

Transcriptional profiling of long non-coding RNAs in mantle of *Crassostrea gigas* and their association with shell pigmentation

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Figure S1 TE components in three subtypes of oyster lncRNAs and mRNAs. The main TE families were identified using RepeatMasker. Percentage of TE components in mRNAs and individual subtype of lncRNAs were represented.

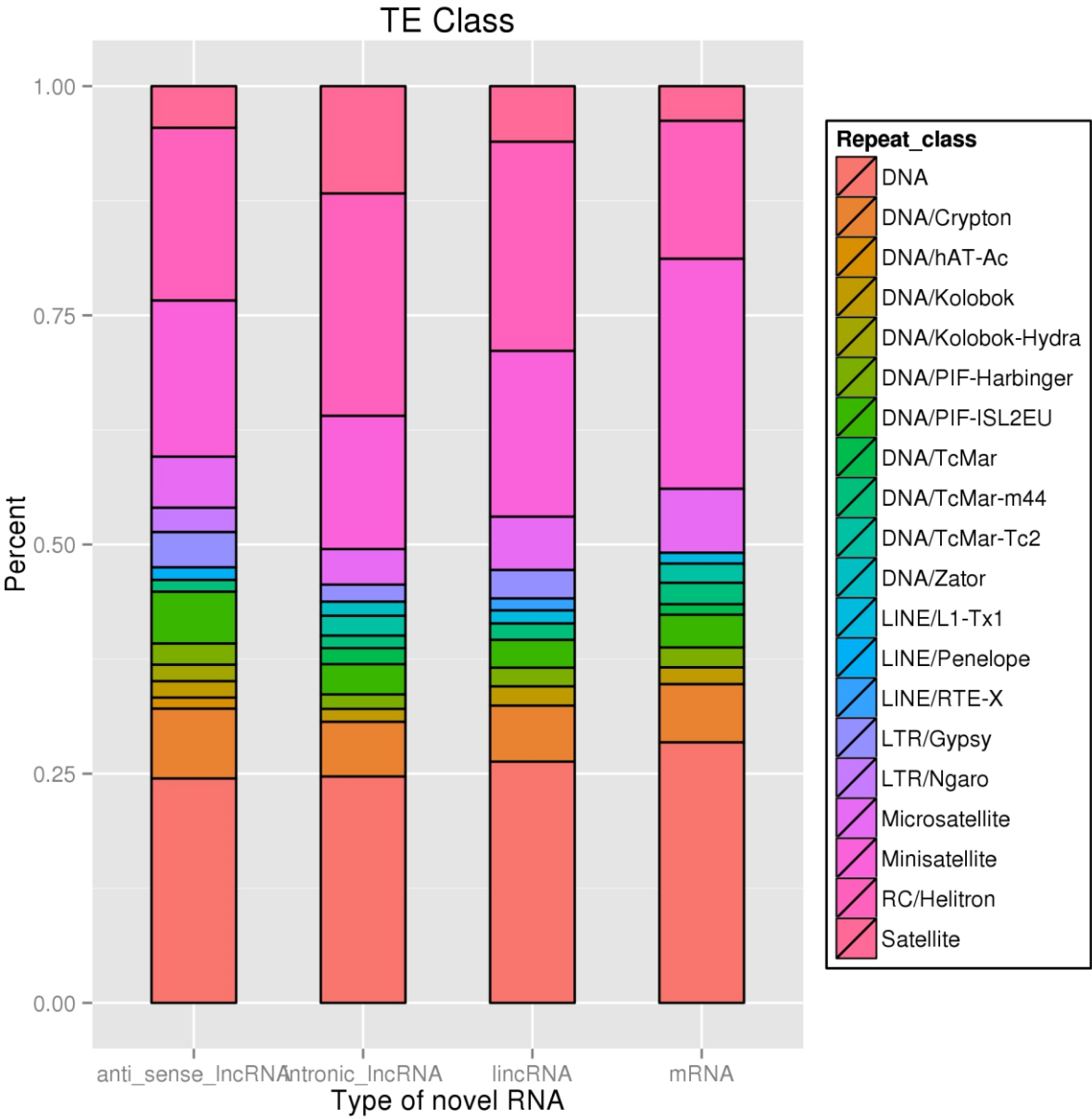


Figure S2 Cluster analysis of differentially expressed lncRNA and mRNA transcripts | a. Cluster analysis of differentially expressed lncRNA transcripts. b. Cluster analysis of differentially expressed mRNA transcripts.

