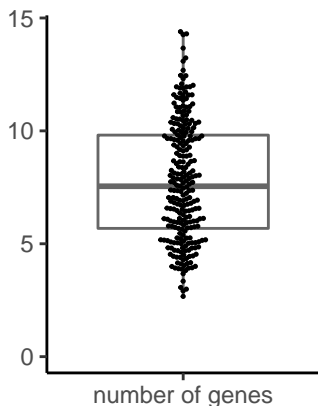
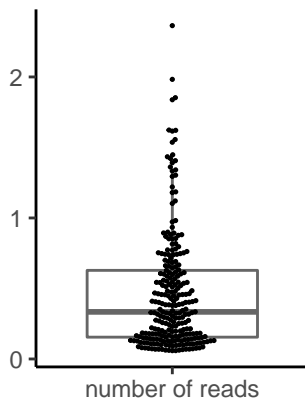


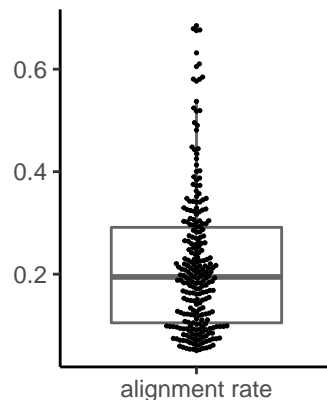
number of genes x
median:
7547



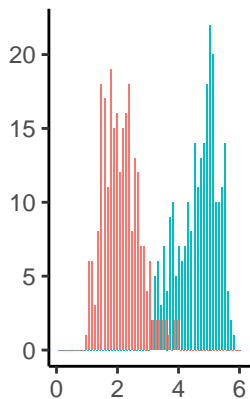
unique M reads
median:
333923



alignment rate (unique)
median:
0.19

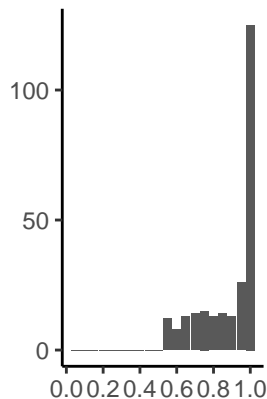


NMS scores
cells



group
cont
neuron

bootstrap score
#cells



Pearson correlation
cells

