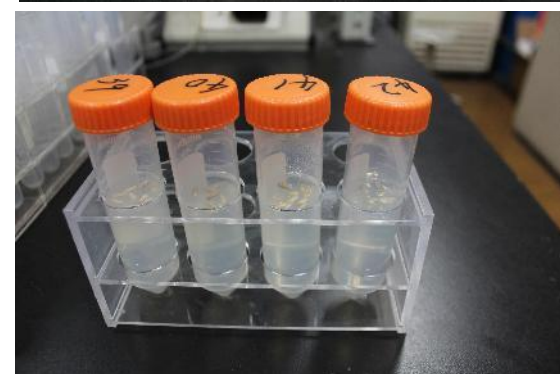
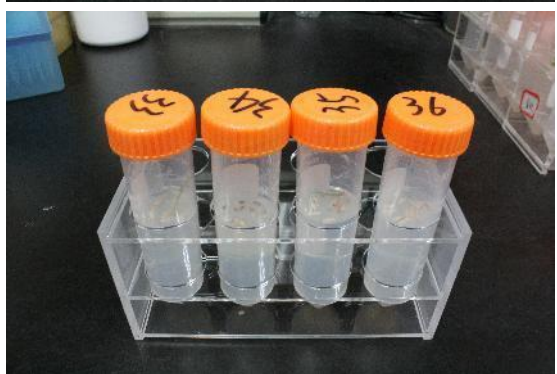
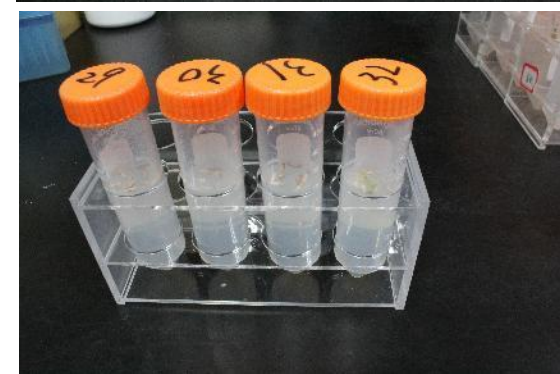
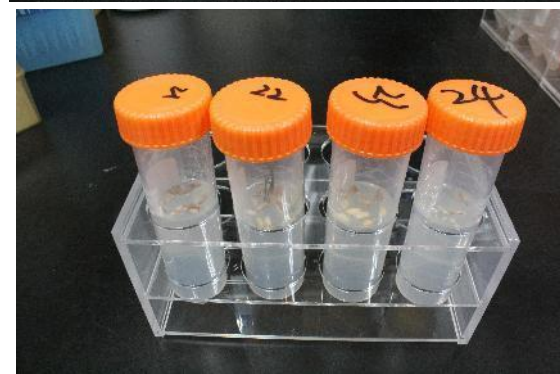
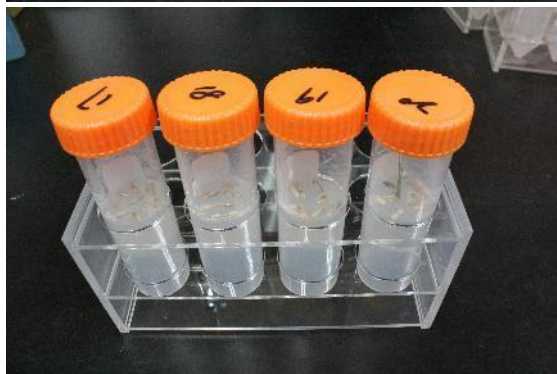
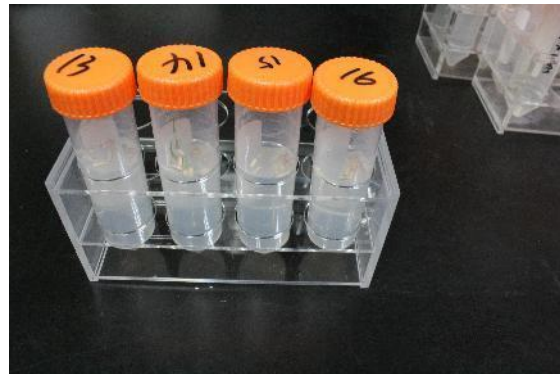
