

Figure S1 Osa-miR444b.2 enhanced rice susceptibility to sheath blight in YSBR1.

A Disease phenotypes at different time points after infected by *R. solani*. Scale bar, 10 cm.
B The lesion length of knockout materials at different time points after infected by *R. solani*. The lesion length of overexpression materials at different time points after infected by *R. solani*. a presents the largest average, bcd present significance ($P<0.01$).

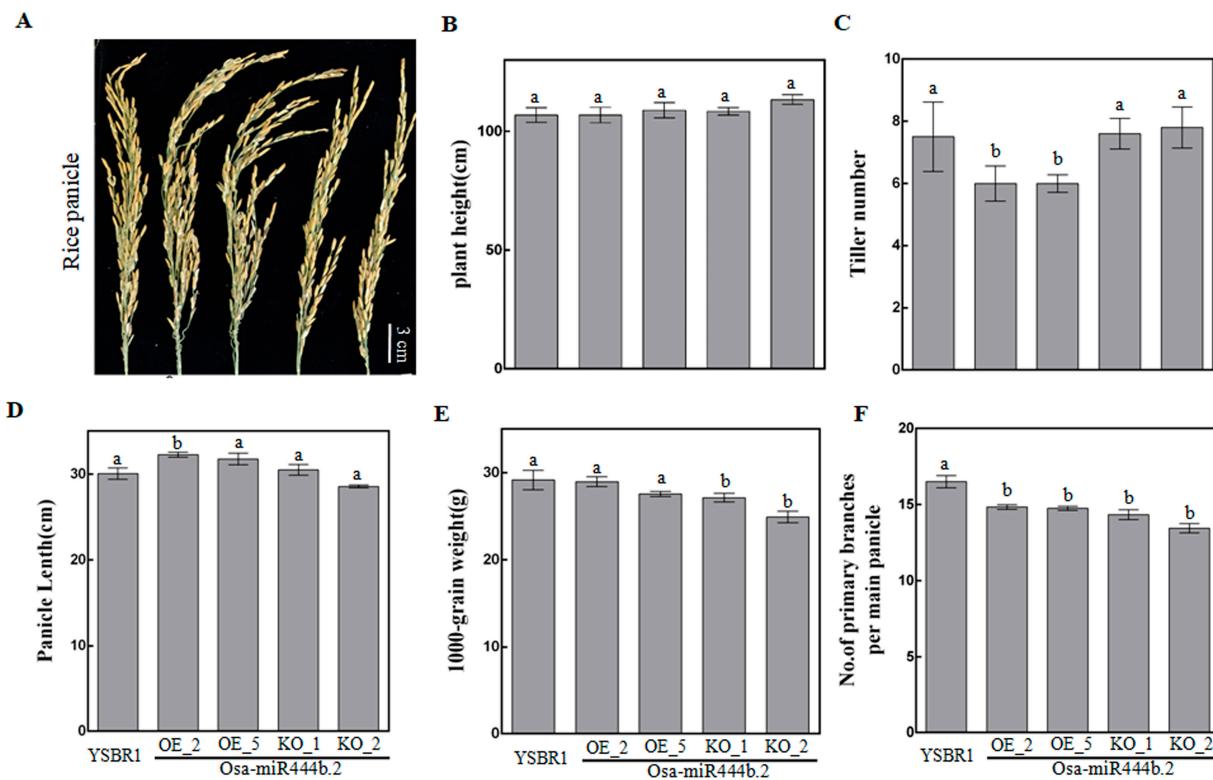


Figure S2 Osa-miR444b.2 expression level affects the agronomic traits in YSBR1.

A Panicle morphology of Osa-miR444b.2 knockout and overexpression materials. **B** Statistics of the plant height. **C** Statistics of the tiller number. **D** Statistics of the panicle length. **E** Statistics of the 1000-grains weight. **F** Statistics of the number of primary branches per main panicle. a presents the largest average, b present significance ($P<0.01$).

