

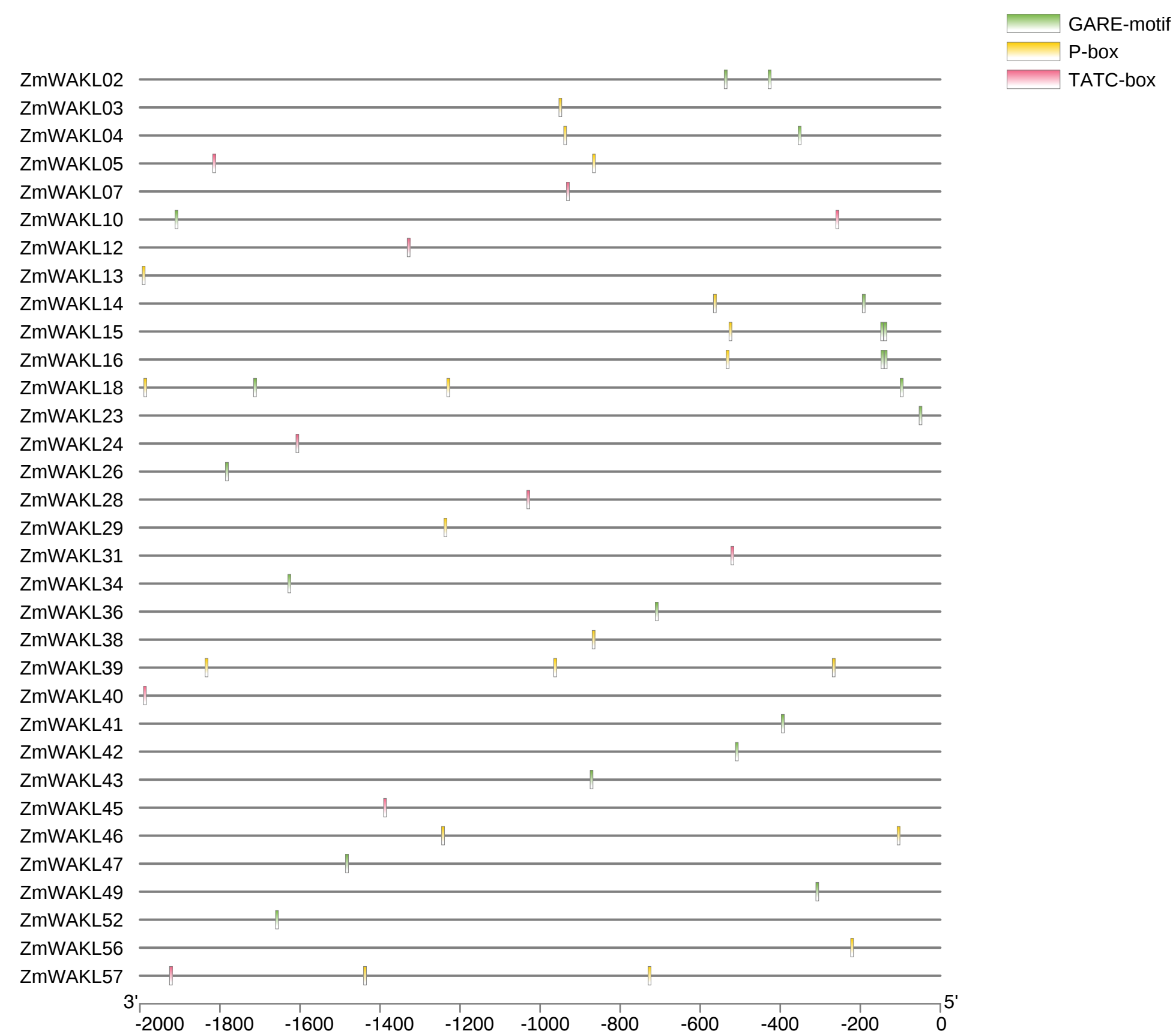


Fig S1.Prediction of GA responsive related *cis*-acting elements in 33 *ZmWAKL* promoters.

Promoter sequences (−2000 bp) of 33 *ZmWAKL* genes are analyzed by PlantCARE. The upstream length to the translation starts site can be inferred according to the scale at the bottom.