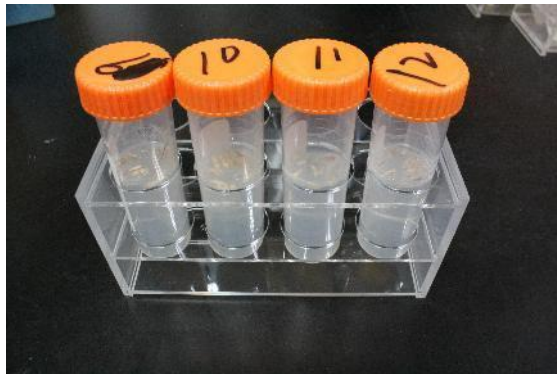
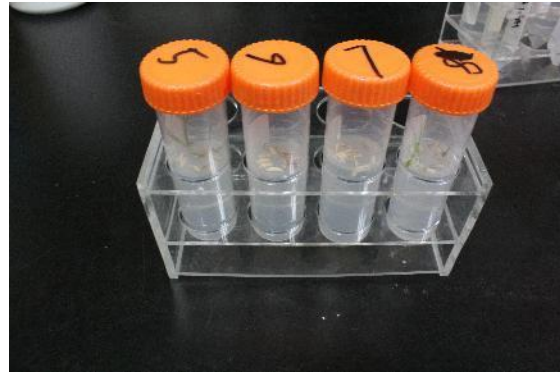
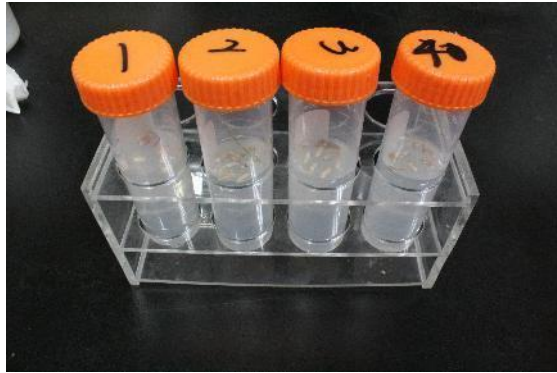
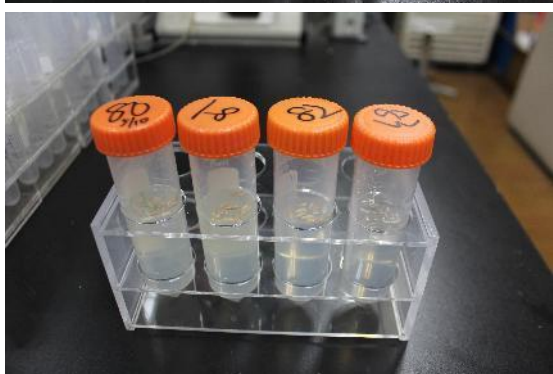
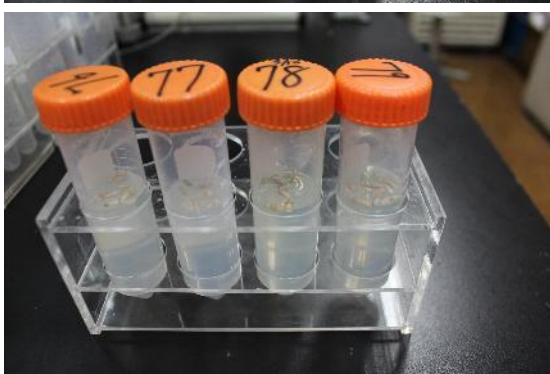
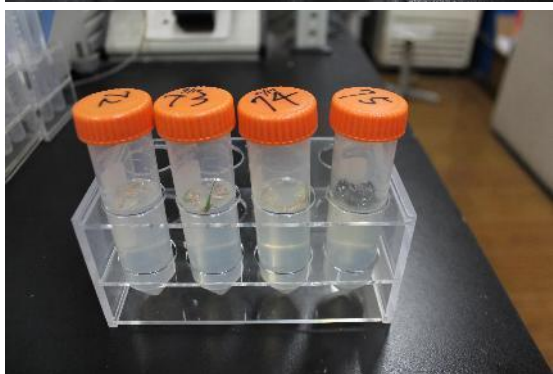
