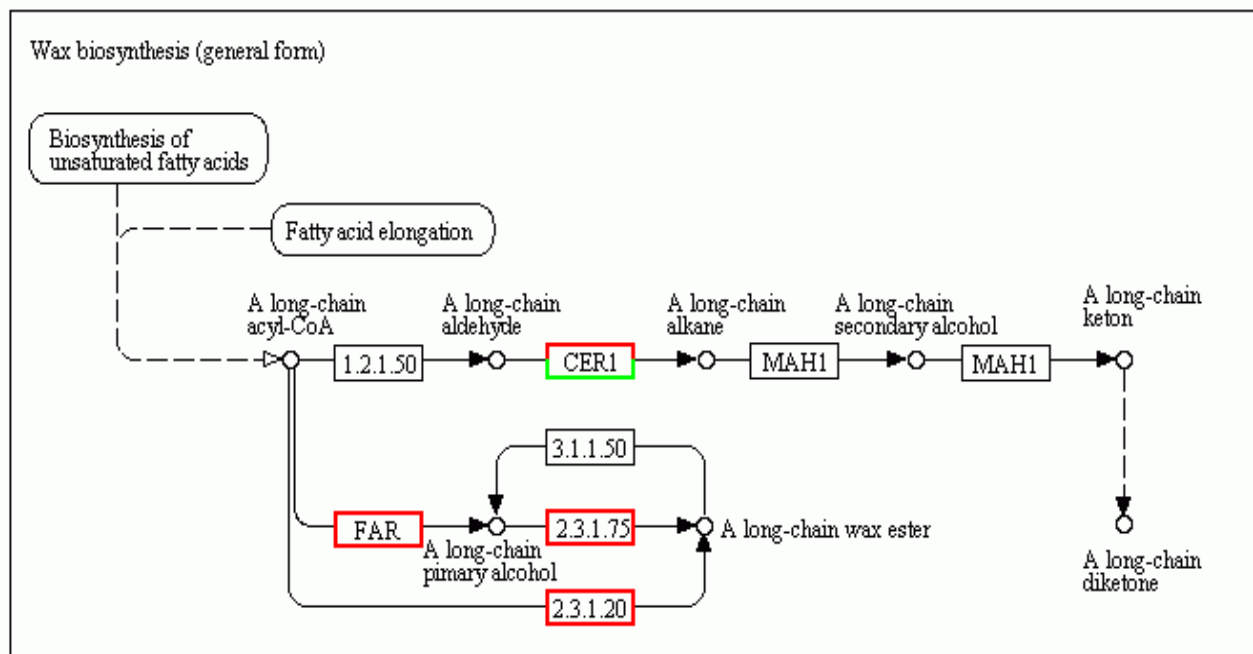
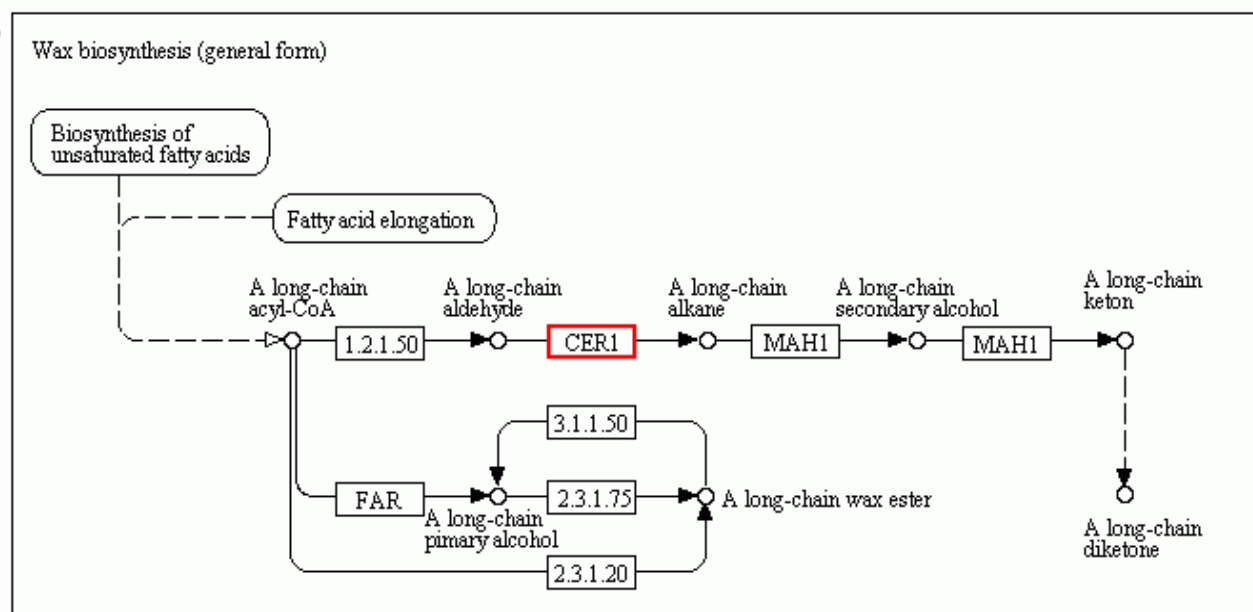


a



b



00073 8/23/11
(c) Kanehisa Laboratories

Supplementary Material 4 KEGG pathway visualization of Wax biosynthesis

a: A-VS-B; b: D-VS-E. KEGG pathway analysis of significant differentially expressed transcripts in Wax biosynthesis in Tibetan hulless barley under detached water-deficit stress. In the figure, up-regulated genes are marked with red borders and down-regulated genes with green borders. Non-change genes are marked with black borders.