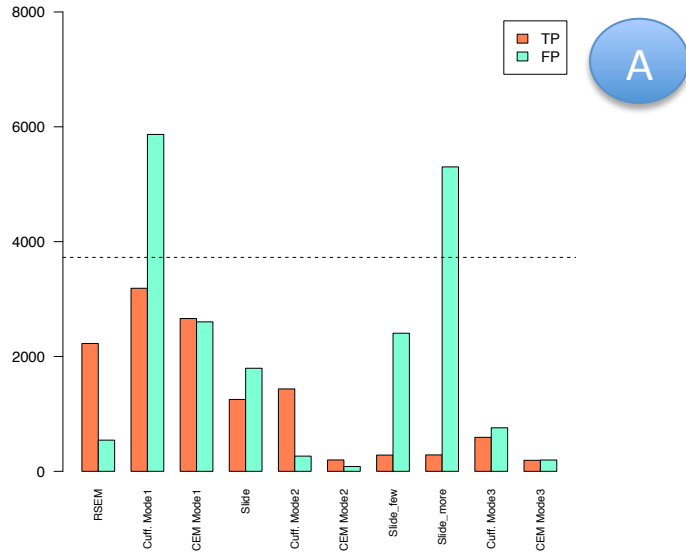


# SE 100 bp – 0.25 M (Set-up 1)

Alignment with transcriptome

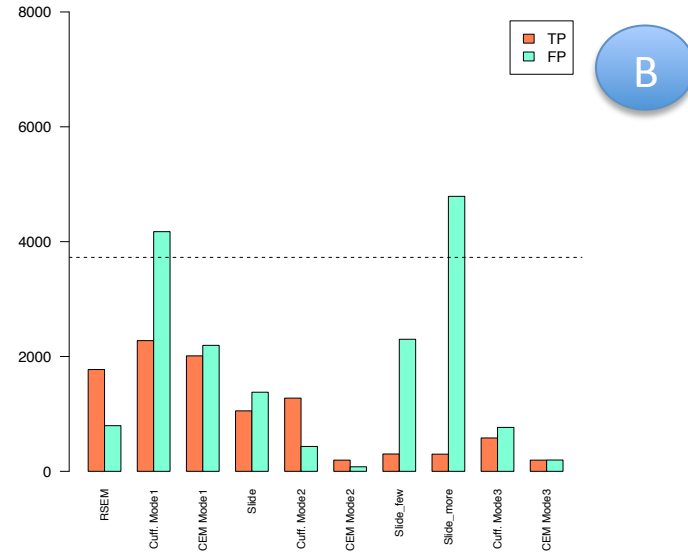
CA

TP and FP (CA– 100 read length)



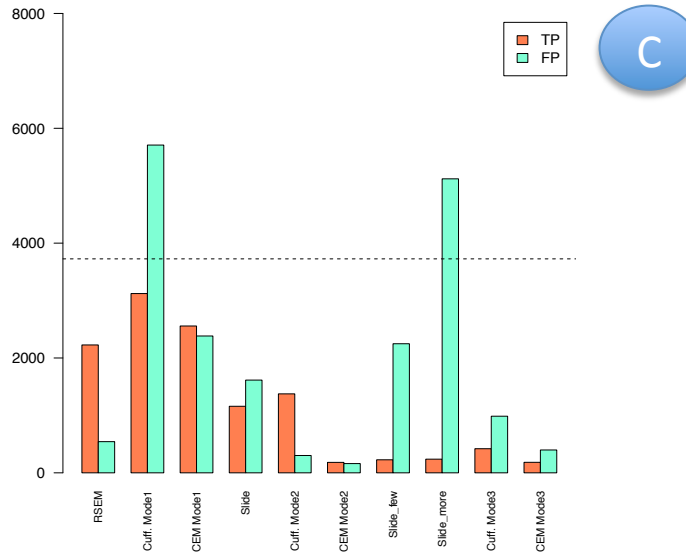
IA

TP and FP (IA– 100 read length)



Alignment data driven

TP and FP (CA– 100 read length)



TP and FP (IA– 100 read length)

