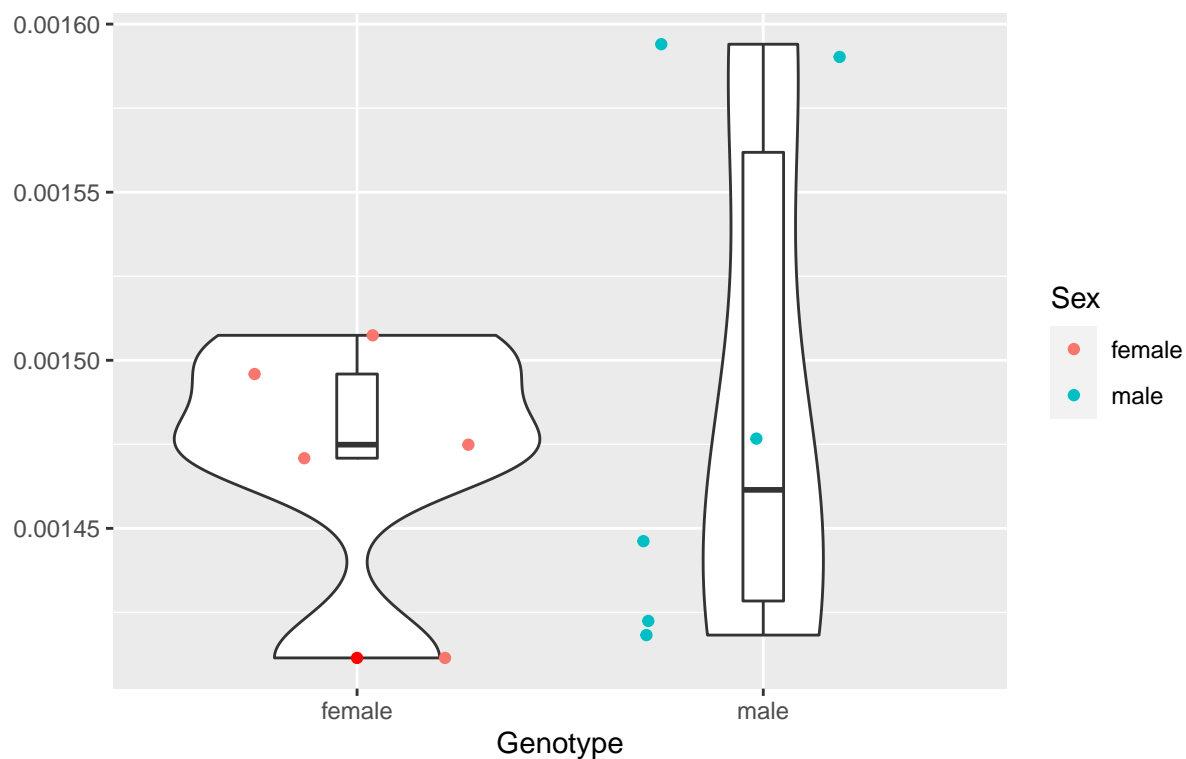
Red points denoting outliers



```
##          Df    Sum Sq  Mean Sq F value Pr(>F)
## Sex          1 2.276e-08 2.276e-08   5.55 0.0429 *
## Residuals    9 3.691e-08 4.101e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

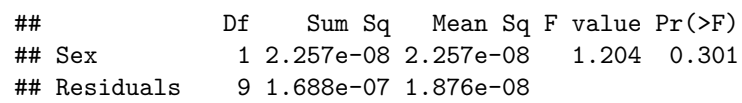
