

Table 1 Suppl. PCR primer sequences used for cloning and gene expression assays. FW - forward, RV - reverse.

Primer name	Sequence (5'-3')
<i>Apx1</i> real time FW	ATTCAGATGCCCGAGAAGTCTTGTTCC
<i>Apx1</i> real time RV	ACCACCGATCCAGACACTGTACTTCC
<i>Actin</i> real time FW	GTCCATTTCGGAACAATGAGGTTTGAC
<i>Actin</i> real time RV	GTGGGCACCAGATAAAGCGACAAT
<i>Apx1 Pst I</i> -FW	AACTGCAGATGACGAAGAAGTACCCA
<i>Apx1 Bln I</i> -RV	ATCCTAGGAGCATCAGCAAACCCGAG
OE <i>AtApx1</i> -FW	GCCCTGACATTCTTTCCAC
OE <i>AtApx1</i> -RV	AGTTCACCTTGATGCCGTTTC
OE <i>TsApx1</i> -FW	TGAGAAATACGCTGCTGACGA
OE <i>TsApx1</i> -RV	AAGTTCACCTTGATGCCGTTTC
Hyg-FW	GAAAAAGCCTGAACTCACCGC
Hyg-RV	TGCTCCATACAAGCCAACCAC

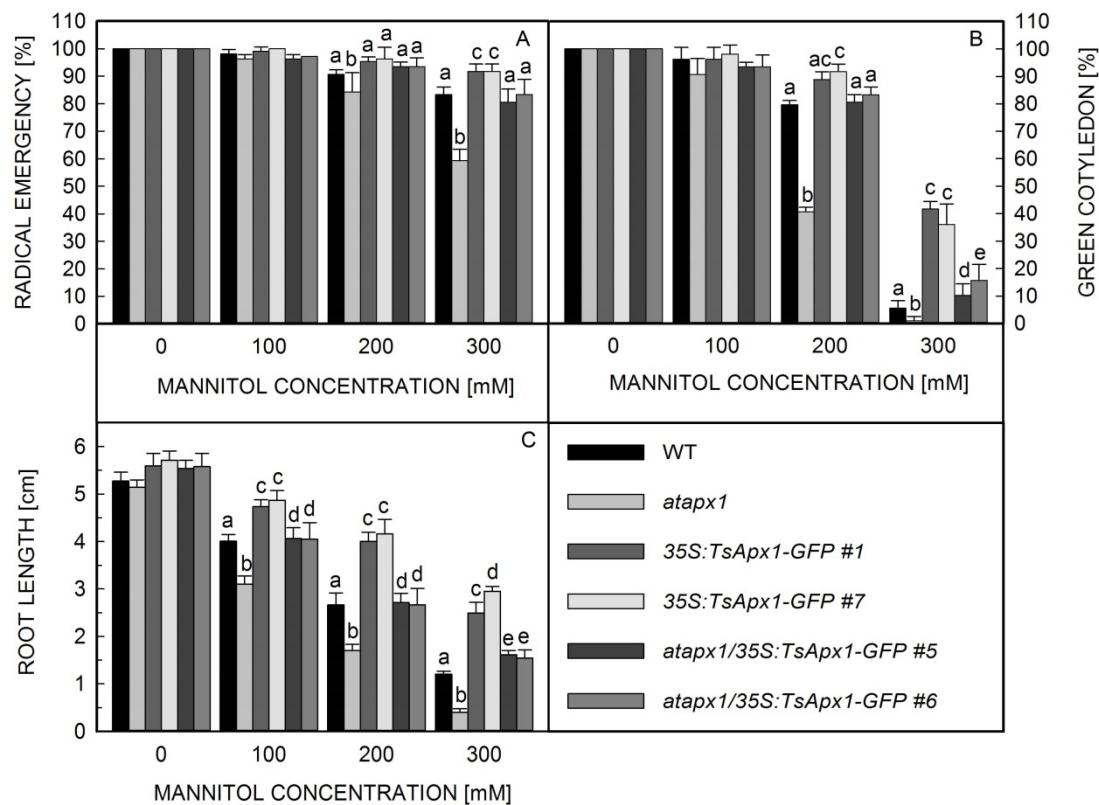


Fig. 1 Suppl. Effect of mannitol (0, 100, 200, and 300 mM in MS medium) on radicle emergence (A), cotyledon greening (B) and root length (C) of wild-type (WT), *atapx1* loss-of-function mutant, and 35S:*TsApx1*-GFP and *atapx1*/35S:*TsApx1*-GFP transgenic plants at the germination stage (10-d-old). Means $\pm$ SD, $n = 144$ for germination and greening and 30 for root length. Means followed by different letters are significantly different at $P < 0.05$.