## **Description of Additional Supplementary Files**

File Name: Supplementary Data 1 Description: Number of duplicated ancestral core-eudicot genes in grape and Panax stipuleanatus genomes.

File Name: Supplementary Data 2

Description: Proportion of the gene retention of the six duplicated ancestral core-eudicot chromosomes in Panax stipuleanatus.

File Name: Supplementary Data 3 Description: Collinear matrix of the protein-coding genes of the four Panax species.

File Name: Supplementary Data 4 Description: Matrix of the collinear protein-coding genes of the three tetraploid species.

File Name: Supplementary Data 5 Description: PCA scores of the compartment A/B in Panax stipuleanatus genome.

File Name: Supplementary Data 6 Description: Topological associated domains (TADs) in Panax stipuleanatus genome.

File Name: Supplementary Data 7 Description: Pattenrs of the gene expression and cytosine methylation in all 42 homologous genomic regions.

File Name: Supplementary Data 8 Description: Identification of the CYP450 gene family and their duplciation modes in Panax species.

File Name: Supplementary Data 9 Description: Information of the CYP450 superfamily in Panax stipuleanatus genome.