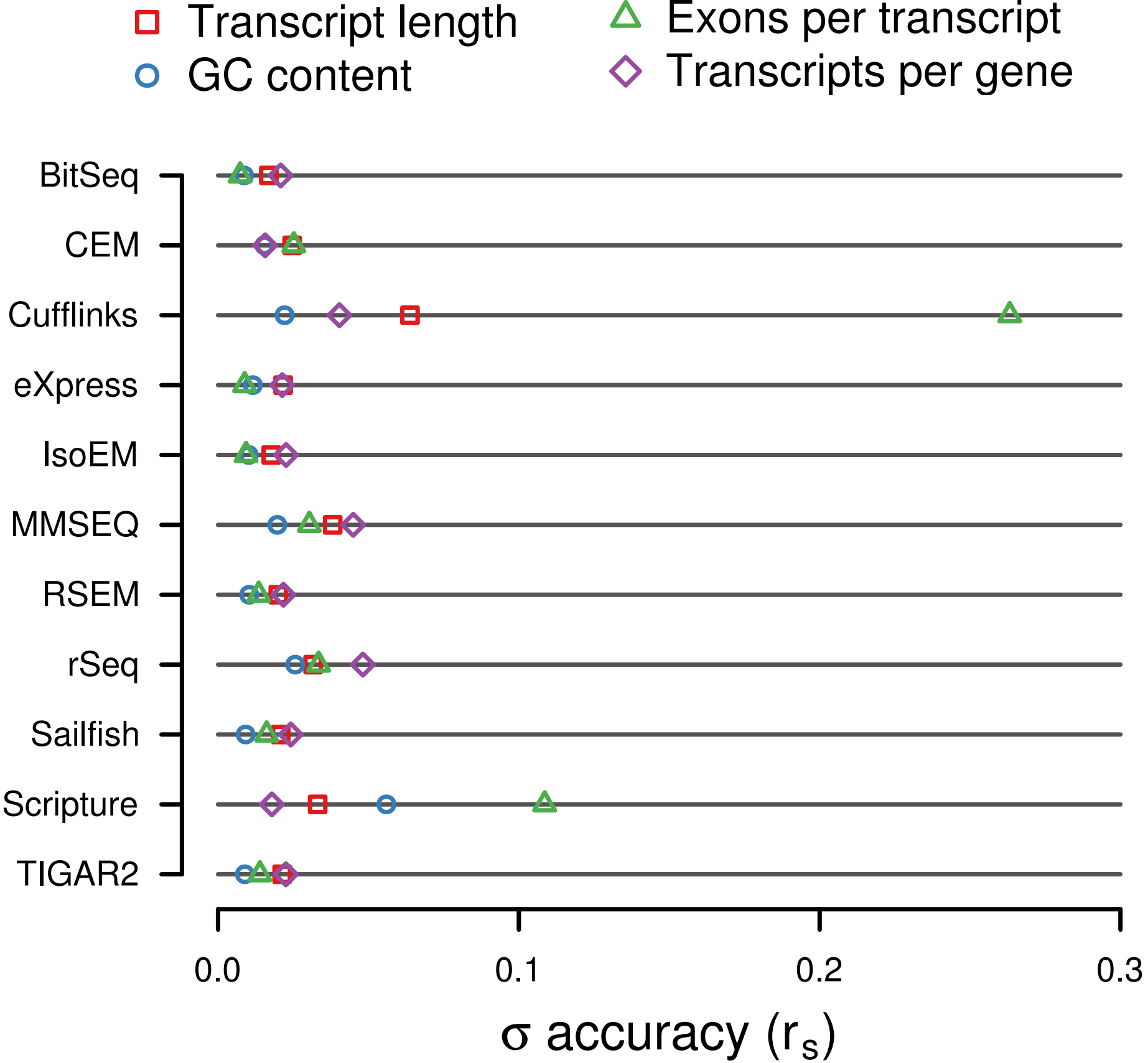
Transcript length
GC content
Exons per transcript
Transcripts per gene
BitSeq
CEM
Cufflinks
eXpress
IsoEM
MMSEQ
RSEM
rSeq
Sailfish
Scripture
TIGAR2
0.0
0.1
0.2
0.3
σ accuracy (r_s)