

# Proportional Volume Distributions by Region

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11/14/2020

## New File Plots

### Dorsolateral Orbital Cortex

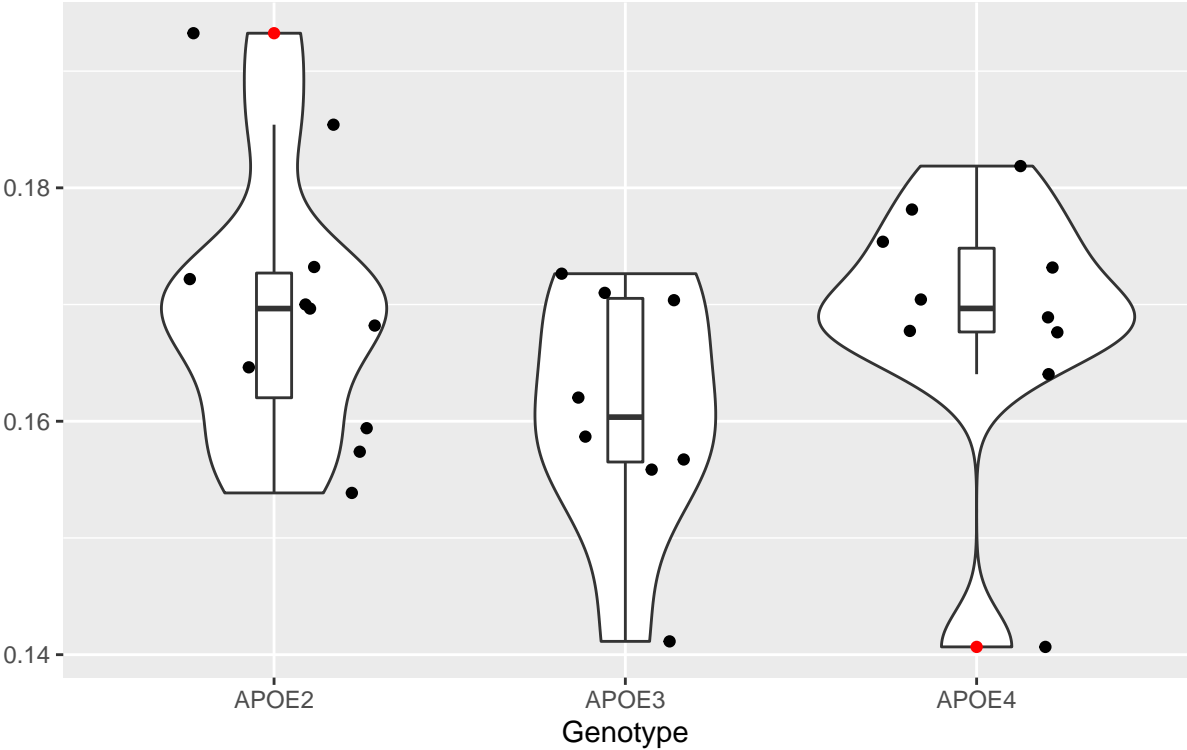
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## geno	2	0.000183	9.141e-05	0.591	0.561
## Residuals	26	0.004021	1.547e-04		

Secondary Auditory Cortex Ventral Part

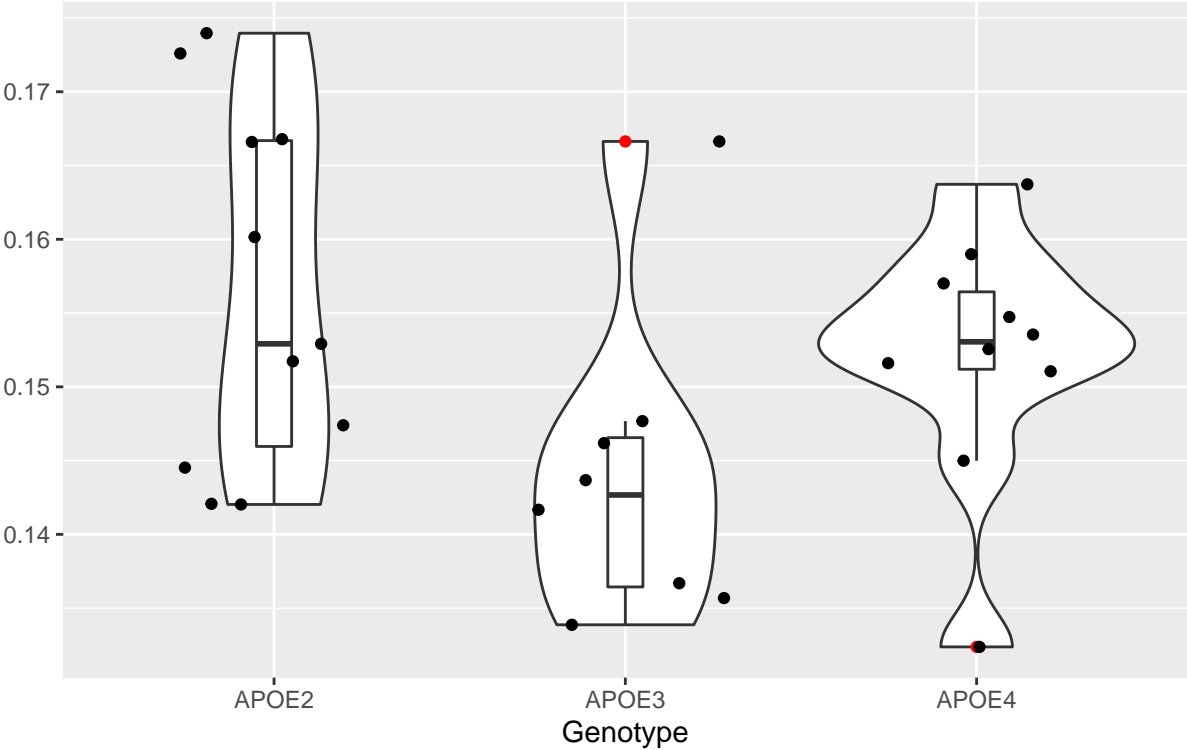
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## geno	2	0.000183	9.141e-05	0.591	0.561
## Residuals	26	0.004021	1.547e-04		

