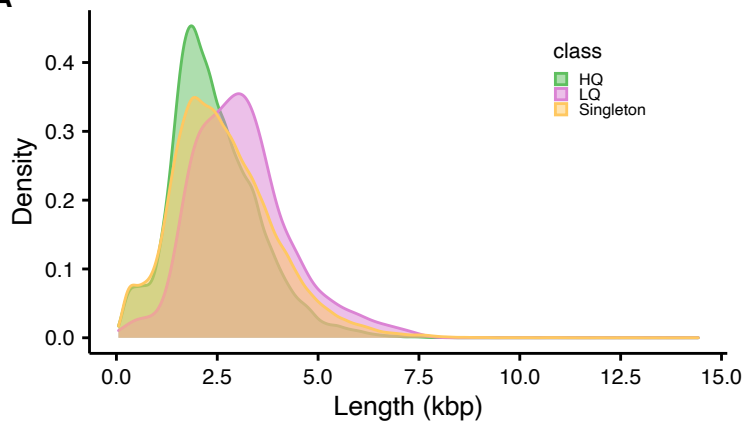
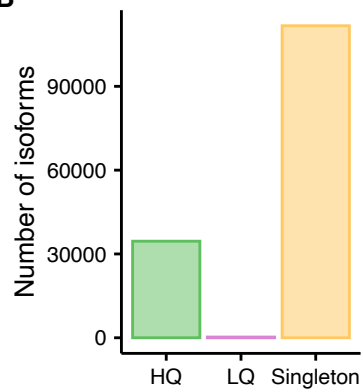
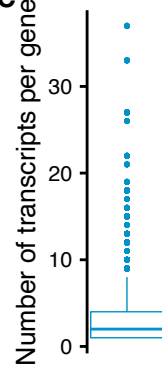


A**B****C**

Supplemental Figure S6: Summary statistics on the full-length cDNA sequences.

Full-length cDNA were sequenced to improve the accuracy of the gene annotation and identify spliced variants. The distribution of the isoform length (**A**) and the total number of isoforms (**B**) are represented by class. **C.** Summary of the number of transcripts per gene after isoform prediction.