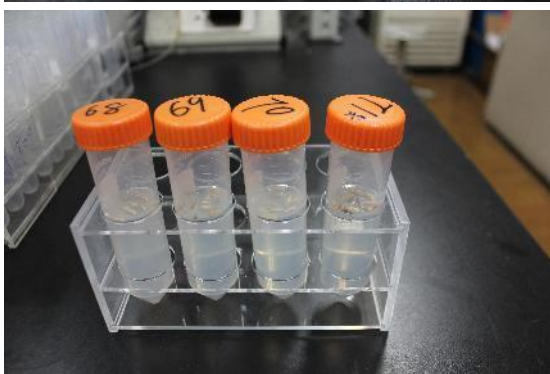
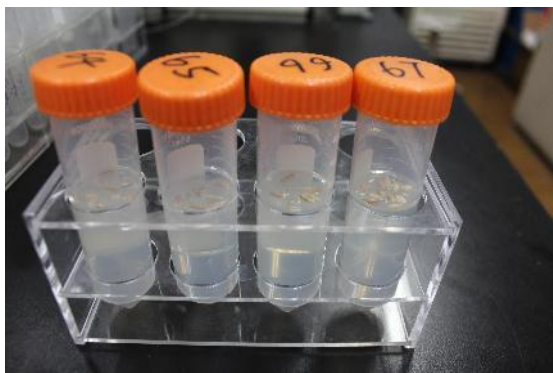
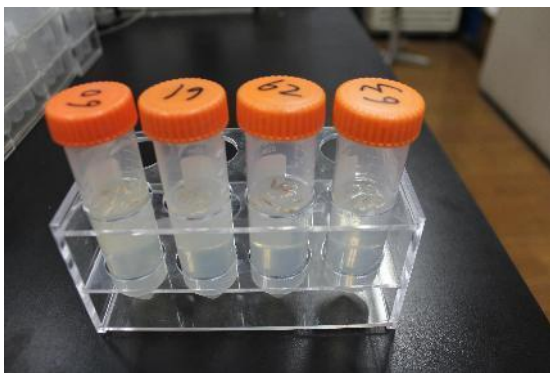
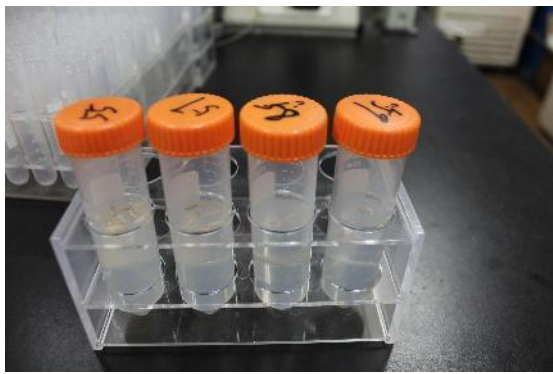
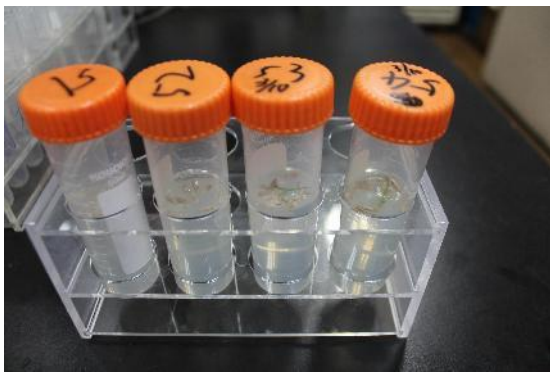