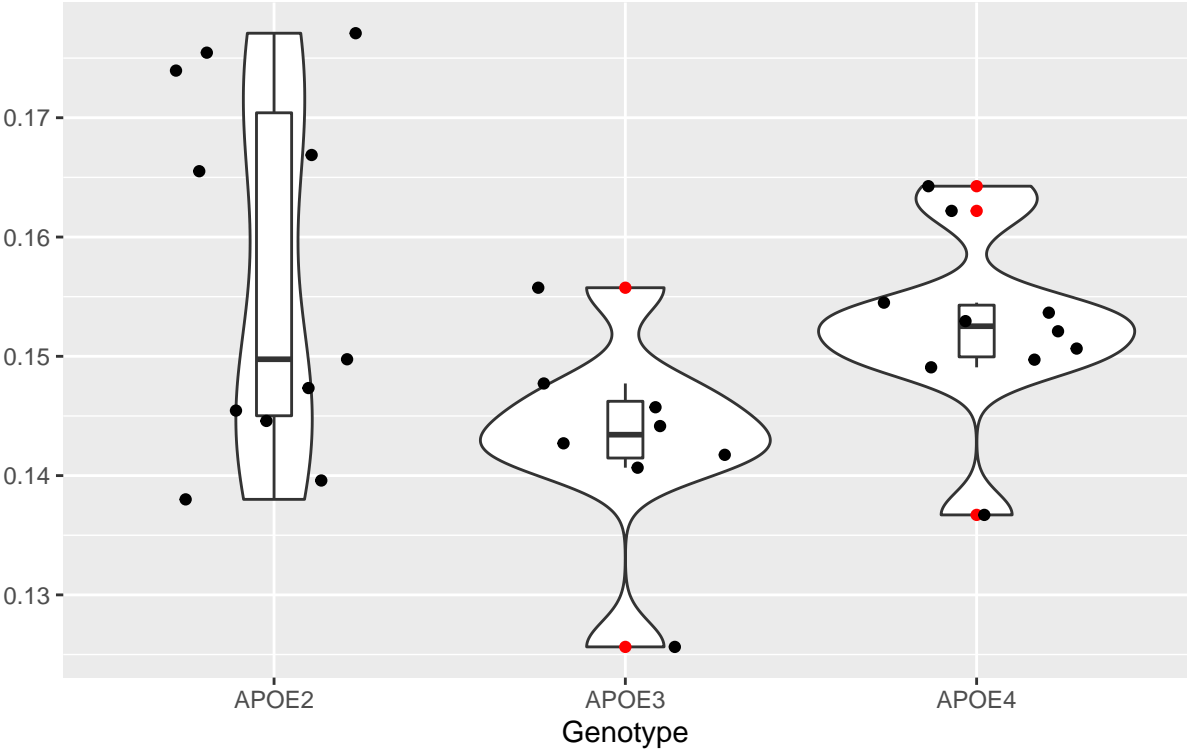
Secondary Auditory Cortex Dorsal Part

Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## geno	2	0.000183	9.141e-05	0.591	0.561
## Residuals	26	0.004021	1.547e-04		

