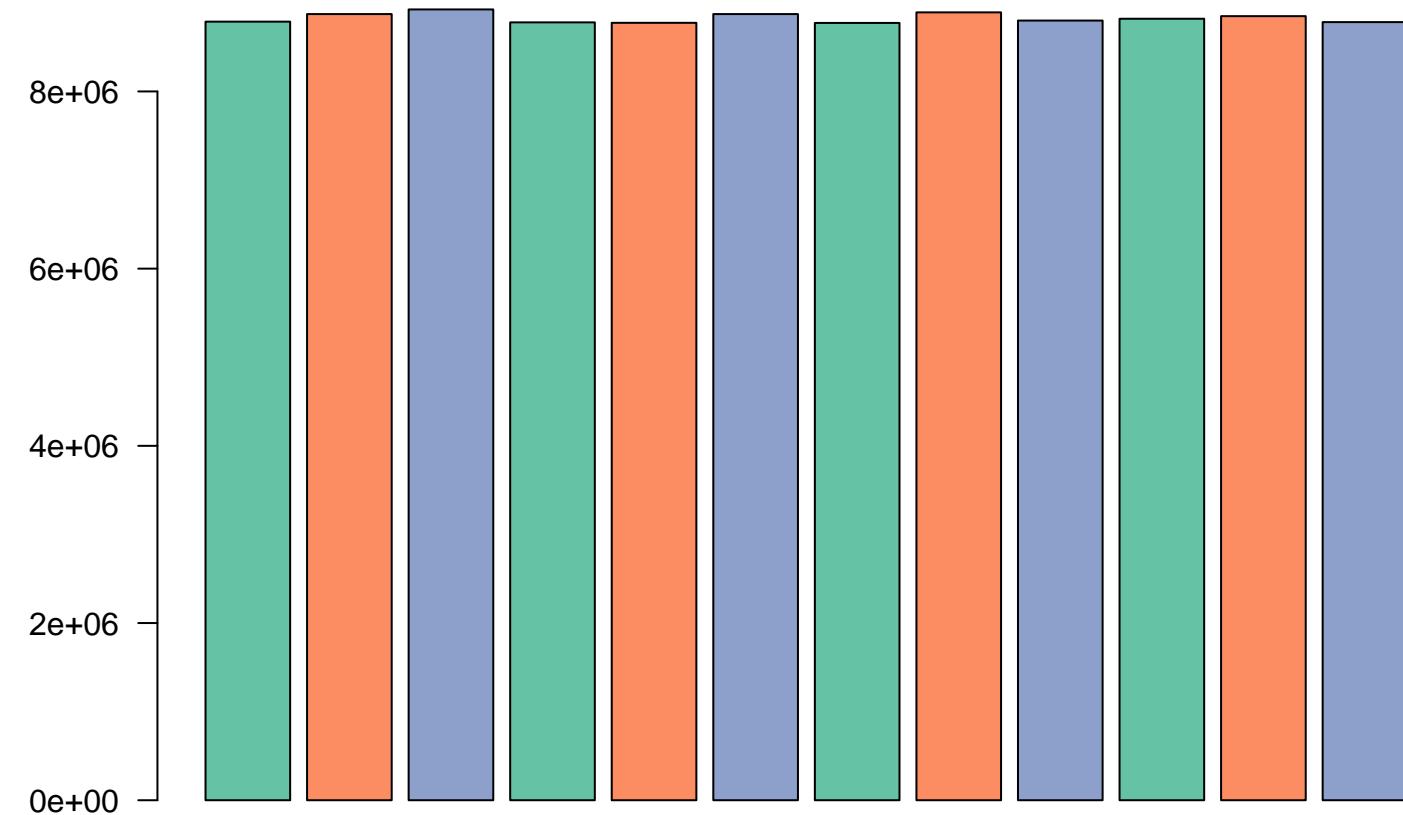
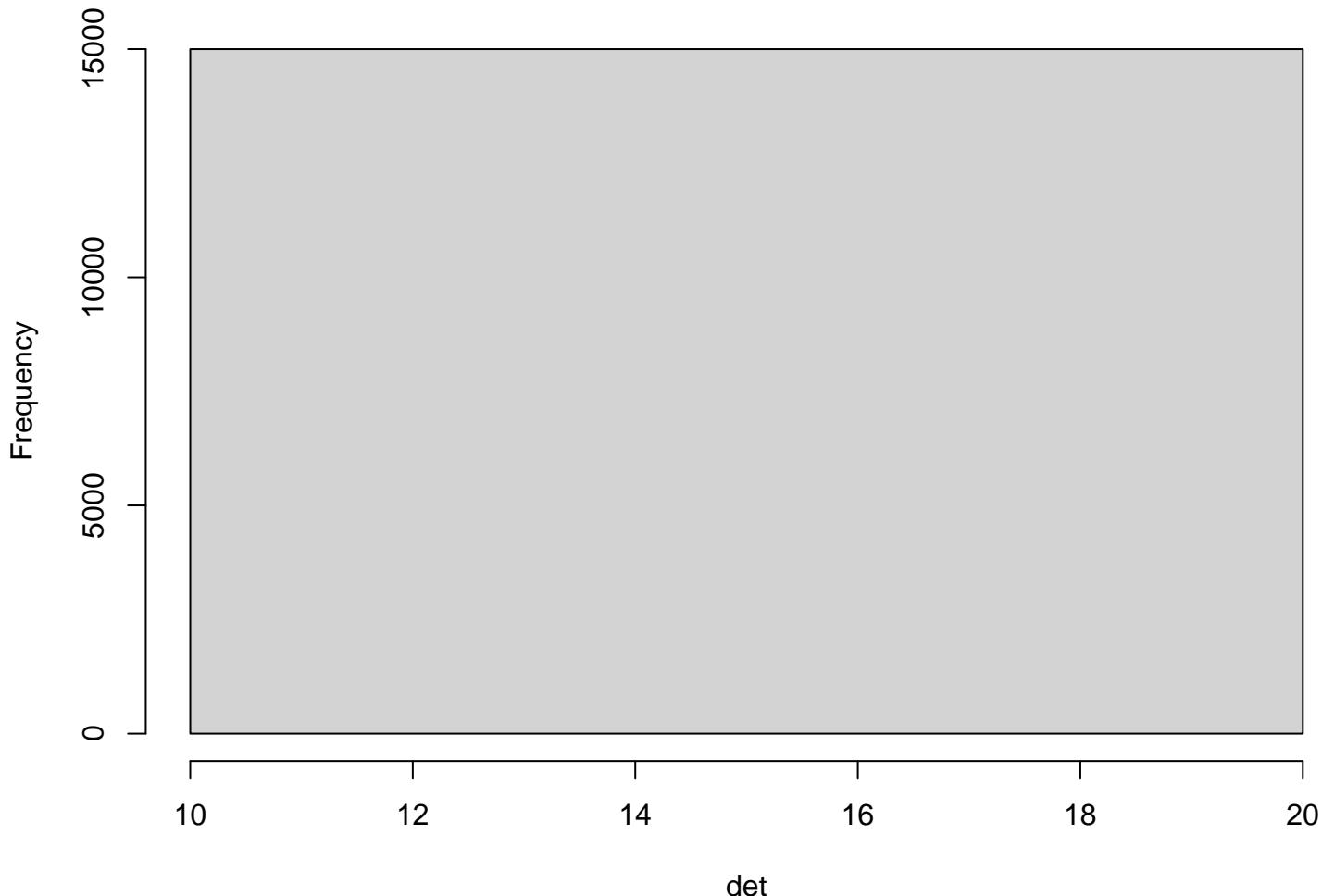


Library sizes



Genes detected (count > 0)



$\log_2(\text{CPM})$ per sample

