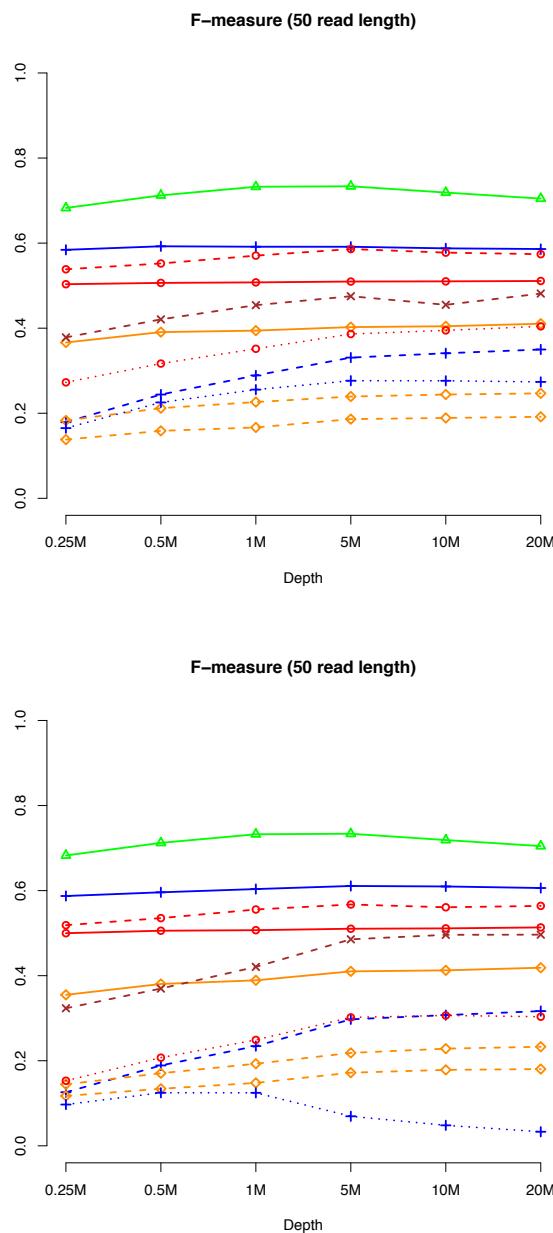


## Alignment with transcriptome

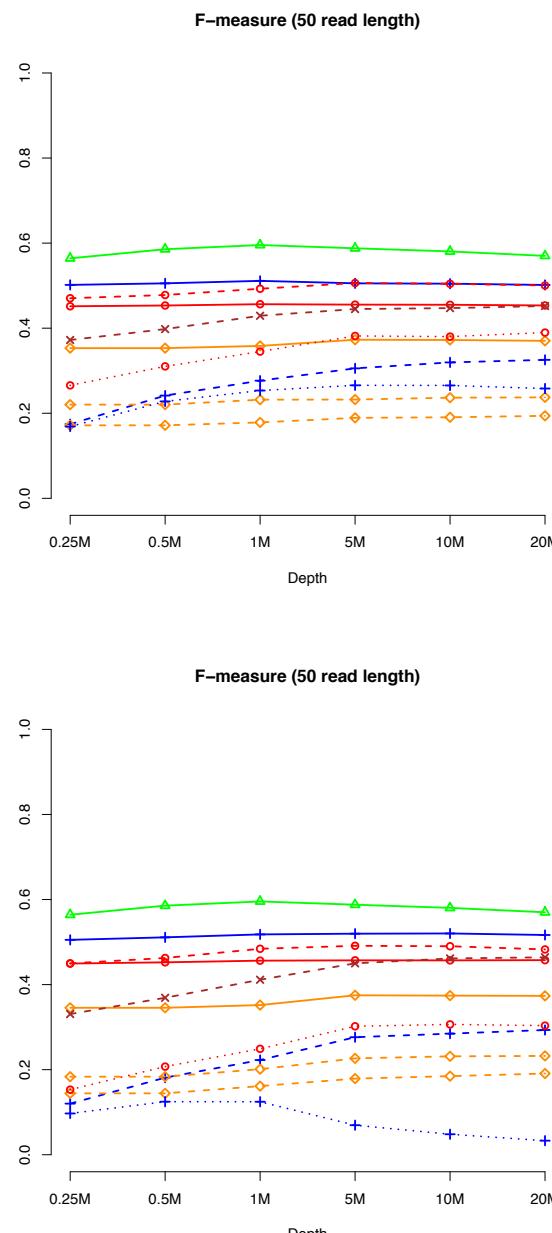
### PE 50 bp (Set-up 1)

CA



A

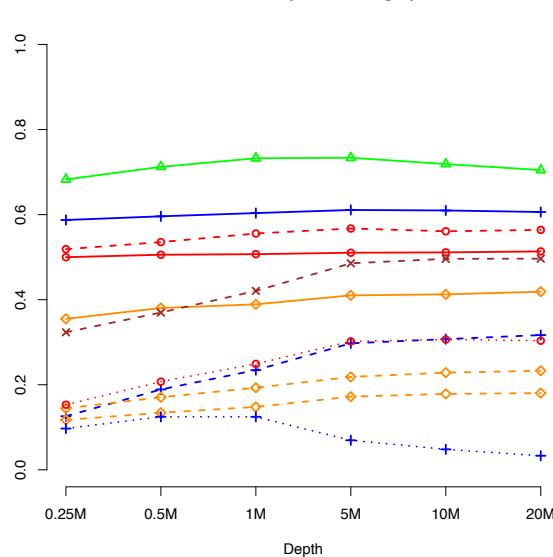
IA



B

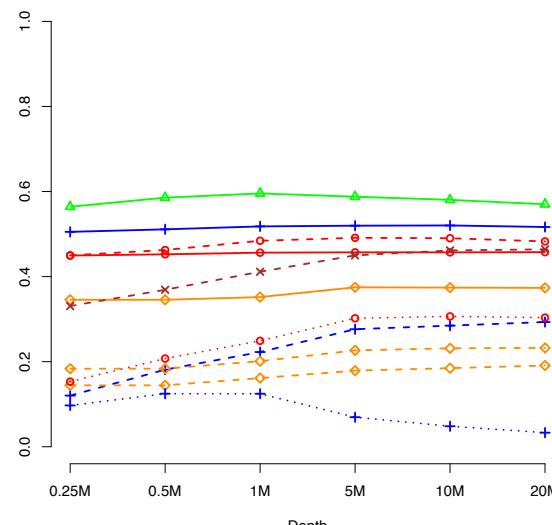
## Alignment data driven

### F-measure (50 read length)



C

### F-measure (50 read length)



D