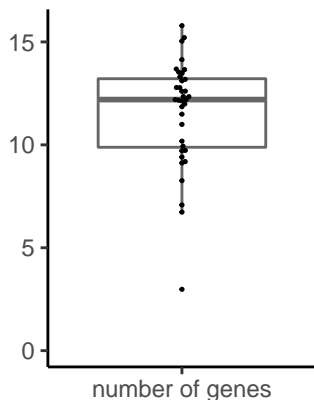
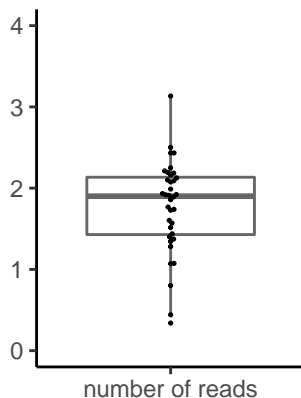


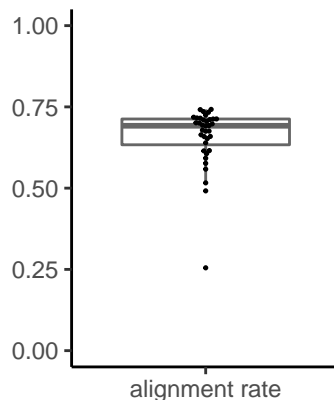
number of genes x
median:
12198



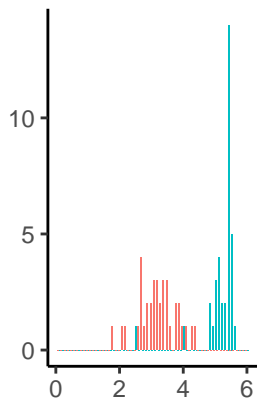
unique M reads
median:
1900807.5



alignment rate (unio
median:
0.69

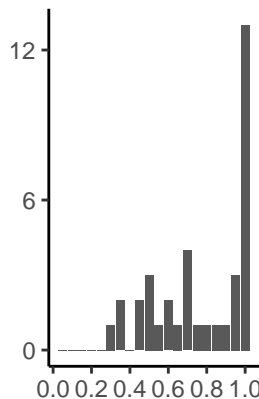


NMS scores
cells



group
cont
neuron

bootstrap score
#cells



Pearson correla
cells

