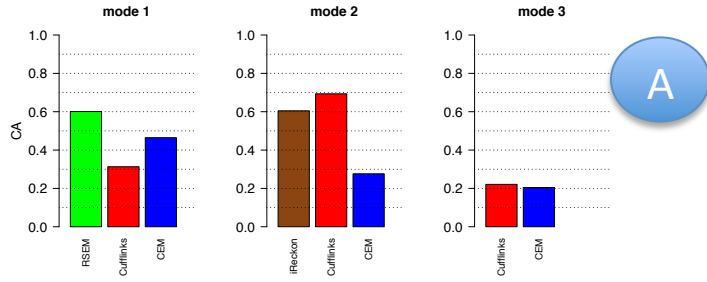


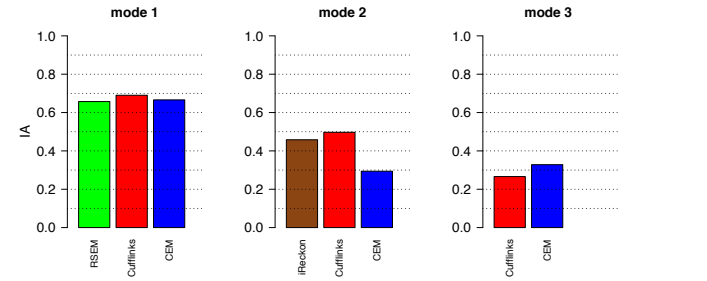
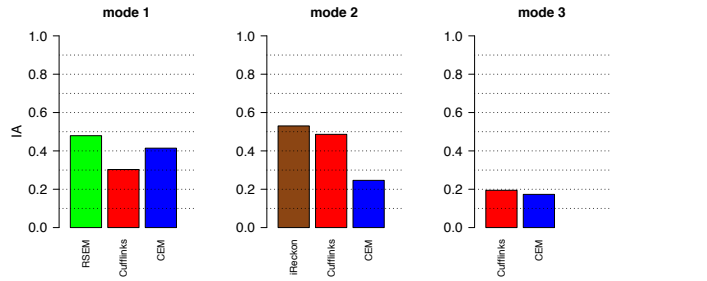
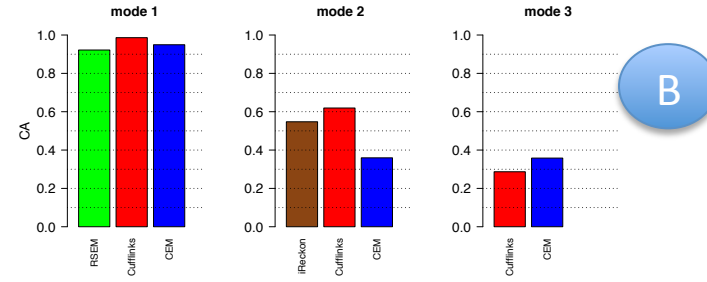
# PE 50 bp (Set-up 2)

Alignment with transcriptome

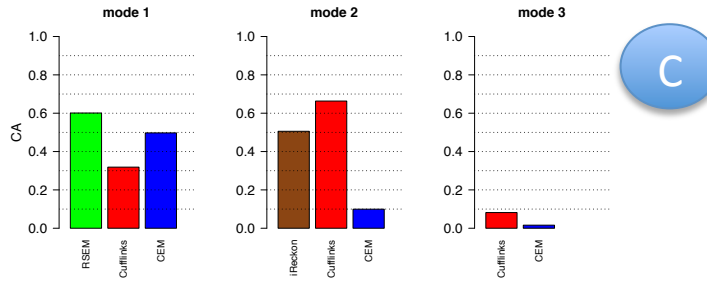
Precision (50 read length)



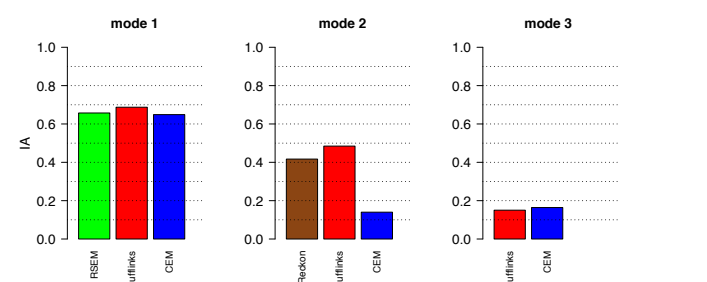
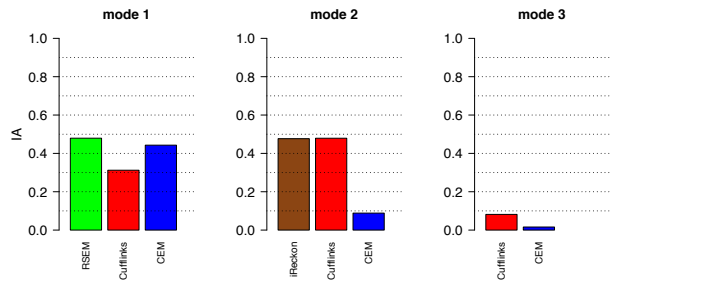
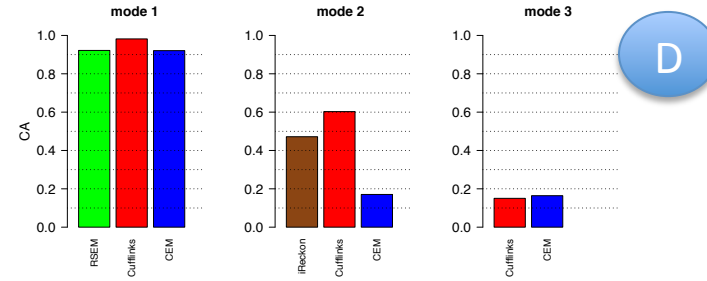
Recall (50 read length)



Precision (50 read length)



Recall (50 read length)



Alignment data driven