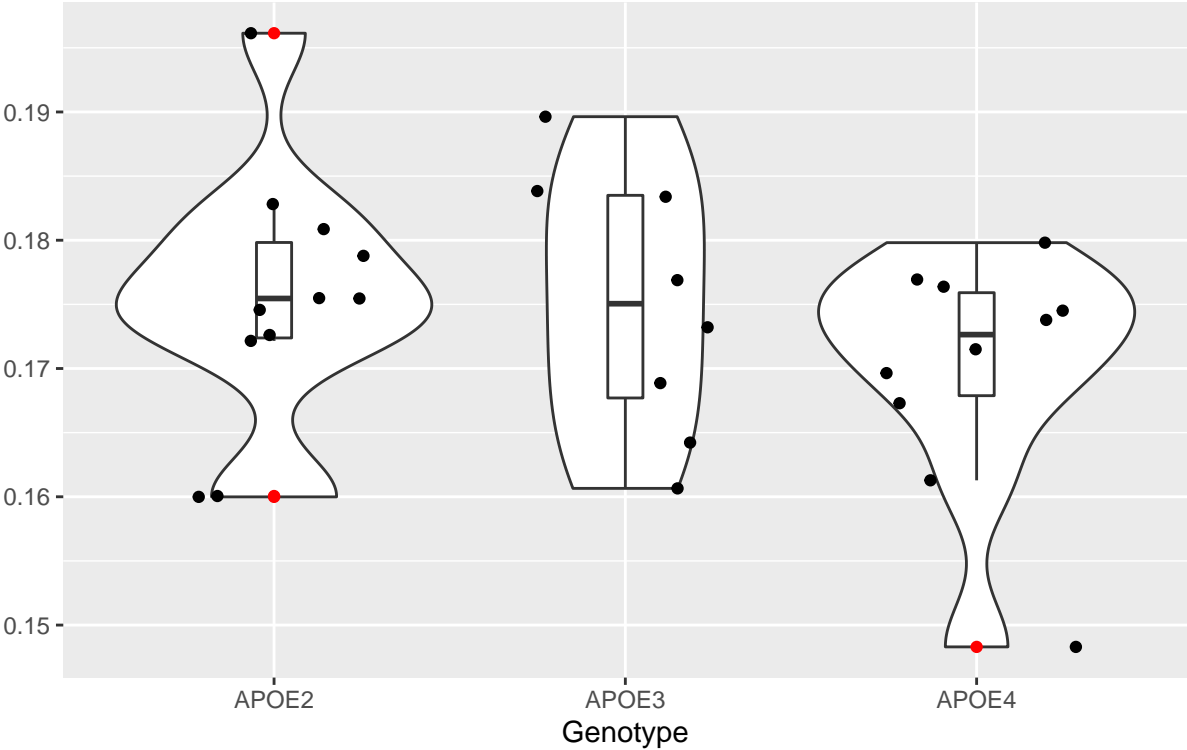
Primary Auditory Cortex  
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## geno	2	0.000183	9.141e-05	0.591	0.561
## Residuals	26	0.004021	1.547e-04		

Cingulate Cortex Area 32

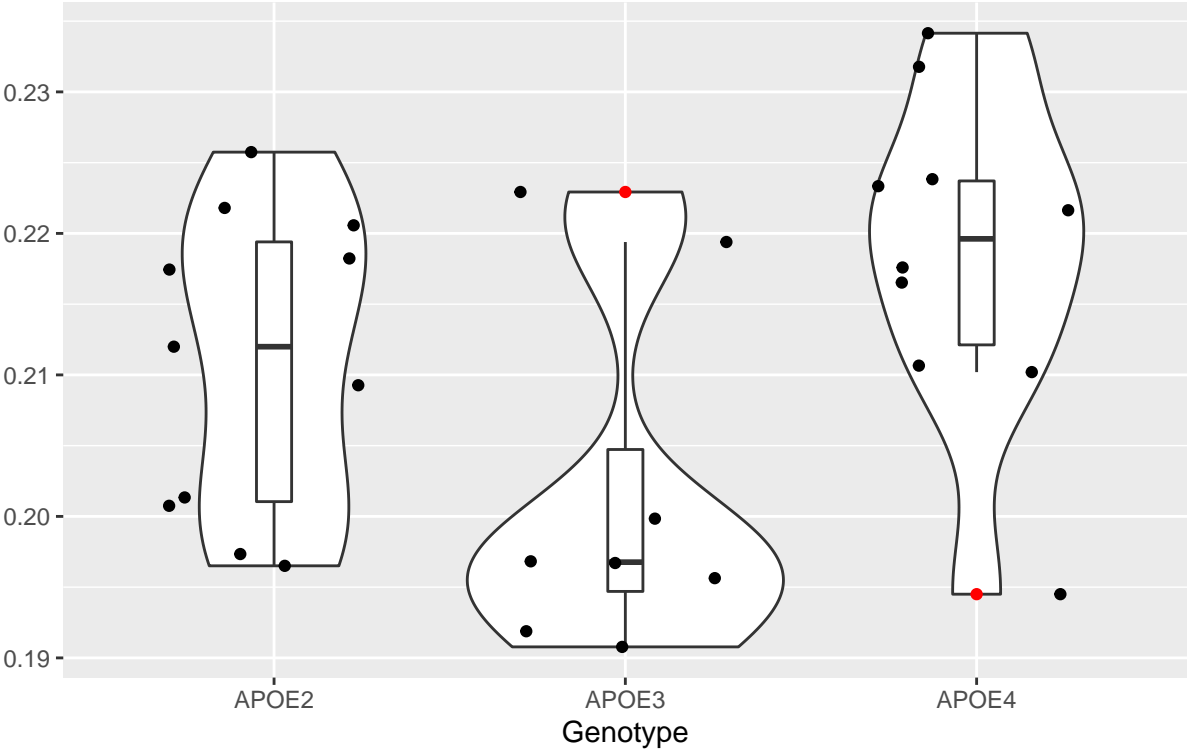
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## geno	2	0.000183	9.141e-05	0.591	0.561
## Residuals	26	0.004021	1.547e-04		

