

**Supplementary Table 1:** List of proteins used for phylogenetic analysis in this study.

Species	Protein name	Accession No. in NCBI
<i>Oryza sativa</i>	Osj3g1BTF3	NP_001048709
	Osj3g2BTF3	NP_001051911
	Osj10gBTF3	NP_001064883
<i>Sorghum bicolor</i>	SbBTF3	EER93008
<i>Arabidopsis thaliana</i>	AtBTF3	AEE29647
<i>Nicotiana benthamiana</i>	NbBTF3	ABE01085
<i>Capsicum annuum</i>	CaBTF3	ABM55742
<i>Ricinus communis</i>	RcBTF3	EEF34688
<i>Solanum lycopersicum</i>	SlBTF3	NP_001234229
<i>Triticum aestivum</i>	TaBTF3	AFV31408
<i>Zea mays</i>	ZmBTF3	ACG28870

**Supplementary Table 2:**List of Q-PCR primers, which were used in the experiment.

Gene name (Accession No.)	Primer	Sequence (5'→3')
Osj10gBTF3 (LOC_Os10g34180)	BTF3-10-F	CTCCTGGTGCTTCTGGT
	BTF3-10-R	GGCAATTCGCAACTTAT
Osj3g1BTF3 (LOC_Os03g01910)	BTF3-3-1-F	TTGCTGAGCAGTTCCAGA
	BTF3-3-1-R	CTAGGACGACTCTTTCTTCTC
Osj3g2BTF3 (LOC_Os03g63400)	BTF3-3-2-F	TCCTGGGATCATTAACCAAT
	BTF3-3-2-R	TCAGGCTTTTGTCTCCTGAG
	UBI-F	CTGTCAACTGCCGCAAGAAG
	UBI-R	GGCGAGTGACGCTCTAGTTC