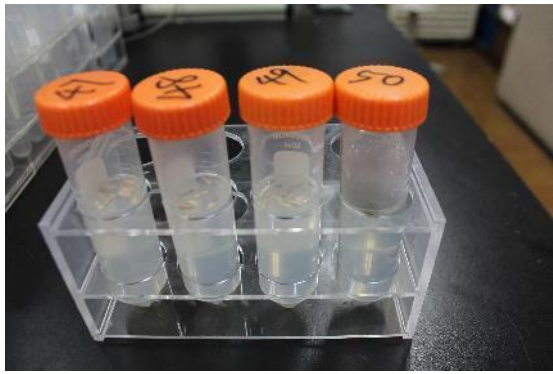
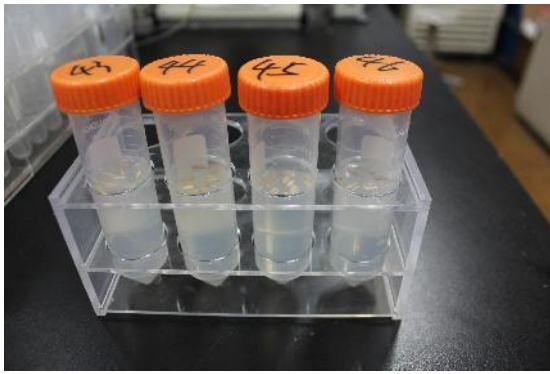
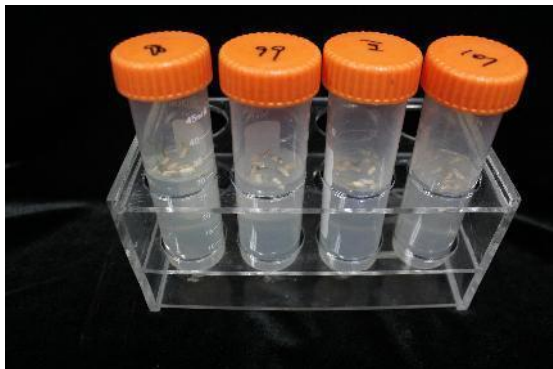
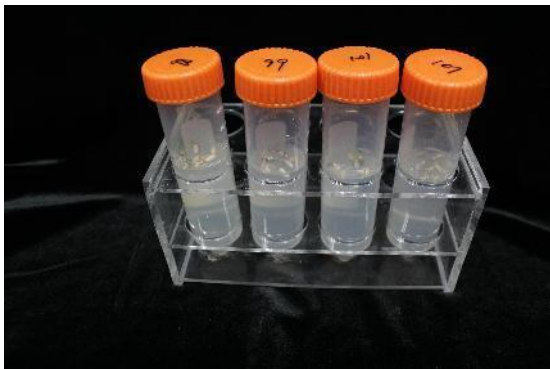
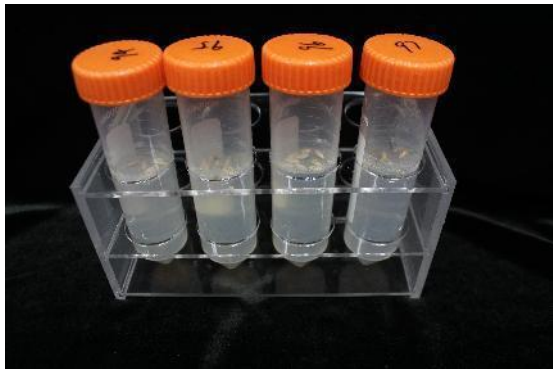
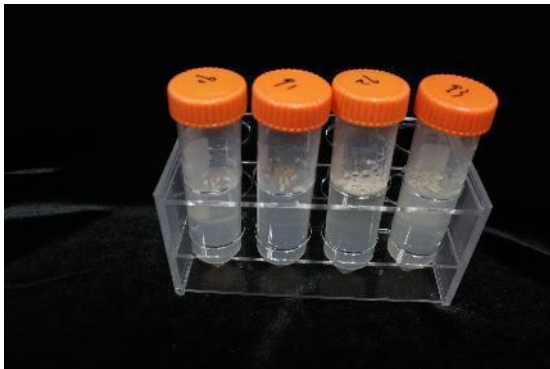