Left Primary Auditory Cortex

Red points denoting outliers

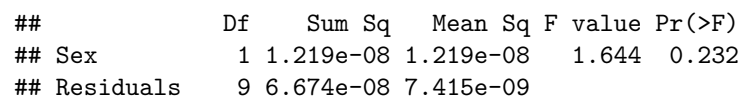


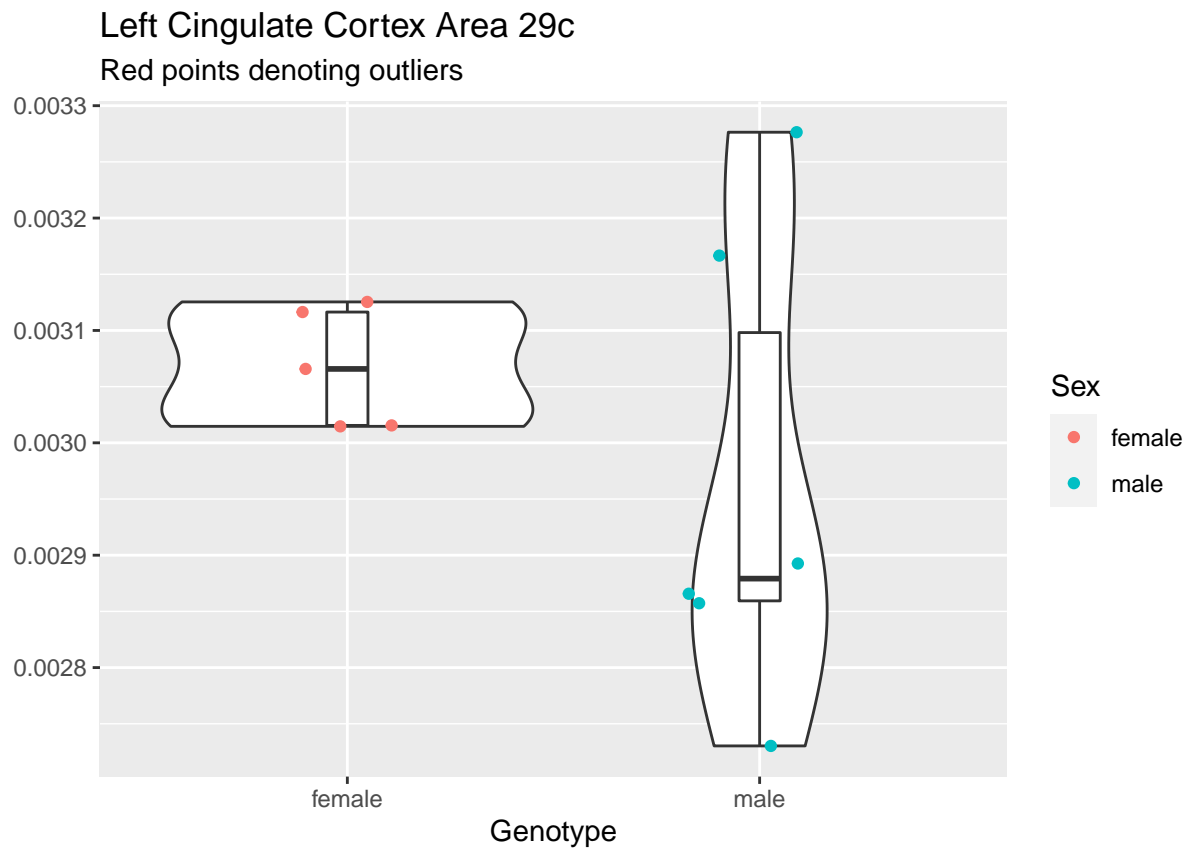
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.010e-09	1.005e-09	0.237	0.638
## Residuals	9	3.816e-08	4.241e-09		

Red points denoting outliers



