

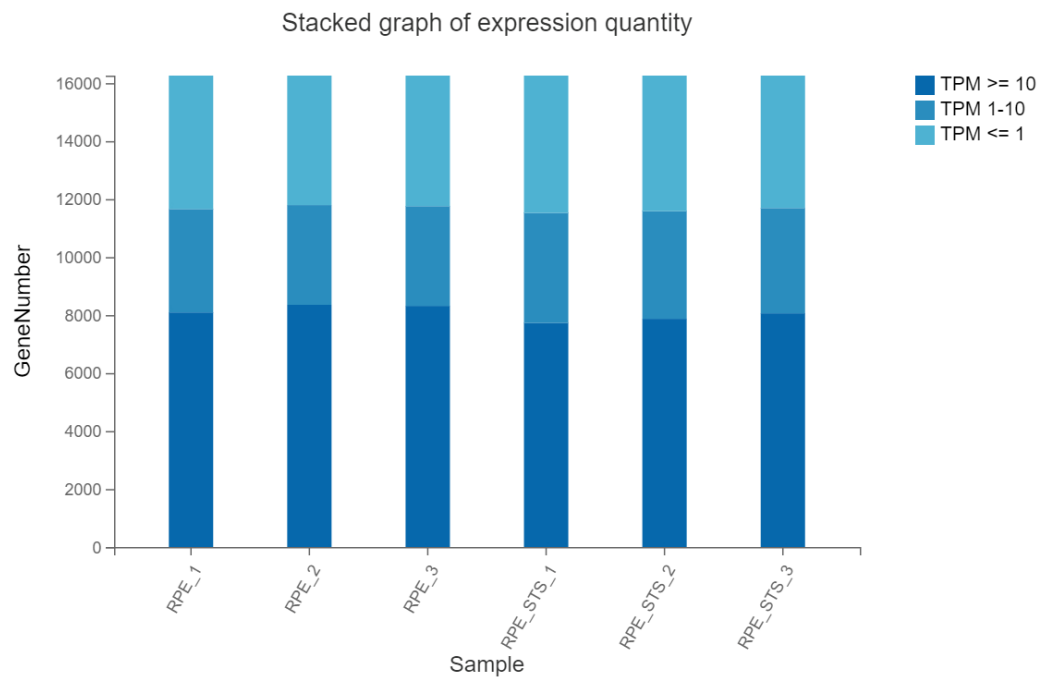
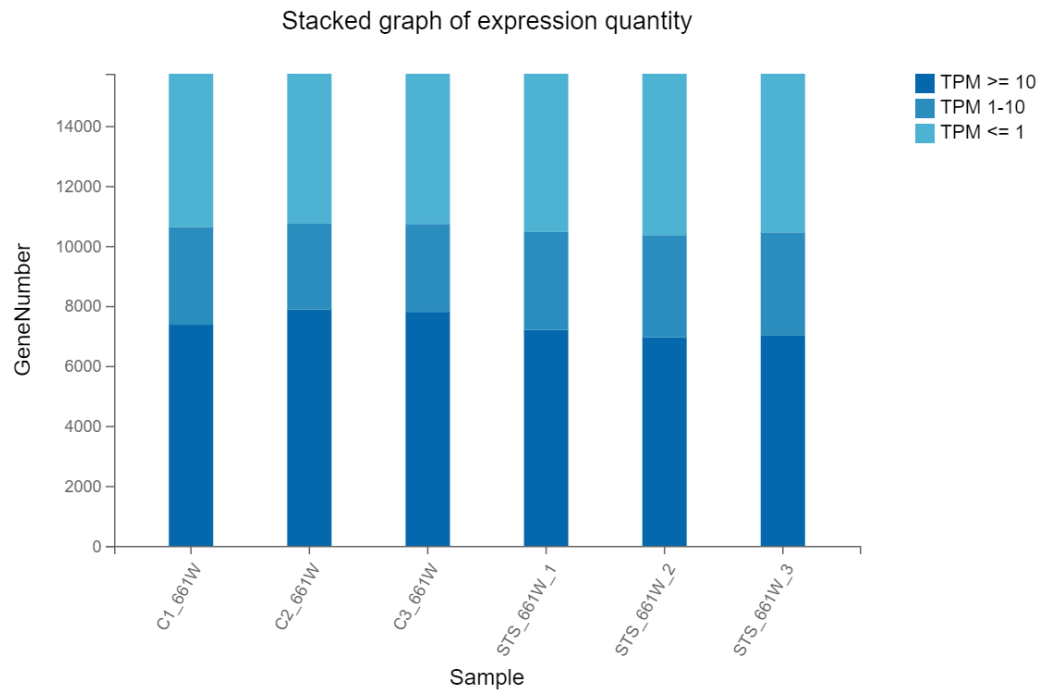


Figure S2. Stacked histogram analysis of transcriptome sequencing data showed that the RNA-Seq data of each sample in each group were highly reproducible. Transcripts Per Million (TPM) ≤ 1 indicates a gene with a very low expression level, TPM between 1 and 10 is a gene with a lower expression level, TPM Genes with ≥ 10 are highly and moderately expressed.