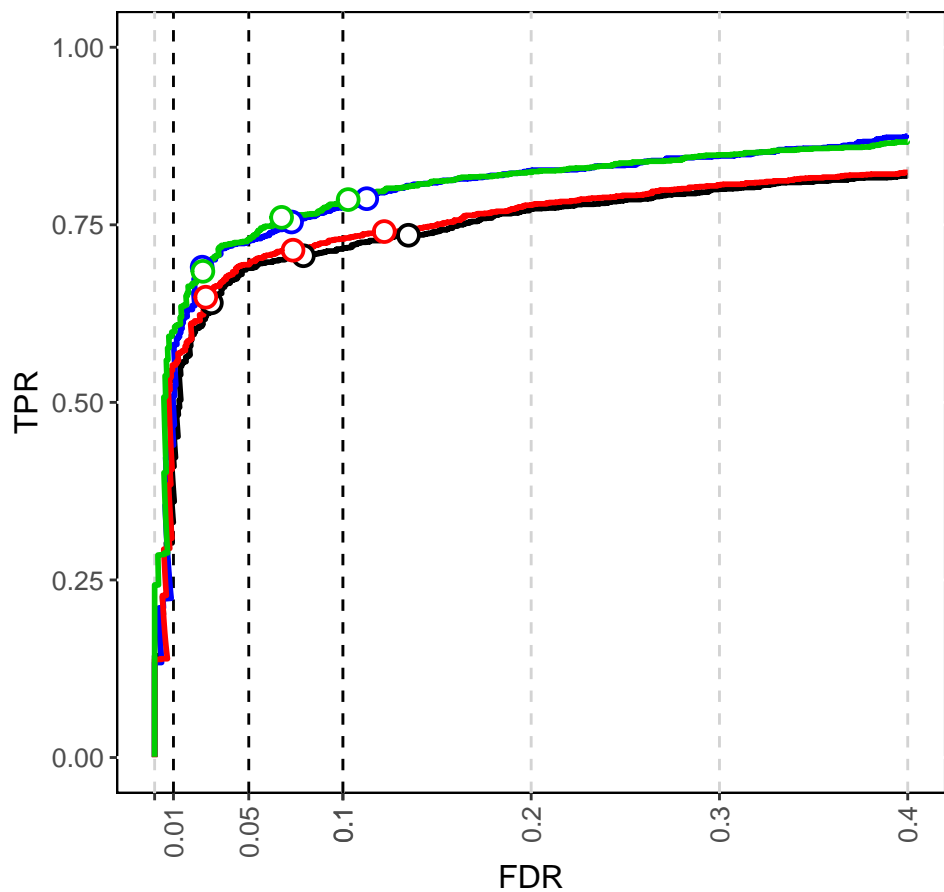
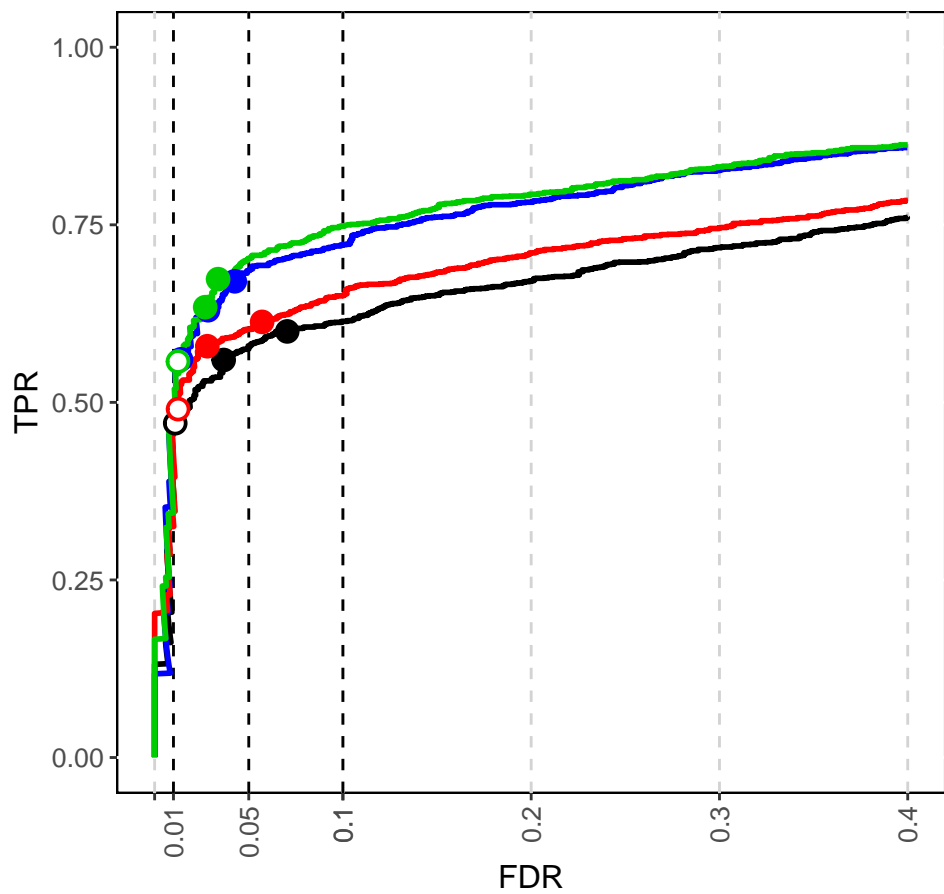


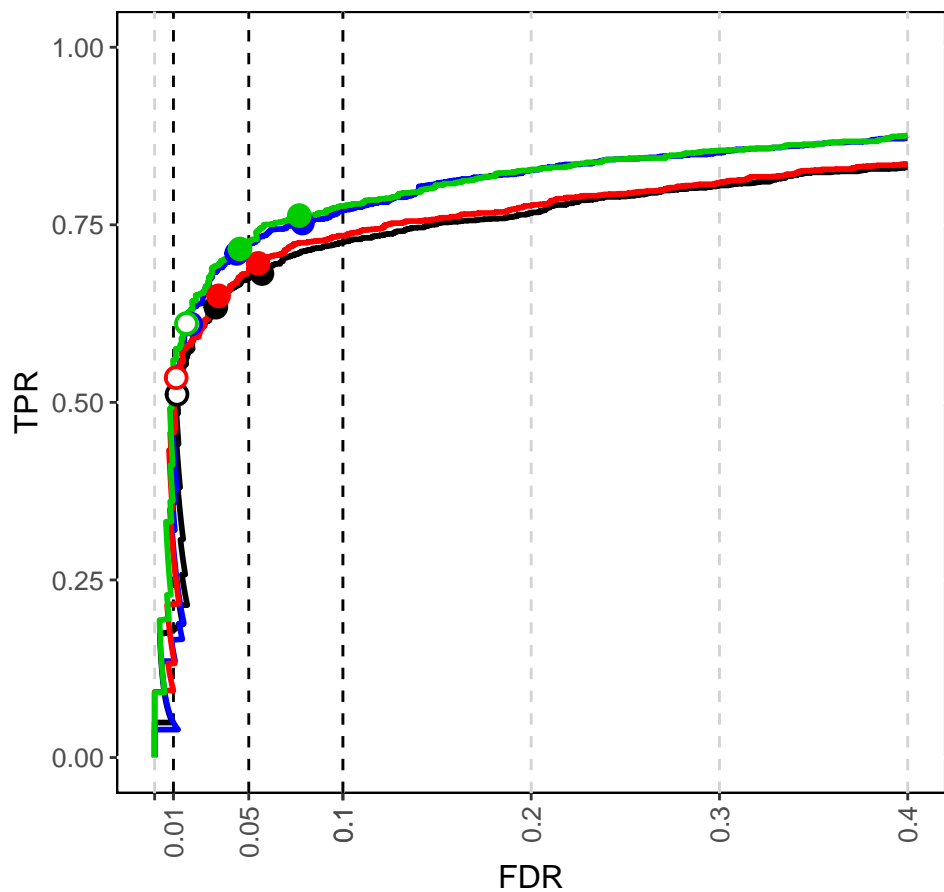
Dmelanogaster: 5 samples (DoubleExpSeq)



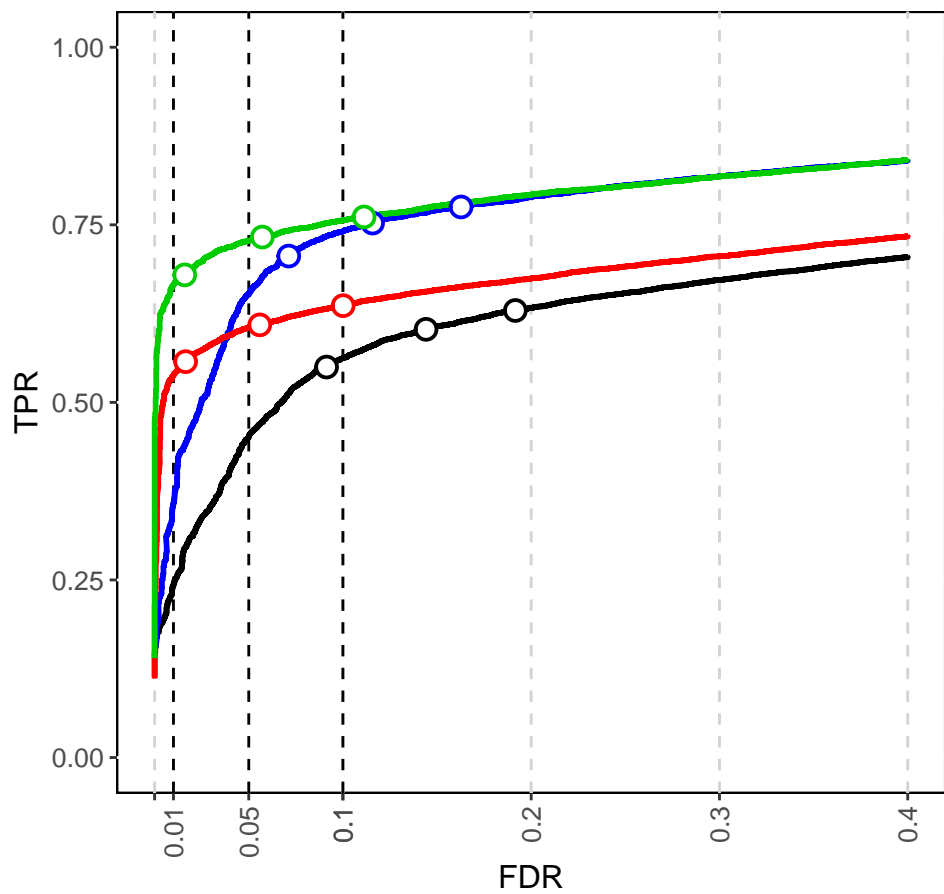
Dmelanogaster: 5 samples (limma_diffsplice)



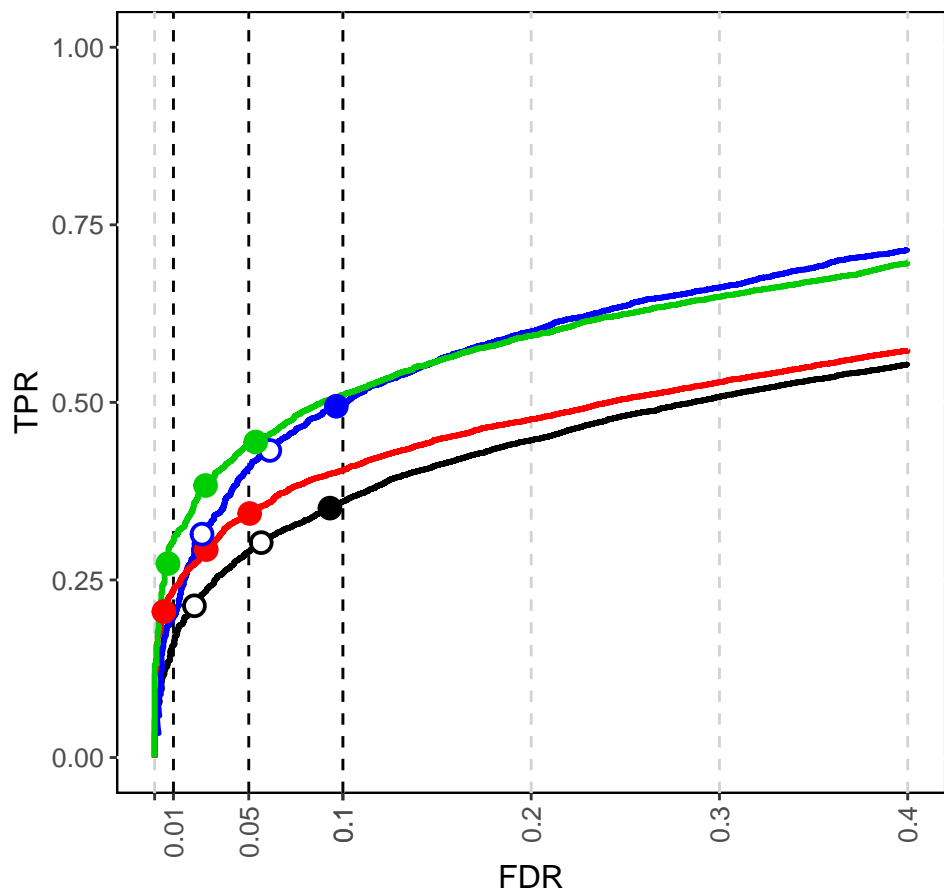
Dmelanogaster: 5 samples (satuRn)



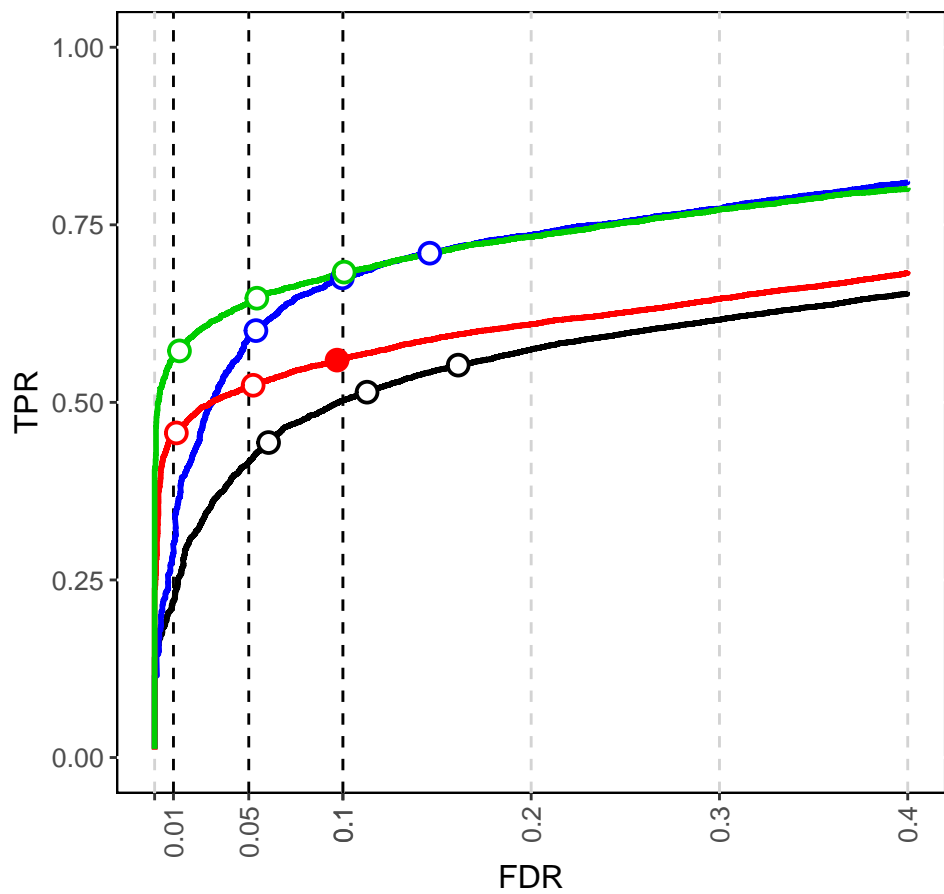
Love: 10 samples (DoubleExpSeq)



