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Supplementary data

Transcriptome analysis of differential expression genes from petals and lips of Phalaenopsis amabilis (*Phalaenopsis aphrodite Rchb*. F) to identify genes associated with floral development

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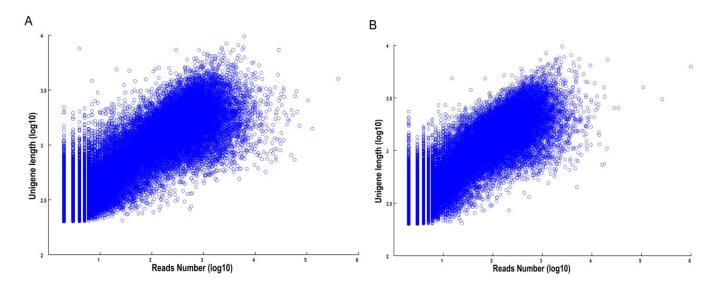
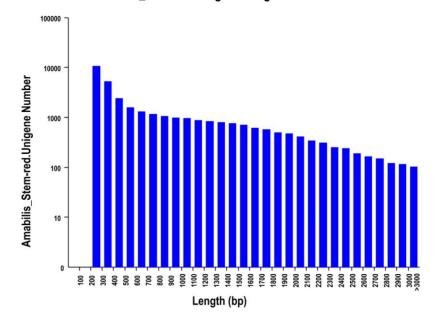
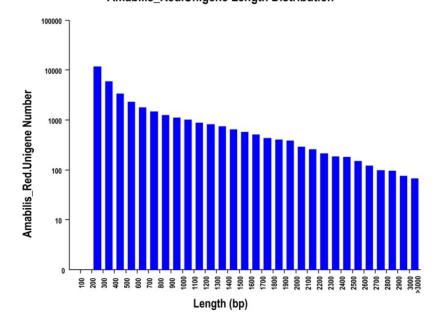


Figure S1. The distributions of unigenes length and read numbers. A is the unigenes length distributions of the Amabilis_Stem-red; B is the unigenes length distributions of the Amabilis_ red.





B Amabilis_Red.Unigene Length Distribution



Amabilis. Unigene Length Distribution

C

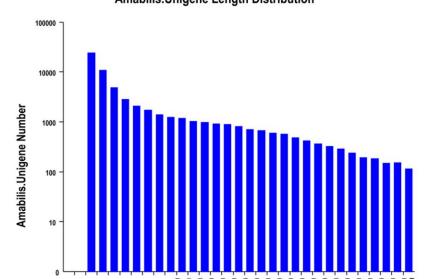


Figure S2. The unigenes length distributions. A is the unigenes length distributions of the Amabilis_Stem-red; B is the unigenes length distributions of the Amabilis_ red; C is all the unigenes length distributions.

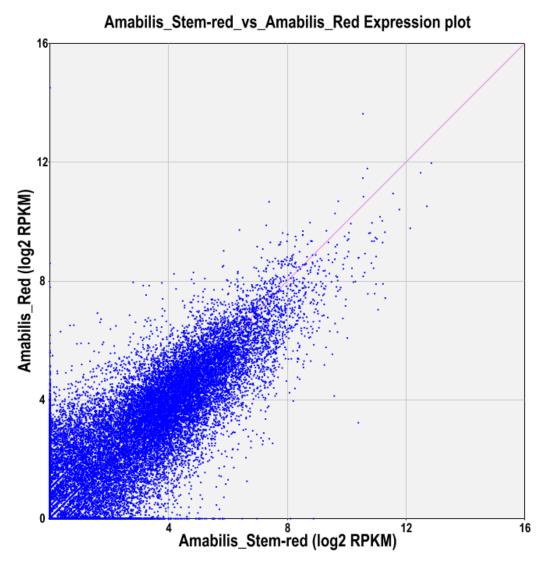


Figure S3. Amabilis_Stem_ red vs Amabilis_ red expression plot. The Y-axis represents Amabilis_red (lip) expression, and the X-axis means the Amabilis_Stem-red (tepal).