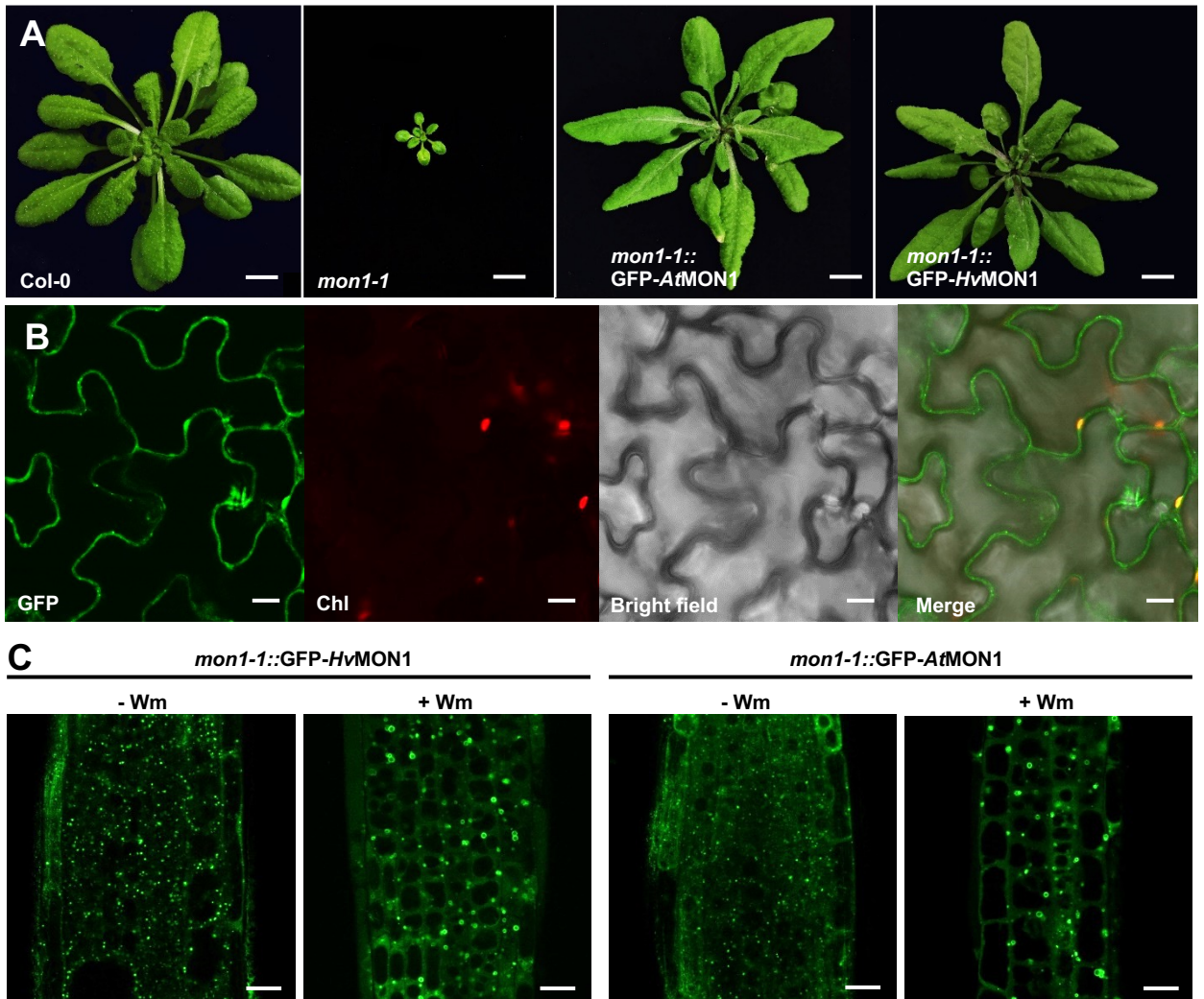
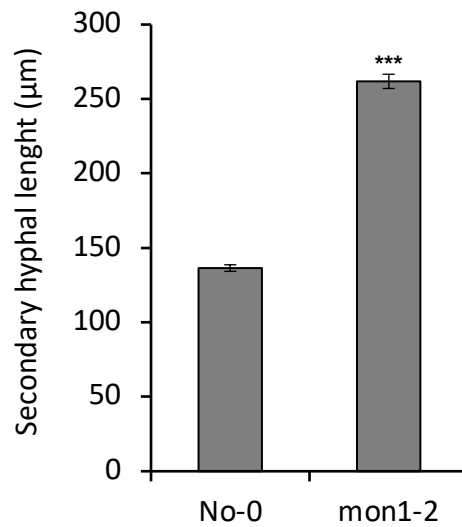


Supplementary Figure S1. Callose-containing encasements around *Bh* haustoria in barley induced by tetraconazole. (A) *Bh* colony and conidia chains developed on barley 4 days after inoculation. (B,C) Tetraconazole-induced encasements around *Bh* haustoria in barley epidermal cells. Inoculation was made 2 h after treatment. Images taken 4 dpi. UV-fluorescence microscopy after aniline blue treatment. Size bars, 100 μm in A and B, 10 μm in C.



Supplementary Figure S2. Barley *HvMON1* complements the function of Arabidopsis *AtMON1*. (A) Transformation of Col-0 *mon1-1* with *35S::HvMON1* rescues it from its lethality phenotype. Size bars, 1 cm, (B) CSLM subcellular localization of GFP-*HvMON1* expressed in *mon1-1* leaf cells epidermal. (C) Wortmannin-induced ring-shaped structures of GFP-*HvMON1* and GFP-*AtMON1* resolved by CSLM. Roots were treated with 33 μ M wortmannin for 1 hour. Size bars, 10 μ m in B and 10 μ m C.



