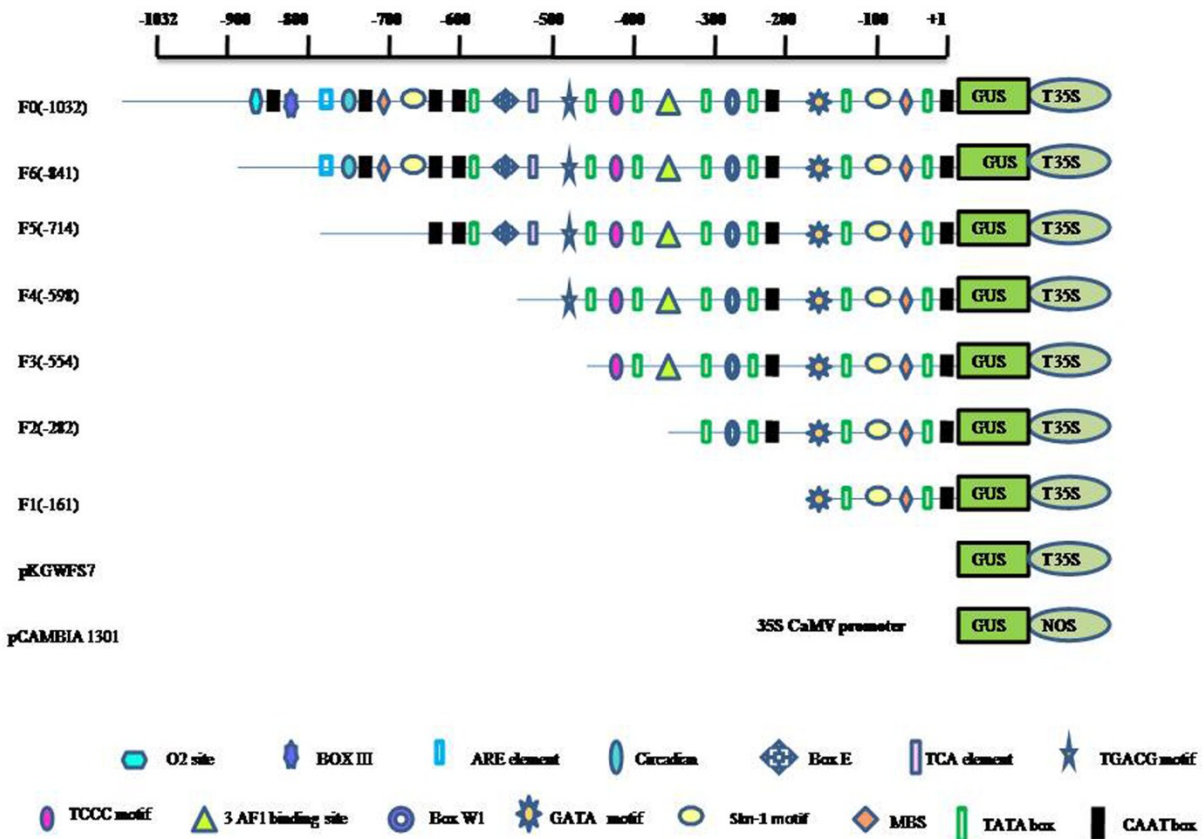


Supplementary Table I: GUS activities of different deletion fragments of promoter *PEaMYBAS1* in transient tobacco leaf discs.

Fragment	GUS activities in control and test samples				
	Mock(control)	Drought	Cold	Salt	Wounding
F0	51.96±8.97	307.00±7.68	254.12±1.99	253.51±7.98	211.11±19.84
F6	40.74±3.92	177.37±17.97	183.43±7.70	183.13±1.89	159.48±15.05
F5	35.50±21.01	124.30±21.38	132.16±8.97	148.55±4.68	121.47±8.64
F4	29.23±22.95	97.99±21.12	105.04±15.49	114.41±3.34	73.63±11.80
F3	22.17±15.73	55.72±27.40	44.58±7.47	58.59±4.68	50.16±18.43
F2	11.23±6.49	15.13±3.73	15.50±19.47	26.09±1.89	22.31±16.25
F1	8.49±2.09	11.22±9.34	9.22±15.53	17.01±7.98	10.83±9.87

Values are mean of three experiments ± SD;

Where, F1, F2, F3, F4, F5, F6 and F0 are deletion fragments of promoter *PEaMYBAS1*
GUS activity in nmol 4-PNP/mg protein/min



Supplementary Fig. I: Schematic representation of *PEaMYBAS1* promoter constructs for assaying GUS expression in tobacco leaves. The serially 5'-deleted promoter constructs of the *PEaMYBAS1* were fused to the GUS reporter gene in the vector pKGWFS7