Supplemental Materials: A methodology and tools for evaluating the performance of RNA-Seq bioinformatic pipelines

Supplementary Figure 1: Memory and time of mapping softwares
Performances were measured with unix /usr/bin/time command. Mappers were run with 10 cores. Time is set to user time (in minutes), while memory is set to max memory usage (in Gb).

Supplementary Figure 2: Accuracy sensitivity for alignments
Supplementary Figure 3: Accuracy sensitivity for splices

Supplementary Figure 4: Accuracy sensitivity for snps

Supplementary Figure 5: Accuracy sensitivity for insertions
Supplementary Figure 6: Accuracy sensitivity for deletions

Supplementary Figure 7: Accuracy sensitivity for colinear_fusions
Supplementary Figure 8: Accuracy sensitivity for noncolinear fusions