

Supplementary Table S1. Shoot length of LYP9 and two parents under control and salt stress conditions

Traits	CK			7 days			14 days		
	9311	LYP9	PA64s	9311	LYP9	PA64s	9311	LYP9	PA64s
SL(cm)	30.3± 2.0	34.5± 2.4**	28.6± 1.9	23.0± 1.4	27.8± 2.6**	24.6± 1.9	17.5± 1.0	24.1± 1.5**	19.8± 1.4
SFW(mg)	1230.3 ±10.5	1704.7 ±12.5**	1089.7 ±10.4	853.5± 10.1	978.0± 9.6**	881± 7.4	376.4± 5.0	510.0± 5.1**	400.4 ± 4.8
SDW(mg)	411.6± 9.3	479.7± 10.3*	356.5± 9.5	110.4± 5.7	190.3 ± 7.2**	166.6± 7.1	56.2± 6.8	91.4± 7.2**	68.9± 5.5
RL(cm)	10.9± 0.5	13.9 ± 0.3**	11.2± 0.3	8.4± 0.4	12.2± 0.5**	10.1± 0.8	6.7± 0.4	9.1± 0.7**	7.8± 0.3
RFW(mg)	546.5± 11.8	630.2± 13.1**	526.7± 13.5	306.3± 11.3	460.4± 9.1**	436.7± 9.4	213.6± 4.1	360.3± 4.2**	273.3± 3.3
RDW(mg)	66.7± 2.8	83.3± 5.7**	60.2± 3.7	50.7± 2.1	61.7± 3.7*	56.7± 3.4**	34.2± 1.2**	50.5± 3.0*	46.7± 1.7

Values are denoted as mean ± SD (n = 5). Differences between mean values of treatments and controls were compared using t-tests (*P < 0.05, **P < 0.01) according to LSD test.

Supplementary Table S3. Primers for RT-qPCR are listed

Marker Name	Sequence
1F	GGTCCAGAGCGGTATAGTGAAC
1R	GTCATTCTGCTTGTCCCGCT
2F	AGGGAGAACTTAAGCGGGAC
2R	GCCTGTCAGGTTCTAGCAAGTT
3F	TGGGAACTTGCTAGAACCTGAC
3R	ATGCTGTTATCGTCGTGCAG
Histone F	GGTCAACTTGTTGATTCCCCTCT
Histone R	AACCGCAAAATCCAAAGAACG