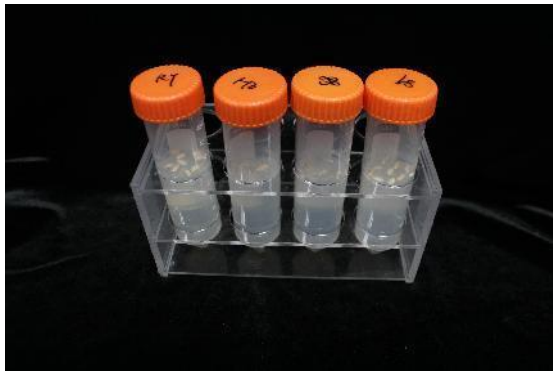
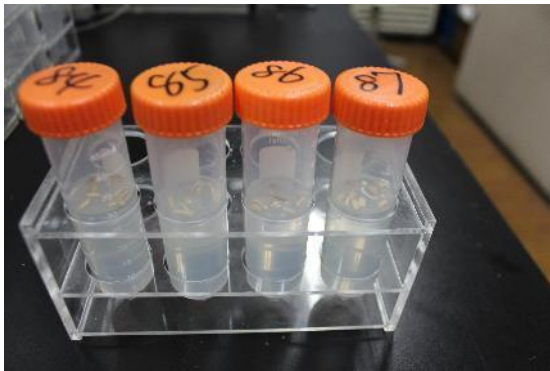


ADDITIONAL FILE FIGURE TABLE







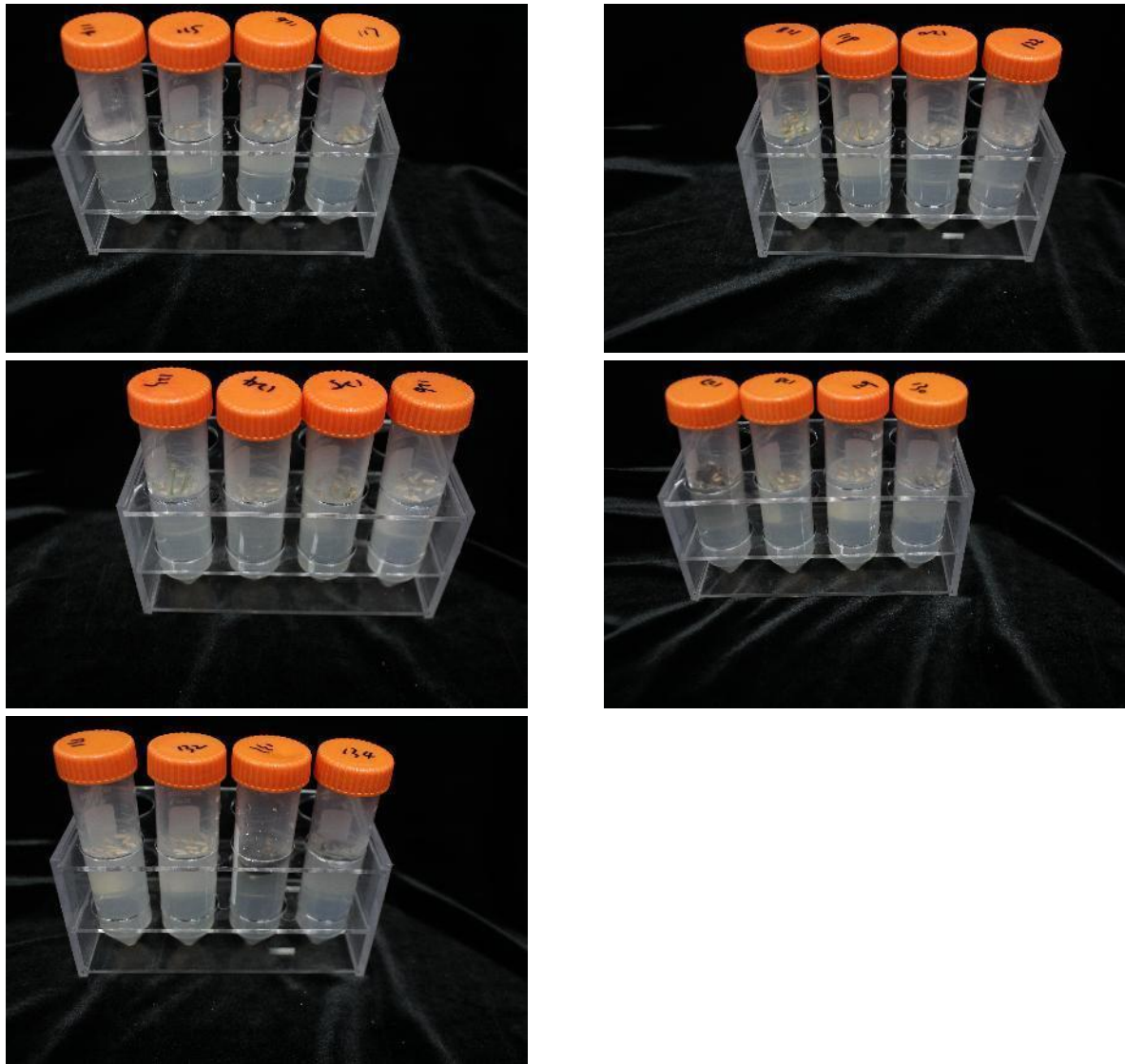


Figure S1. Phenotypes of both parents and RIL populations treated with 80mM NaCl.

