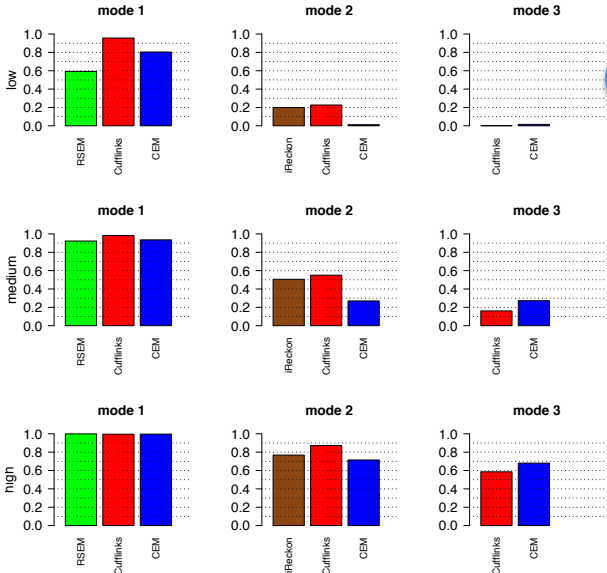


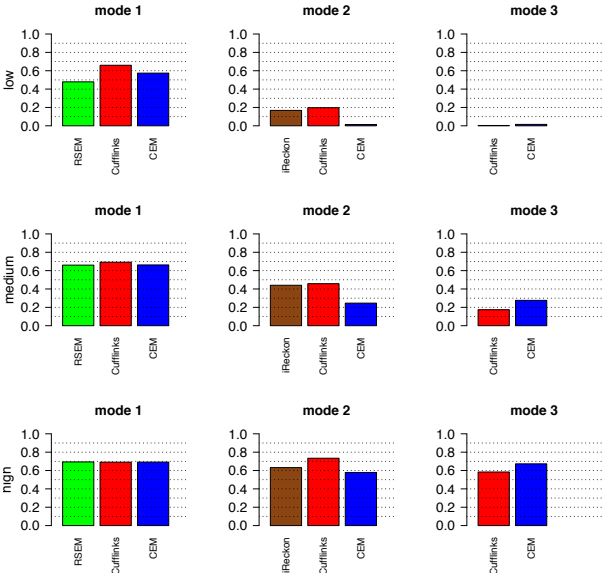
PE 75 bp (Set-up 2)

Alignment with transcriptome

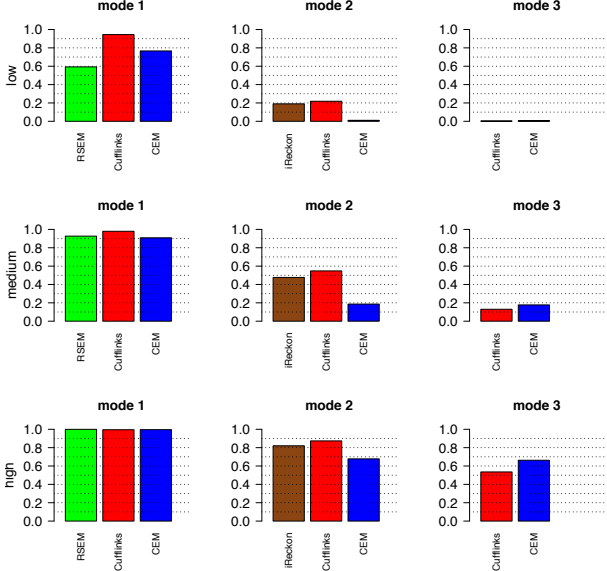
Recall (CA – 75 read length)



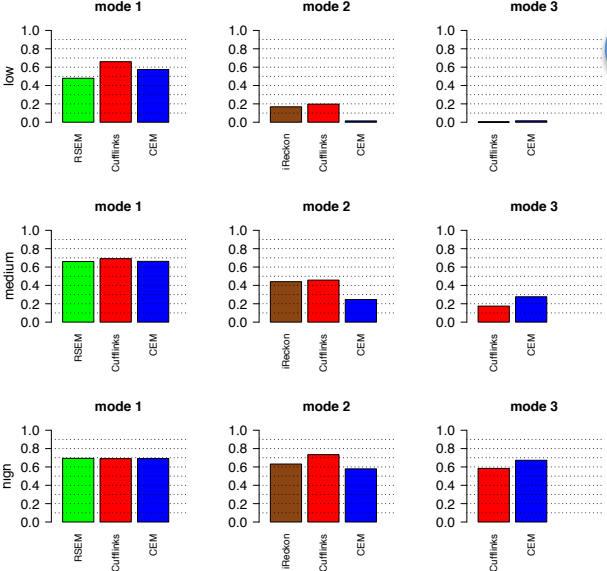
Recall (IA – 75 read length)



Recall (CA – 75 read length)



Recall (IA – 75 read length)



Alignment data driven