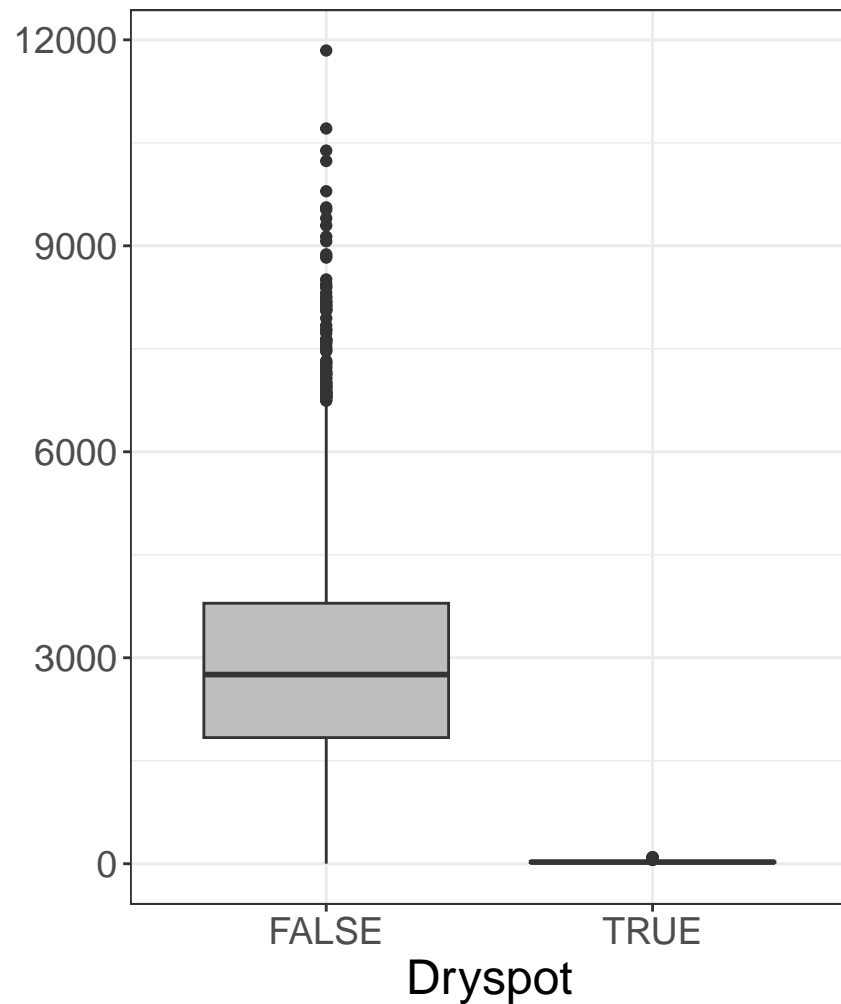
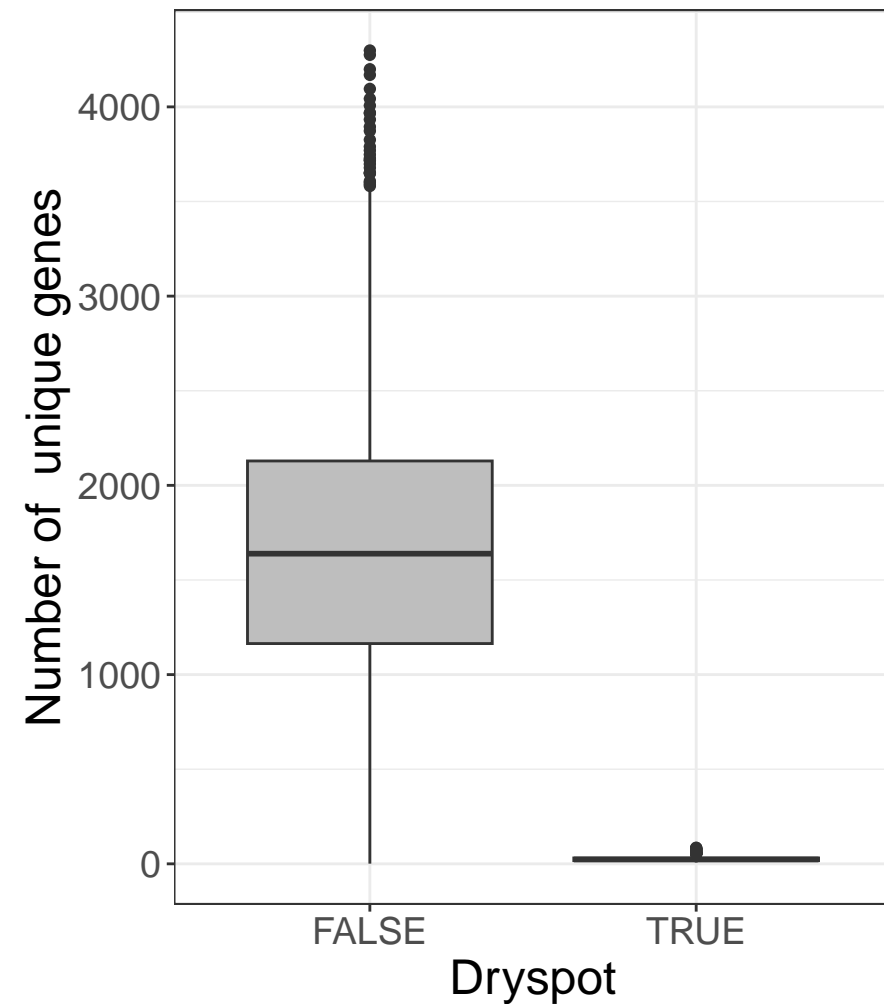


