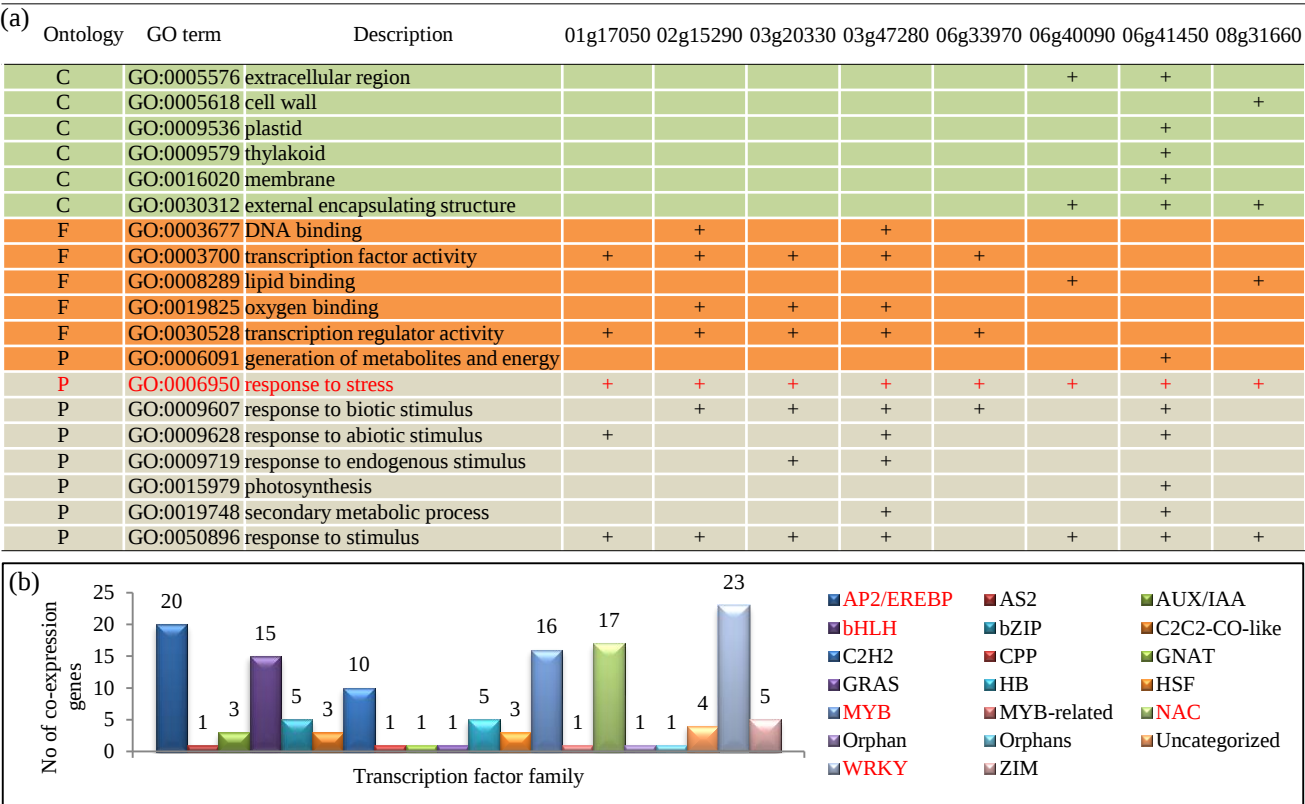




Additional file 11: Figure S5. Gene set enrichment analysis of co-expressed genes with VQs. (a) Over-representative (enrichment) GO terms among co-expressed genes with 8 VQs. The prefix “LOC_Os” in each locus name was omitted for convenience. (b) A general overview of co-expressed genes encoding transcription factors and their family names. The five family names were highlighted with red fonts, which contained 15 or more co-expressed members in each family.