Cingulate Cortex Area 30

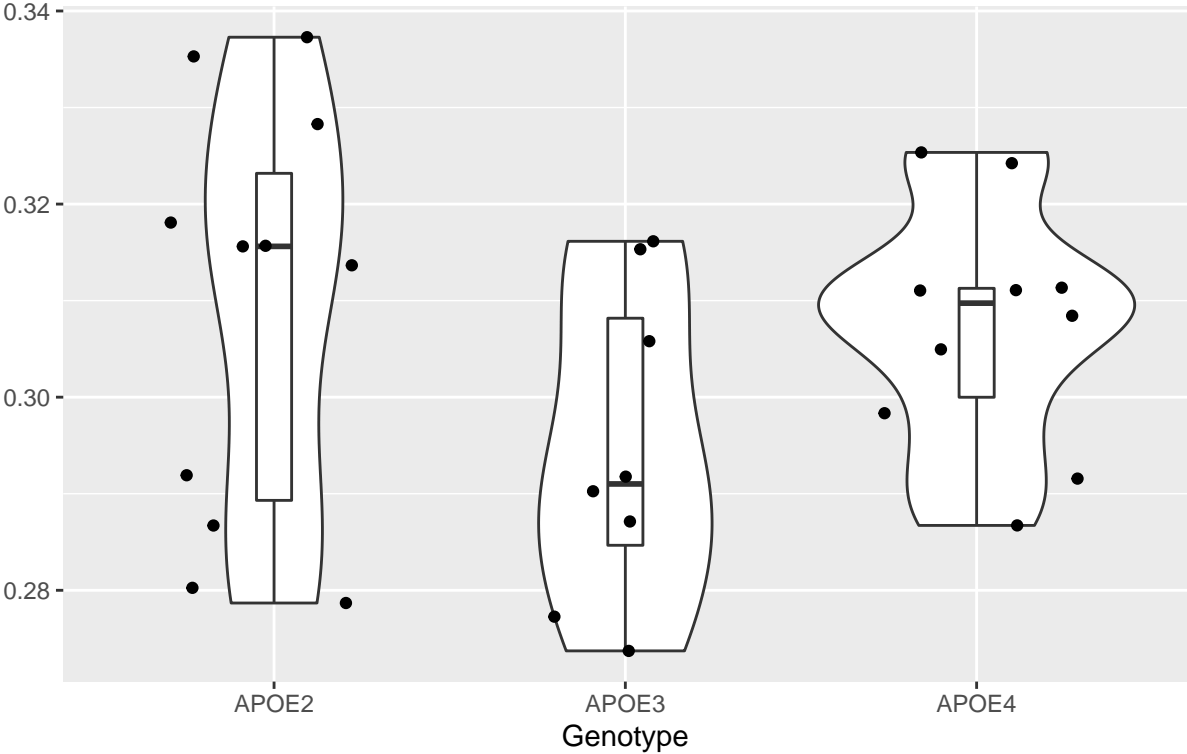
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## geno	2	0.000183	9.141e-05	0.591	0.561
## Residuals	26	0.004021	1.547e-04		

Cingulate Cortex Area 29c

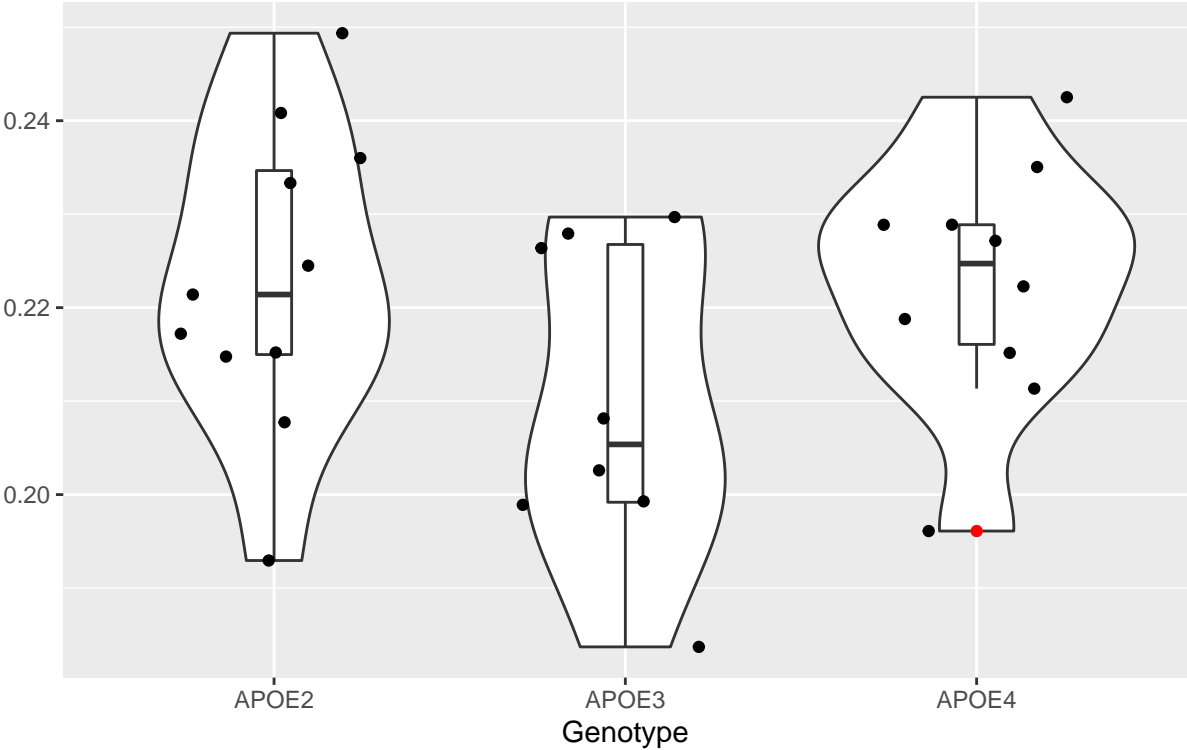
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## geno	2	0.000183	9.141e-05	0.591	0.561
## Residuals	26	0.004021	1.547e-04		