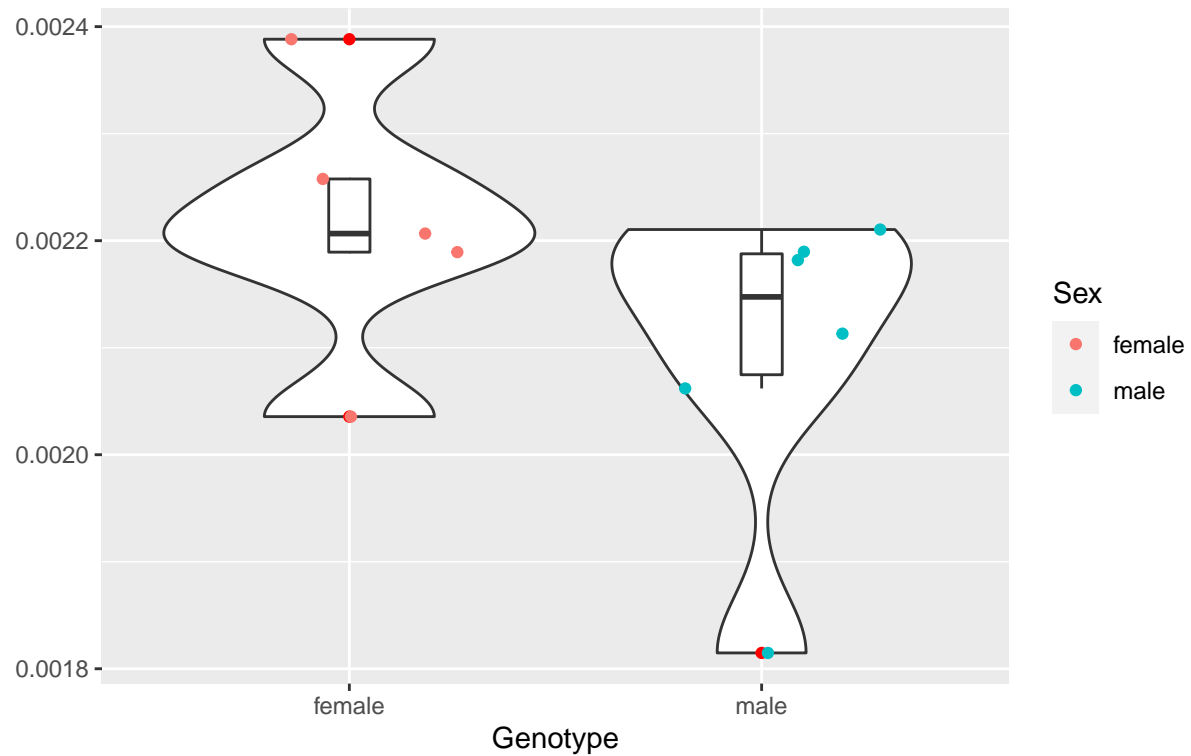
Red points denoting outliers





Left Cingulate Cortex Area 29b

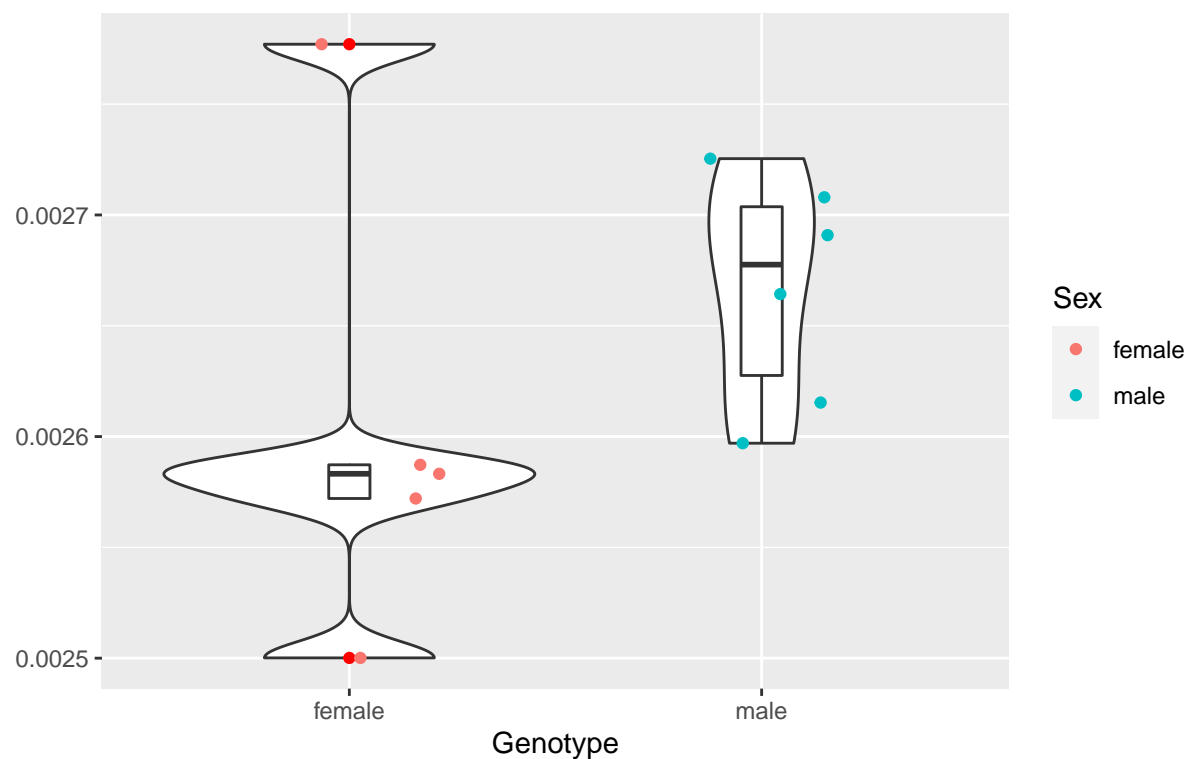
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	3.940e-08	3.940e-08	2.032	0.188
## Residuals	9	1.745e-07	1.939e-08		

Left Cingulate Cortex Area 29a

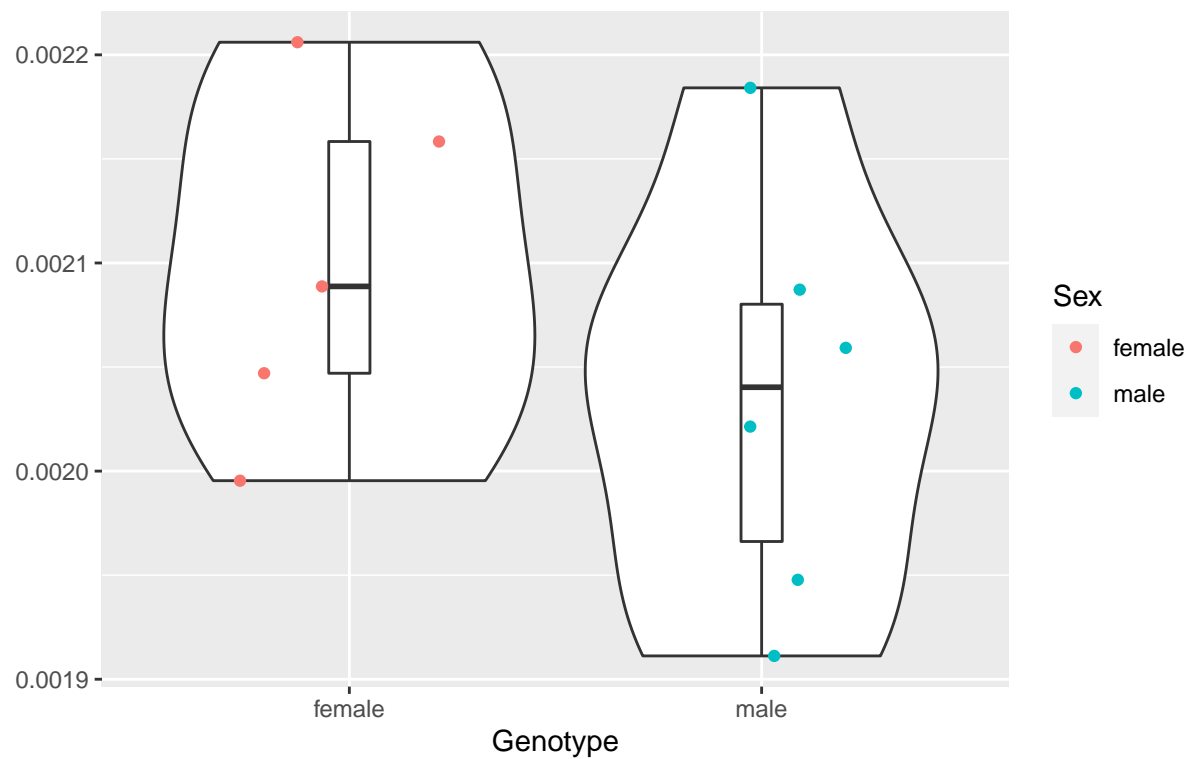
