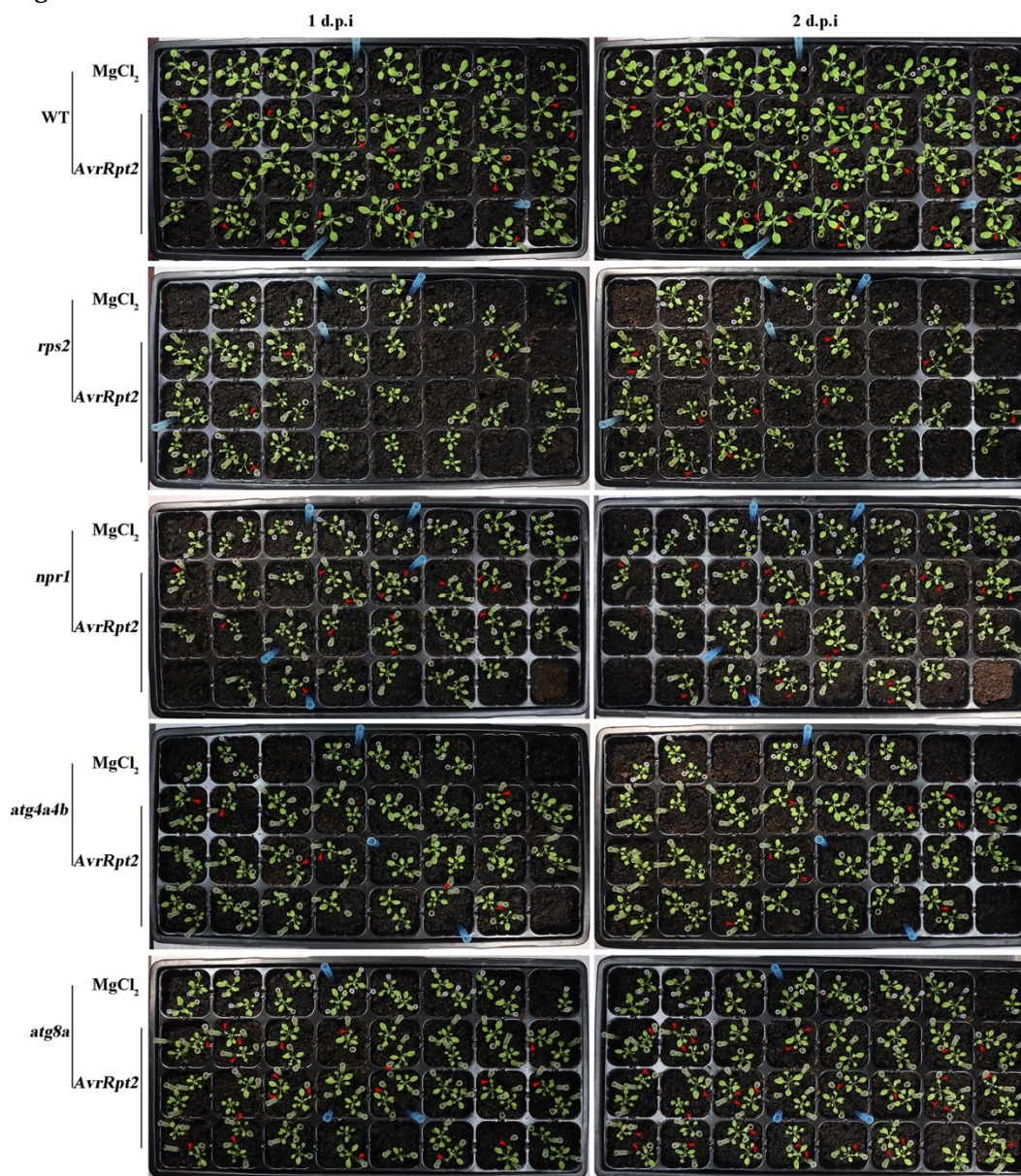




Supplementary data

Figure S1



phenotypes of WT, *rps2*, *npr1*, *atg4a4b* and *atg8a* after *Psm ES4326/avrRpt2* infiltration for 1 or 2 days. The red arrow showed the leaves with HR phenotype.

Table S1 Primers for several genes.

gene name	sequence (5'→3')
<i>AtATG1</i> (AT3G61960)	F: ATTTGAATTCACCTGCACCA
	R: GGCAGCACTTGTTTCGTTTA
<i>AtATG6</i> (AT3G61710.1)	F: TTGCAAATTCAAAGGACCAA

	R: TGGTCCAACCTCTCTTGCTTG
<i>AtATG8a</i> (AT4G21980.1)	F: AAGCTTGGAGCTGAGAAAGC
	R: GGAACCCATCCTCATCTTTG
<i>ICS1</i> (AT1G74710.2)	F: GGCAGGGAGACTTACG
	R: AGGTCCCGCATAACATT
<i>NPR1</i> (AT1G64280.1)	F: CAATTCATCGGAACCTGTTG
	R: GAGGAGTCGGTGTTATCGGT
<i>PR1</i> (AT2G14610)	F:CTCAAGATAGCCCACAAGAT
	R:GCGTAGTTGTAGTTAGCCTTCT
<i>EDS1</i> (AT3G48090)	F:CCAATTGGATCCCAGAAAGT
	R:AACAGCTTGGTTTGCAACAG
<i>ubiquitin 5</i>	F:GCCGAAGAAGATCAAGCACAAGCA
	R:TCGATGGATCTGGAAAGGTTTCAGCGT
<i>AtACTIN2</i>	F: GGTAACATTGTGCTCAGTGGTGG
	R: AACGACCTTAATCTTCATGCTGC
