Table 1 Suppl. Primer sequences used in the experiments.

Primer	Sequence (5'-3')
P1	5'-CGGGGGACTCTTGACCATGGATGGGGTCTGGTATGGAGGA-3'
P2	5'-CGGGGAAATTCGAGCTGGTCACCTCATTCCCATCCATGAACAC-3'
P3	5'-CTCGAGCTCAAGGAATTCATGGGGTCTGGTATGGAGGA-3
P4	5'-GGTGGCGACCGGGTACCTTCCCATCCATGAACAC-3
P5	5'-ATACGTGGCACAATGTGACAGAG-3'
P6	5'-GATAGGAAAGACAGCGGGAAGAC-3'
P7	5'-ATTTGCTGGATTGCTTAGTGCGTA-3'
P8	5'-ACTTGCTGGTTGAAGGTGATTTTT-3'
A1	5'- TATGCTAGTGGTCGTACAACTG -3'
A2	5'- AACAACCTTAATCTTCATGCTG -3'

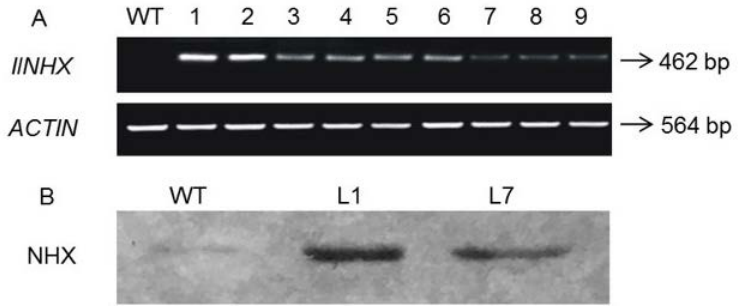

Fig. 1 Suppl. *A* - Reverse transcription (RT-PCR) analysis of *Iris lactaea tonoplast* Na^+/H^+ *antiporter* (*IlNHX*) in wild type (WT) and transgenic tobacco lines (1 - 9) overexpressing *IlNHX*. Specific PCR products of *IlNHX* of 462 bp were detected in the leaves of the tobacco plants. A 564 bp *ACTIN* fragment was amplified by RT-PCR as an internal control. *B* - Western blot analysis of IlNHX in WT and transgenic (L1 and L7) tobacco plants under 200 mM NaCl treatment for 10 d.