Figure S3 A gene network derived from co-expression analysis based on the rice functionally related gene expression network database (RiceFREND; http://ricefrend.dna.affrc.go.jp) using CYP96B3, CYP96B4, and CYP96B5 as inputs. The strength of co-expression is indicated by thickness of the lines. Detailed descriptions of the genes indicated in the network are give in Table S5.