

**Supple. Fig.1** The evaluation of the annotation of *Camellia chekiangoleosa* transcriptome. **A**, The distribution of unigenes that are annotated in the Gene Ontology (GO) database. **B**, The distribution of unigenes that are annotated in the eggNOG database. **C**, The distribution of unigenes that are annotated in the KEGG database.

