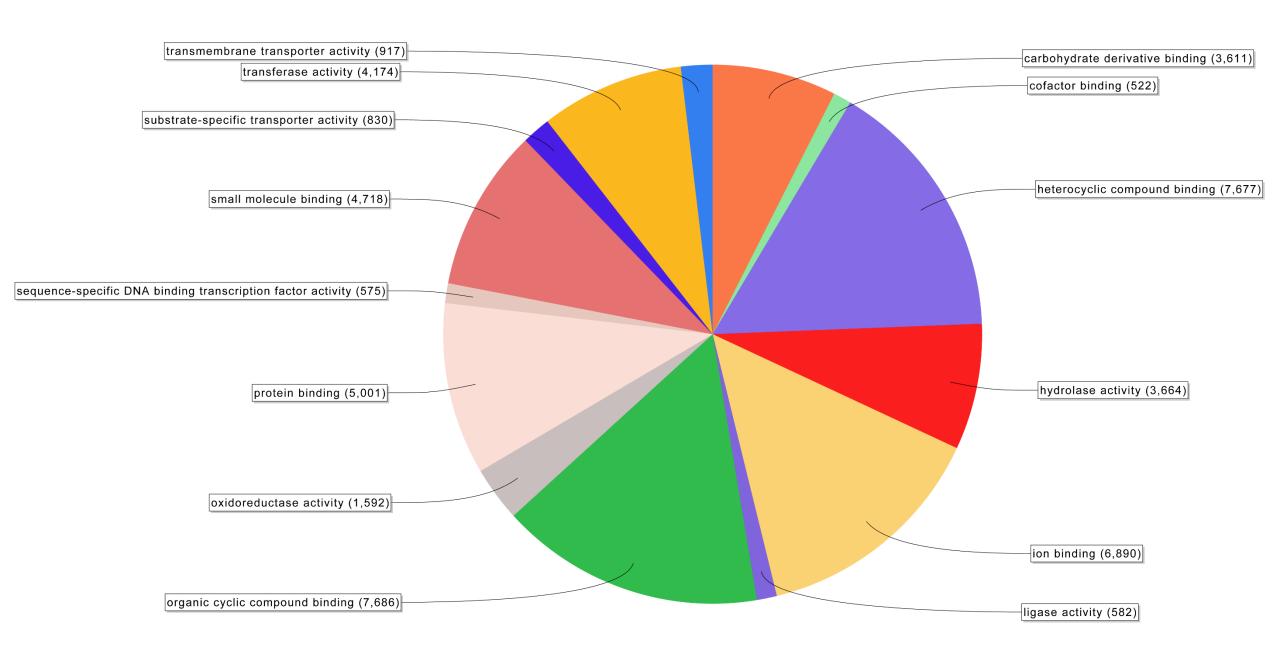
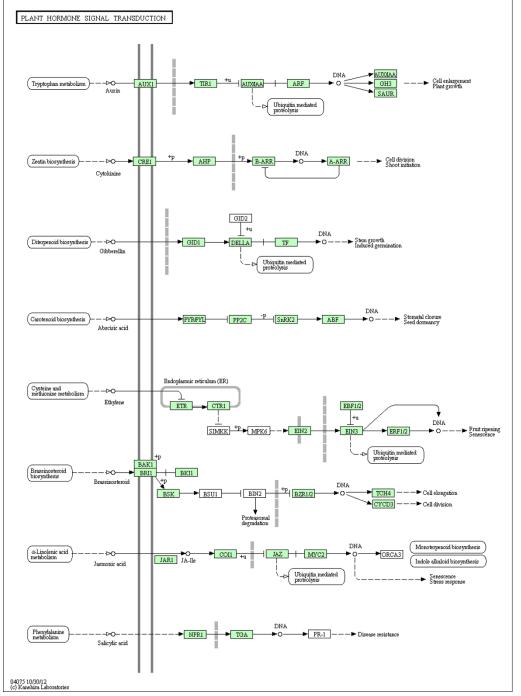