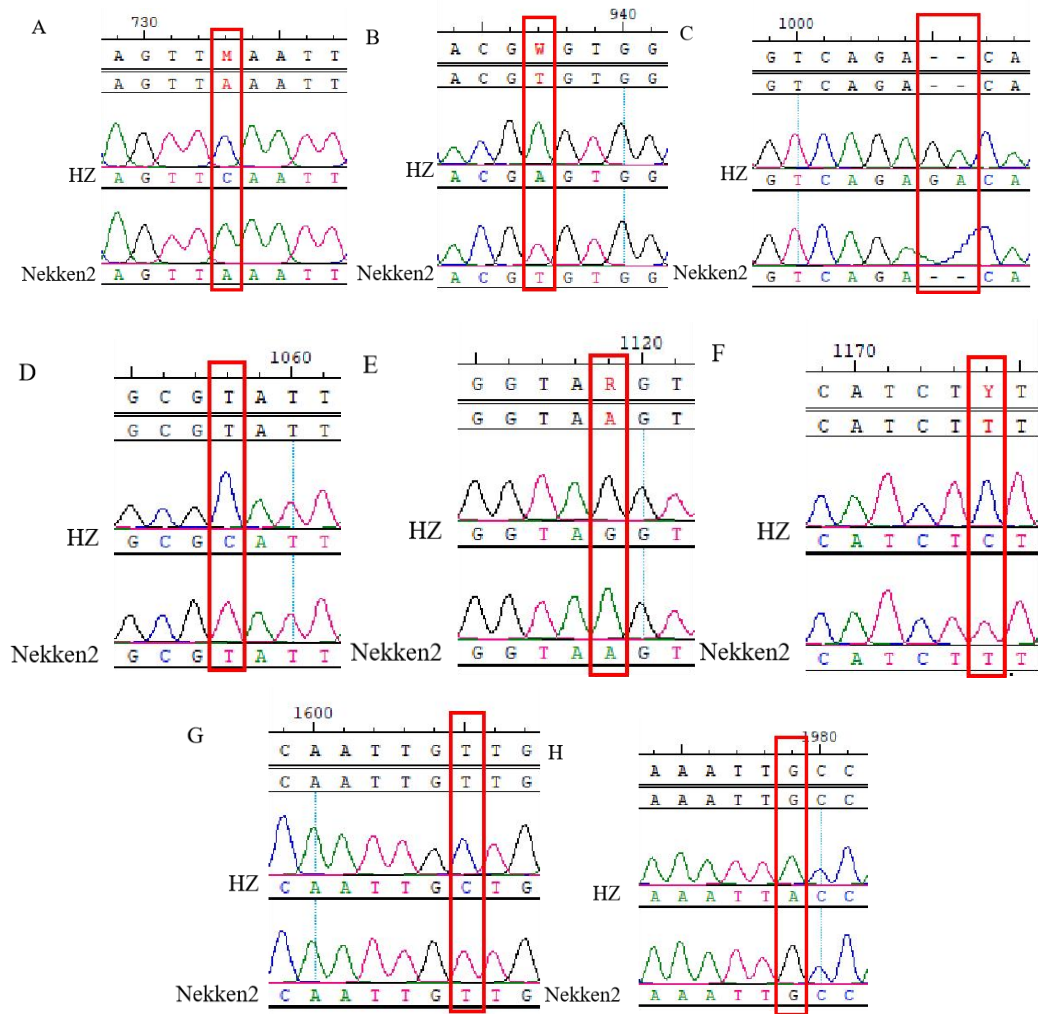


Figure S2. Sequencing of *LOC_Os12g25200* promoter region.

Table S1. SSR markers selected to identify the CSSLs.