Cingulate Cortex Area 29b

Red points denoting outliers

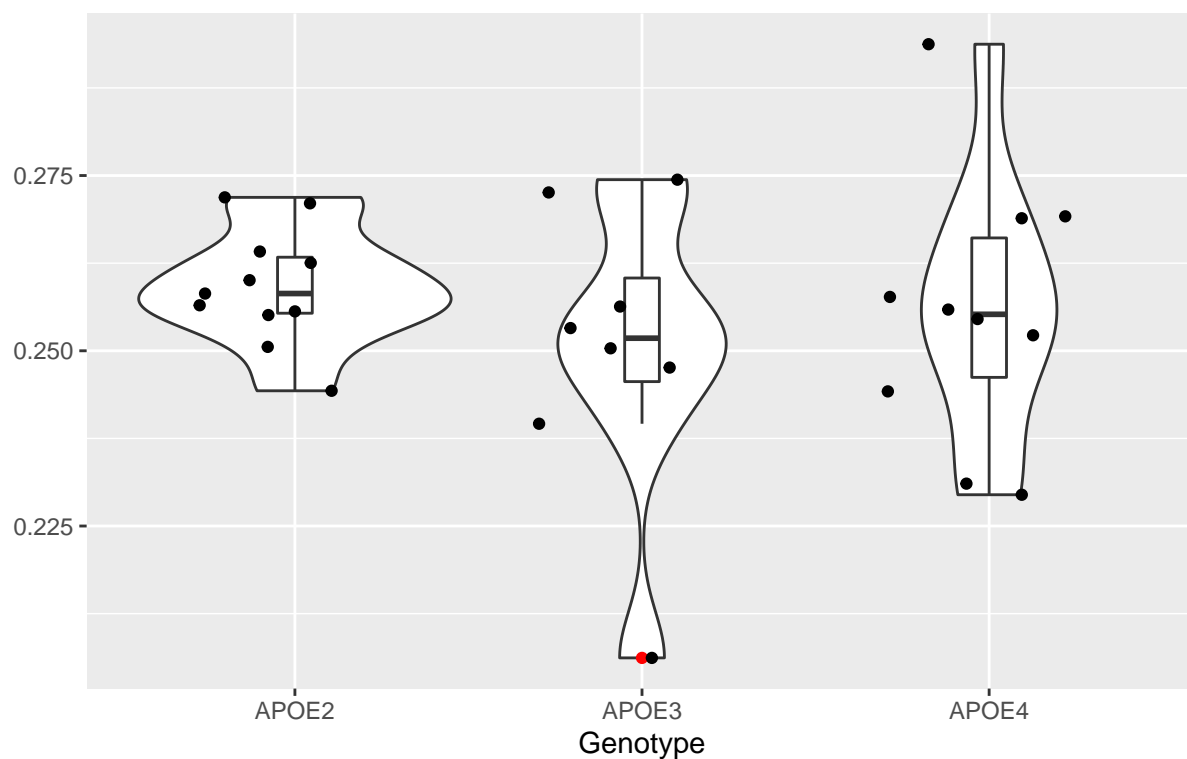


##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## geno	2	0.000183	9.141e-05	0.591	0.561
## Residuals	26	0.004021	1.547e-04		



