

Transcriptome-Wide Analysis Revealed the Potential of the High-Affinity Potassium Transporter (*HKT*) Gene Family in Rice Salinity Tolerance via Ion Homeostasis

Table S1. Primers information of *OsHKT* genes for qRT-PCR expression analysis in rice.

Genes	Primer sequences
<i>OsHKT1</i>	F: TGAAGCCAAGCAACCCAGAA R: AGGCTGGAAAGTGTCTCAGAGC
<i>OsHKT3</i>	F: AGAGGCGACTCCCCAAACT R: TGATCATGTATGCCAGCACCA
<i>OsHKT4</i>	F: GGTGCCTCGGATCTATTGGG R: TCAGAGCAAGGAAACCAGCA
<i>OsHKT6</i>	F: GTCCTCAAGCCACGAGACAA R: CCAGAGTTGAGCGCTTGAGA
<i>OsHKT7</i>	F: ACCAGGGAATCAGACTGCTC R: CCATATGCACTGACAACTTCGAC
<i>OsHKT8</i>	F: CACCCATTCTGGCTCCAACT R: ATCGGCAGAGCTTTCAGCAT
<i>OsHKT9</i>	F: AGTGCCTACGGGAACATTGG R: CACCCCGAGAAGCTGTATGG