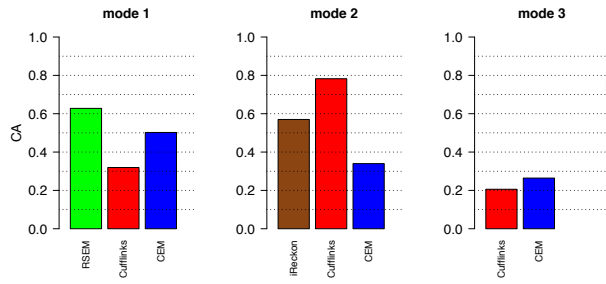


# PE 75 bp (Set-up 2)

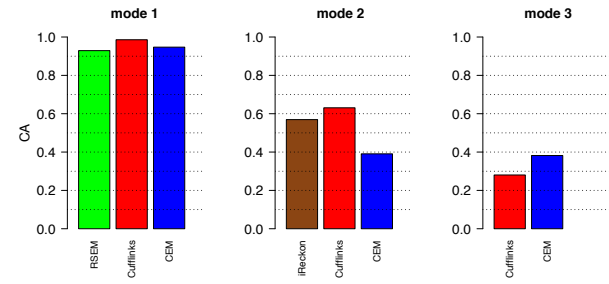
Alignment with transcriptome

Precision (75 read length)

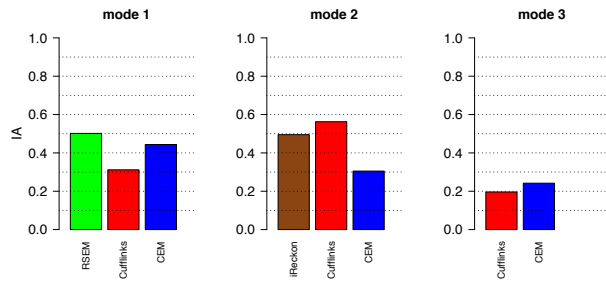


A

Recall (75 read length)

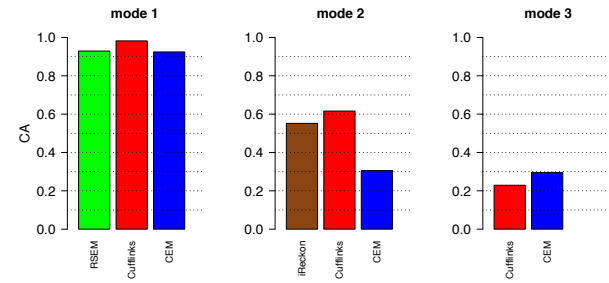


B



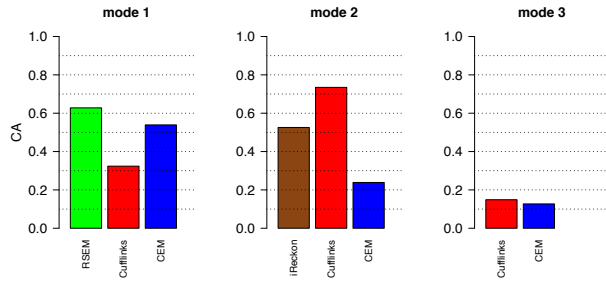
C

Recall (75 read length)



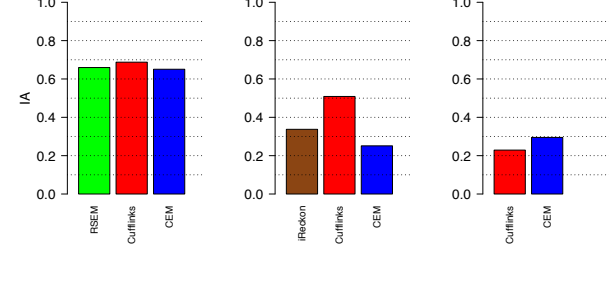
D

Precision (75 read length)



E

Recall (75 read length)



Alignment data driven