# Cingulate Cortex Area 29a

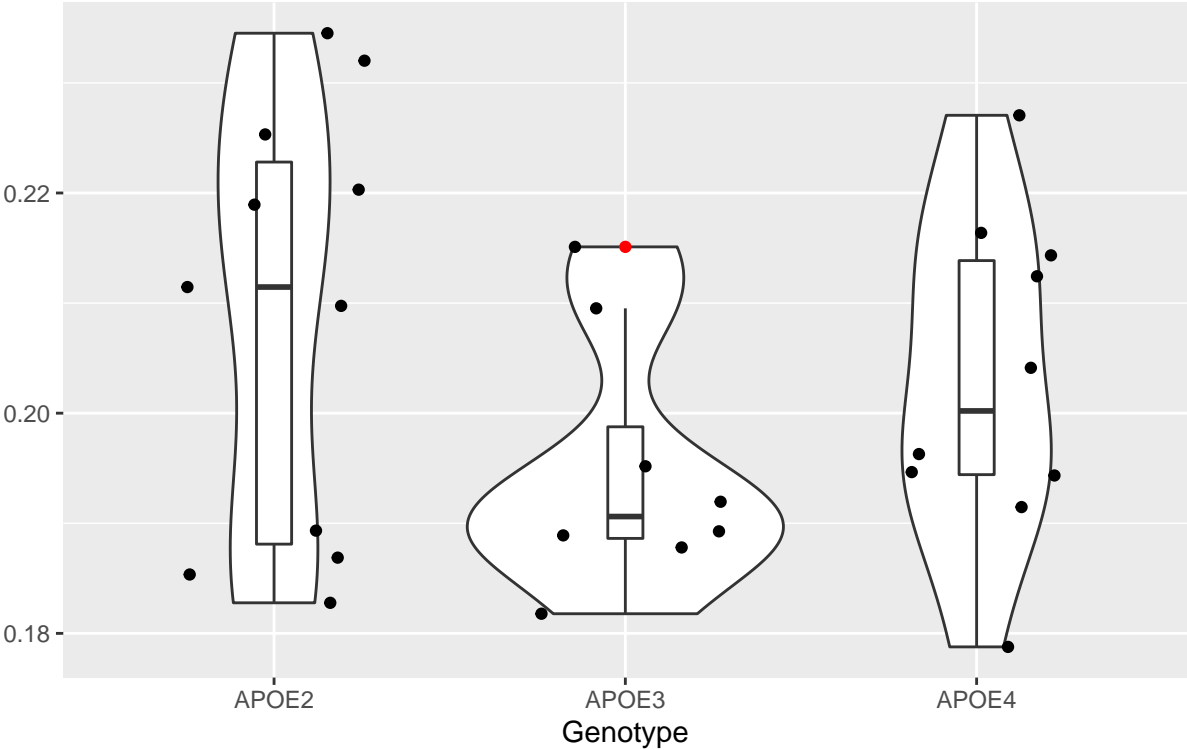
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## geno	2	0.000183	9.141e-05	0.591	0.561
## Residuals	26	0.004021	1.547e-04		

Cingulate Cortex Area 24b Prime

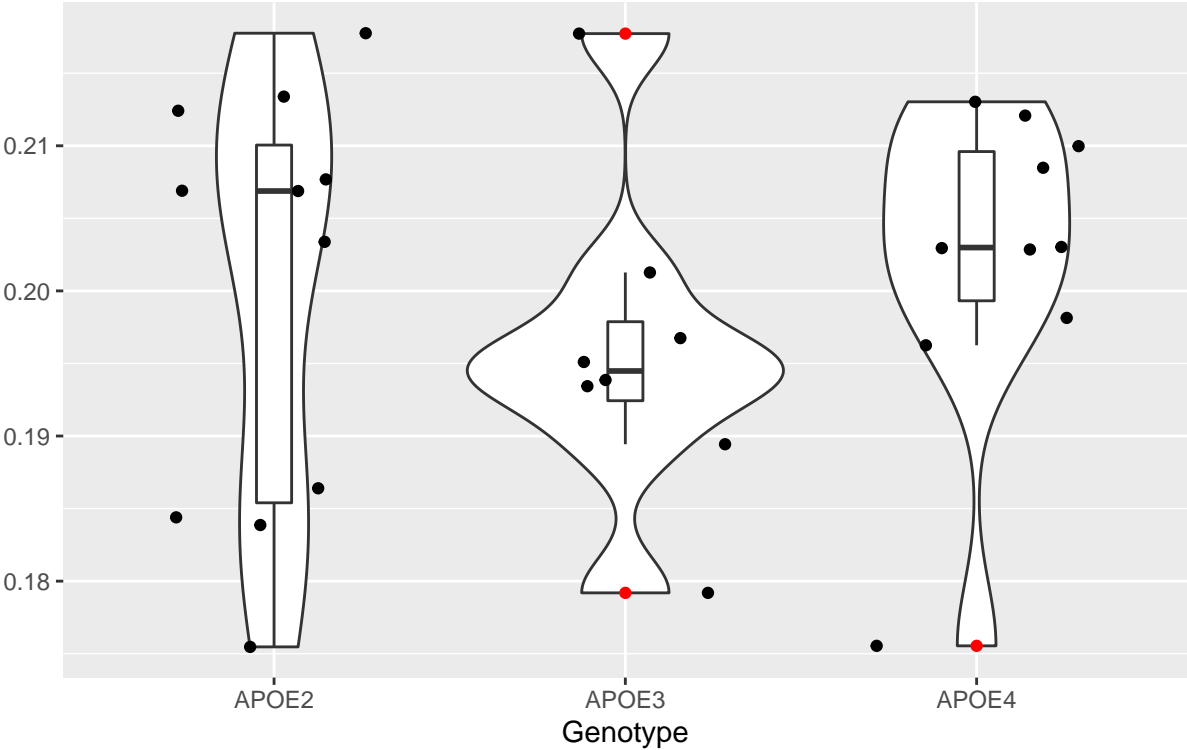
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## geno	2	0.000183	9.141e-05	0.591	0.561
## Residuals	26	0.004021	1.547e-04		