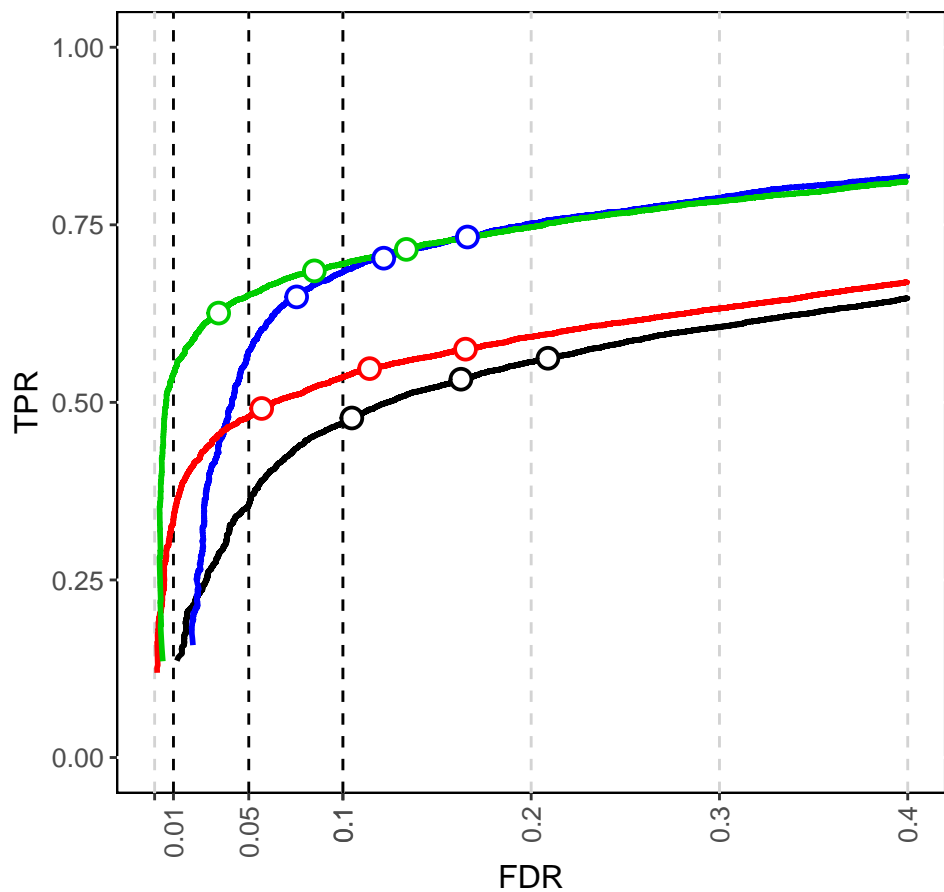
Love: 3 samples (DoubleExpSeq)



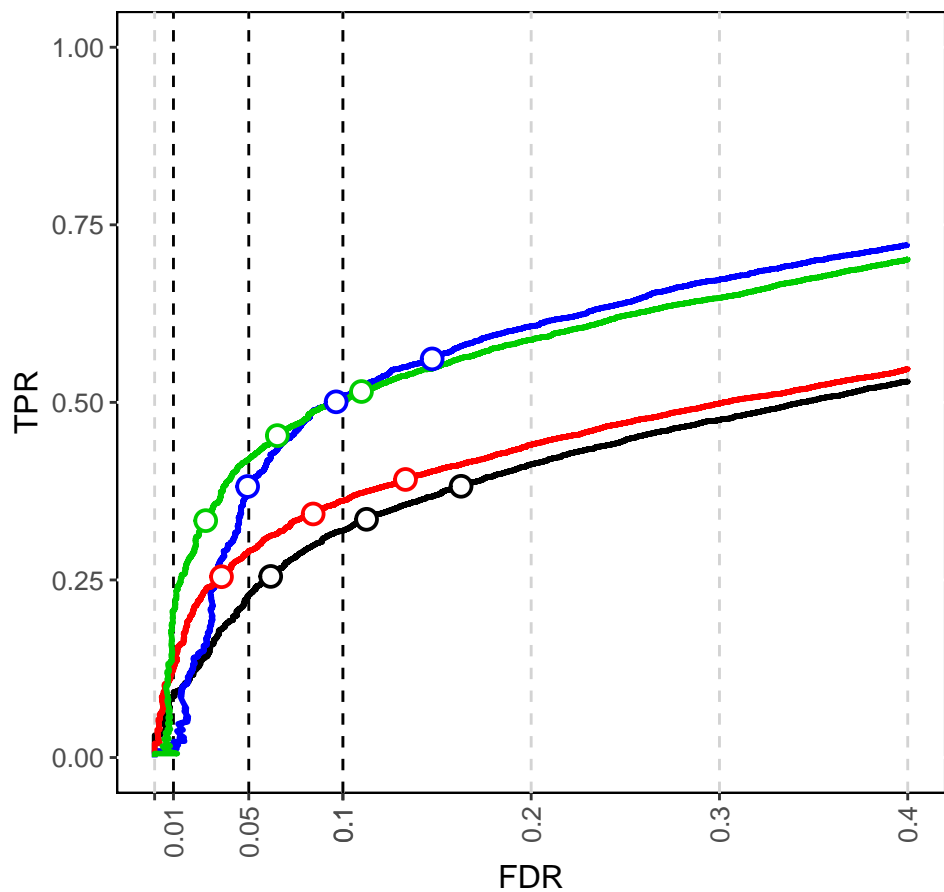
Love: 6 samples (DoubleExpSeq)



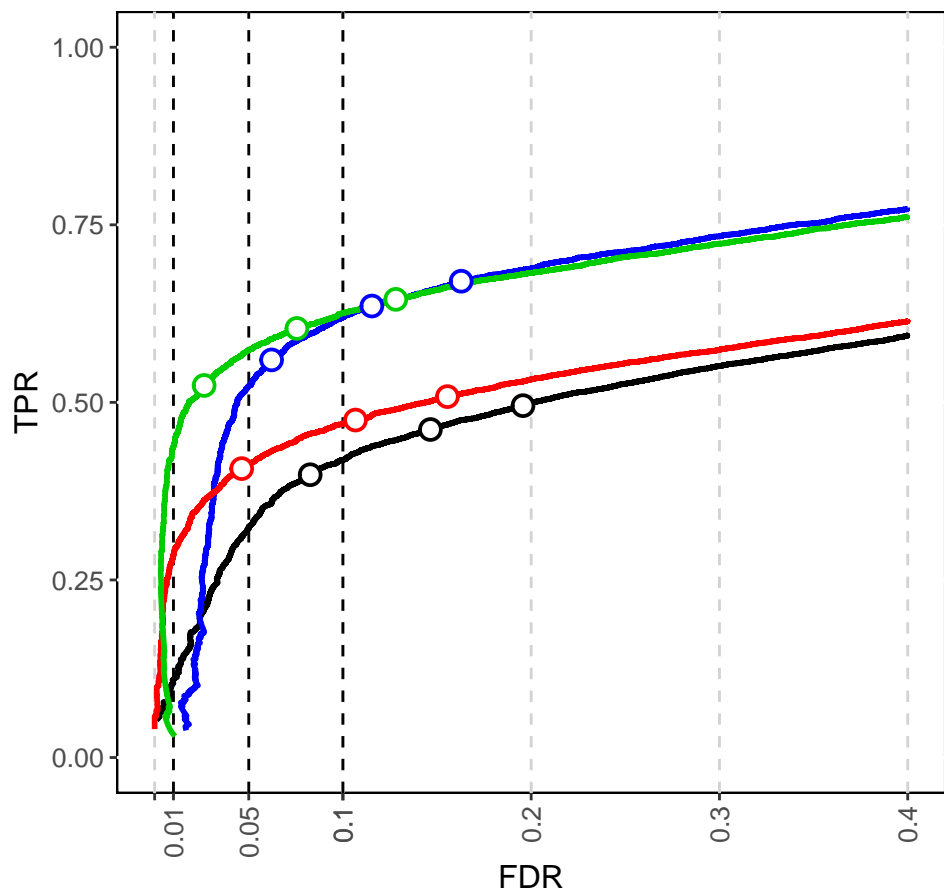
Love: 10 samples (limma_diffsplice)



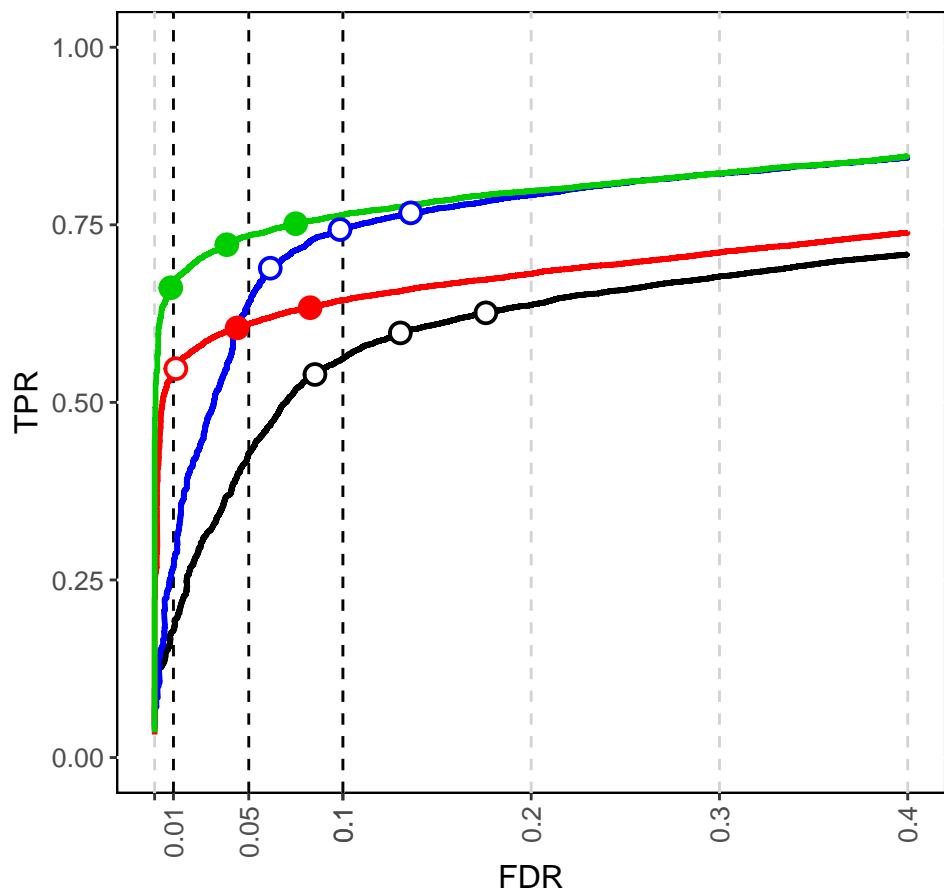
Love: 3 samples (limma_diffsplice)



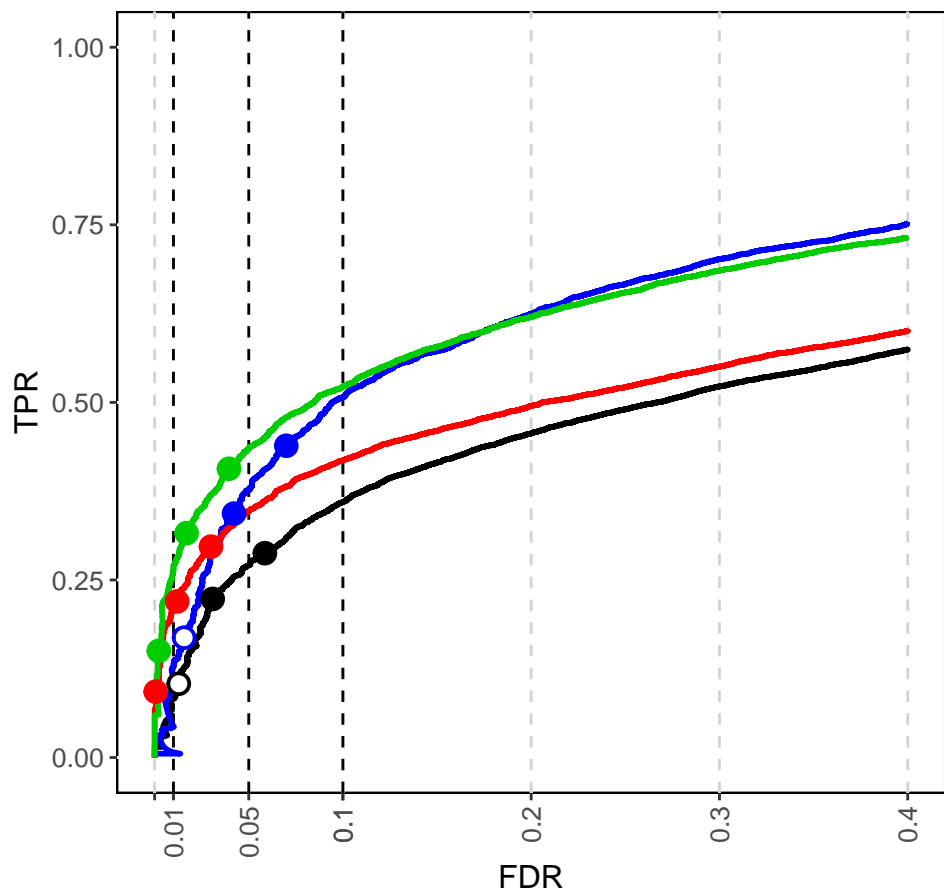
Love: 6 samples (limma_diffsplice)



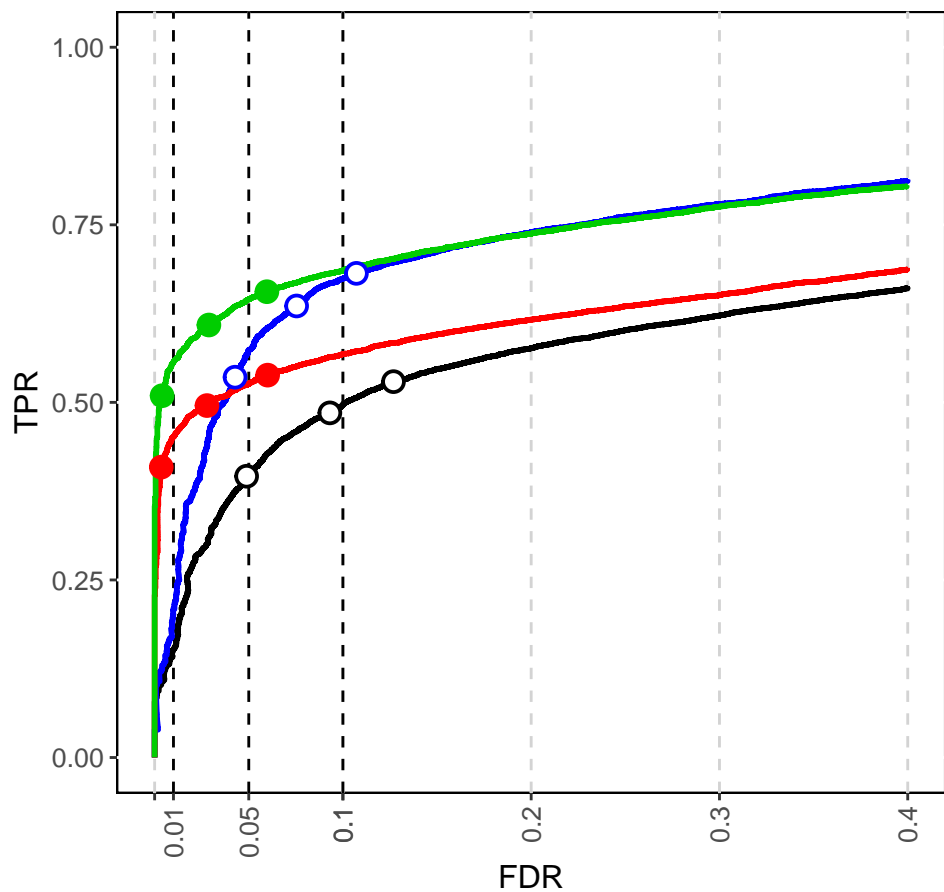
Love: 10 samples (satuRn)