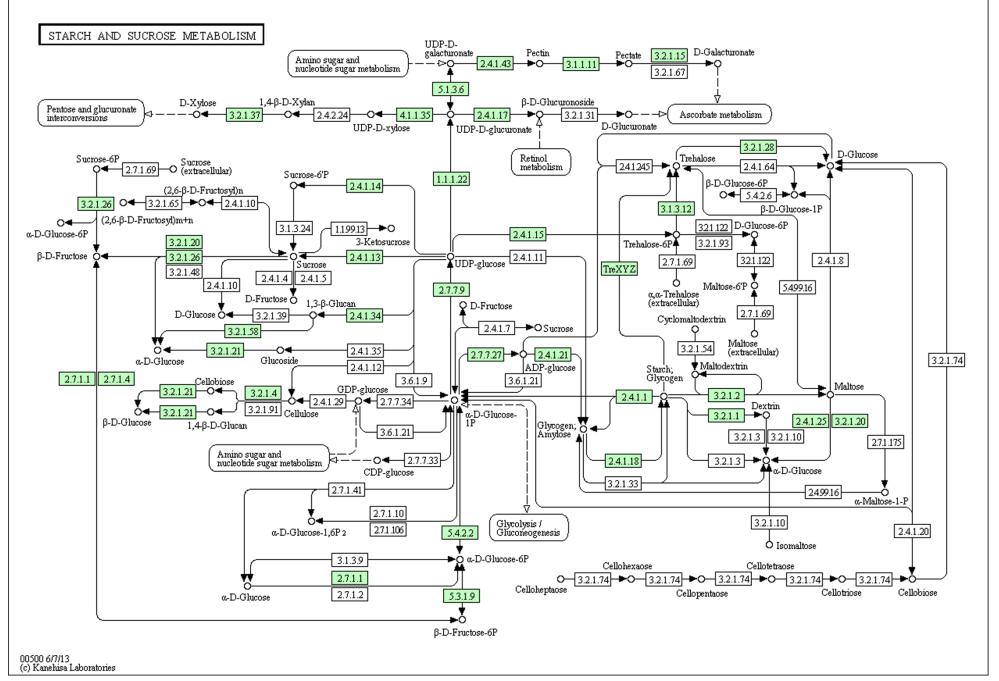
Supplementary Figure 2. Gene Ontology classification according to Biological Process



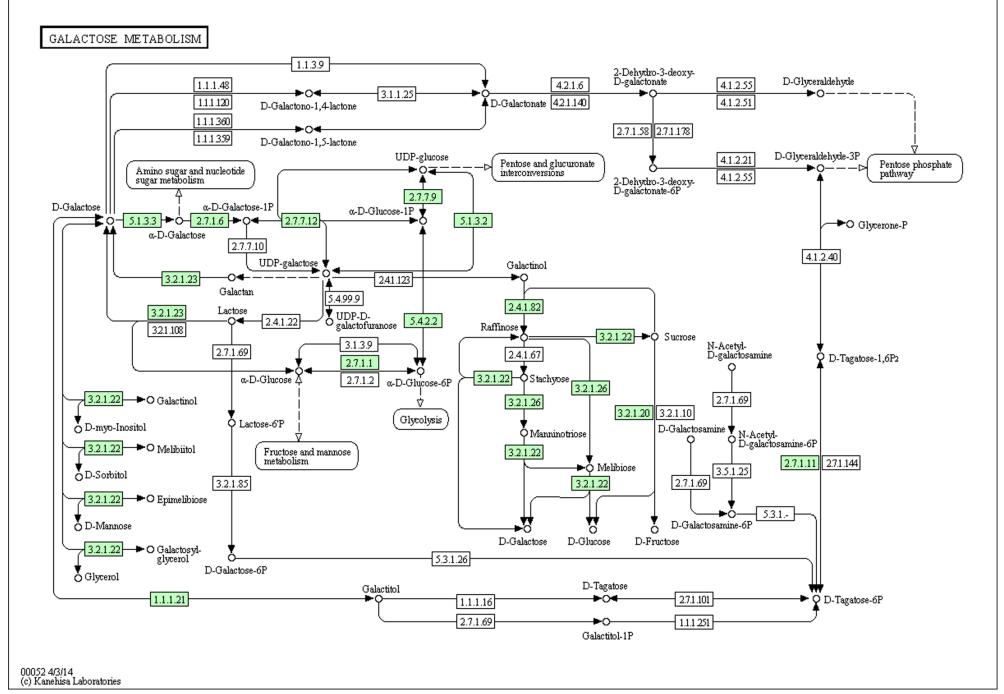
Supplementary Figure 3. Gene Ontology classification according to Molecular Function



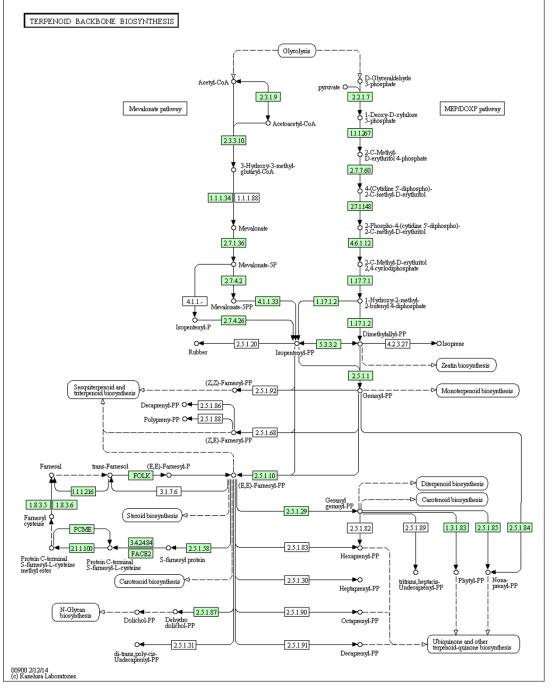
Supplementary Figure 4. KEGG metabolic pathway of plant hormone signal transduction obtained with mango Kent mesocarp transcriptome data.



