

Table S1. Comparison of the partial sequences of genes in *Arabidopsis* and *Thellungiella* and among *Thellungiella* ecotypes

Expected real-time PCR product and its flanking nucleotide sequences were amplified by regular PCR from cDNA of *Arabidopsis* and *Thellungiella* ecotypes. Sequences were aligned and similarities between species are shown.

Gene	Product size (bases)	Similarity in <i>Arabidopsis</i> and <i>Thellungiella</i> (%)	Similarity in <i>Thellungiella</i> ecotypes Shandong and Yukon (%)
<i>NRT1.1</i>	400	90	100
<i>NRT2.1</i>	650	90	99
<i>NRT3.1</i>	550	80	100
<i>CLCa</i>	440	90	100
<i>NLA</i>	800	98	100
<i>NR2</i>	405	90	100
<i>GS1.1</i>	540	93	100
<i>GS2</i>	524	100	100
<i>UBQ10</i>	506	100	100