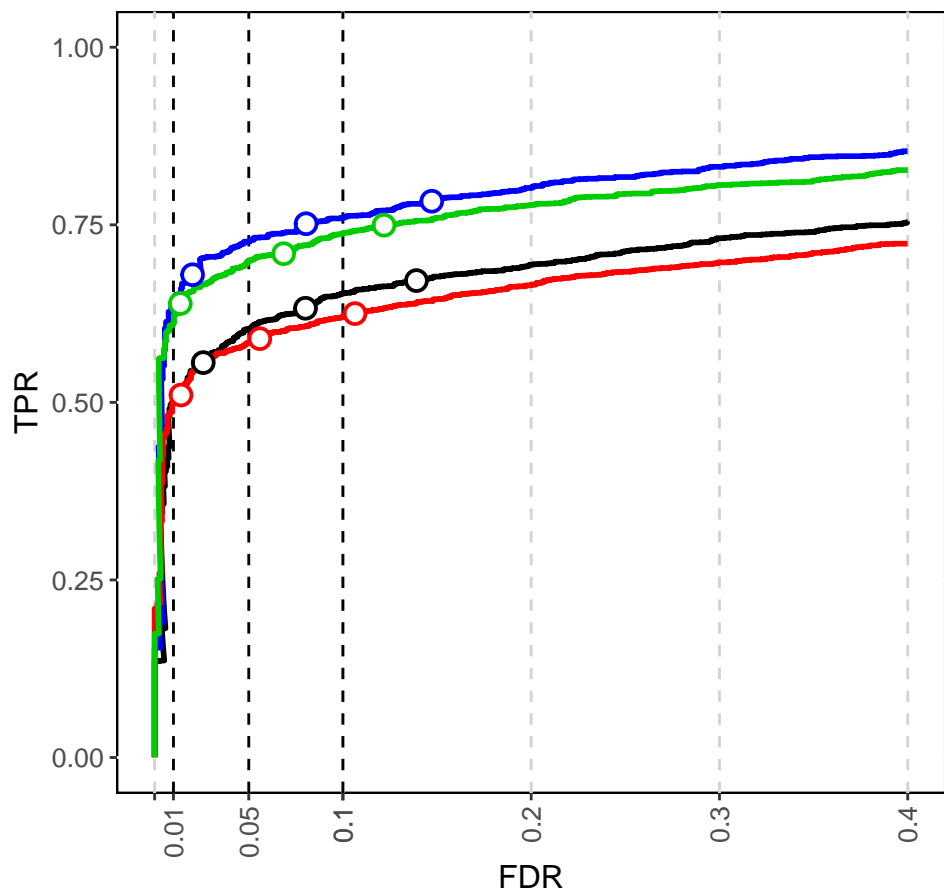
Love: 3 samples (satuRn)



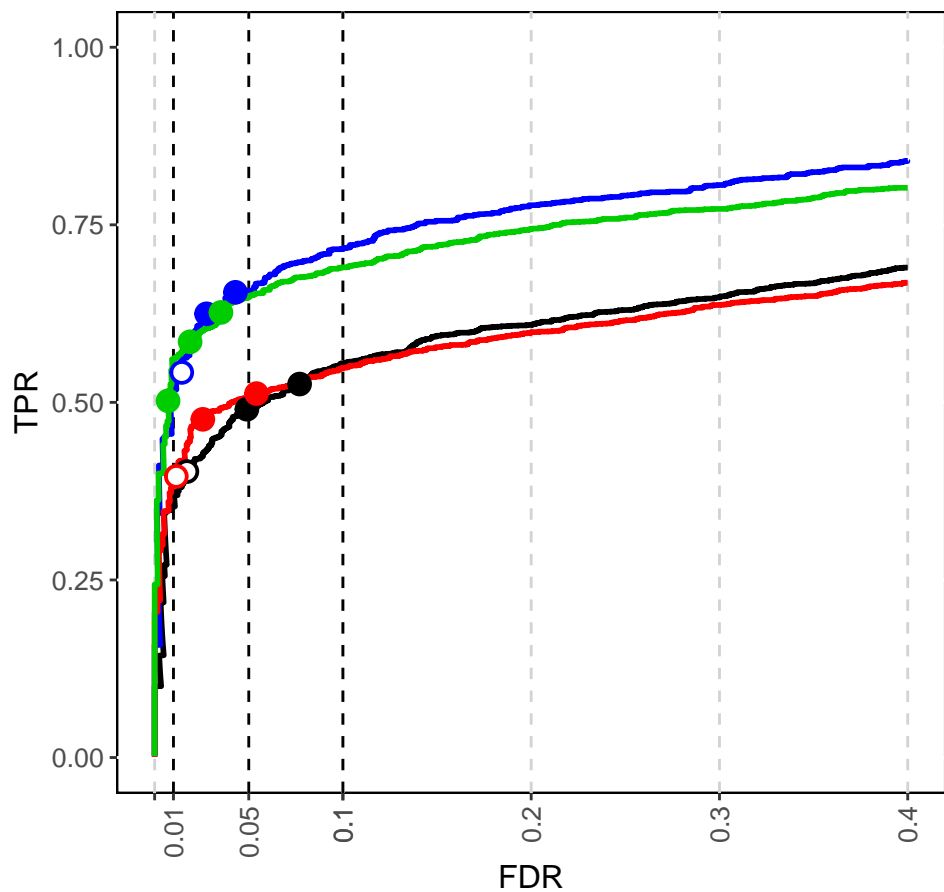
Love: 6 samples (satuRn)



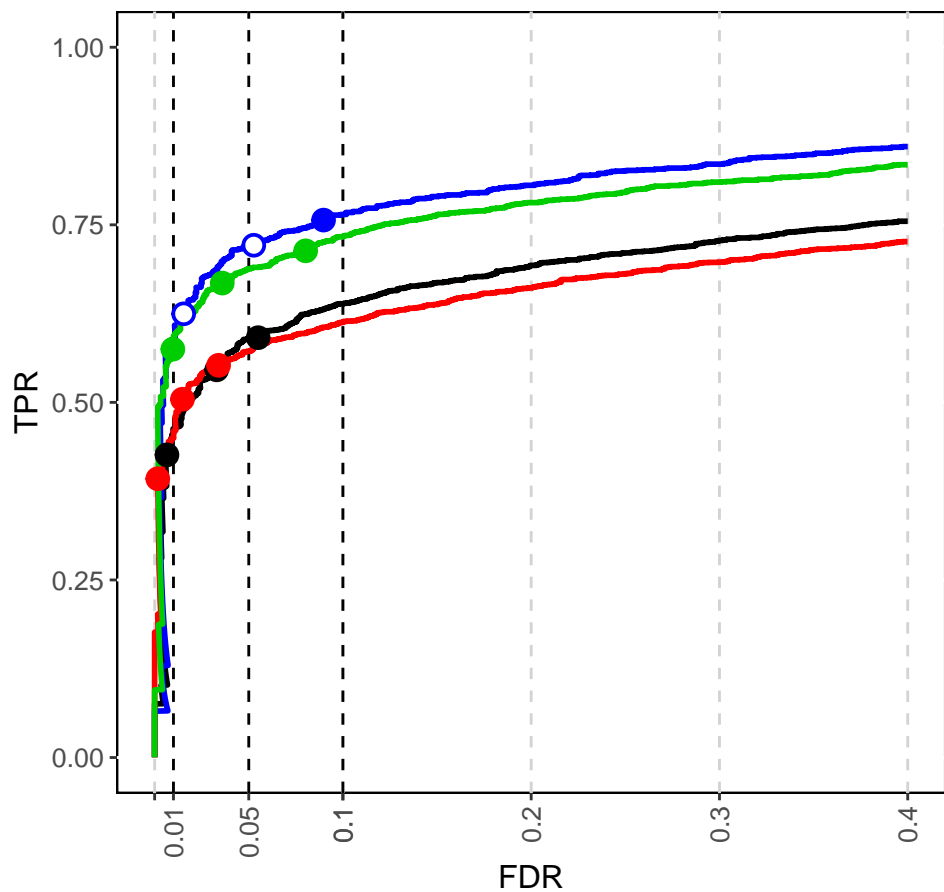
Hsapiens: 5 samples (DoubleExpSeq)



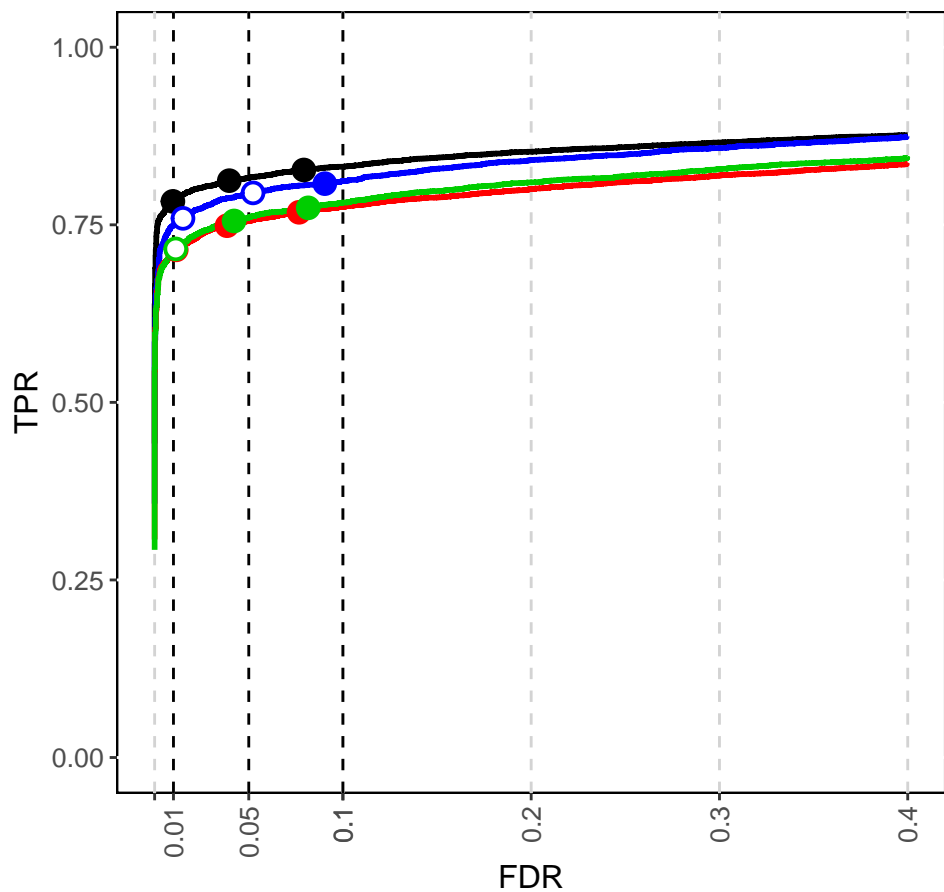
Hsapiens: 5 samples (limma_diffsplice)



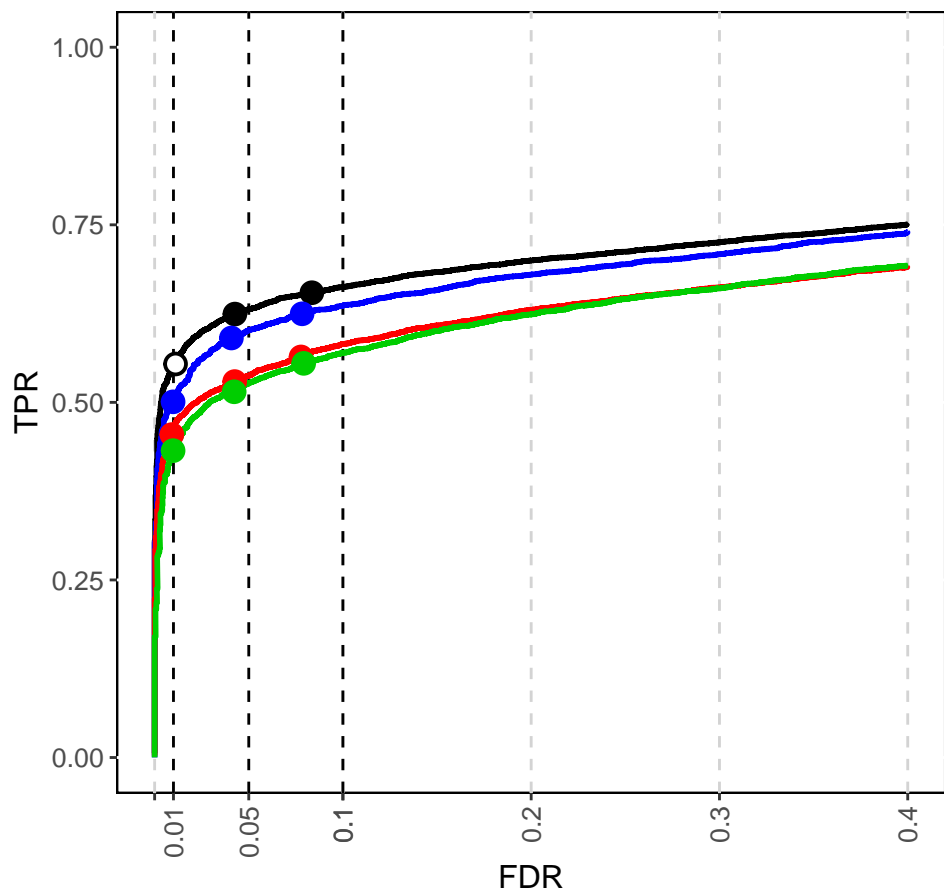
Hsapiens: 5 samples (satuRn)



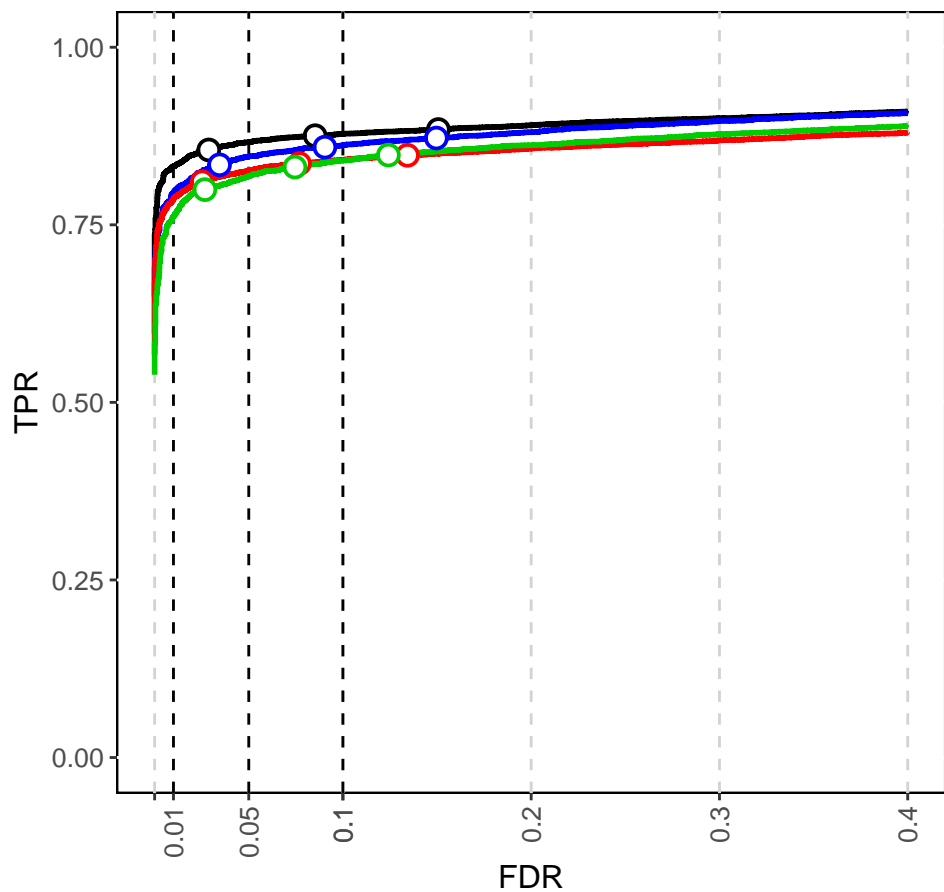
Gtex: 20 samples (DoubleExpSeq)



