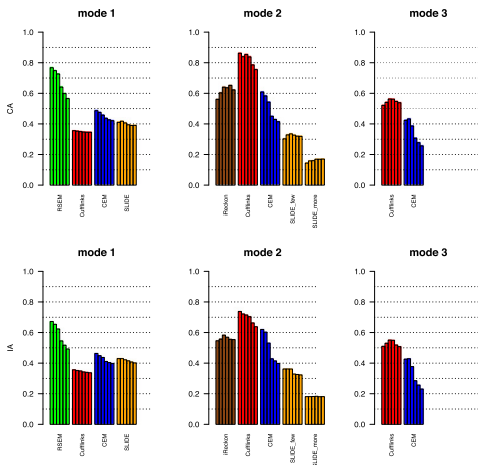


PE 50 bp (Set-up 1)

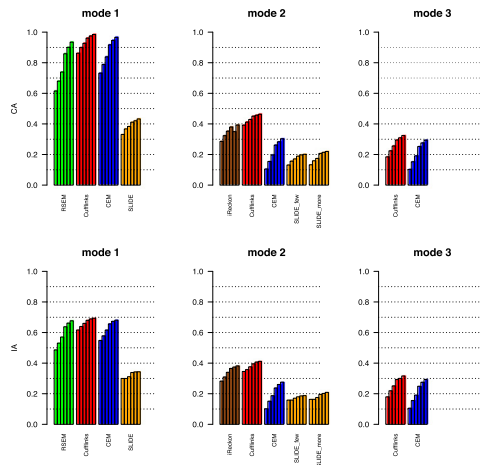
Alignment with transcriptome

Precision (50 read length)



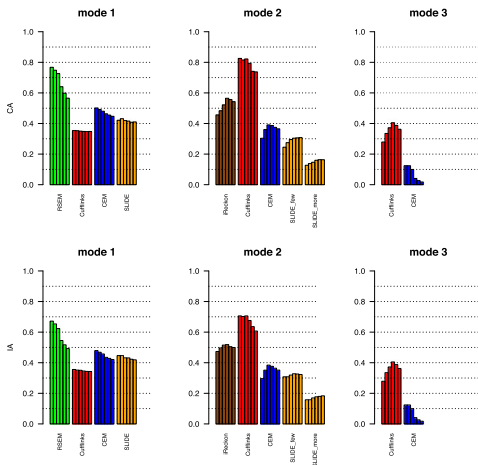
A

Recall (50 read length)



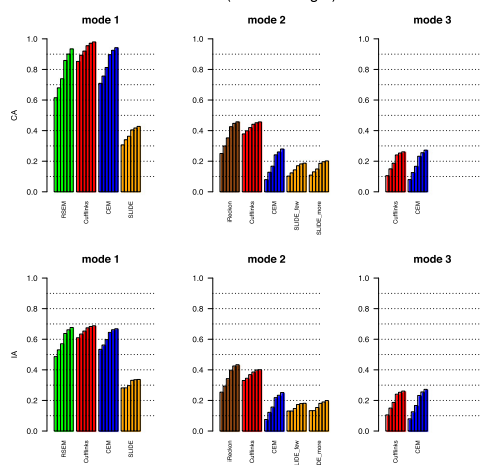
B

Precision (50 read length)



C

Recall (50 read length)



D

Alignment data driven