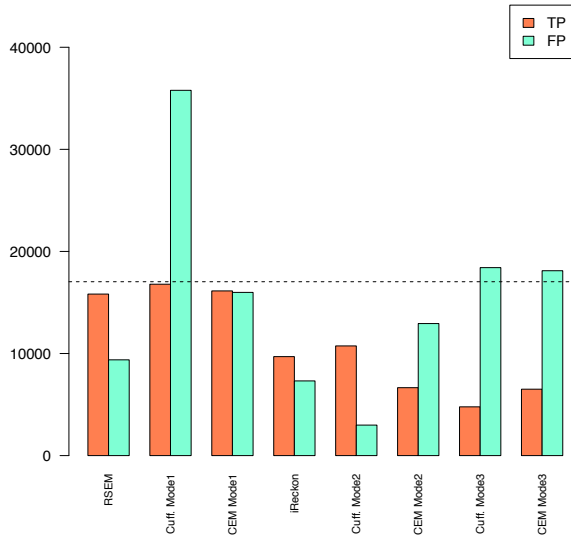


PE 75 bp – 60 M (Set-up 2)

Alignment with transcriptome

CA

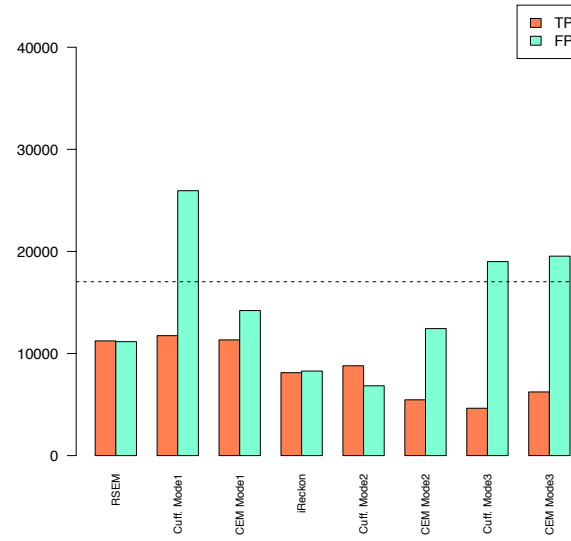
TP and FP (CA– 75 read length)



A

IA

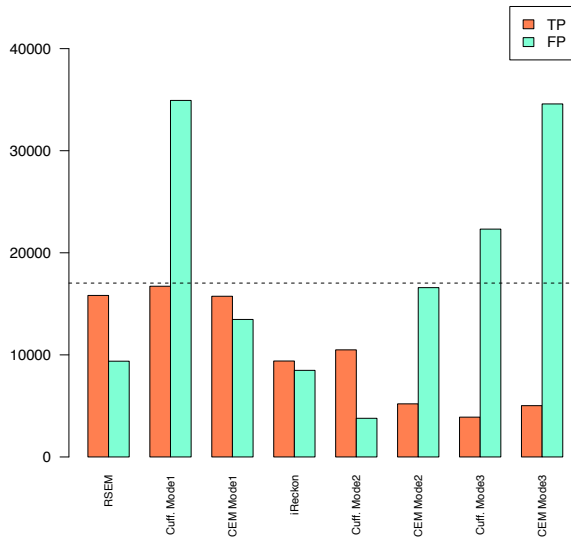
TP and FP (IA– 75 read length)



B

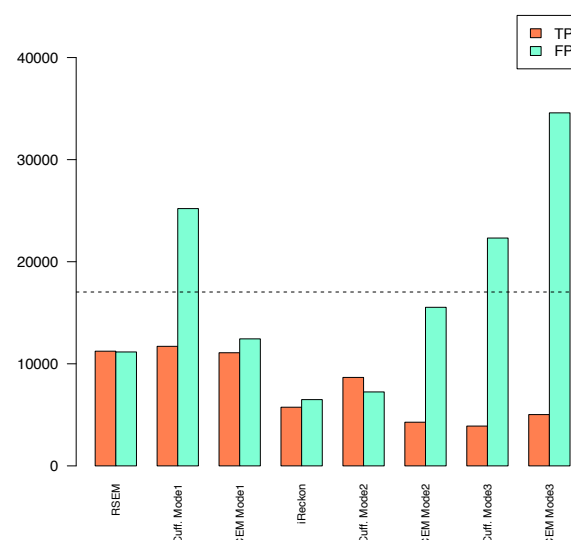
Alignment data driven

TP and FP (CA– 75 read length)



C

TP and FP (IA– 75 read length)



D