

**Table S1. Primers for all manuscript.**

Experiment	Annotation	Forward sequence	Reverse sequence
P1	Reverse Transcription-qPCR (RT-qPCR)	CGGTGAAGTTGGGATGTG	TTTTGTTTCTCCGTAAGGC
P2	Full length of sequence	GGATAACAACATAGAAAC AGT	TAAACATCAGAAACGAAAAA GA
P3	<i>In situ</i> hybridization	AAACTGCAGAAGTCGGAGG ACAGAACAA	GGGGGTACCCAGCGGTCACA AGAACAG
P4	Subcellular localization	CAGTGGTCTCACAACATGA AAATCTTTAACCCATC	CGATGGTCTCAAATGATGGTT TTAATTTTGTG