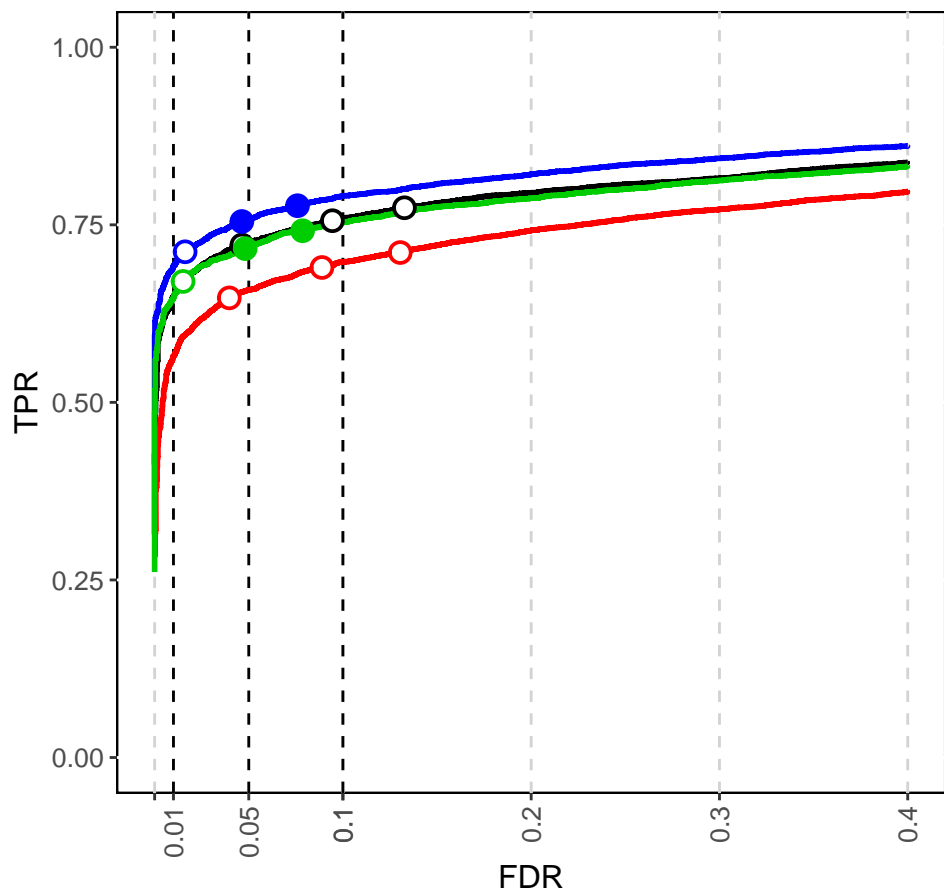
Gtex: 5 samples (DoubleExpSeq)



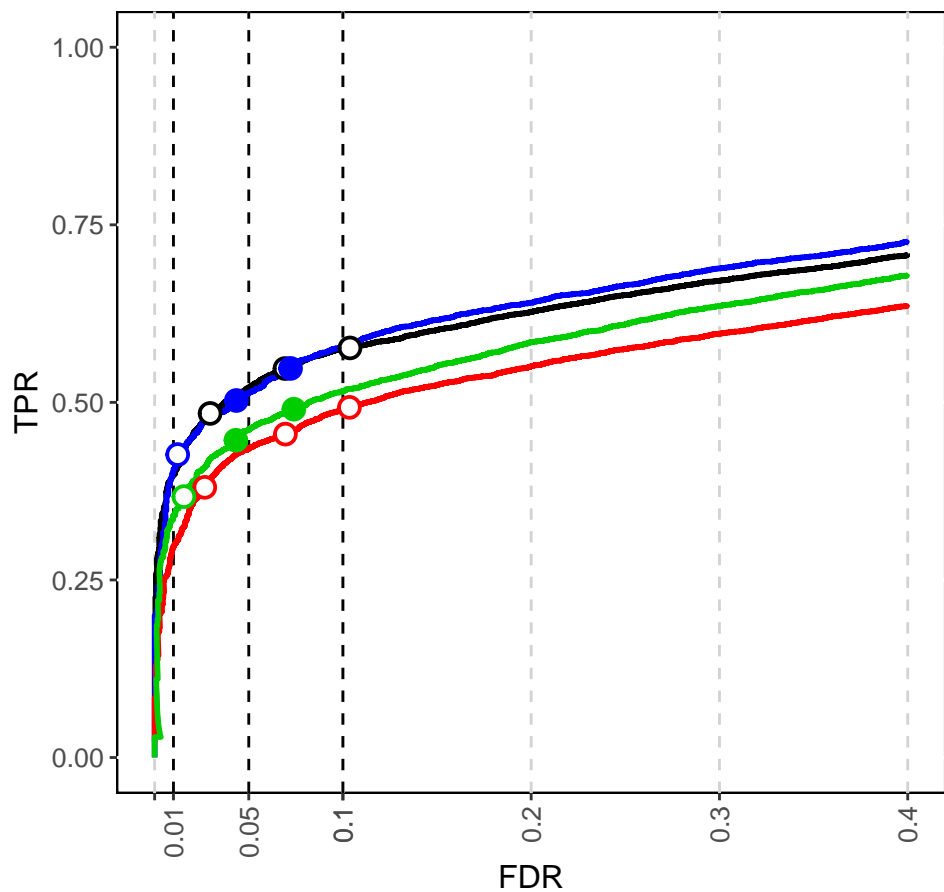
Gtex: 50 samples (DoubleExpSeq)



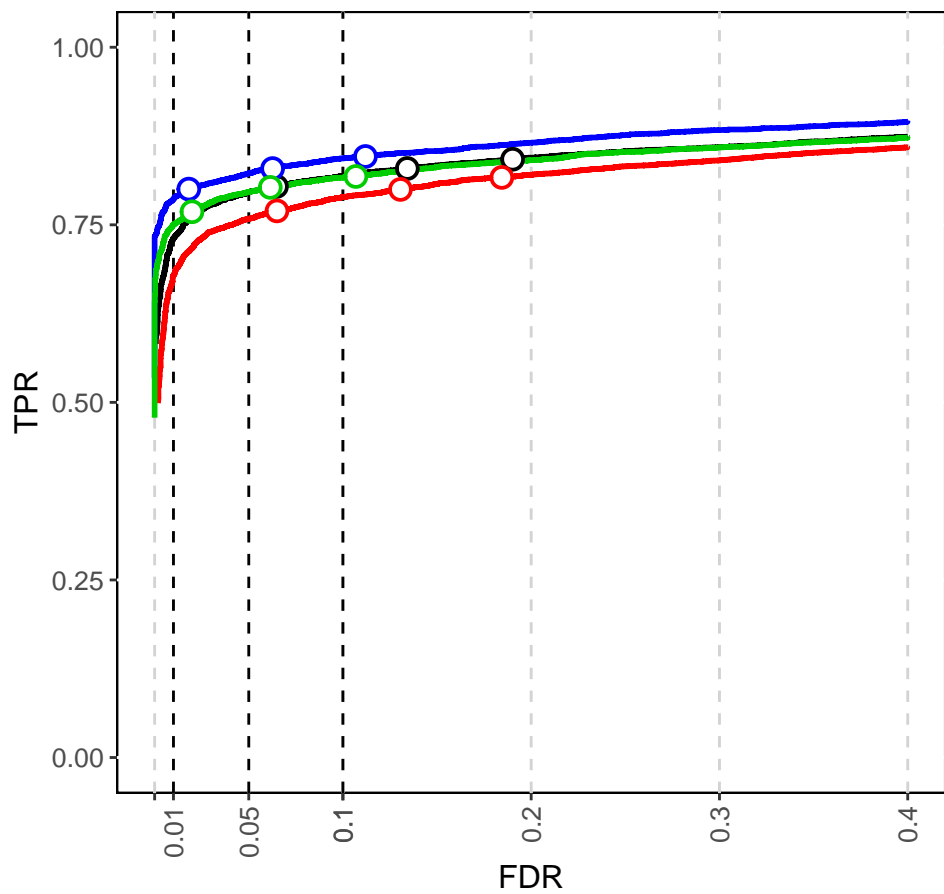
Gtex: 20 samples (limma_diffsplice)



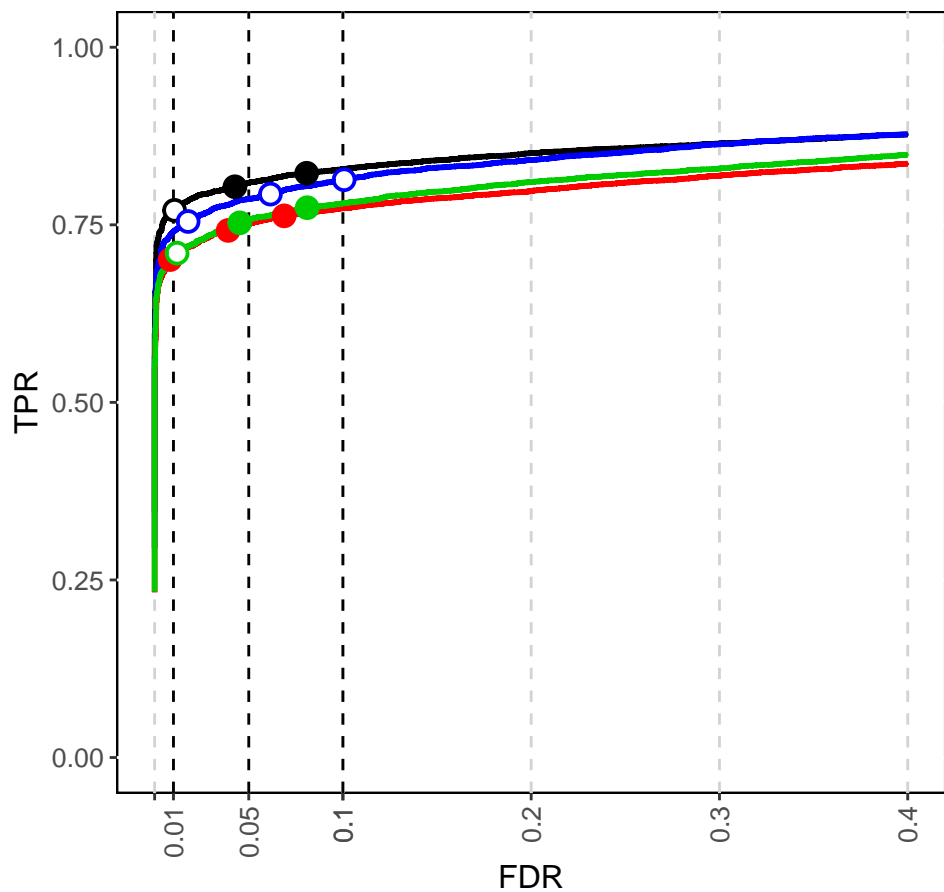
Gtex: 5 samples (limma_diffsplice)



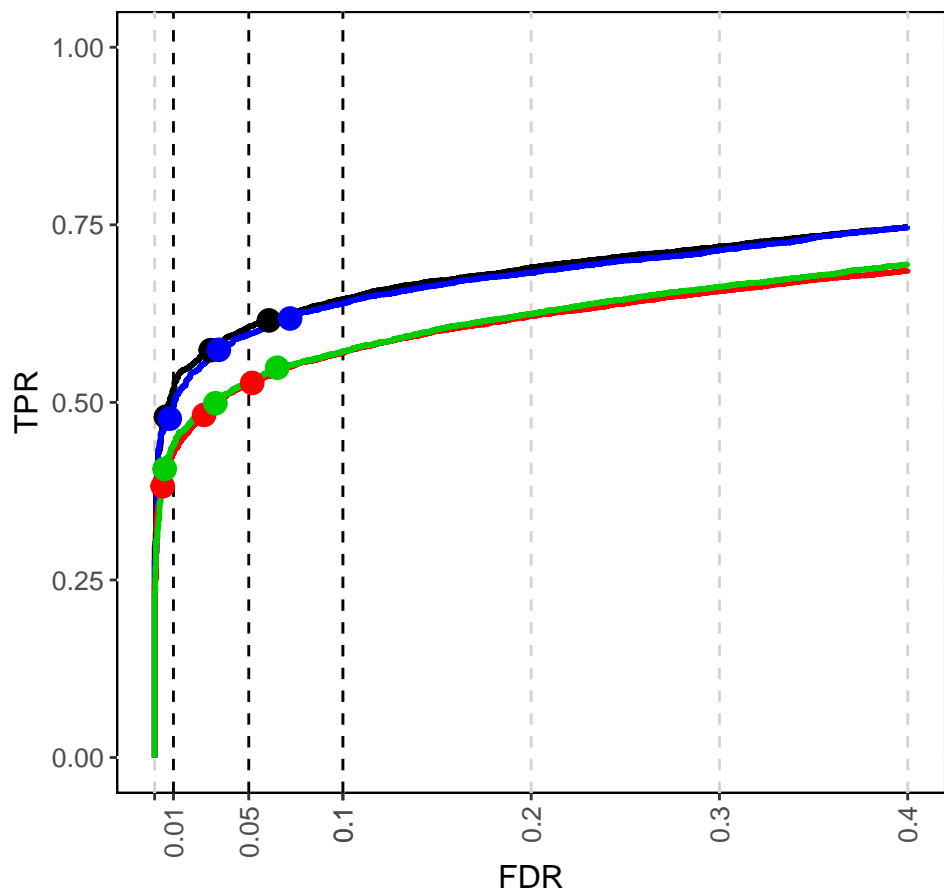
Gtex: 50 samples (limma_diffsplice)



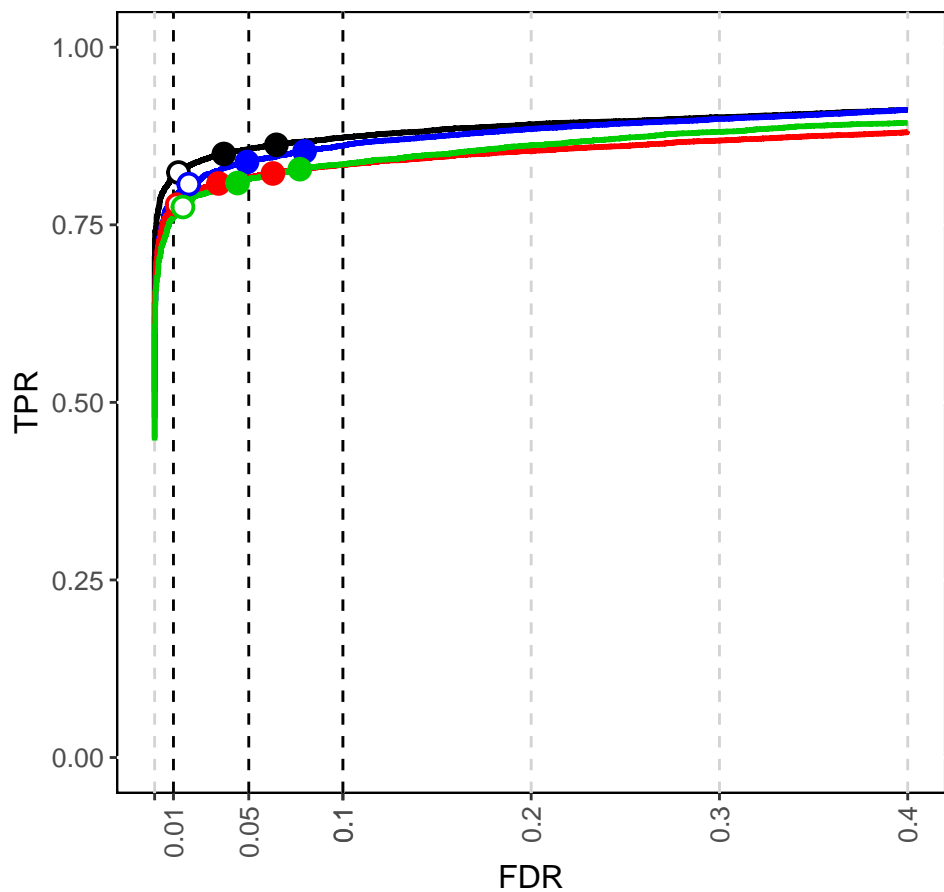
Gtex: 20 samples (saturn)