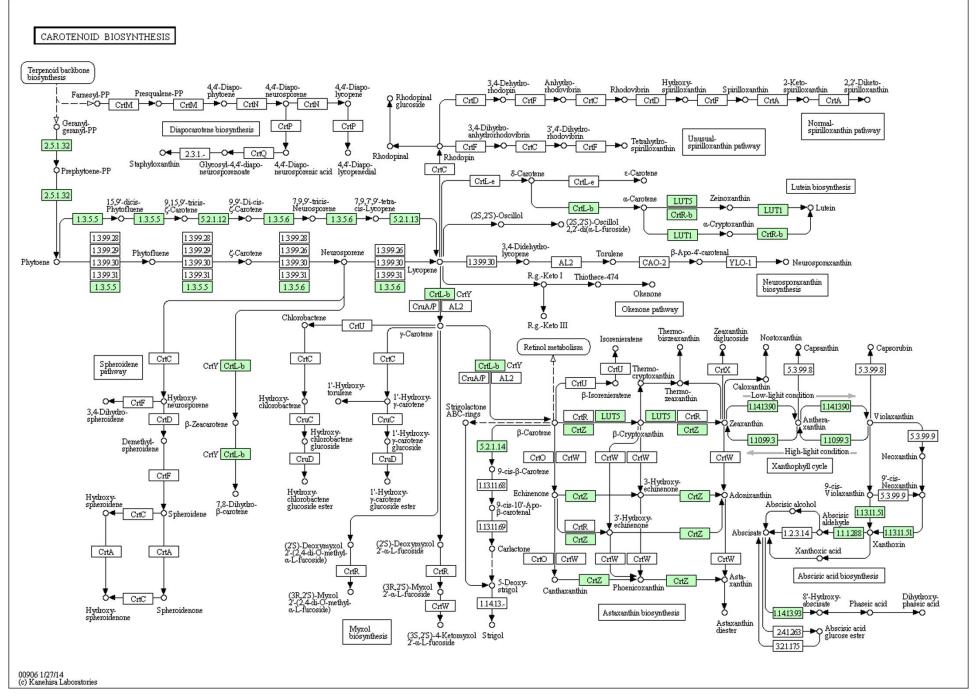
Supplementary Figure 5. KEGG metabolic pathway of terpenoid backbone biosynthesis obtained with mango Kent mesocarp transcriptome data.



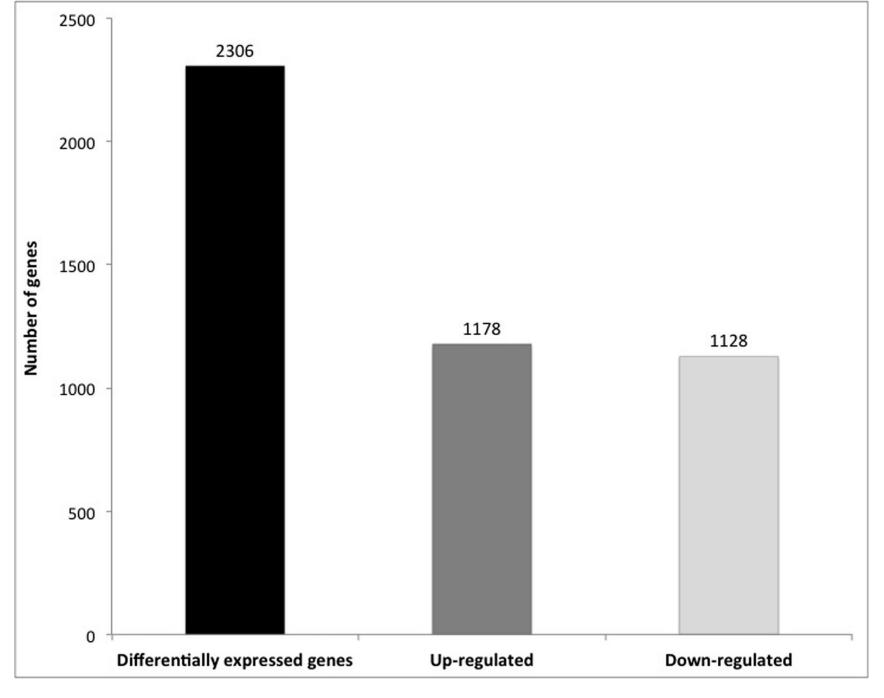
Supplementary Figure 6. KEGG metabolic pathway of galactose metabolism obtained with mango Kent mesocarp transcriptome data.



Supplementary Figure 7. KEGG metabolic pathway of terpenoid backbone biosynthesis obtained with mango Kent mesocarp transcriptome data.



Supplementary Figure 8. KEGG metabolic pathway of carotenoid biosynthesis obtained with mango Kent mesocarp transcriptome data.



Supplementary Figure 9. Differential expressed genes in mango Kent mesocarp transcriptome