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.078e-08	1.078e-08	1.742	0.219
## Residuals	9	5.571e-08	6.190e-09		

Left Cingulate Cortex Area 24b Prime

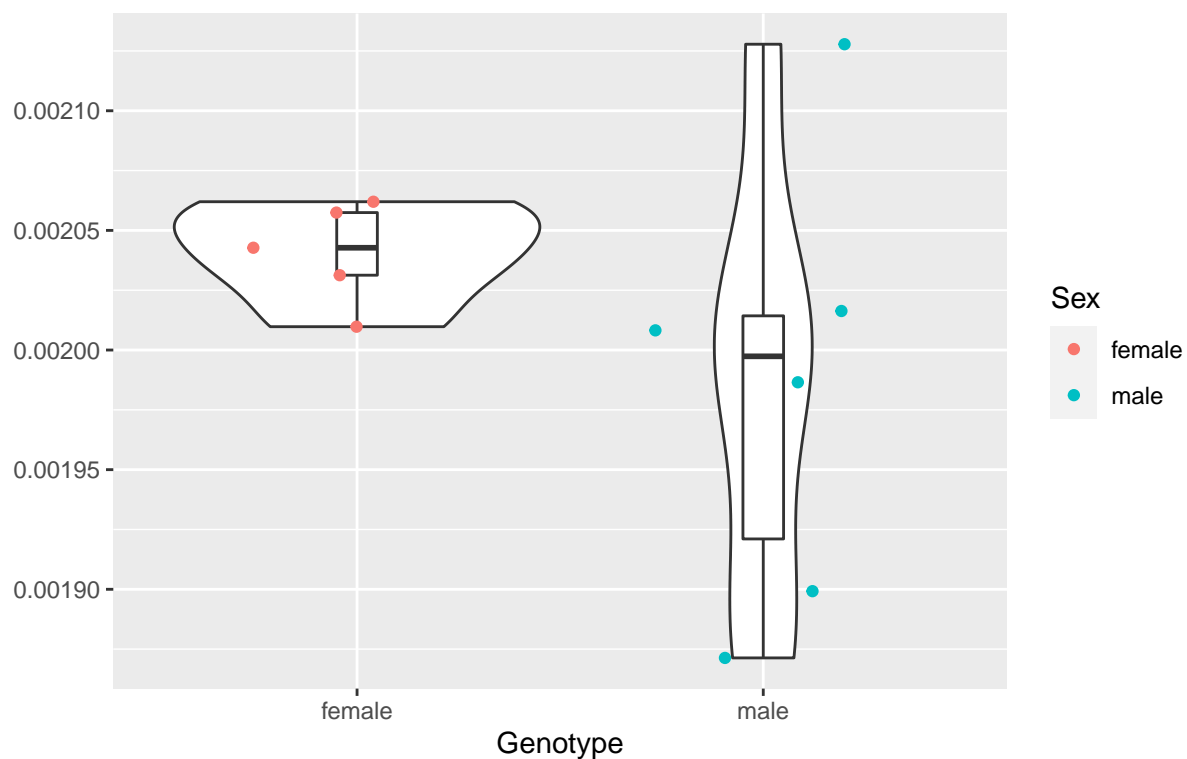
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	1.116e-08	1.116e-08	1.301	0.283
