

CHLOROPHYLL METABOLISM

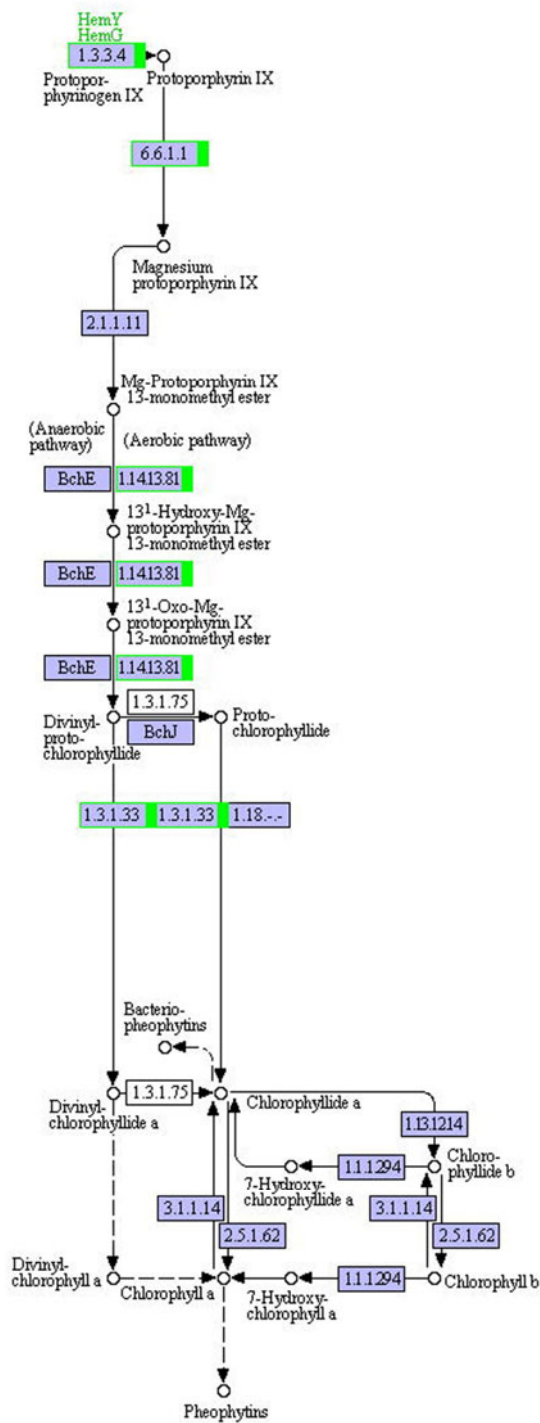


Fig. 5S. Kyoto encyclopedia of genes and genomes (KEGG) schematic of the chlorophyll metabolism pathway. Differentially expressed genes (DEGs) were compared between salt-treated group and the control. Boxes with a green frame indicate the corresponding DEGs were down-regulated in the NaCl-treated samples and those without any colored frame indicate the expression level of corresponding genes were not changed, as determined by RNA-Seq.