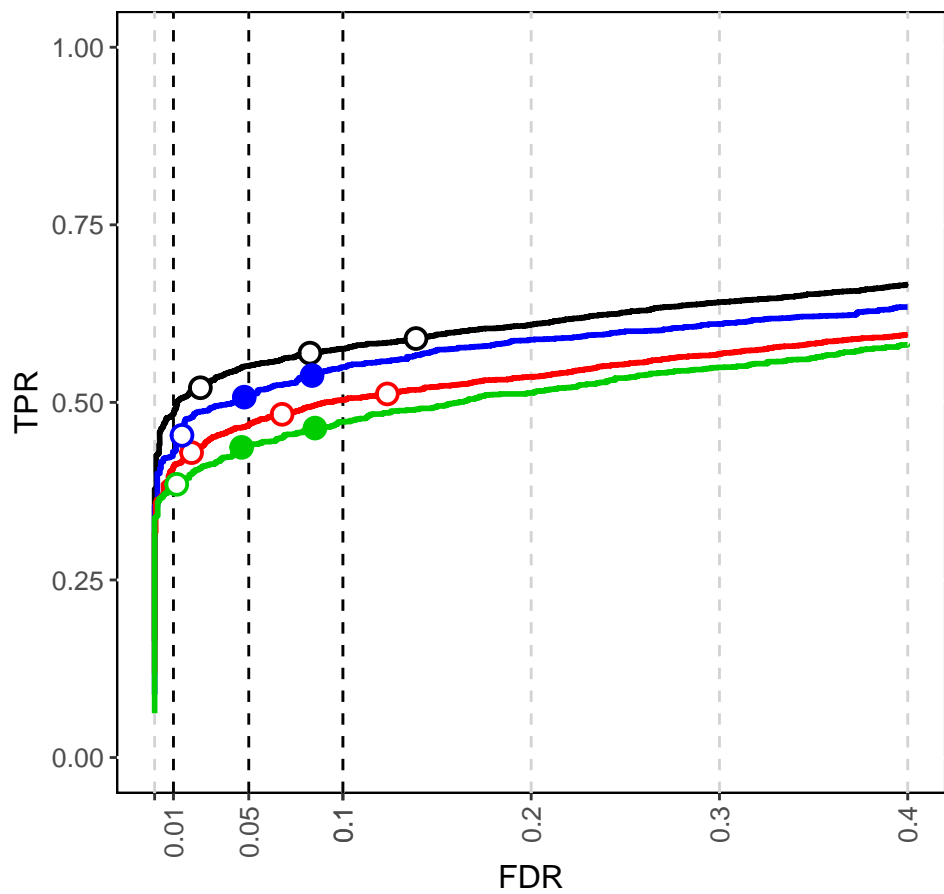
Gtex: 5 samples (satuRn)



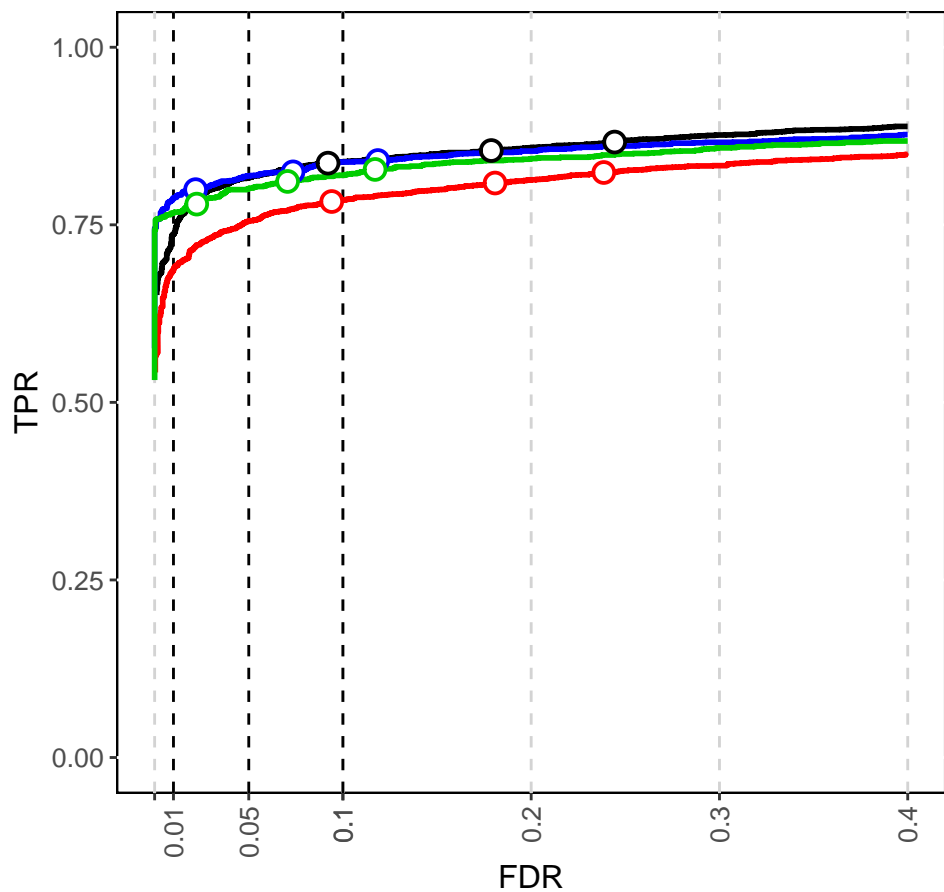
Gtex: 50 samples (saturn)



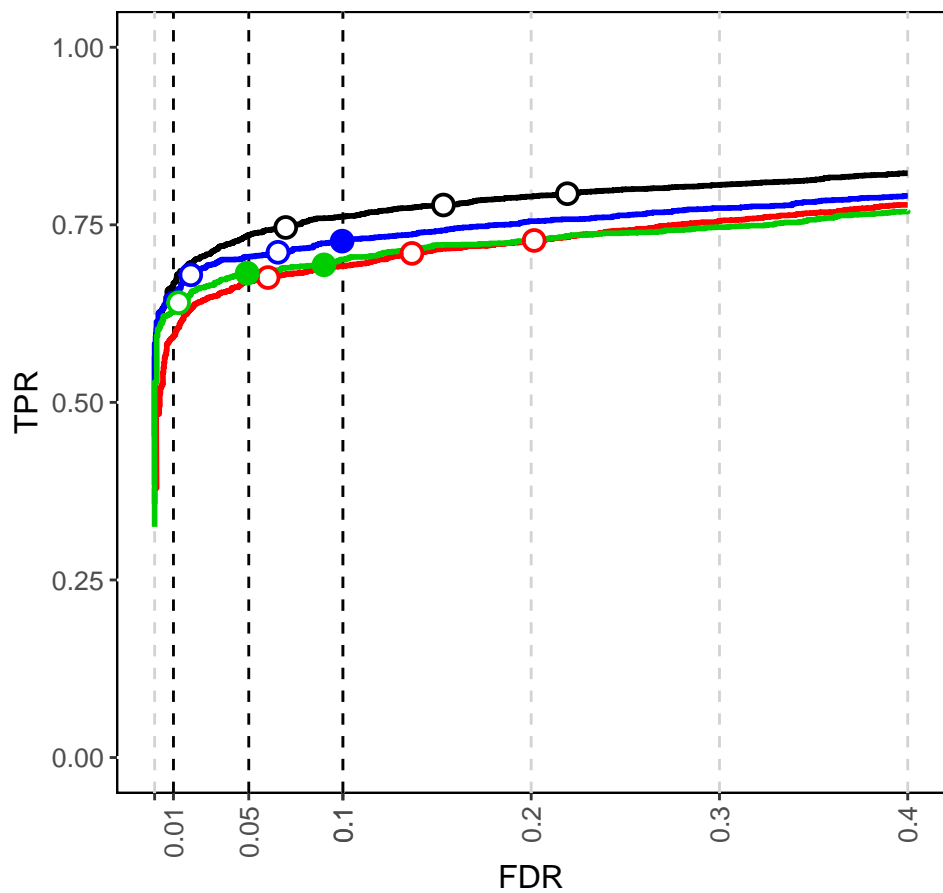
Tasic: 20 cells (DoubleExpSeq)



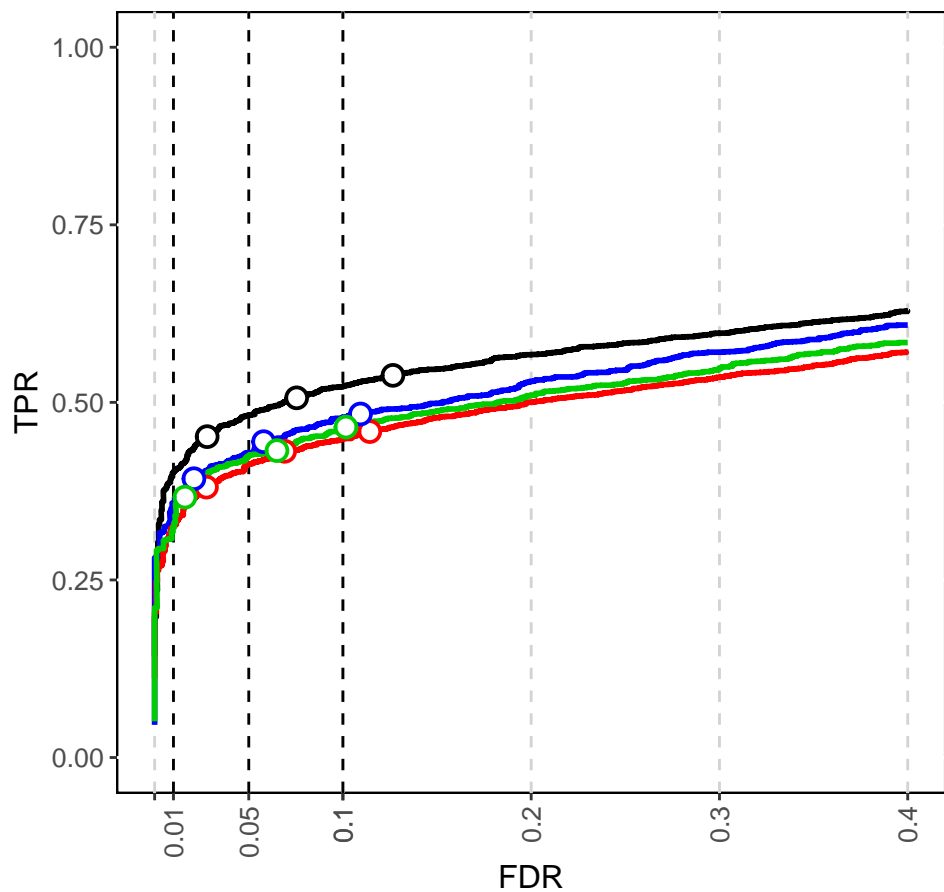
Tasic: 200 cells (DoubleExpSeq)



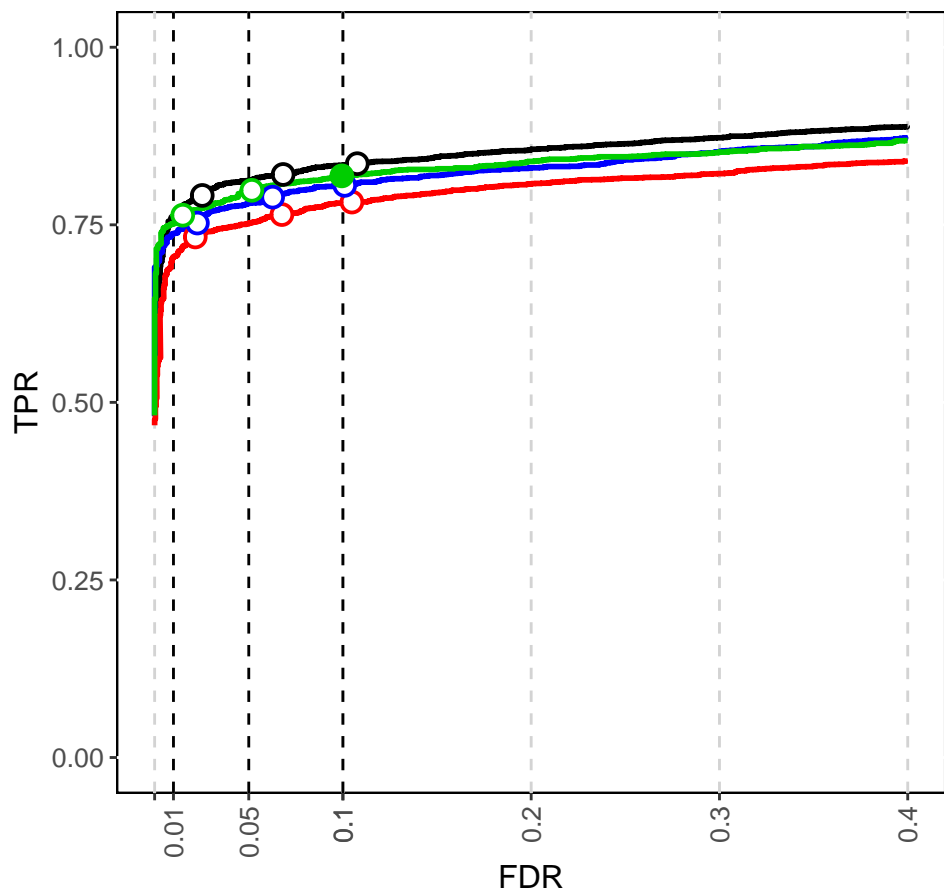
Tasic: 75 cells (DoubleExpSeq)