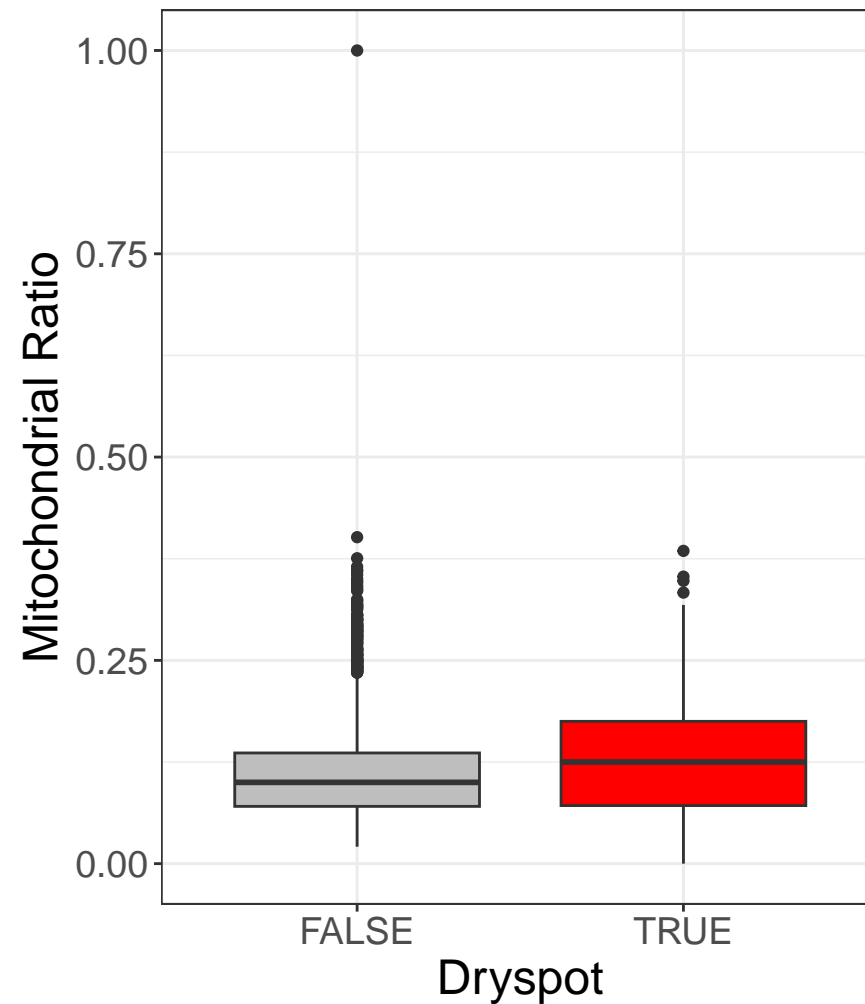
Library size



Unique Genes



Mito Ratio



Multiscale variance

