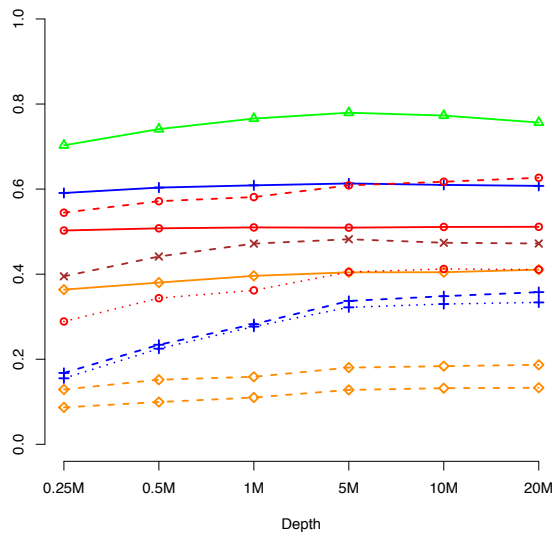


# PE 100 bp (Set-up 1)

Alignment with transcriptome

CA

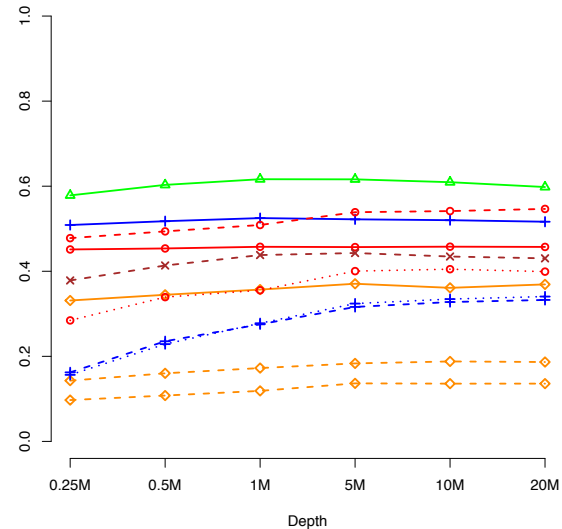
F-measure (100 read length)



A

IA

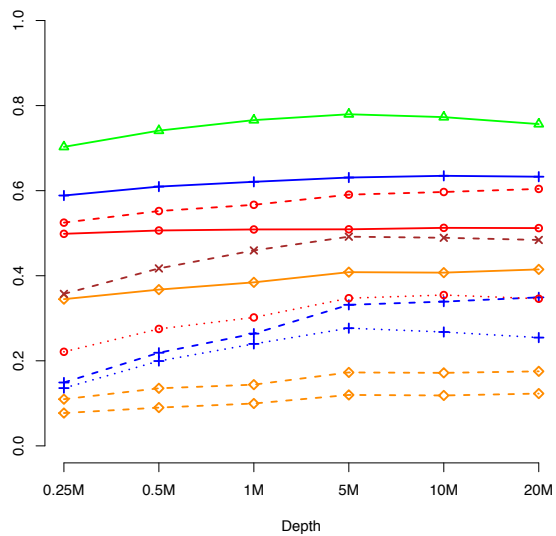
F-measure (100 read length)



B

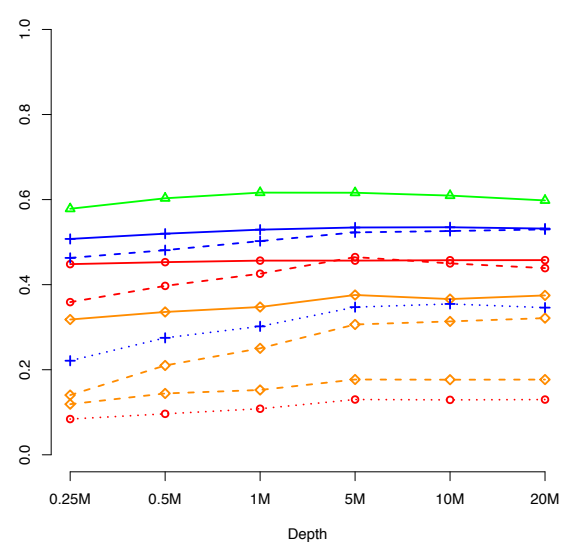
Alignment data driven

F-measure (100 read length)



C

F-measure (100 read length)



D