

Supplemental Table 3. Conserved motif compositions of CsWRKY proteins.

Motif	Consensus sequence	E-value	Width	Description
1	ILDDGYR <u>WRKYGQK</u> PVKGSPFP RSYYRCT	1.4E-1756	29	WRKY domain
2	QGCPVRKQVZRSEDPSIVITTYE GKHNH	4.7E-1228	29	
3	PSDDGYN <u>WRKYGQK</u> QVKGSEF PRSYYKCTHPNCPVKK	4.70E-260	37	WRKY domain
4	KKGEKKIREPRVAFQTRSEVD	1.20E-229	21	
5	KVERSJDGQITEIVYKGTHNHPK PQPNRRSTSG	5.10E-174	33	
6	KNELEVLQAE LERMNAENKRLR EMLNQICKNYNTLQMHLMDLM QQNQ	7.60E-170	47	
7	PLPPAAMAMASTTSAAASMLLS GSMSSADGLIN	1.50E-122	33	
8	VEAATKAJTADPNFTAALAAAS SIIGGG	1.00E-114	29	
9	SGRCHCSKKRKSrvkrtirvpa isklad	4.10E-109	29	
10	VEVFRQTIAGNILVGSYCAFSNK GGLVHPYTSIEDLDELSTLLQVPL VAG	3.30E-87	50	