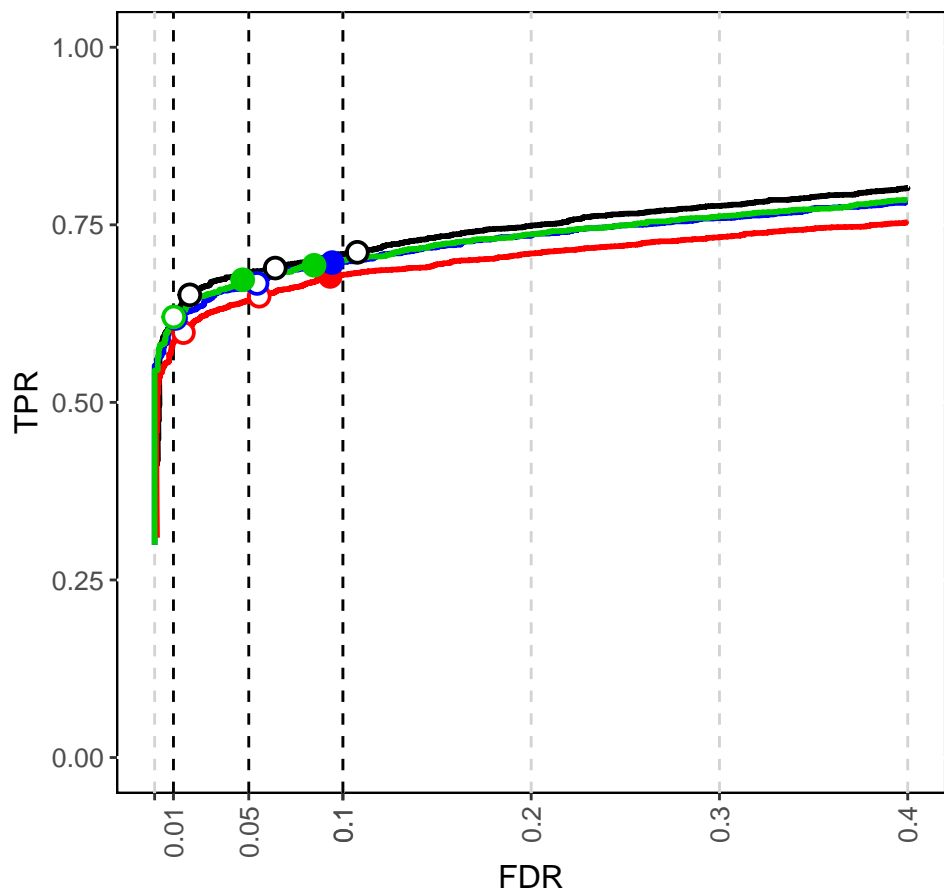
Tasic: 20 cells (limma_diffsplice)



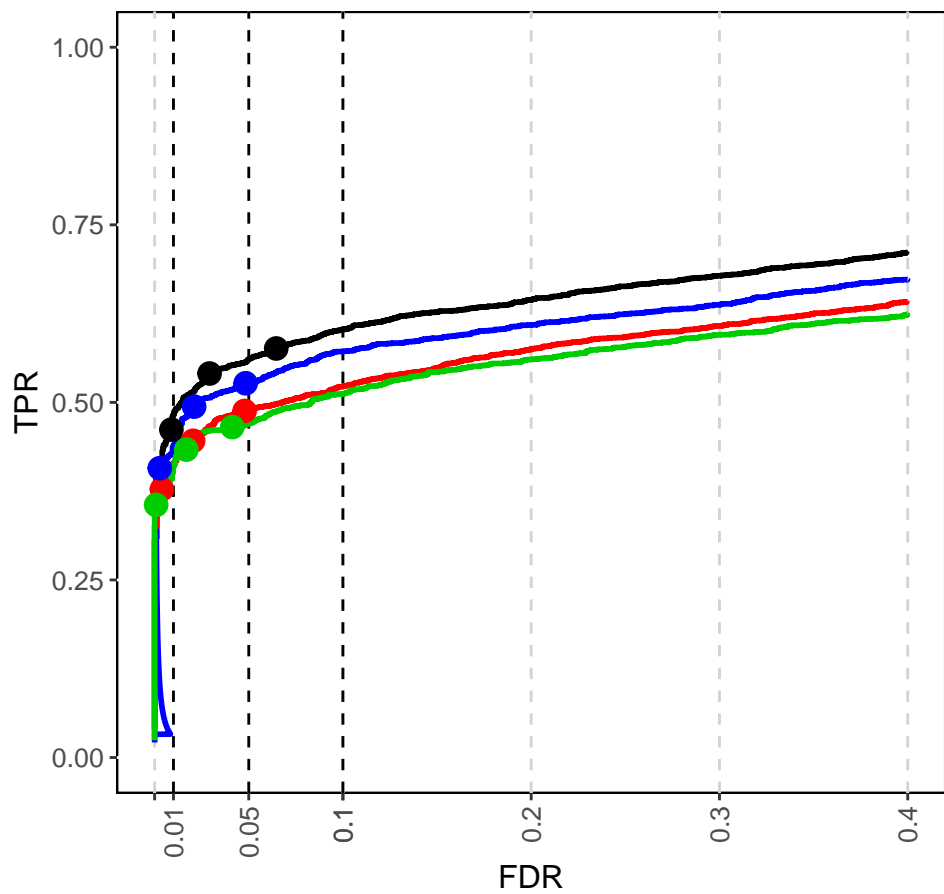
Tasic: 200 cells (limma_diffsplice)



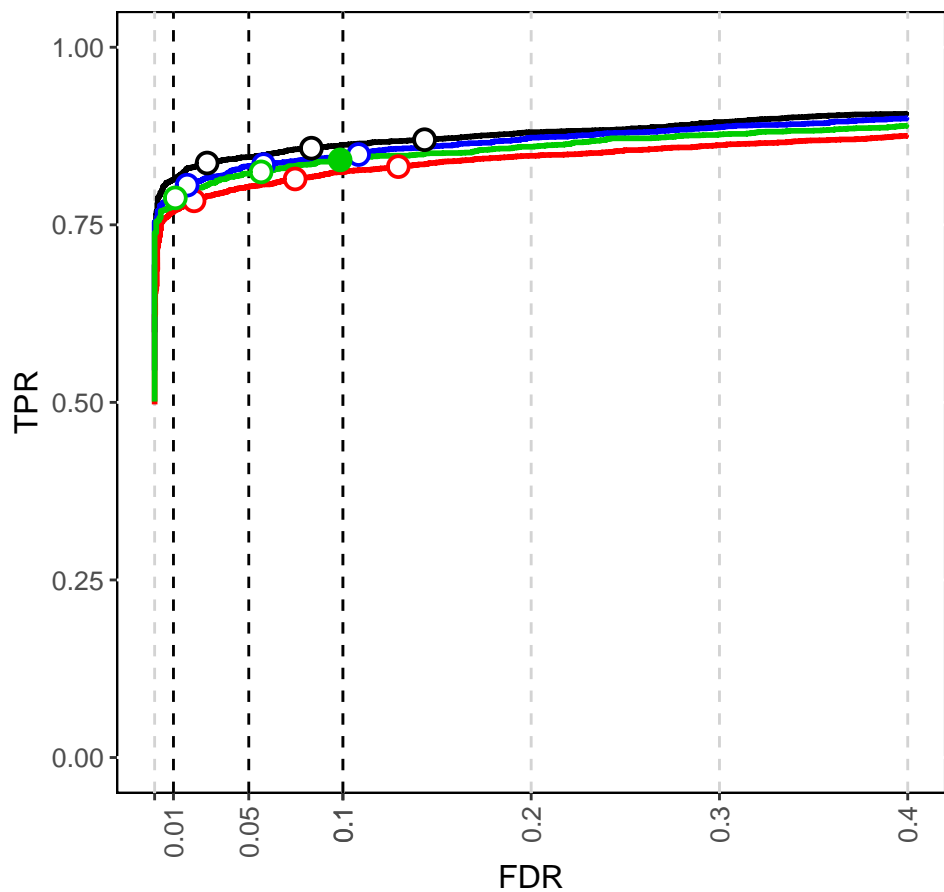
Tasic: 75 cells (limma_diffsplice)



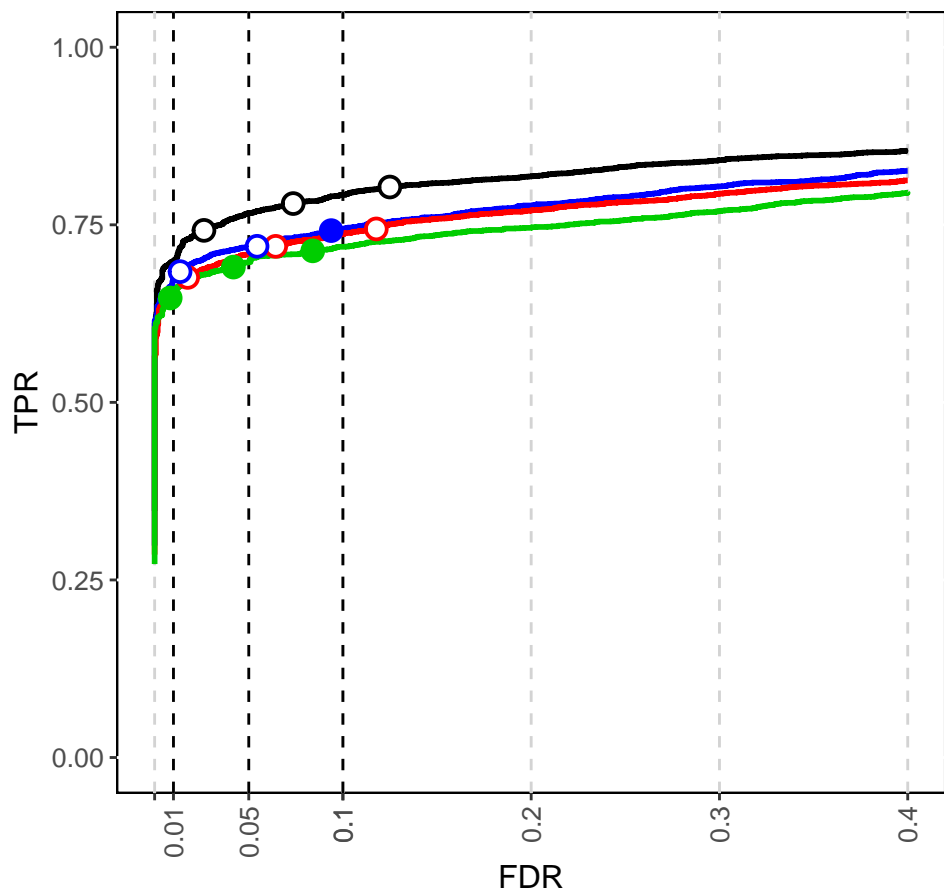
Tasic: 20 cells (satuRn)



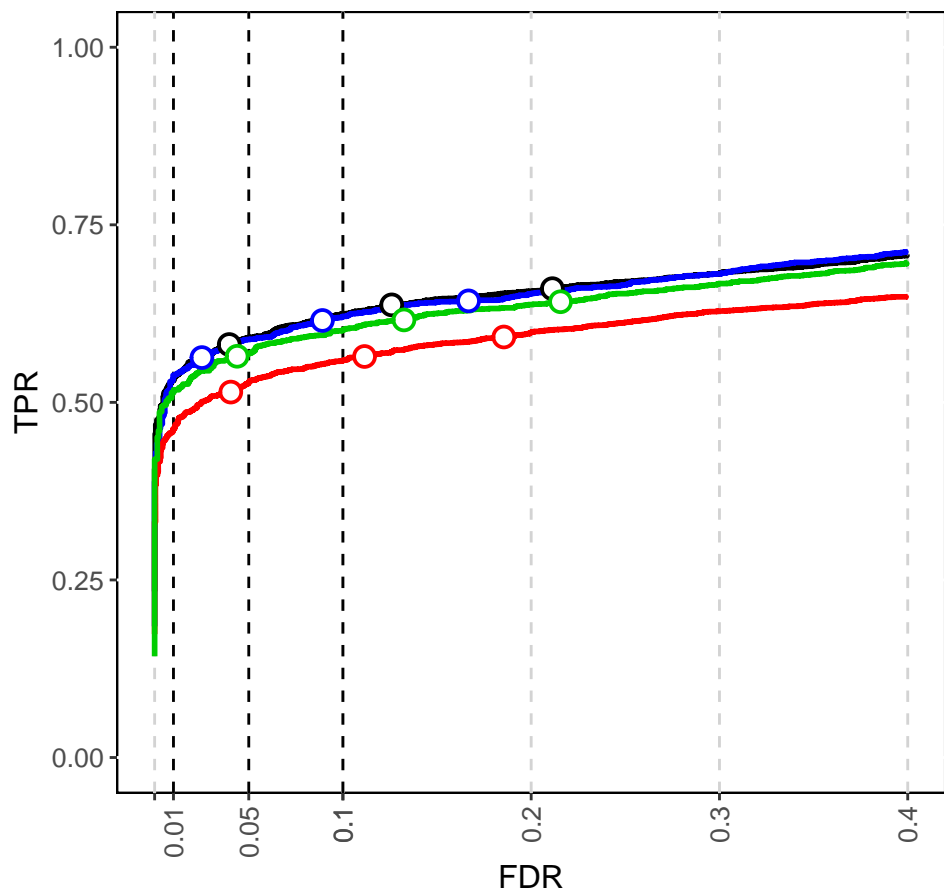
Tasic: 200 cells (satuRn)



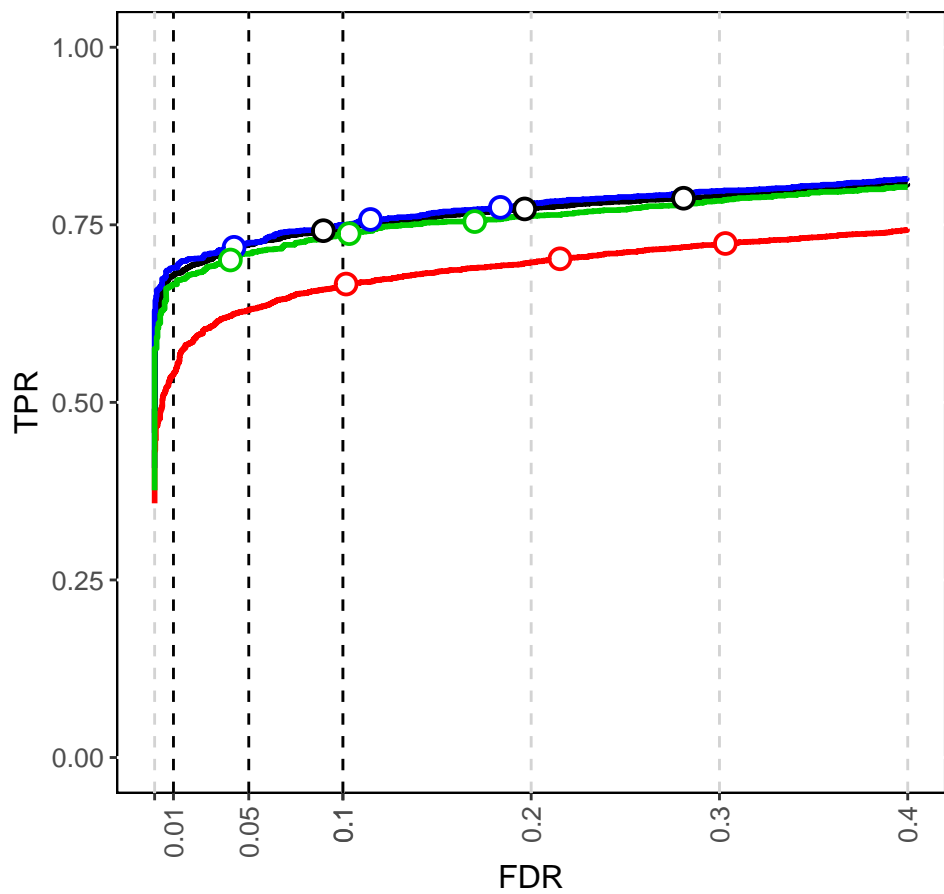
Tasic: 75 cells (satuRn)



