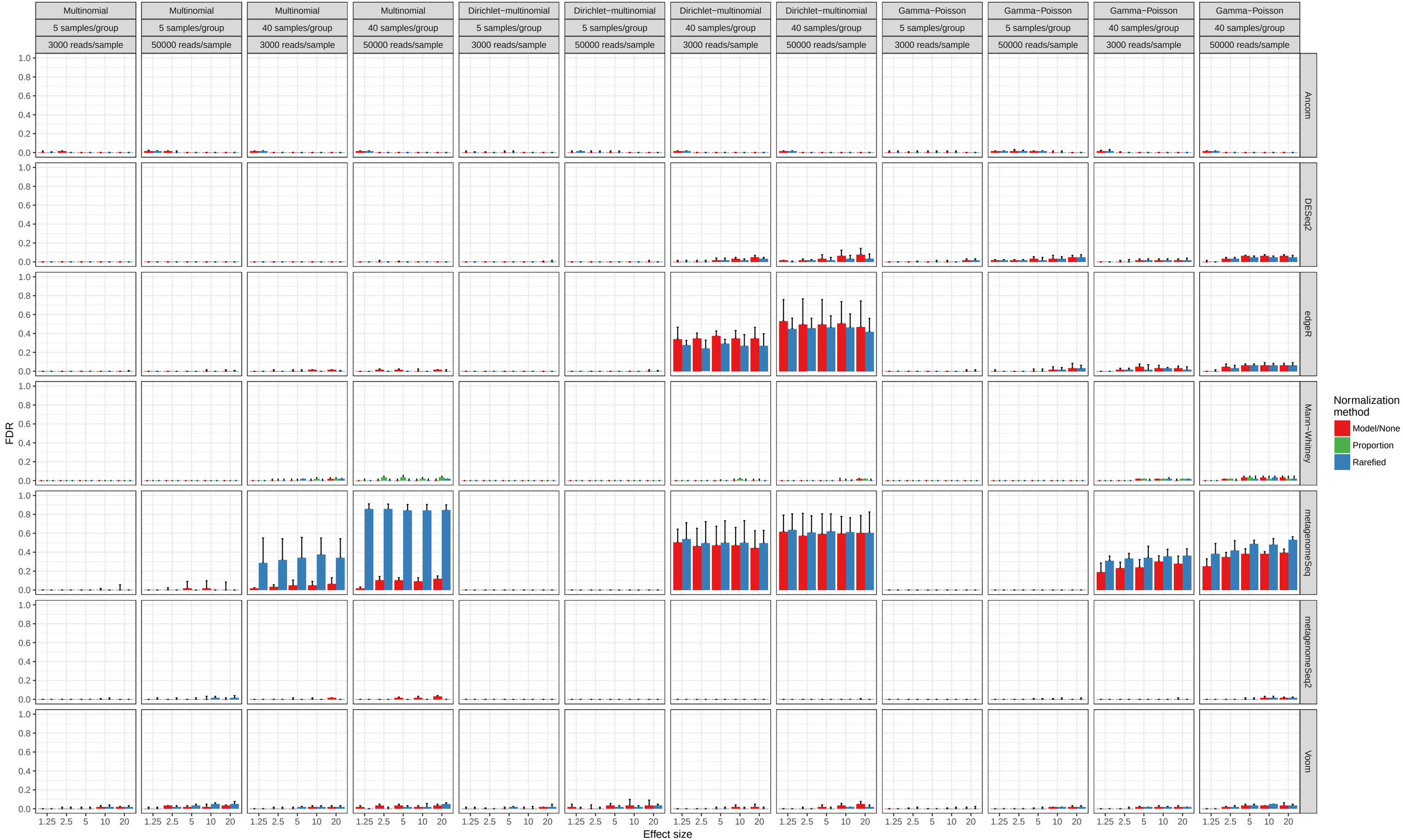
Multinomial
Dirichlet−multinomial
Gamma−Poisson
5 samples/group
40 samples/group
3000 reads/sample
50000 reads/sample
Ancom
DESeq2
edgeR
Mann−Whitney
metagenomeSeq
metagenomeSeq2
Voom
FDR
Effect size
1.25
2.5
5
10
20
Normalization method
Model/None
Proportion
Rarefied