Supplemental Fig S6. Examples of the effect of TransRate contig filtering on *de novo* transcriptome assemblies. **A)** Comparison of the number of genes that had full length or partial length matches in *de novo* assembled transcripts before and after TransRate filtering. **B)** Histogram of lengths of genes encompassed by the *de novo* assembled transcripts in A that were discarded by TransRate filtering. **C)** As in A, but for genes that had their entire length contained within a single contig.