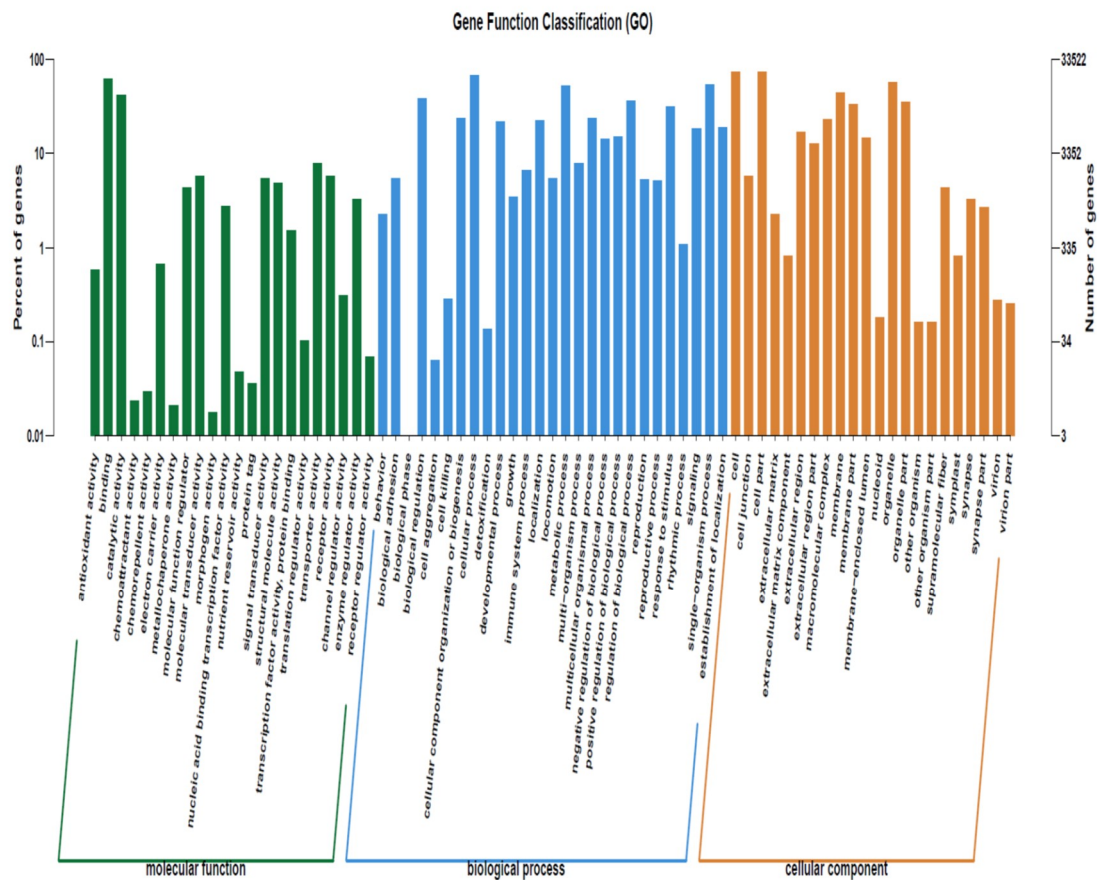
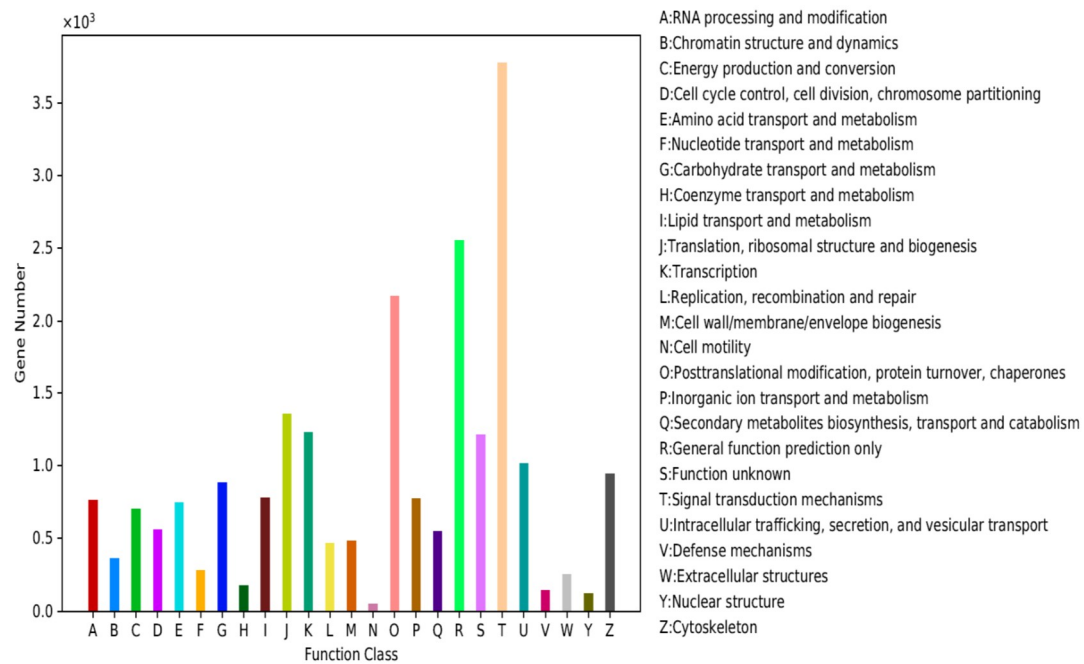


