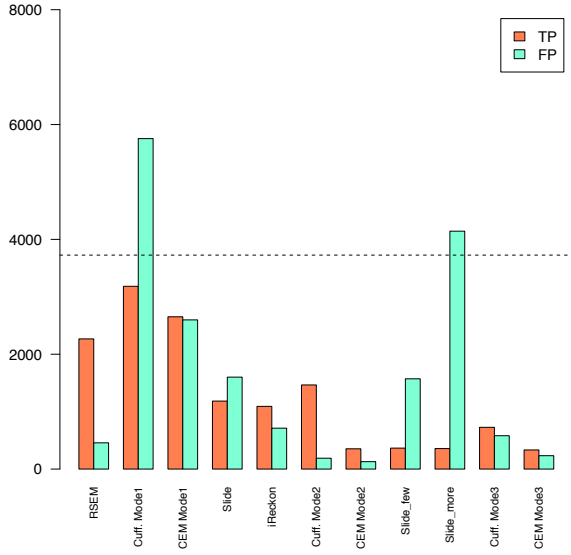


PE 100 bp – 0.25 M (Set-up 1)

Alignment with transcriptome

CA

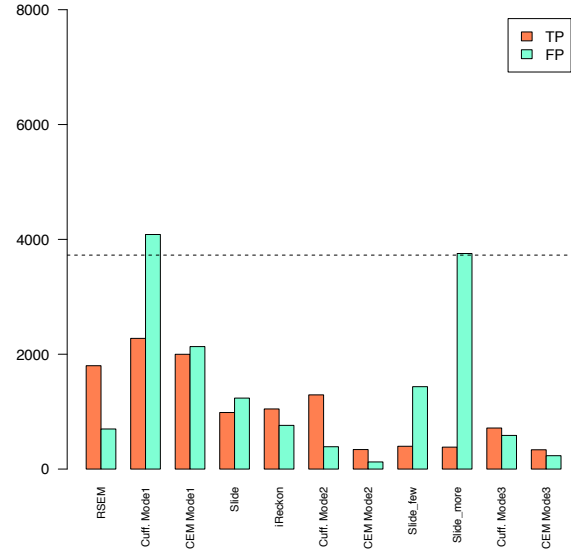
TP and FP (CA- 100 read length)



A

IA

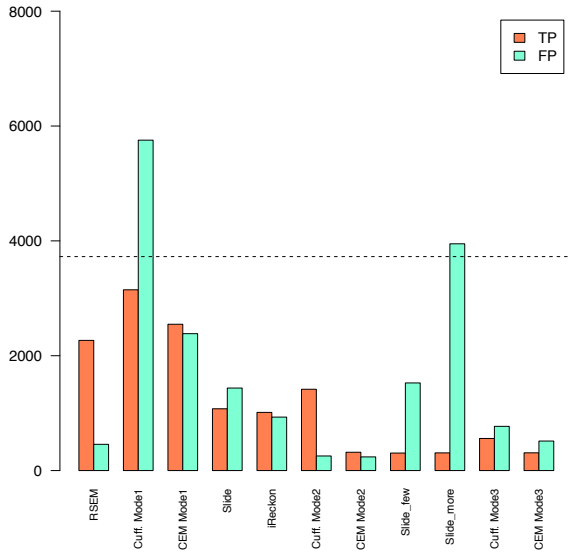
TP and FP (IA- 100 read length)



B

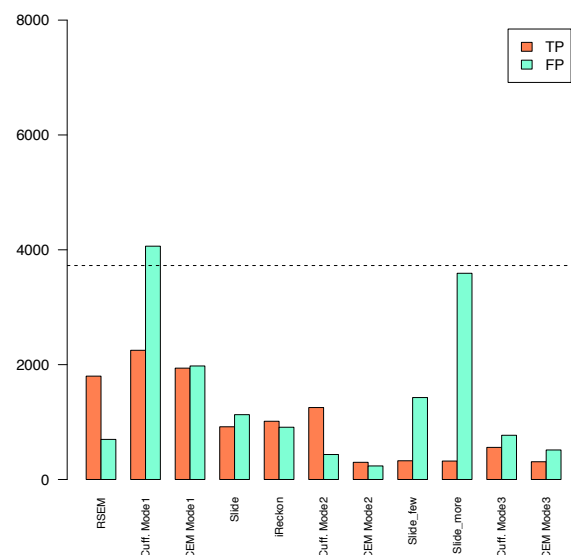
Alignment data driven

TP and FP (CA- 100 read length)



C

TP and FP (IA- 100 read length)



D