

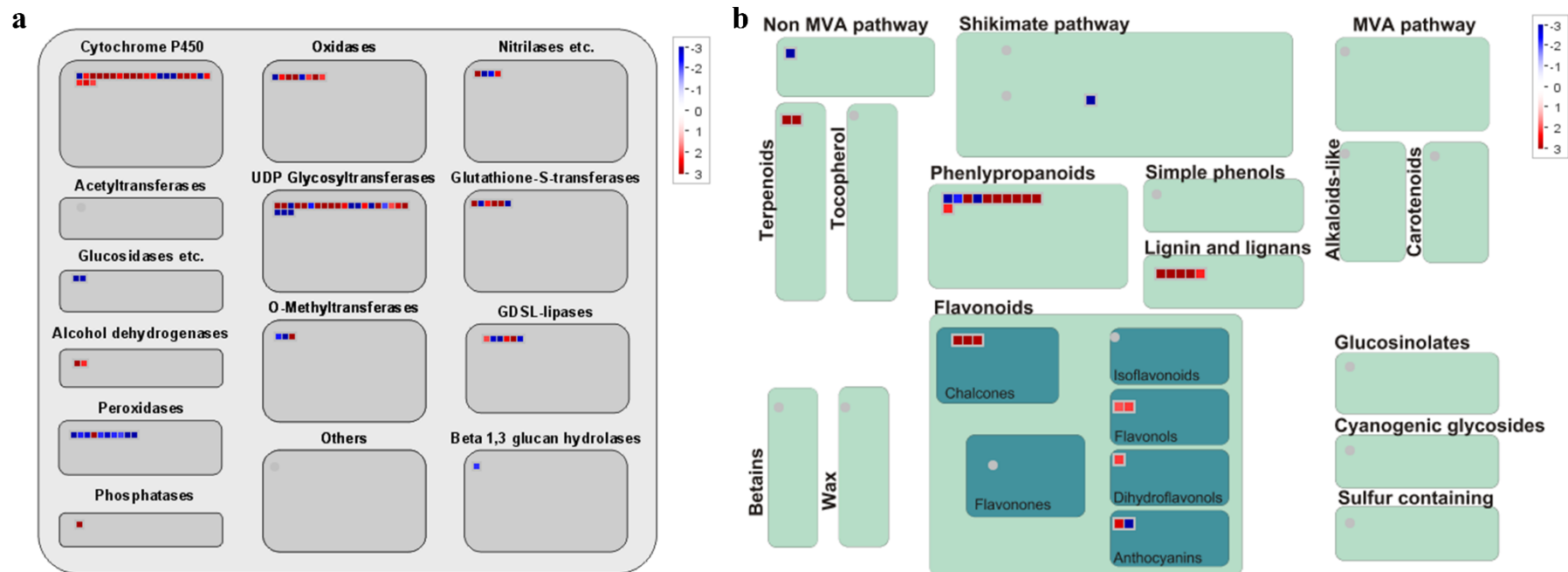


Fig. S1 MapMan overview of changes in the expression of differentially expressed genes (DEGs) in *yel-hc* developing seeds. **a** Large enzyme families. **b** Secondary metabolism. The DEGs were binned into MapMan functional categories. Data represent log₂FC values. Red and blue colors represent upregulated and downregulated genes, respectively.