



Figure 5. The Venn diagram of DEGs. (A) The Venn diagram of DEGs between C1306, W1227 and T801. (B) Kyoto Encyclopedia of Genes and Genomes (KEGG) analysis of 2816 DEGs, which were mapped to the KEGG database and the pathway names were extracted. The enrichment was conducted for DEGs involved in each pathway using the genome-scale expressed genes as the background with a hypergeometric test. The pathways with the p value under 0.05 were used to draw the graph; the ordinate is the pathway and the abscissa is the gene percent (the redder the color, the smaller the p value).