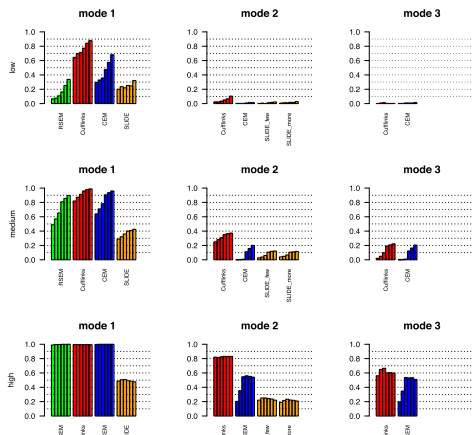


SE 100 bp (Set-up 1)

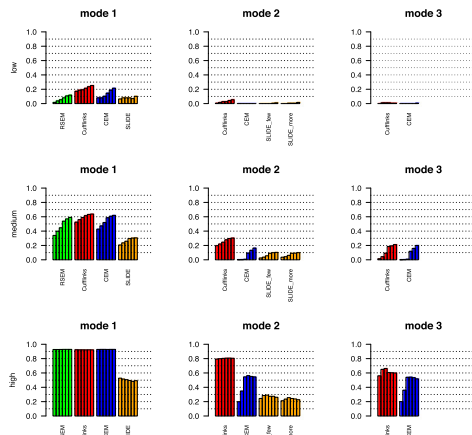
Alignment with transcriptome

Recall (CA – 100 read length)



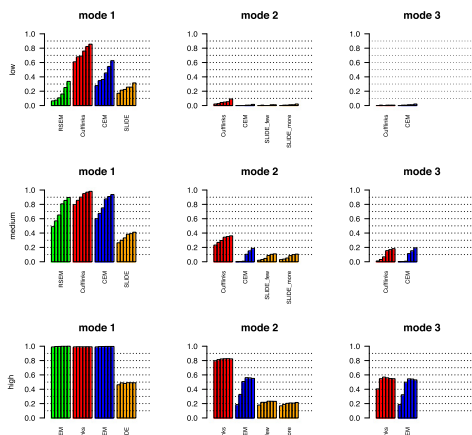
A

Recall (IA – 100 read length)



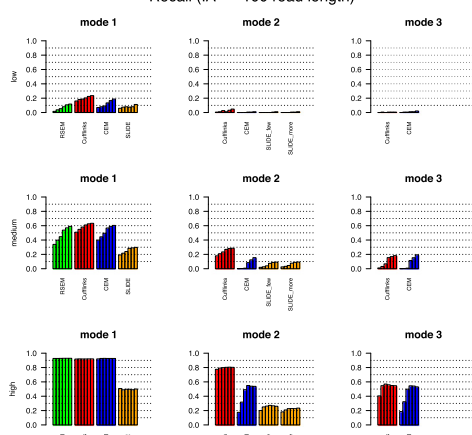
B

Recall (CA – 100 read length)



C

Recall (IA – 100 read length)



D

Alignment data driven