

Figure S1. Results for Ex90N50 statistic.

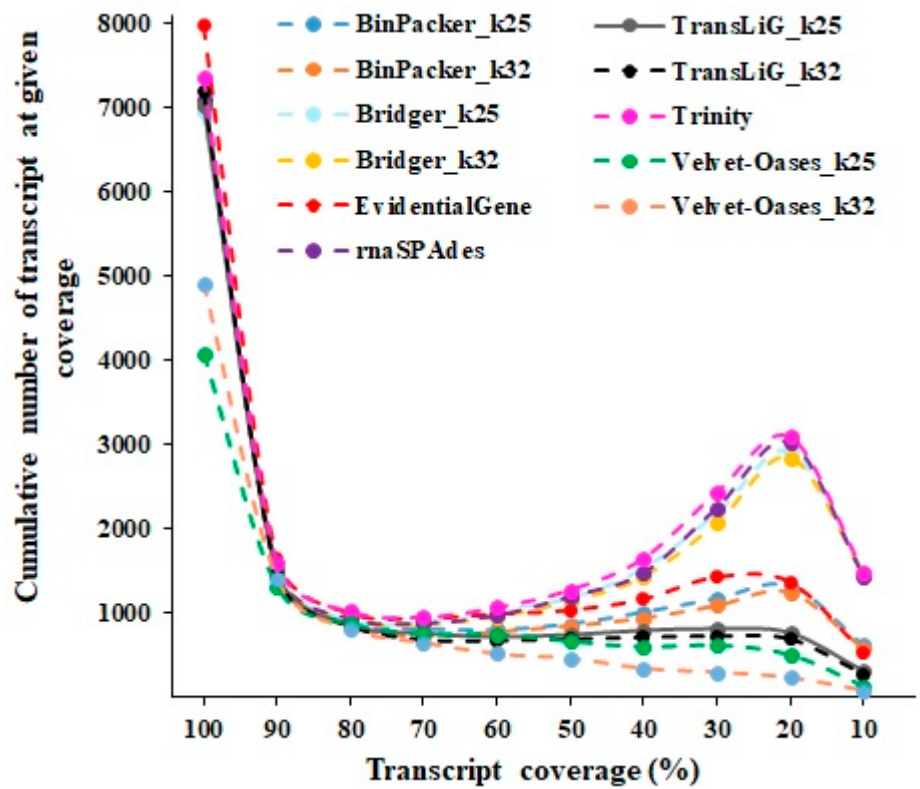


Figure S2. Cumulative number of genes that have been aligned to the Swiss-Prot database at a given coverage.

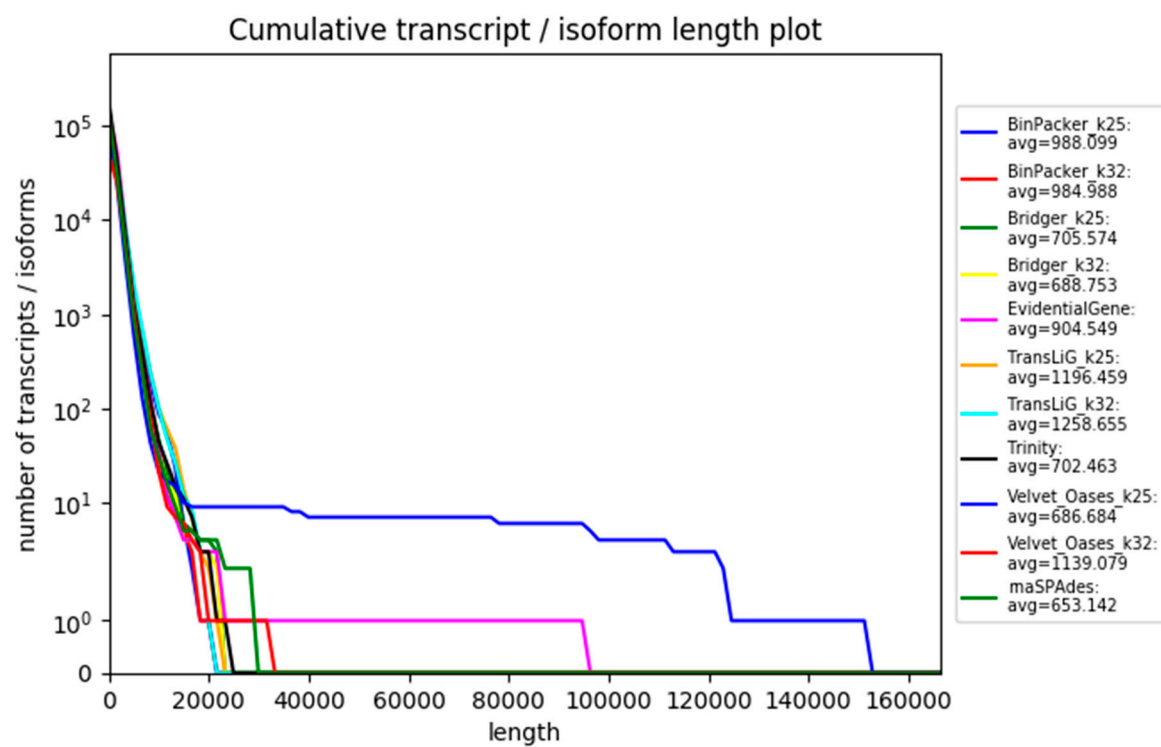


Figure S3. Transcripts length and the number of transcripts per isoform among assemblies.