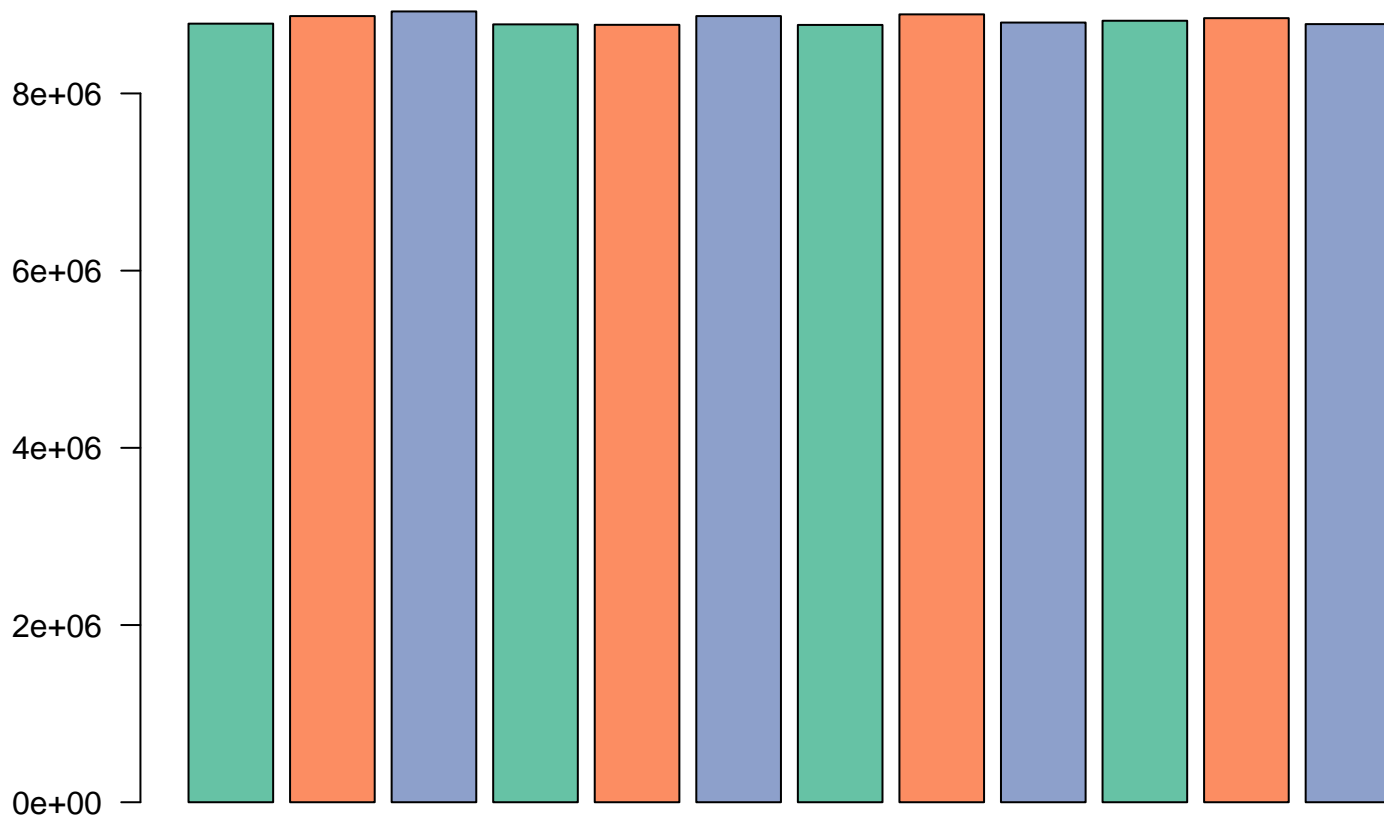
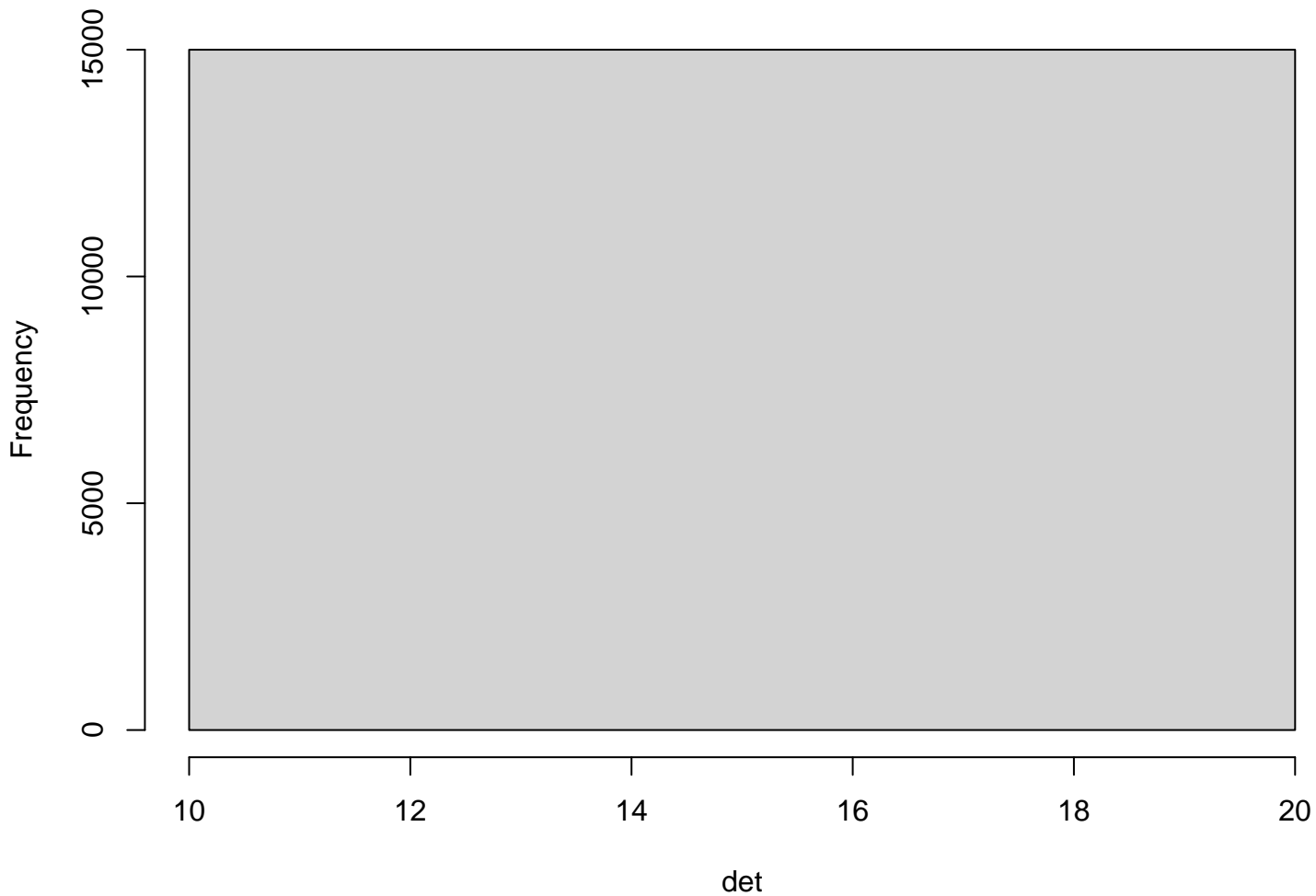


Library sizes



Genes detected (count > 0)



log2(CPM) per sample