Chromosome	SSR markers
Chr.01	RM3252, RM1247, RM3740, RM6902, RM8111, RM6786, RM583, RM580, RM259, RM493, RM6880, RM8097, RM6703, RM3738, RM1361, RM8100, RM6321
Chr.02	RM7581, RM300, RM262, RM3515, RM6318, RM1920, RM3763, RM6933, RM3316
Chr.03	RM14795, RM6676, RM3513, RM1350, RM3856, RM5172
Chr.04	RM8213, RM307, RM16874, RM1155, RM6997, RM3524, RM3839, RM303, RM317, RM349, RM1113
Chr.05	RM17713, RM17798, RM13, RM17954, RM18384, RM3437, RM430, RM3695, RM18751, RM3476, RM6972, RM3616, RM3809, RM334, RM19221
Chr.06	RM19234, RM585, RM217, RM276, RM19715, RM2523, RM3183, RM3496, RM162, RM5371, RM6395, RM3343, RM412
Chr.07	RM5490, RM5055, RM5711, RM6574, RM3755, RM6835, RM11, RM3826, RM1132, RM234, RM3555, RM1306
Chr.08	RM1019, RM6356, RM3819, RM5556, RM7285, RM5808, RM556, RM80, RM3571
Chr.09	RM444, RM219, RM1328, RM6051, RM3700, RM242, RM201, RM3744
Chr.10	RM6370, RM2125, RM4455, RM5689, RM1375, RM1873, RM6737, RM3451, RM4771
Chr.11	RM286, RM167, RM4504, RM6272, RM6094
Chr.12	RM5795, RM247, RM101, RM1337, RM1986, RM7376, RM1103, RM5479, RM3739, RM1226

Table S2. Primers used in the study.

QTL	Gene ID	Forward primer (5'→3')	Downstream primer (5'→3')
	<i>OsActin</i>	TGGCATCTCAGCACATTCC	TGCACAATGGATGGGTCAGA
<i>qST3</i>	<i>LOC_Os03g12820</i>	CCGTCGACAACCTTGAGAAT	AGGGAACATGTCGCGAGTAG
<i>qST3</i>	<i>LOC_Os03g12840</i>	ACGAGGAGGAAGTGGTGATG	ATCAAATGGGCCTTGTTCTG
<i>qST9</i>	<i>LOC_Os09g37949</i>	TCCACGAGATTGAGTTGCTG	ACACTGATCGGCAGATTTCC
<i>qST9</i>	<i>LOC_Os09g38000</i>	TCCGGAGTGGTATTTCTTCG	GACCAGCGTCTTCTTCATCC
<i>qST10.2</i>	<i>LOC_Os10g01480</i>	ACTAACGTTGCCCTTGGTTG	TGCCATCATTGTTTTCCAGA
<i>qST10.2</i>	<i>LOC_Os10g01760</i>	TTCCTCCACCTCGTCTCTTC	GGCACGTCTCGTCGTAGAAT
<i>qST11.1</i>	<i>LOC_Os11g10590</i>	CACTAAGATGGCGGTGGTG	GGACGAAACATGCAAAAACC
<i>qST11.1</i>	<i>LOC_Os11g10640</i>	TGGAACATCACCTGTTTGGGA	TGTATGAATCGTGCGGGTTA
<i>qST12.1</i>	<i>LOC_Os12g10660</i>	GTCATCGTCAGCCCAACAAG	GCTGGAGGAGTGAGGTTTCAG
<i>qST12.1</i>	<i>LOC_Os12g10740</i>	CCTTGAGCAAGCTCCAAAAC	GGAAGTCCCTCCAGTGAGCTG
<i>qST12.2</i>	<i>LOC_Os12g12860</i>	CTGGGAGAAGCAGTCAGGTC	AAGTGCCCCTTCTTTTTGGT
<i>qST12.2</i>	<i>LOC_Os12g13570</i>	AGTGCTGCCAAAAGCTTCAT	TCGCTCAATACGCAAACTG
<i>qST12.3</i>	<i>LOC_Os12g24800</i>	ATGGAAACGAGGATAGTGGT	CTTATTGTTGTGCGAGAAGT
<i>qST12.3</i>	<i>LOC_Os12g25200</i>	AGGAGATGGGGATGTACGTG	ATCTCGGGTCCTTGGA ACTT