## Residuals	9	7.720e-08	8.578e-09		

Left Cingulate Cortex Area 24b

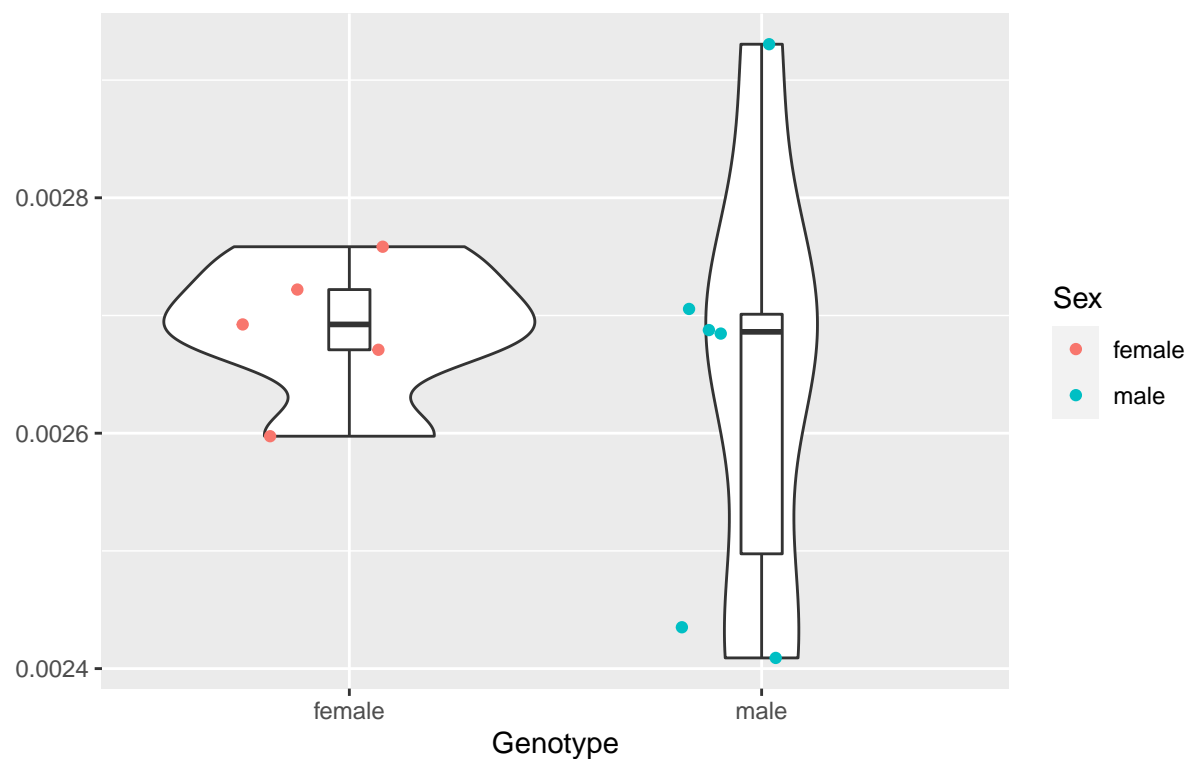
Red points denoting outliers



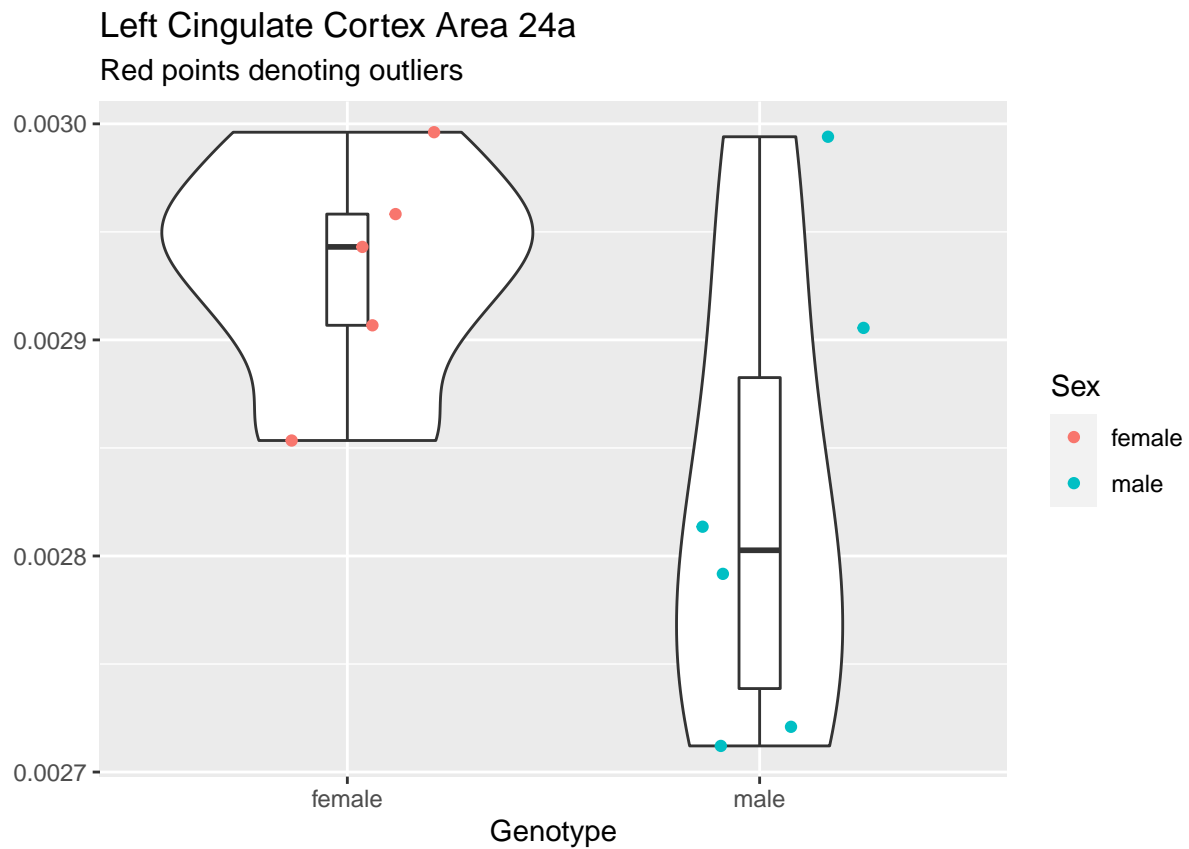
##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	8.470e-09	8.473e-09	1.734	0.22
## Residuals	9	4.399e-08	4.887e-09		

Left Cingulate Cortex Area 24a Prime