Supplementary Figure S3. Secondary hyphal length of *Go* on No-0 and *mon1-2*. Length of secondary hyphae developed from penetrated appressoria at 2 days after inoculation. Error bars, SE. ***, $P < 0.001$ assessed by Student's T-tests. $n=3$.

Supplementary Table S1. Primers used in this work.

Name	Information	Sequence (5'-3')
attbHvMON1-F	HvMON1	AAAAAAGCAGGCTACATGGATCCGGCCCCCGAT
attbHvMON1-R	Cloning	AGAAAGCTGGGTCTCACCAGGCAATGGTGCT
attBCSEP0162-F	CSEP0162	AAAAAAGCAGGCTACATGGCCCAATATTCTAGACATATTTAA
attBCSEP0162-R	Cloning	AGAAAGCTGGGTCTCCCGAGCCAAGTGGC
attbAtMON1-F	AtMON1 Cloning	AAAAAAGCAGGCTACATGGCGACTTCAGATTCCG
attbAtMON1-R		AGAAAGCTGGGTCTCACCAGGCAATGGTGCT
pGB9-FP	pDEST-AS2-1 sequencing	AGTGCACATCATCATCG
pGB9-RP		CGTTTTAAAACCTAAGAGTCAC
pACT2-FP	pDEST-ACT2 sequencing	GATGATGAAGATACCCAC
pACT2-RP		CAGTTGAAGTGAAGTGC
attbHvMON1i-F	HvMON1 RNAi	AAAAAAGCAGGCTACCACCACTTCTAGGTGGCACA
attbHvMON1i-R	Cloning	AGAAAGCTGGGTCTCGCAAGCATATGGGTGAAAA
qRT-VIGS-Mi-F1	qRT-PCR for HvMON1 RNAi	CCGCACAAAACACAATTGAG
qRT-VIGS-Mi-R1		CCCTGATCCACTGGCATACT
HvUBC2-F		GGATCCAGGGGCACCTCAC
HvUBC2-R		CGTCCAAGCTTTTGGAGGAC
Bgh-GPD-F		<i>Blumeria hordei</i> quantification
Bgh-GPD-R	CATTTCCGGCGGCAATCTTT	
qPCR-HvM-F	TTCTGATGAGGCAATTGGTG	
qPCR-HvM-R	TTCTTATGGGCAAGGCAAAC	
BS10	VIGS vector sequencing	GGTGCTTGATGCTTTGGATAAAGG
BS32		TGGTCTTCCCTTGGGGGAC
At3g21215-RBD-F	<i>Golovinomyces orontii</i> quantification	GAATCCACCCATACCACCAG
At3g21215-RBD-R		GAGGAGGAGGATGGTGATGA
GoATPase1-F		TCGCCGCTATATTGGAGTC
GoATPase1-R		CTGGGTCAGATGGTTCACCT
mon1-mutant-LP	mon1 mutant	CGGTTTGCTGAGTTACTCAG
mon1-mutant-RP		AAAAGCCCAACAATATGGGTC
DS5-3	<i>mon1-2</i> (No-0)	TACCTCGGGTTCGAAATCGAT
LBb1.3	<i>mon1-1</i> (Col-0)	ATTTTGCCGATTCGGAAC
eds1-F760	<i>eds1-2</i> mutant	ACACAAGGGTGATGCGAGACA
eds1-R1458		GGCTTGATTCATCTTCTATCC
eds1-R2333		GTGGAAACCAAAATTTGACATTAG
ndr1-F3129	<i>ndr1-1</i> mutant	GGTTGTGAAATCAAGAATTAATGTGGA
ndr1-F4443		TTGCCTAATGGATCGGCTG
ndr1-R4928		GTTCTTGATTTGAAACCAACA

Supplementary Table S2. Gateway destination vectors used in this work.

Name	Information	Reference
pDEST-ACT2-GW	Yeast two-hybrid	Robertson. 2004
pDEST-AS2-GW		
pB4GWnG	BiFC	Kamigaki et al. 2016
pB4GWcCG		
pB4nGGW		
pB4cCGGW		
p35S-mCherry-GW	Protoplast transformation and particle bombardment	Kwaaitaal et al. 2010
pUbi-GW-YFP		
pUbi-GW-nos		
pIPKTA30N-GW		
pCaBS-γ-GW	VIGS	pCaBS- γ (Yuan et al. 2011) were adapted to Gateway compatible
pUBN10-GFP-GW	Arabidopsis overexpression	Grefen et al. 2010

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Supplementary Table S3. Overview of *HvMON1*/CSEP0162 bifluorescence complementation (BiFC) results.

Combination	Fluorescence signal
<i>HvMON1</i> -nGFP + CSEP0162-cCFP	No
<i>HvMON1</i> -nGFP + cCFP-CSEP0162	No
nGFP- <i>HvMON1</i> + cCFP-CSEP0162	No
nGFP- <i>HvMON1</i> + CSEP0162-cCFP	Yes
<i>HvMON1</i> -cCFP + CSEP0162-nGFP	No
<i>HvMON1</i> -cCFP + nGFP-CSEP0162	No
cCFP- <i>HvMON1</i> + nGFP-CSEP0162	No
cCFP- <i>HvMON1</i> + CSEP0162-nGFP	No