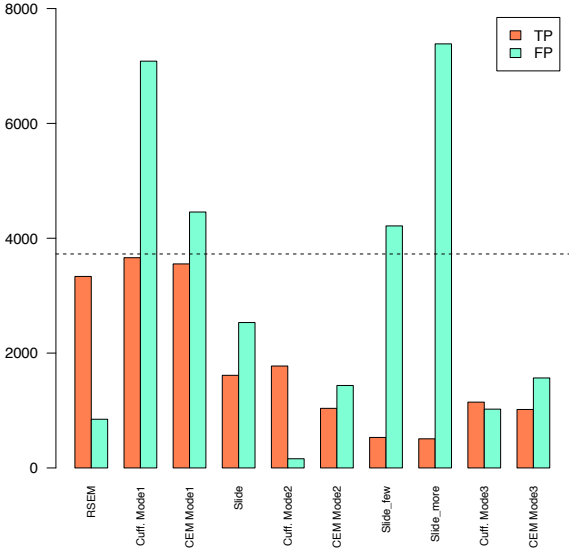


SE 100 bp – 20 M (Set-up 1)

Alignment with transcriptome

CA

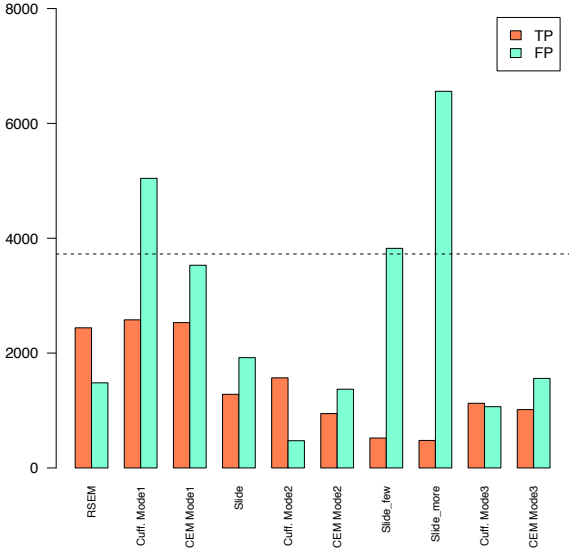
TP and FP (CA– 100 read length)



A

IA

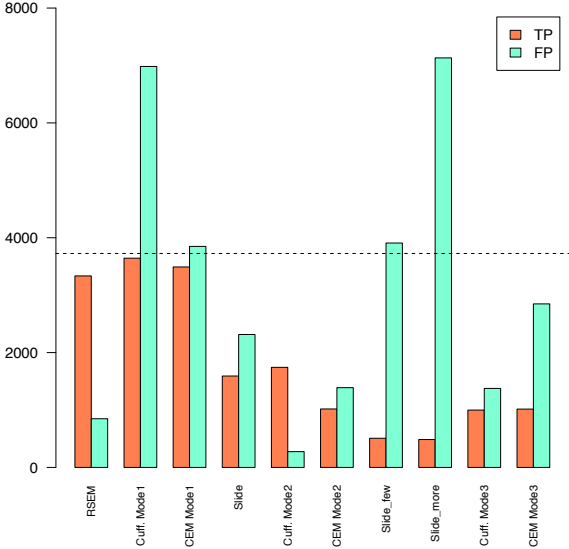
TP and FP (IA– 100 read length)



B

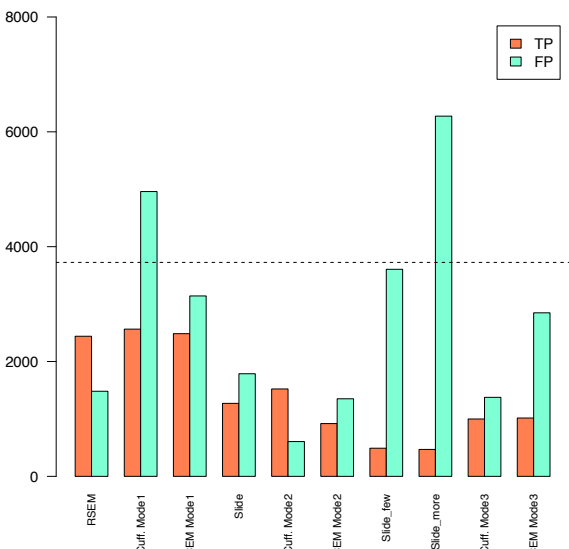
Alignment data driven

TP and FP (CA– 100 read length)



C

TP and FP (IA– 100 read length)



D