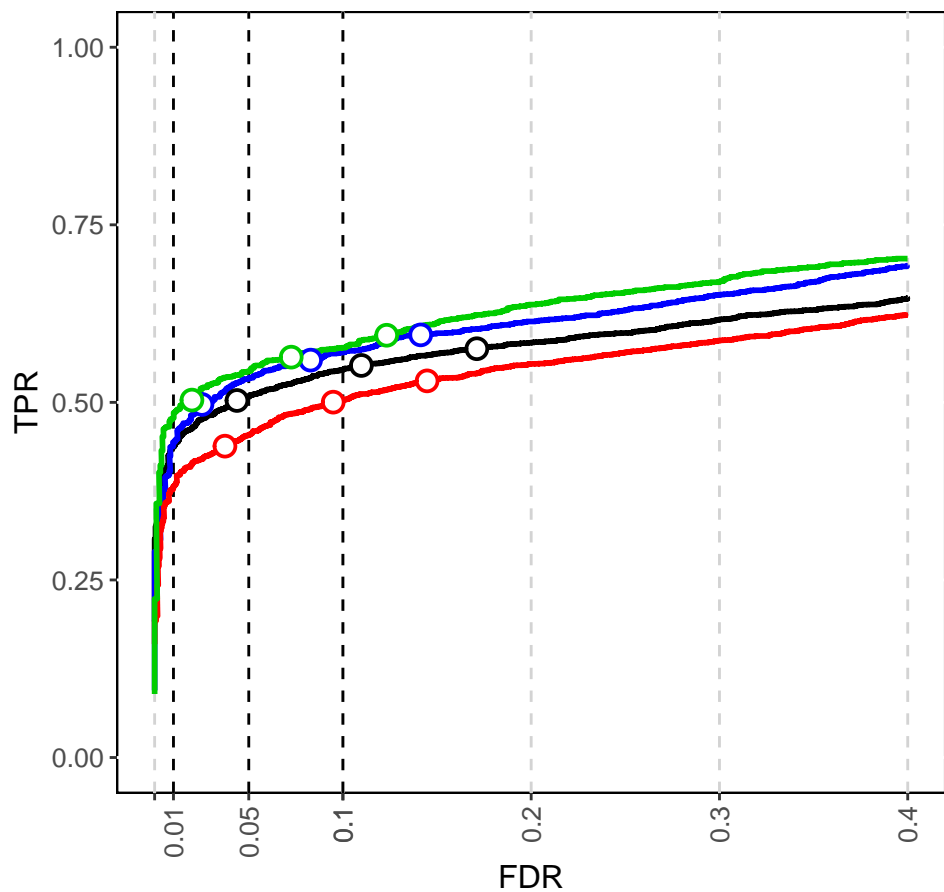
Chen: 20 cells (DoubleExpSeq)



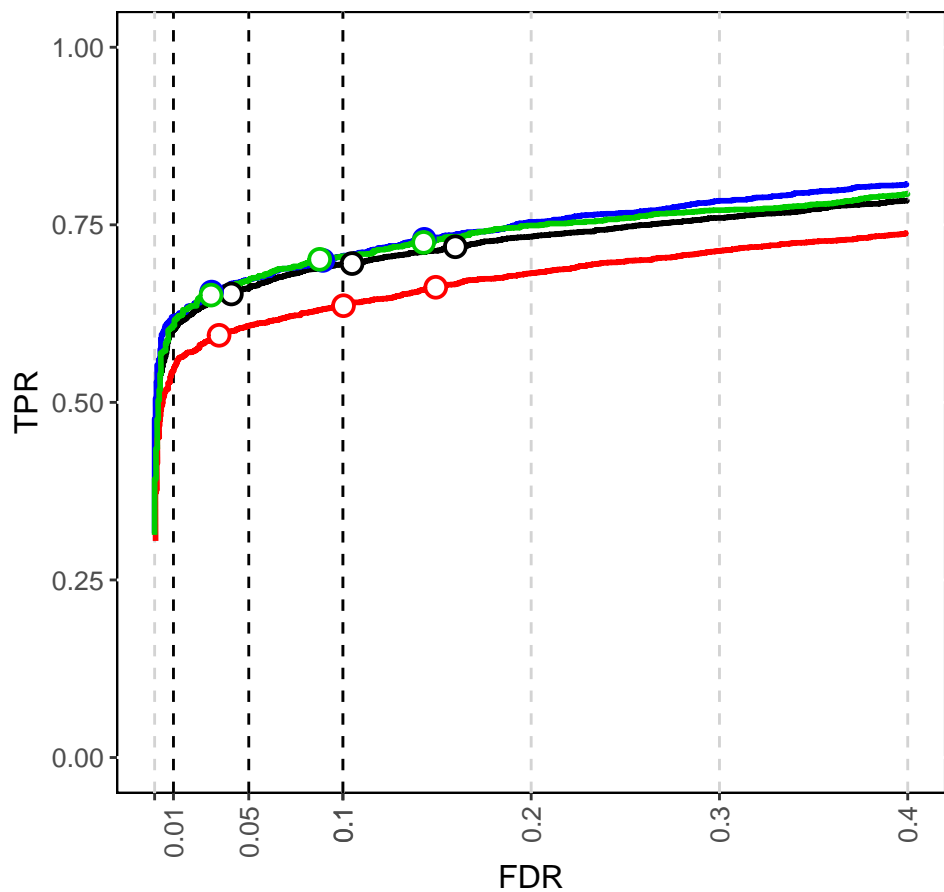
Chen: 50 cells (DoubleExpSeq)



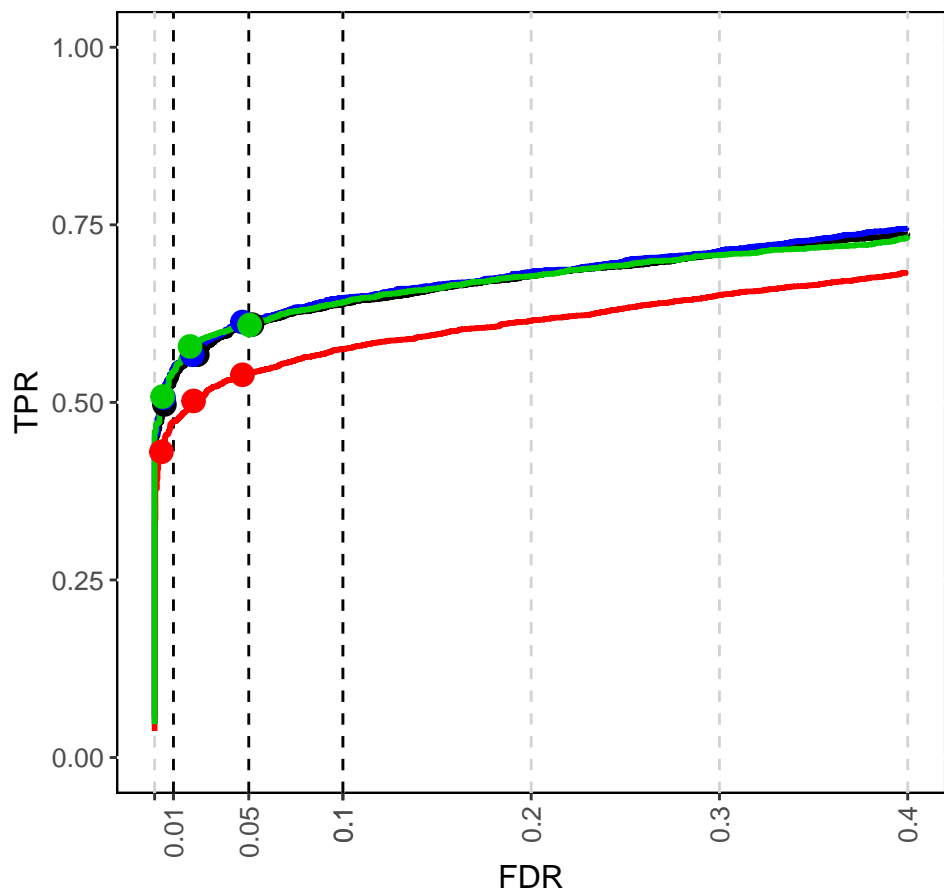
Chen: 20 cells (limma_diffsplice)



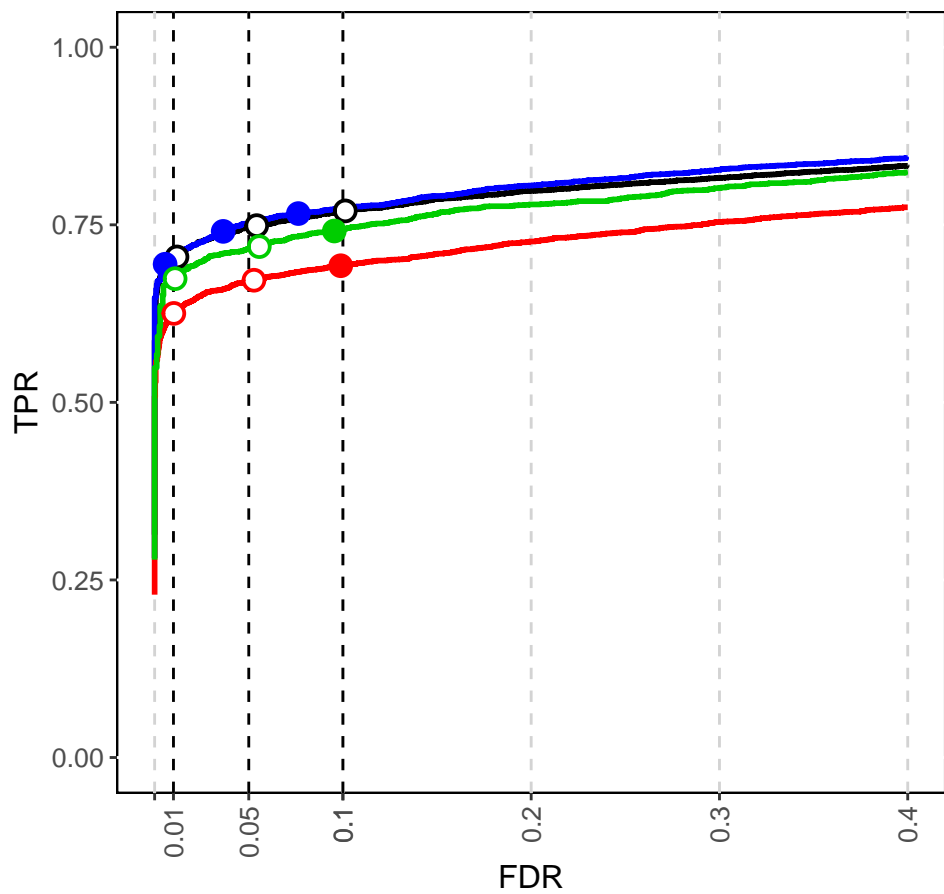
Chen: 50 cells (limma_diffsplice)



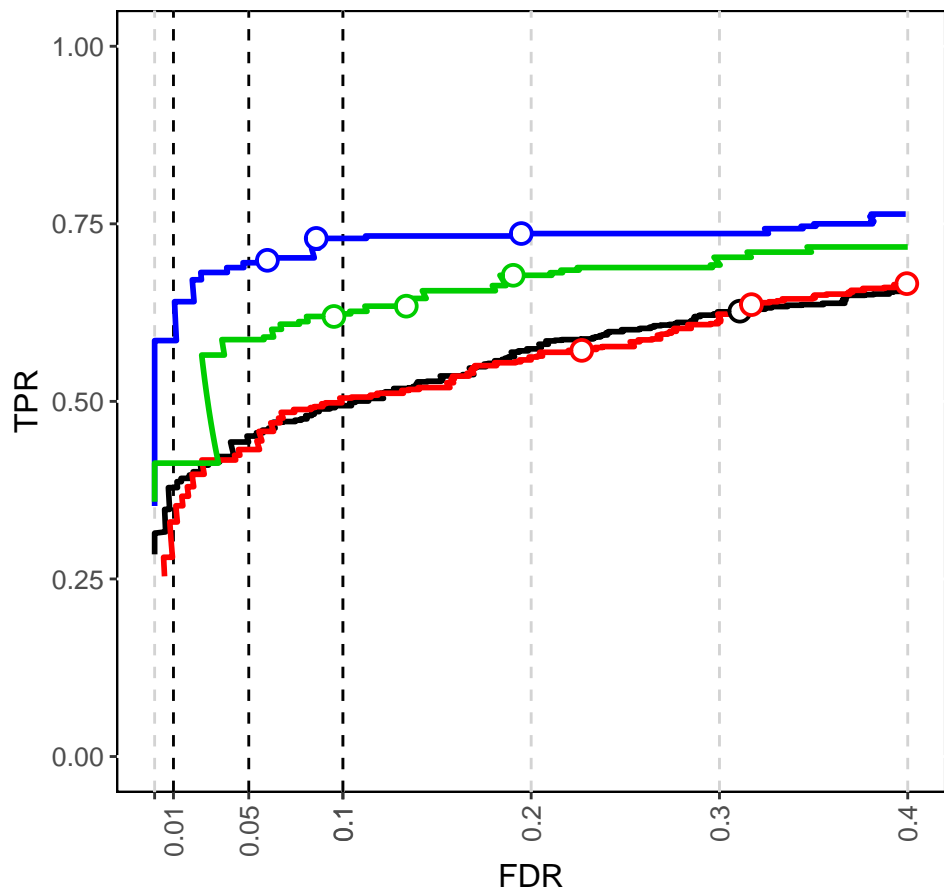
Chen: 20 cells (satuRn)



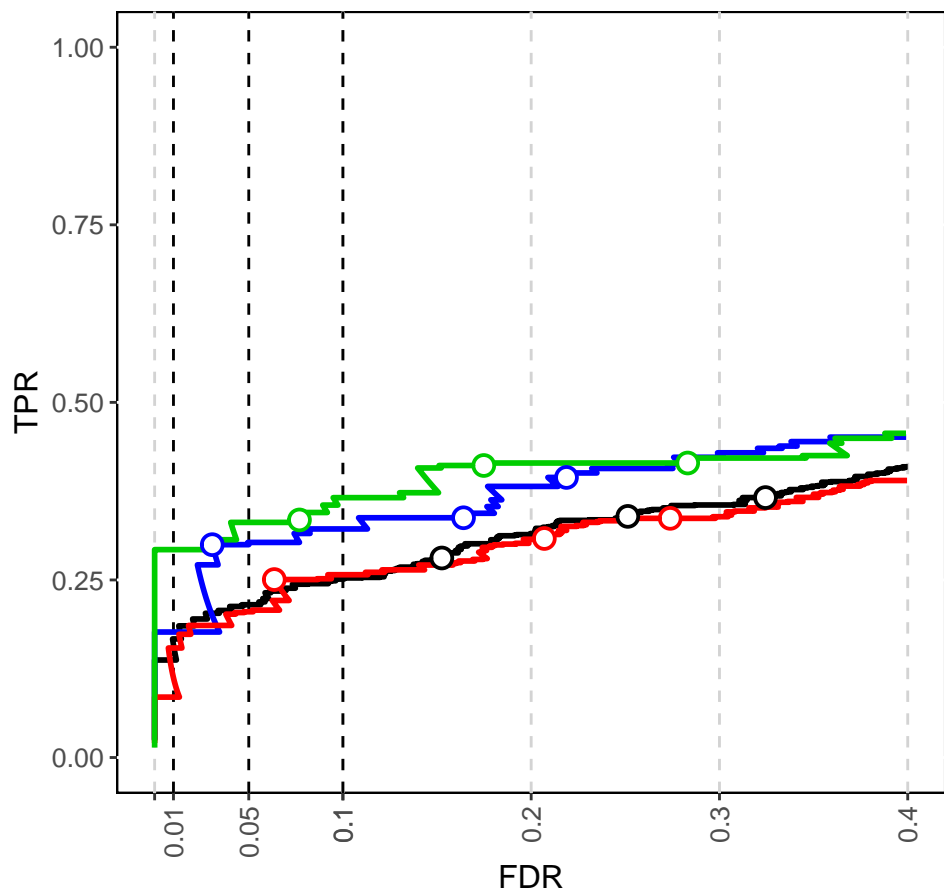
Chen: 50 cells (satuRn)