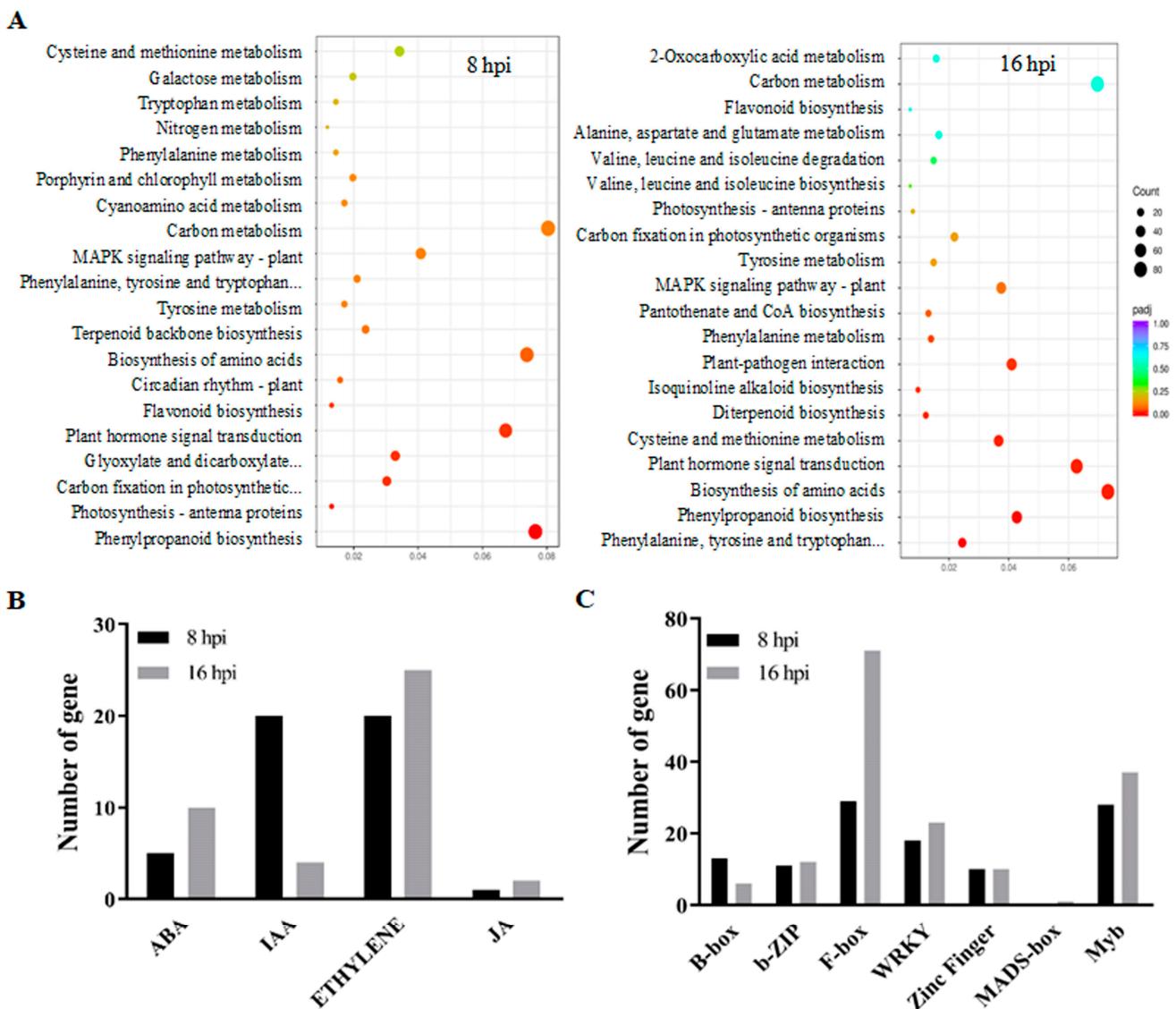


Figure S3 Osa-miR444b.2 participates in the response to *R. solani* infection by regulating plant hormone signal transduction pathway and transcription factors.

A Top 20 enriched KEGG pathways of differentially expressed genes in differential time.
B The number of differentially expressed genes in plant hormone signal transduction. **C** The number of differentially expressed genes of TFs.

Table S1 Table Primers used in this study.

Purpose	ID	Sequence (5'→3')
Reverse	miR444- stemloop	GTCGTATCCAGTGCAGGGTCCGAGGTA
Transcription primer		TTCGCACTGGATACGACAAGCTT
qRT-PCR	miR444-RT-F	GCGGCGGTGCAGTTGTTGTC
	Universal reverse primer	GTGCAGGGTCCGAGGT
	Os18s RNA-F	CTACGTCCCTGCCCTTGTACA

<i>Os18s</i> RNA-R	ACACTTCACCGGACCATTCAA
<i>Os12g26290</i> RT-F	GACCTCCGTATGTTTCTC
<i>Os12g26290</i> RT-R	CAGCATTATGGGGTCGTG
<i>Os07g03730</i> RT-F	CGGAGAAGCAGTGGTACGAC
<i>Os07g03730</i> RT-R	GCGAGTAGTTGCAGGTGATG
<i>Os08g38990</i> RT-F	GTGCCAACCAAGAAATCAAGC
<i>Os08g38990</i> RT-R	CAGTGGTAGGAGAAGGTTGT
<i>Os12g36850</i> RT-F	GCTGTGGAAGGTCTGCTTGG
<i>Os12g36850</i> RT-R	CCAACACCTCAACCTTACGC
<i>Os11g37950</i> RT-F	CTACGTGGACCGCTAACAAAG
<i>Os11g37950</i> RT-R	CTTGGTGAAGACGGTGTCCC
<i>Os09g25070</i> RT-F	CGTGACAACCCCTACCTAG
<i>Os09g25070</i> RT-R	CCGACGAGTTGATGGAGATG
<i>Os02g33820</i> RT-F	GCGTGGTGGACTACGACAAG
<i>Os02g33820</i> RT-R	CGTGCTTCTTGGCGTCCTTC
<i>Os01g72910</i> RT-F	GCAAGGAGGAGAAGCACCAC
<i>Os01g72910</i> RT-R	GTGCTTCTTGGCTTCCTTCT
<i>Os01g32380</i> RT-F	GACGTGGACAGGTTCTACGC
<i>Os01g32380</i> RT-R	GCTGTCCACCTTCTTCTCCG
<i>Os03g18850</i> RT-F	TCAACTCACCTCAGCCATG
<i>Os03g18850</i> RT-R	GTGATCTCGTCCTTCACCTC
