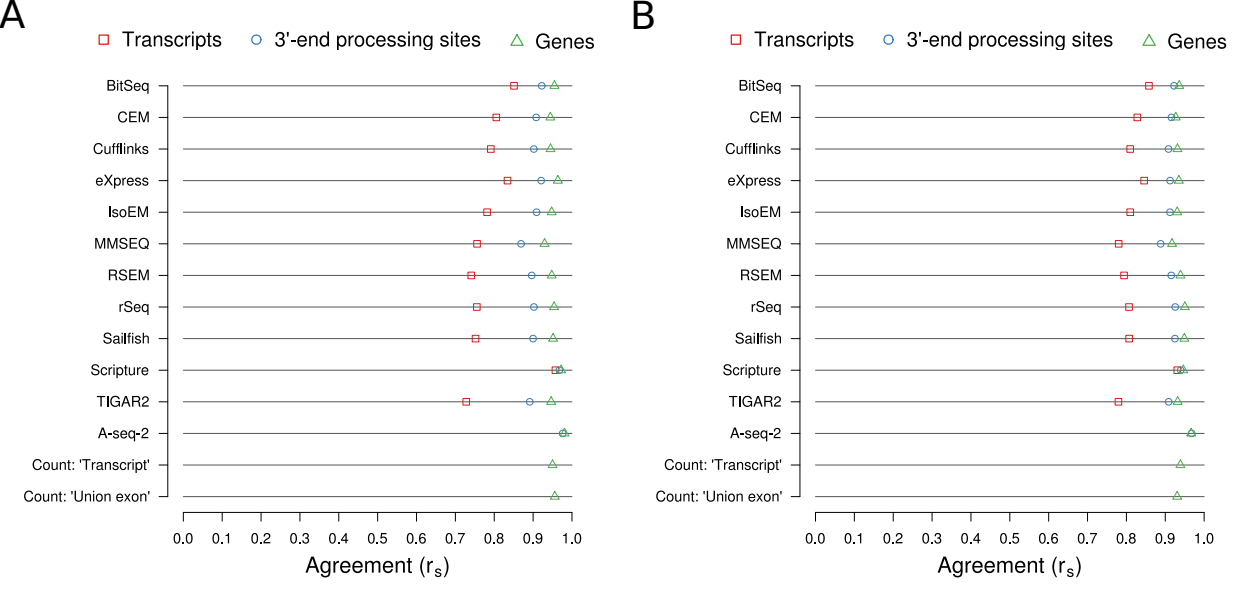
A
Transcripts
3'-end processing sites
Genes
BitSeq
CEM
Cufflinks
eXpress
IsoEM
MMSEQ
RSEM
rSeq
Sailfish
Scripture
TIGAR2
A-seq-2
Count: 'Transcript'
Count: 'Union exon'
0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0
Agreement (r_s)
B
Transcripts
3'-end processing sites
Genes
BitSeq
CEM
Cufflinks
eXpress
IsoEM
MMSEQ
RSEM
rSeq
Sailfish
Scripture
TIGAR2
A-seq-2
Count: 'Transcript'
Count: 'Union exon'
0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0
Agreement (r_s)