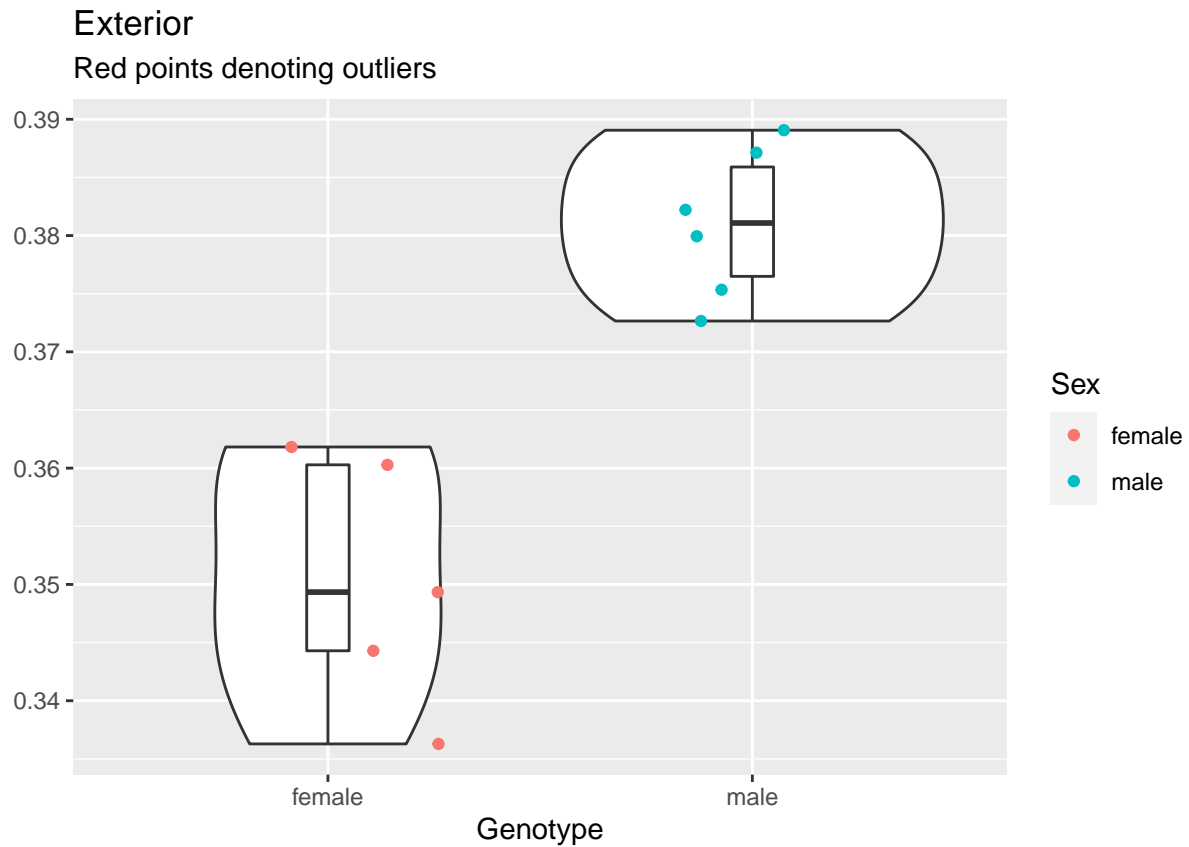
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## Sex	1	5.810e-09	5.809e-09	0.258	0.624
## Residuals	9	2.028e-07	2.254e-08		



```
##           Df    Sum Sq   Mean Sq F value Pr(>F)
## Sex         1 3.213e-08 3.213e-08    4.04 0.0753 .
## Residuals   9 7.158e-08 7.950e-09
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```



```
##          Df    Sum Sq  Mean Sq F value    Pr(>F)
## Sex          1 0.0025623 0.0025623    34.29 0.000242 ***
## Residuals    9 0.0006724 0.0000747
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```