Cingulate Cortex Area 24b

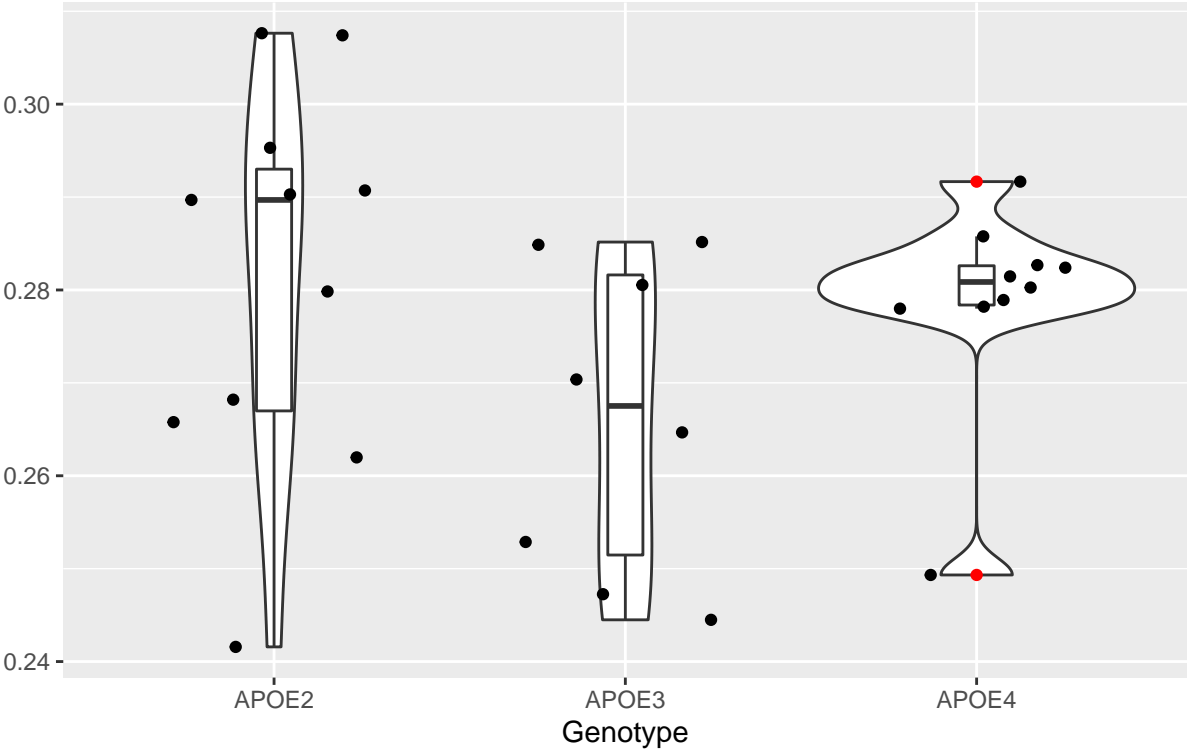
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## geno	2	0.000183	9.141e-05	0.591	0.561
## Residuals	26	0.004021	1.547e-04		

