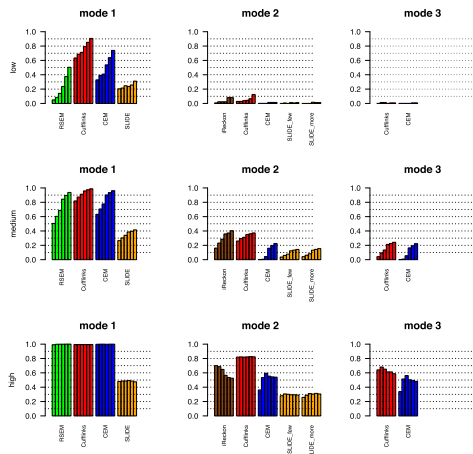


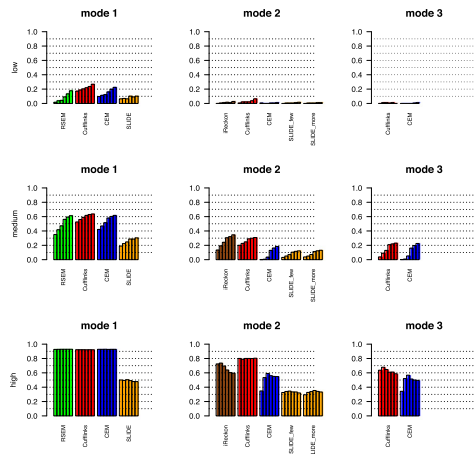
PE 100 bp (Set-up 1)

Alignment with transcriptome

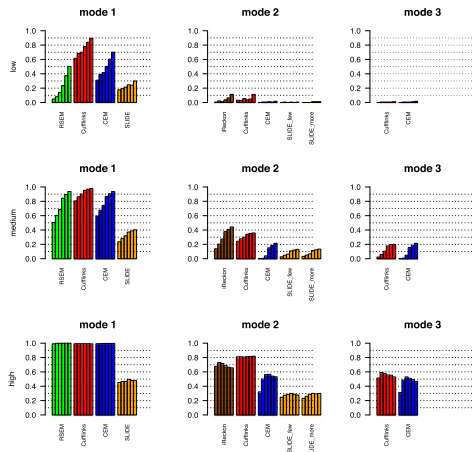
Recall (CA – 100 read length)



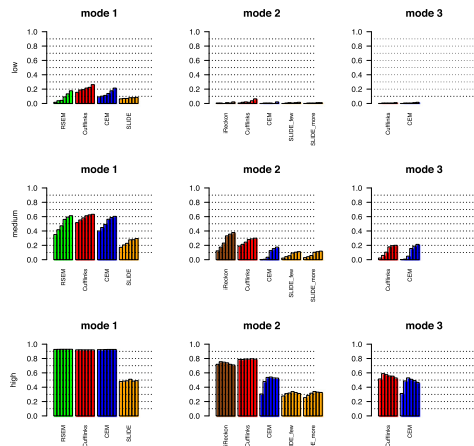
Recall (IA – 100 read length)



Recall (CA – 100 read length)



Recall (IA – 100 read length)



Alignment data driven