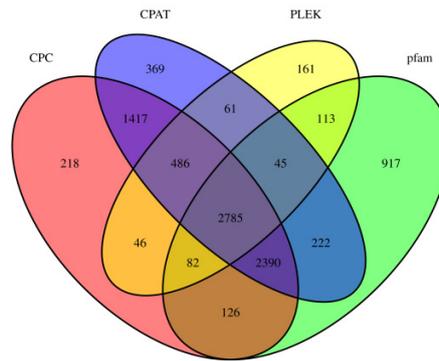


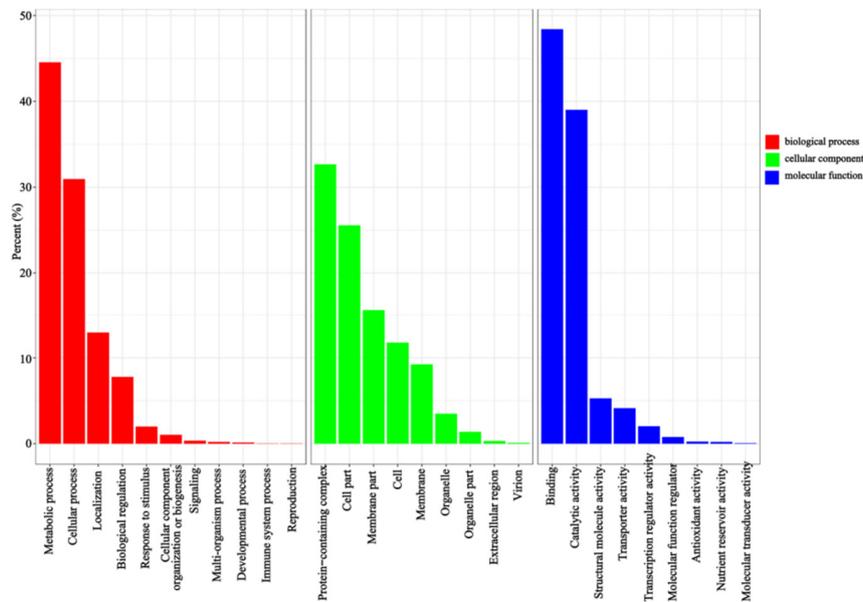
# Supplementary Material

## 1 Supplementary Figures and Tables

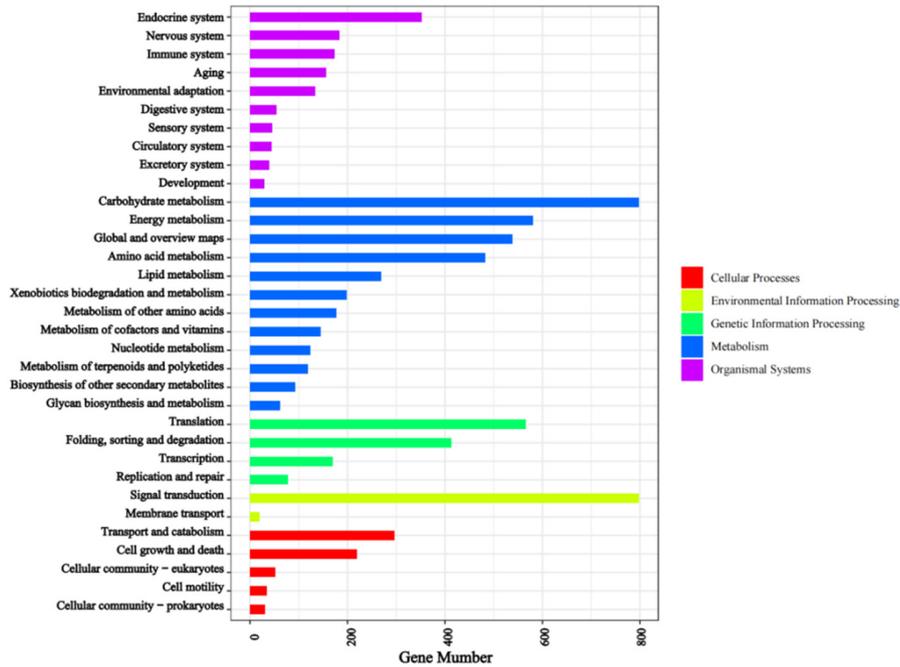
### 1.1 Supplementary Figures



Supplementary Figure S1. Prediction results of lncRNA of the full-length transcript of *C. vietnamensis*



Supplementary Figure S2. GO annotation classification map of the full-length transcriptome sequence



Supplementary Figure S3. KEEG annotation classification map of the full-length transcriptome sequence

## 2.2 Supplementary Tables

Supplementary Table S1. Statistical results for non-redundant high quality full-length transcripts

Seq_ number	Total Length (bp)	<1kb	>1kb&<2kb	>2kb&3kb	>3kb	N50	N90	Mean	Median	Max	Min
14152	12403692	9835	3935	350	32	997	515	876	770	4719	107