

Supplementary data.

Table S1 Primers for qRT-PCR.

Primer name	Primer Sequence	Primer name	Primer Sequence
ZmTIP2;3-qPCR-F	TCGCATTGGAAGCTTCGC	ZmTIP2;3-qPCR -R	CTTCGTCAATTGCGAGTAGGC
Zma-Tubulin-F	TCCTGGACAACGAGGCTATCTAT	Zma-Tubulin-R	TGTGAGATCAGCCTGTTCAAGTT
ZmGAPDH-F	AGCAGGTCGAGCATCTCG	ZmGAPDH-R	CTGTAGCCCCACTCGTTGTC

Table S2 Primers for vector construction.

Primer name	Primer Sequence	Primer name	Primer Sequence
ZmTIP2;3-F	ATGGTGAAGCTCGCATTGG	ZmTIP2;3-R	CGAGTAGGCAATGGCGGA
ZmTIP2;3-GFP-F	ggacagccagatcaactagtATGGTGAAGCTCGCATTGG	ZmTIP2;3-GFP-R	gccttgcacccatggatccTGGGTACTCCTGCTGGCCG
pZmTIP2;3-F	CTACAACCGTACAGGGAAAAAGT	pZmTIP2;3-R	CTTTTTCCCCCCCCTCTCC
pZmTIP2;3-GUS-F	gacctgcaggcatcaagctCTACAACCGTACAGGGAAAAAGTATT	pZmTIP2;3-GUS-R	ttaccctcagatctaccatggCTTTTTCCCCCCCCTCTCC

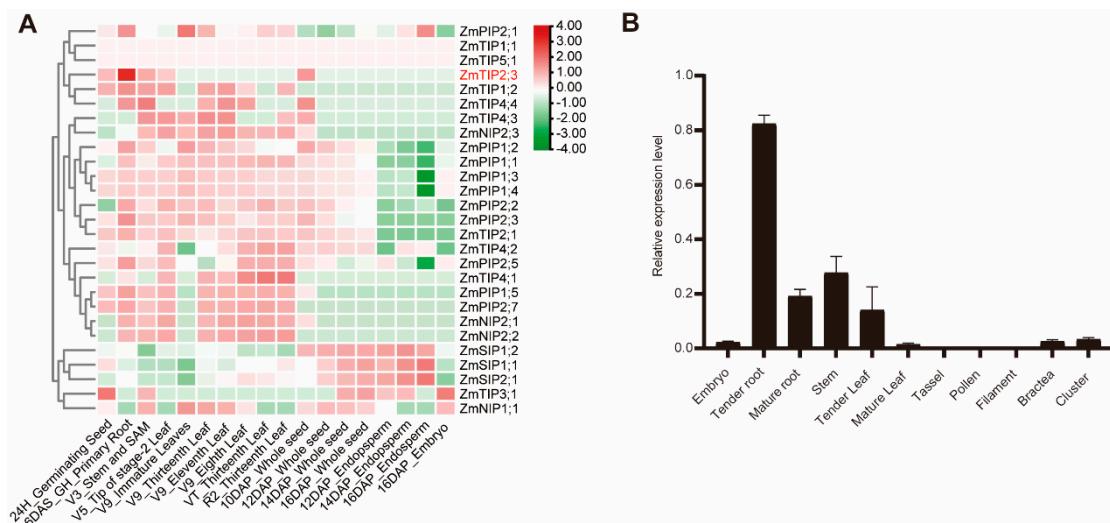


Figure S1. Tissue expression pattern analysis of *ZmTIP2;3*. (A) Heatmap of the tissue expression pattern of *ZmTIP2;3*. (B) The relative expression level of *ZmTIP2;3* in different tissues of maize was measured by qRT-PCR.