

Evolutionary diversification of cytokinin-specific glucosyltransferases in angiosperms and enigma of missing *cis*-zeatin O-glucosyltransferase gene in Brassicaceae

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Figure S3. Maximum likelihood phylogenetic tree of *cis*ZOG and *trans*ZOG (UGT73C for Brassicaceae) based on combined data from Phytozome v12. Numbers below branches indicate bootstrap support values and fifteen conserved motifs are shown for each sequence.

