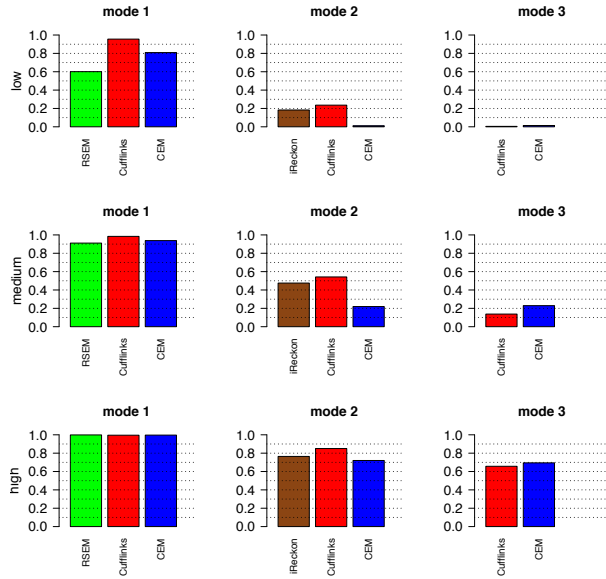


PE 50 bp (Set-up 2)

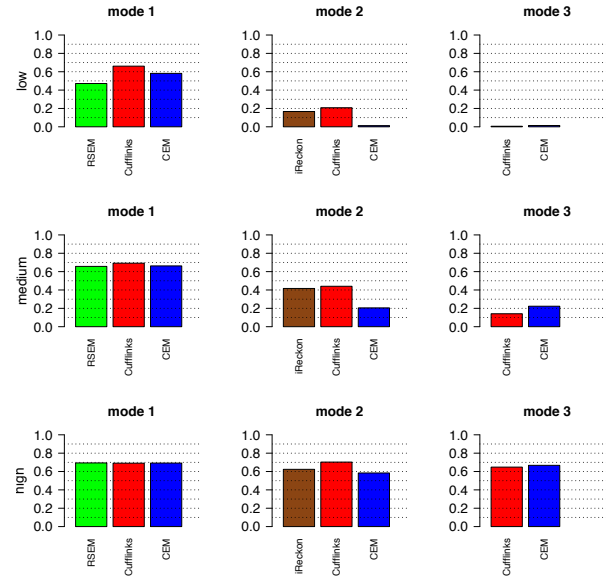
Alignment with transcriptome

Recall (CA – 50 read length)



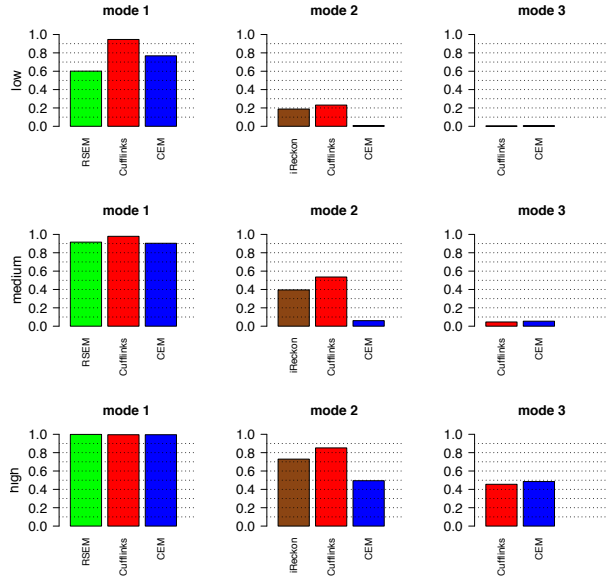
A

Recall (IA – 50 read length)



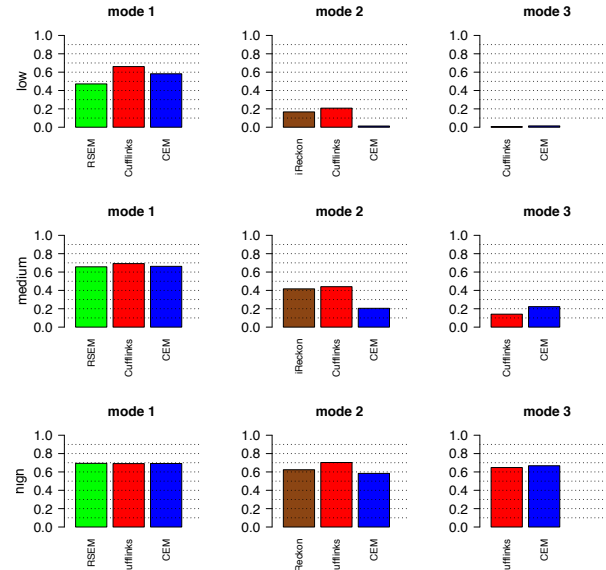
B

Recall (CA – 50 read length)



C

Recall (IA – 50 read length)



D

Alignment data driven