Cingulate Cortex Area 24a Prime

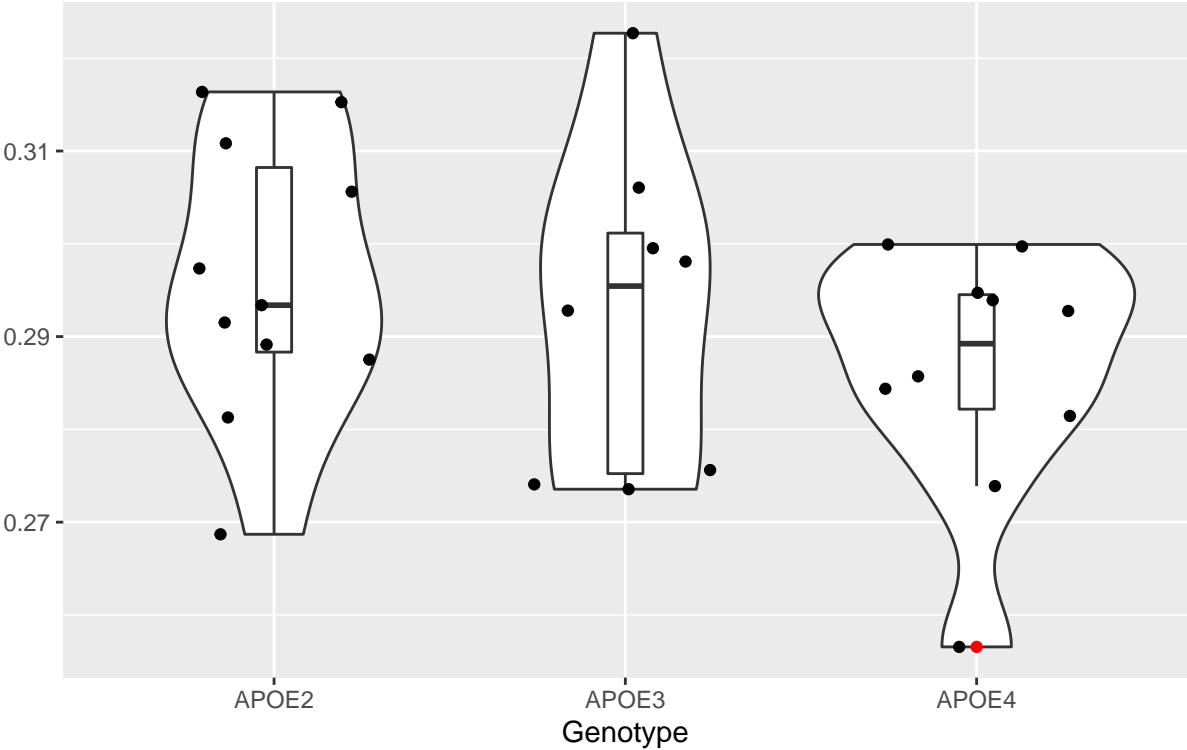
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## geno	2	0.001186	0.0005932	2.131	0.139
## Residuals	26	0.007238	0.0002784		

Cingulate Cortex Area 24a

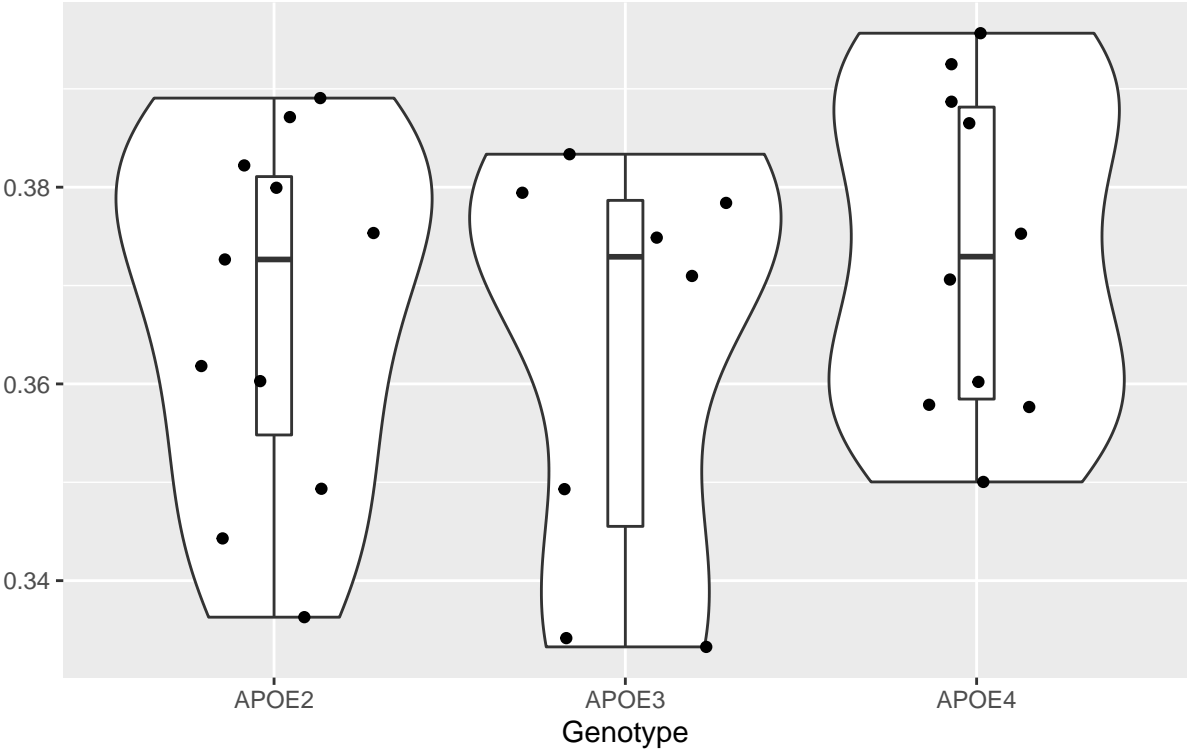
Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## geno	2	0.000512	0.0002562	1.116	0.343
## Residuals	26	0.005970	0.0002296		

Exterior

Red points denoting outliers



##	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## geno	2	0.000512	0.0002562	0.761	0.477
## Residuals	26	0.008750	0.0003365		