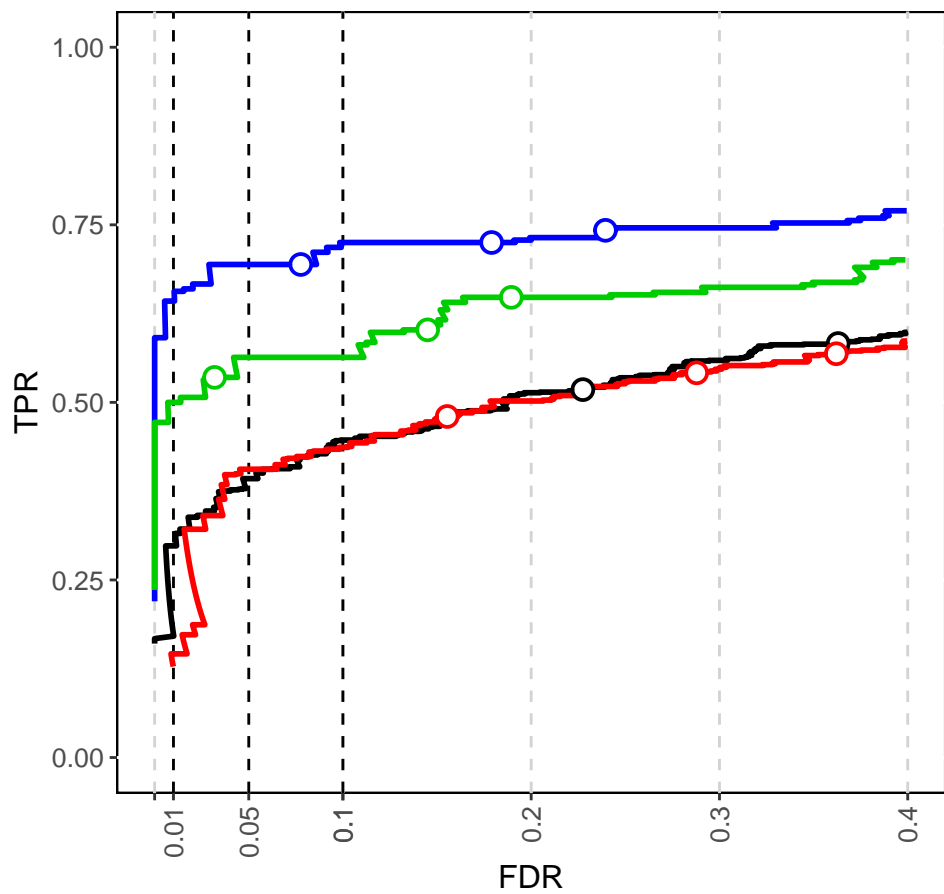
Darmanis: 100 cells (DoubleExpSeq)



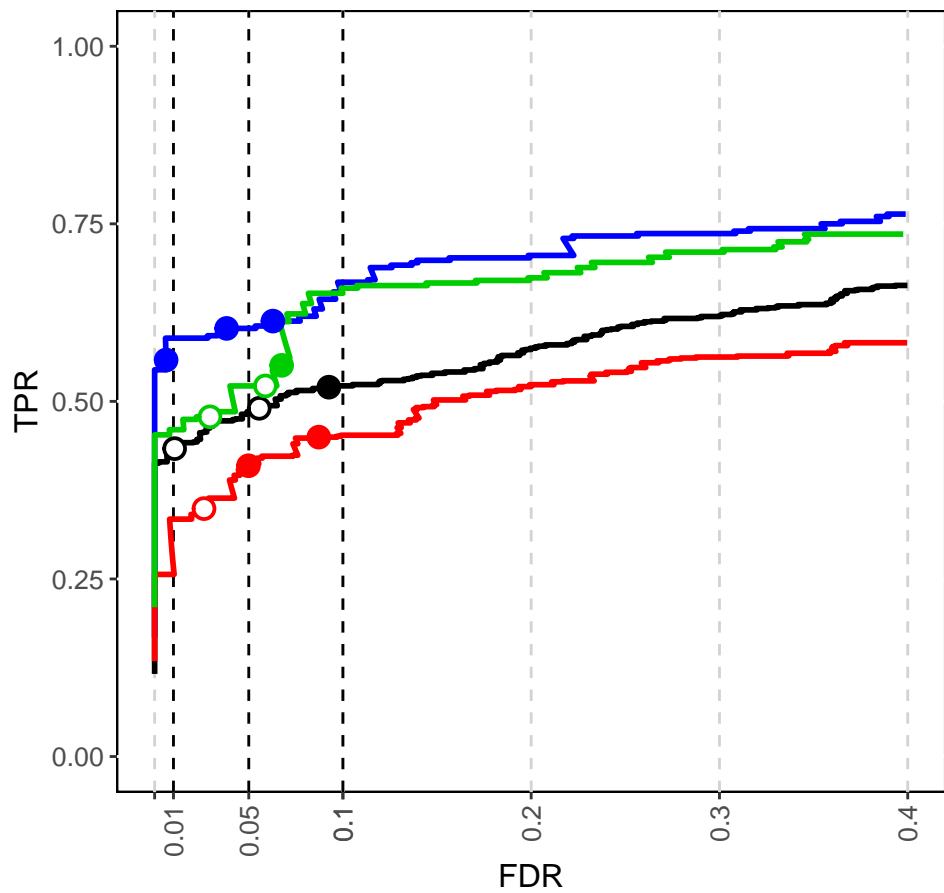
Darmanis: 20 cells (DoubleExpSeq)



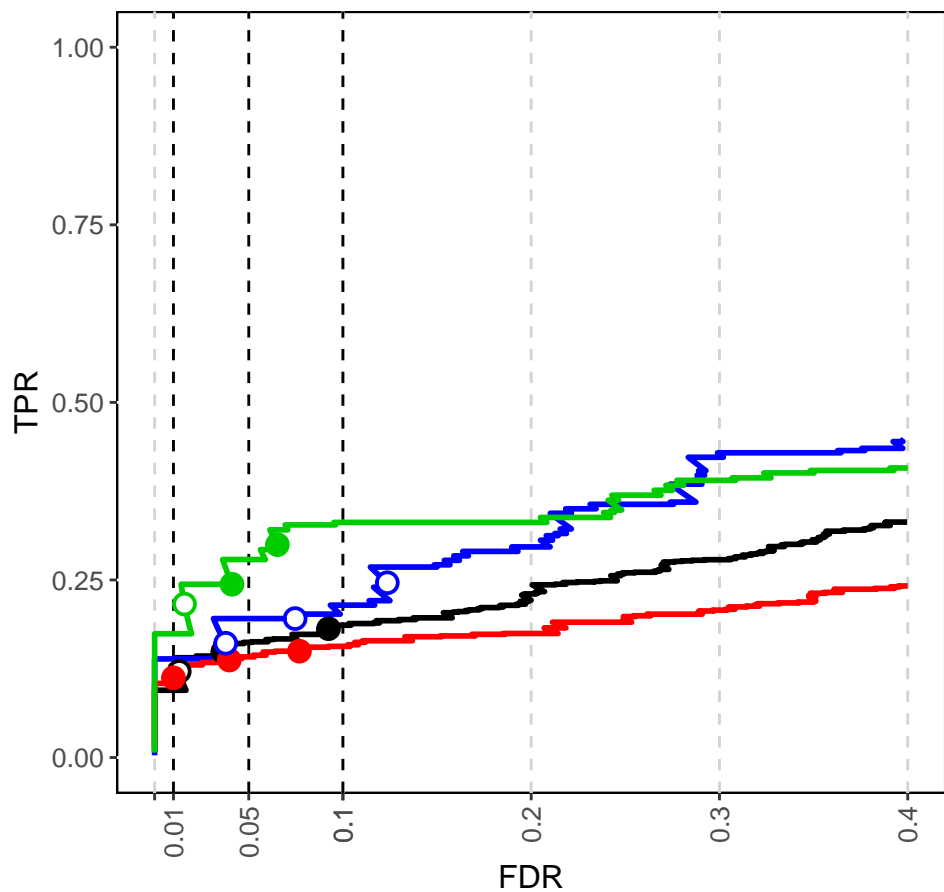
Darmanis: 50 cells (DoubleExpSeq)



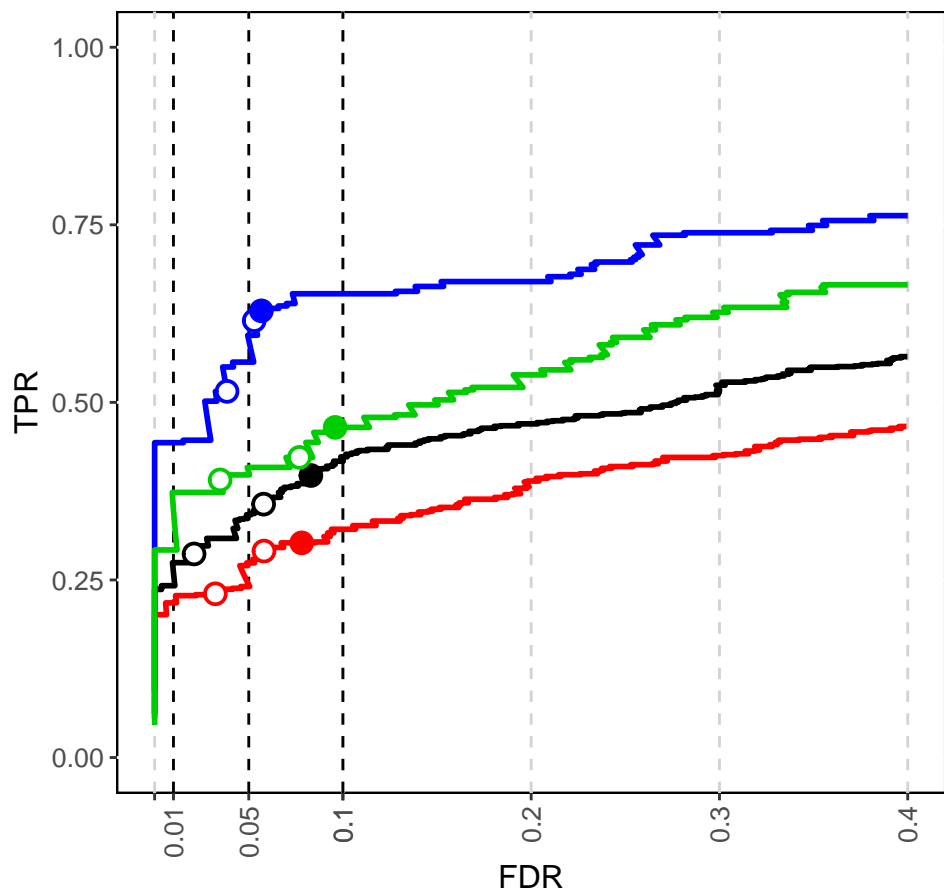
Darmanis: 100 cells (limma_diffsplice)



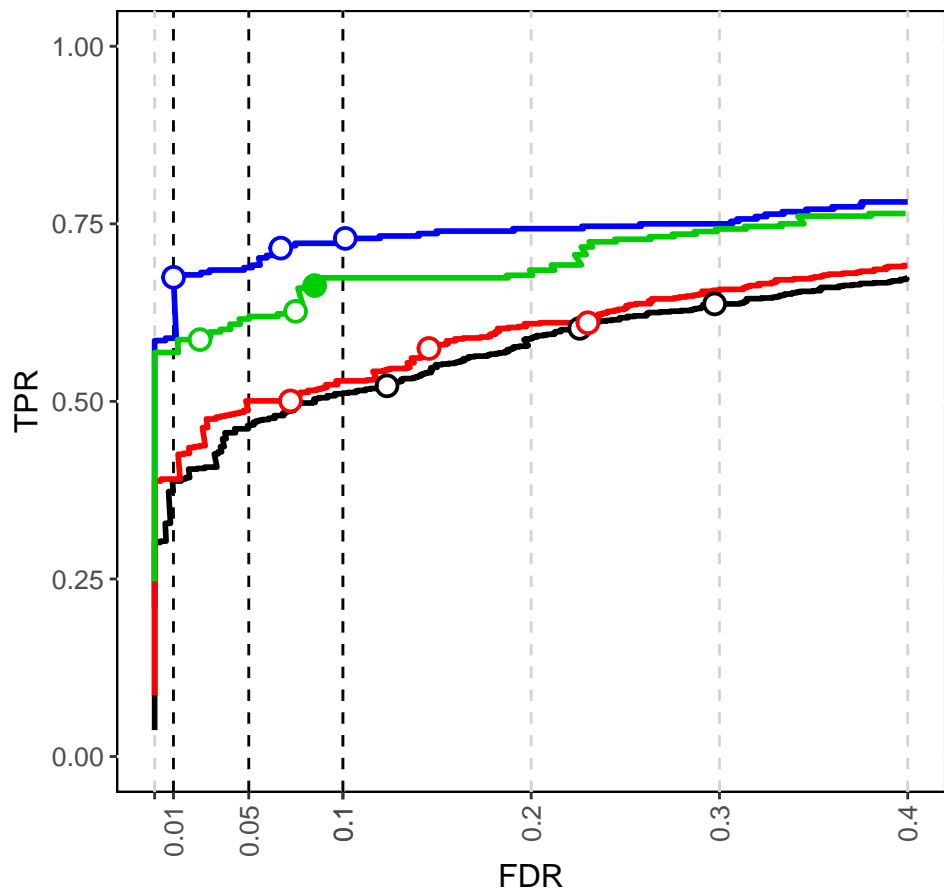
Darmanis: 20 cells (limma_diffsplice)



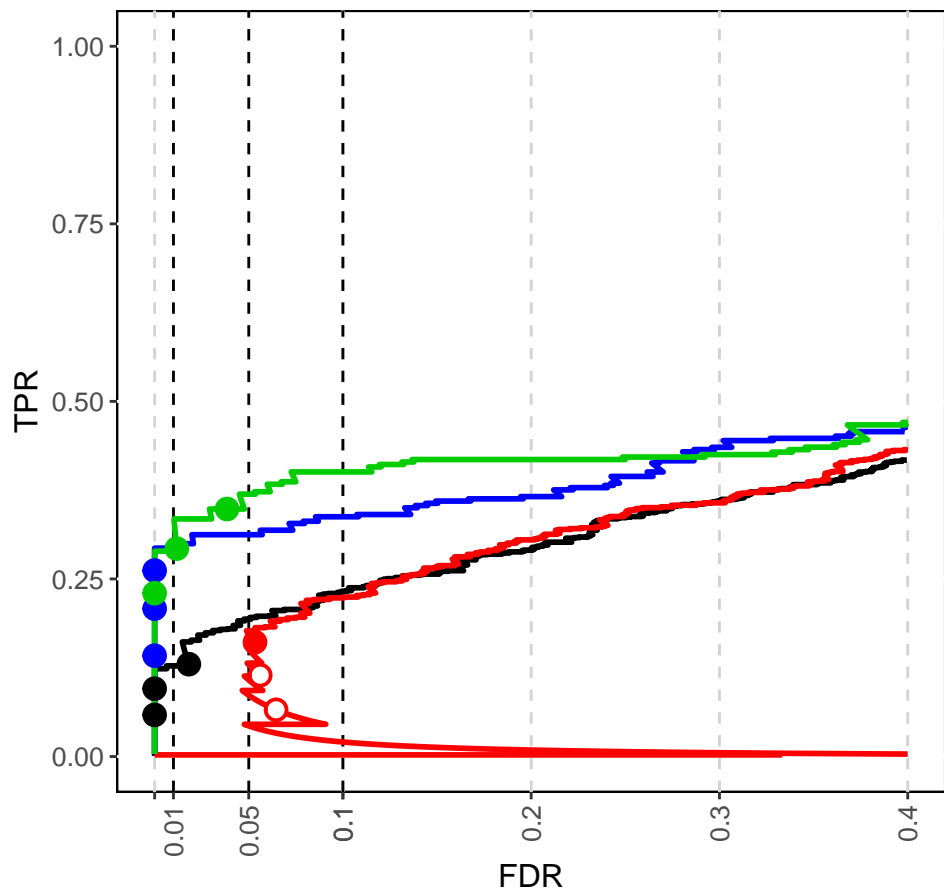
Darmanis: 50 cells (limma_diffsplice)



Darmanis: 100 cells (satuRn)



Darmanis: 20 cells (satuRn)



Darmanis: 50 cells (saturn)

