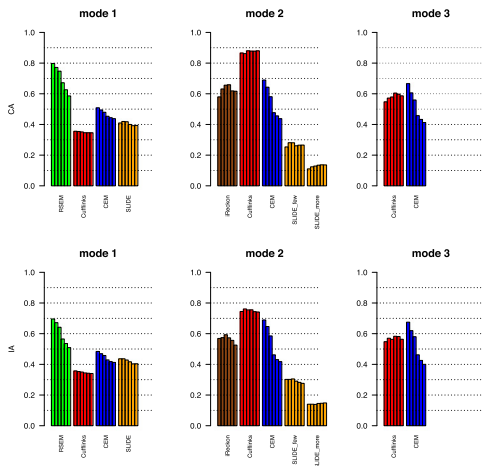


PE 75 bp (Set-1)

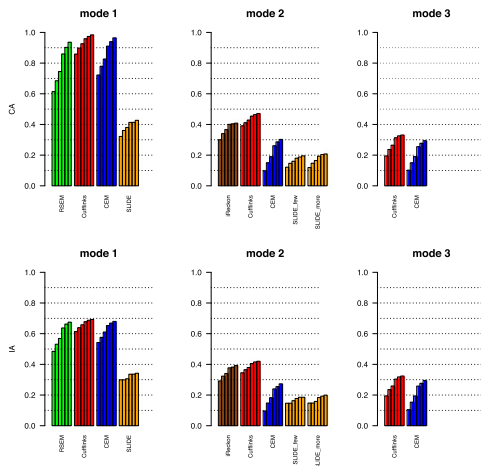
Alignment with transcriptome

Precision (75 read length)



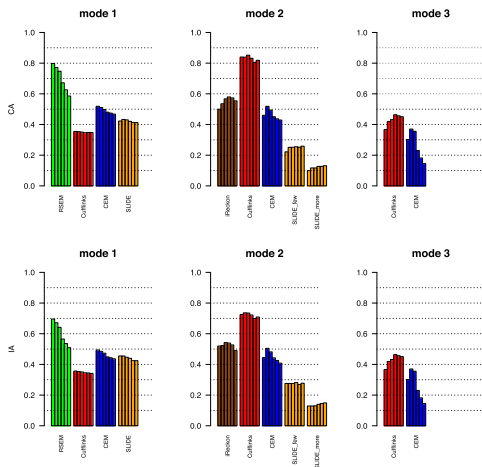
A

Recall (75 read length)



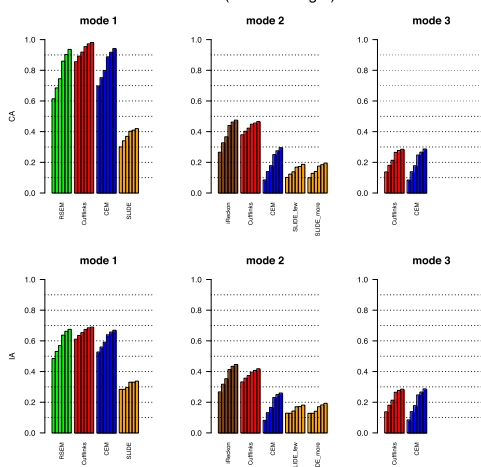
B

Precision (75 read length)



C

Recall (75 read length)



D

Alignment data driven