A



B



C

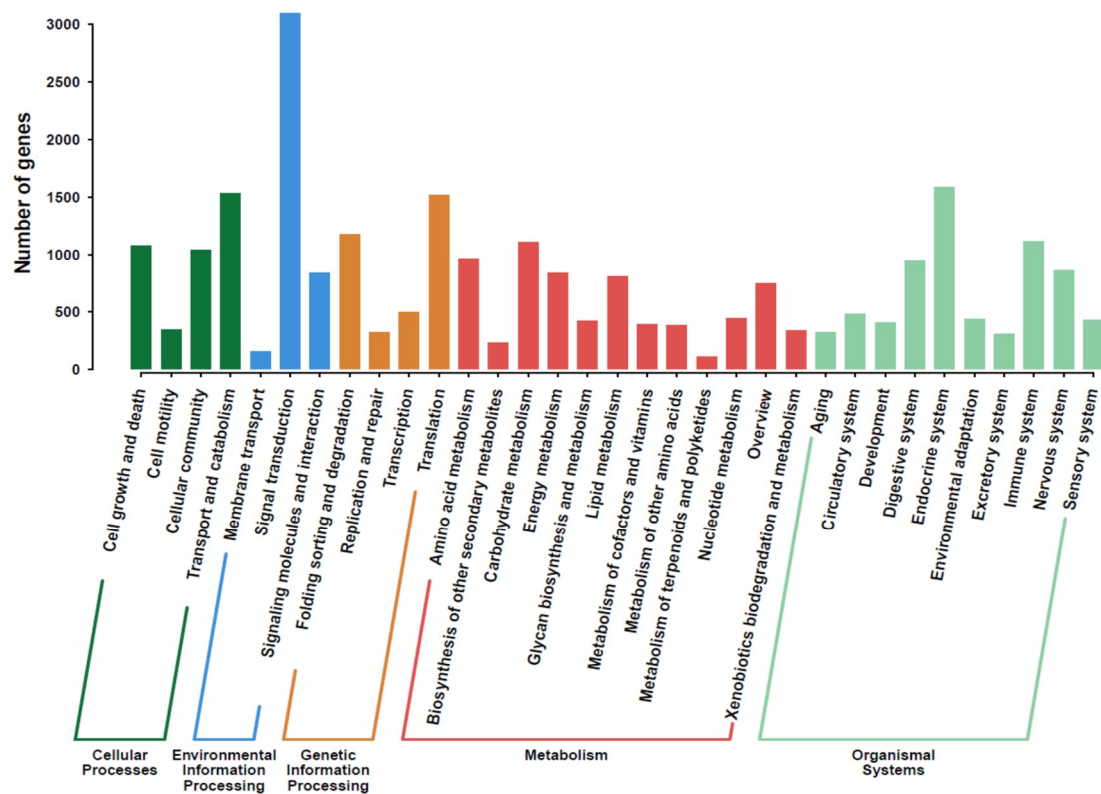


Figure S1. Similarity analysis and gene function annotation in *Anadara kagoshimensis*. A: GO (Gene ontology) classification of the assembled unigenes; B: KOG (EuKaryotic Orthologous Groups) classification of the assembled unigenes; C: KEGG (Kyoto Encyclopedia of Genes and Genomes) annotation analysis of the assembled unigenes.

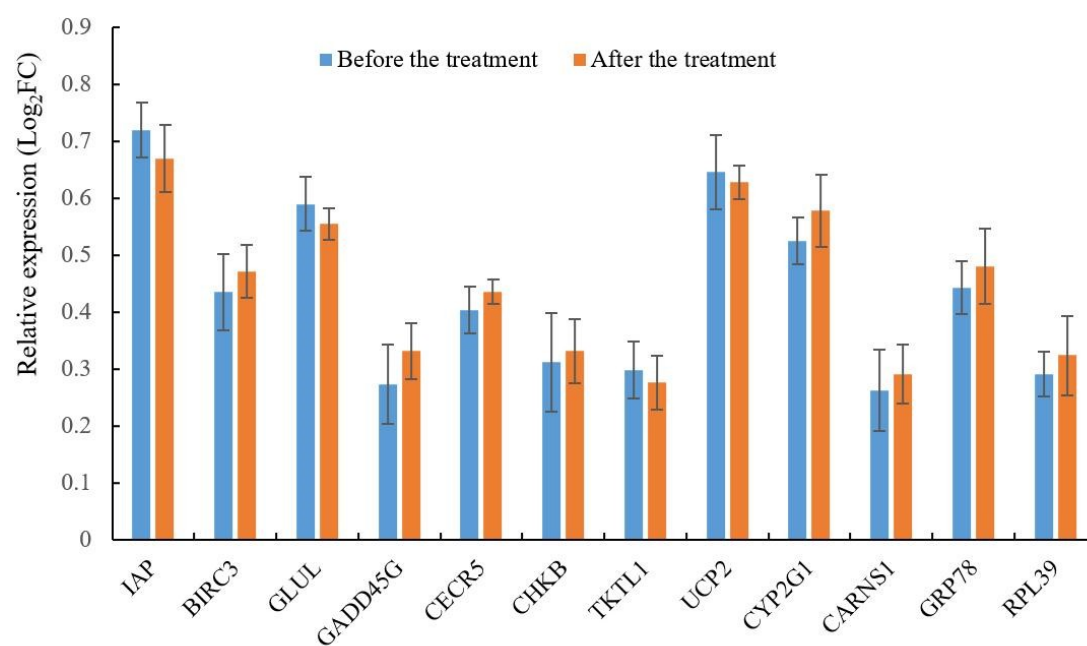


Figure S2. QRT-PCR analysis for 12 DEGs in *Anadara kagoshimensis* before and after 30‰ salinity treatment. The relative expression values were normalized to β -actin gene expression. Vertical bars represent the mean \pm S.D. (n = 3).