

Supplementary material

Article title

Root colonization by fungal entomopathogen systemically primes belowground plant defense against cabbage root fly

Authors

Catalina Posada-Vergara¹, Katharina Lohaus¹, Mohammad Alhussein², Stefan Vidal¹, and Michael Rostás^{1*}.

¹ Agricultural Entomology, Department of Crop Sciences, University of Göttingen, Grisebachstr. 6, 37077 Göttingen, Germany; e-mail@e-mail.com

² Molecular Phytopathology and Mycotoxin Research, University of Göttingen, 37077 Goettingen, Germany

* Correspondence: michael.rostas@uni-goettingen.de

Table S1. List of the fungal strains used in the study (see (Hettlage, 2018)).

Strain	Crop regime	Location
GC1I	Semi-natural permanent grassland for at least 7 years. Grassland harvested twice a year	51°34'10.7"N 10°03'54.1"E
Gd12		
Cb15III		
Cb17b		
CC5		

Table S2. Primers used for qPCR amplification of genes of *Brassica napus* root tissues

ID	Gen/accession	Pathway	Gene description	Sequence 5' --> 3'	Reference
ABA2	LOC106300040	Abscisic acid biosynthesis	Xanthoxin dehydrogenase	GCATCGCTCGTCTGTTCCAC	Karssemeijer et al., 2021
	XM_013736089			CGGCGAACGTCAACAGCGTTA	
ERF2	At5g47220	Ethylene (ET) signaling	Ethylene Response Factor 2	ATGTACGGACAGAGCGAGGT	Yang et al., 2010
				AAGCTTCGAAACCAACAAGTAAC	
ACO	EV102889	ET biosynthesis	ACC oxidase	TCCGTCTGGCTATCACTCT	Maag 2014
				GTGAGTGGGTCGATGTTCT	
PR1	XM_013877950	Salicylic acid (SA) signaling	Pathogenesis-related protein 1	AAAGCTACGCCGACCGACTACGAG	Alkoorane 2017
				CCAGAAAAGTCGGCGCTACTCCA	
PAL	LOC106342153	SA synthesis-Phenylpropanoid pathway	phenylalanine ammonia-lyase 1	TCGCTATGGCTTCTACTGCTCTG	Karssemeijer et al., 2021
	XM_013781008			GAGGTCTTACGAGATGAGATGAGTCC	
AOS	LOC106327419	Jasmonic acid (JA) synthesis	Allene oxide synthase	ACCGCTTGCGACTAGGGATC	Karssemeijer et al., 2021
	XM_013765565			CAAAGTCCTTACCGCGCAC	
MYC2	EV120351	JA signaling	Basic helix-loop-helix (bHLH) DNA-binding family protein	GCAAAGCCCAGACAGAGAAC	Maag et al., 2014
				AGCTCACGCAACACCTTCTT	
TPI	EV144353	JA signaling	Trypsin inhibitor B-like	GTGGTATCACCATGAACCTTG	Maag et al., 2014
				GTTGACCACCTAACCGGAA	
PDF1.2	EV163328	JA signaling	Defensin-like protein 16	TCCATCACCCCTCTCTTGC	Maag et al., 2014
				TTTGGCACCGCATAGTCGTA	
ACTIN	AF111812	Housekeeping gene	Housekeeping gene	ATCGTCCTCAGTGGTGGTTC	Maag et al., 2014
				TTGATCTCATGCTGCTTGG	
GTR1A2	Bra018096	Glucosinolate (GSL) transport	Glucosinolate Transporter 1 A2	ATTCACCTCGGGGAACTGG	(Sontowski et al., 2019)
				TCGCTTGCTCTGCTTGGTC	
CYP79B2	At4G39950	Indole GSLs biosynthesis	CYTOCHROME P450, FAMILY 79, SUBFAMILY B, POLYPEPTIDE 2	AAGAGGTTGTGCTGCTCG	Tytgat et al., 2013 Also in Marthur
				TCCAAGTGAAACCTTGAAGAAGTC	
CYP83A1	At4G13770	Aliphatic GSL biosynthesis	CYTOCHROME P450, FAMILY 83, SUBFAMILY A, POLYPEPTIDE 1	CTCCTTATCCCTCGTGTCTTG	Mathur et al., 2013
				TGTCGTAACCAGCGATCTTG	
BABG	LOC106429220	Myrosinase biosynthesis	Beta-glucosidase 27-like	CCGAGCGAGCTATGGAGTTT	This study
	XM_022718558			CGGCTTGTCTGGATCCACTT	
BnMyr4	LOC106430598	Myrosinase biosynthesis	PREDICTED myrosinase 4-like	TCAACTGCGACAATCCCCTT	This study
	XM_013871387			ATCACAAGCAAGGTCCCGG	
Myr2.Bn1	LOC106382545	Myrosinase biosynthesis	B.napus myrosinase, thioglucoside glucohydrolase	TTGAAGGAGGGAGAGGTCTG	This study
	NM_001316199			AGCATTGAGTTCGCCCCATCA	
DTCMT.a	LOC106392535	Phytoalexin brassinin biosynthesis	Dithiocarbamate S-Methyltransferase	TGTTCCACTGGACCTAACACG	This study
	XM_013833342			GGCCAAAGAAAAGATCCGGGA	

Table S3. Acquisition parameters for phytohormones analysis.

Compound	RT [min]	Polarity	Parent Ion [m/z]	Fragmentor V	Collision Energy V	Product Ion [m/z]
<i>Trans</i> -zeatin	1.93	+	220.1	100	15	136.1
					9	202.1
					22	148.1
<i>Trans</i> -zeatin-d5	1.92	+	225.1	105	16	137.1
					10	207.2
Abscisic acid (ABA)	4.68	-	263.1	85	4	153.1
					5	219.1
					12	204.1
Abscisic acid-d6	4.67	-	269.1	88	4	159.1
					8	225.1
Jasmonic acid (JA)	5.20	+	211.2	85	8	133.1
					8	151.1
					5	193
Jasmonic acid-d5	5.19	+	216.1	85	8	135.1
					9	153.2
					7	198.2
Salicylic acid (SA)	4.05	-	137	140	15	93.1
					35	65.1
Salicylic acid glucoside (SA-Glu)	2.41	-	137	140	15	93.1
					35	65.1

Table S4. Statistical summary of GLM analysis with binomial distribution of the total mortality on in vitro experiments. Mortality includes mycosed larvae, pupae, and adults that developed mycosis after emerging.

Isolate	L3 Larvae			
	Estimate	SEM	z value	Pr(> z)
Intercept	-1.6398	0.3706	-4.425	9.65E-06***
Cb15III	2.339	0.4349	5.378	7.52E-08***
Cb17B	3.0474	0.4525	6.735	1.64E-11***
CC5	3.1991	0.4588	6.972	3.11E-12***
Gc1I	4.1194	0.5248	7.85	4.15E-15***
Gd12	4.1194	0.5248	7.85	4.15E-15***
Sand	-0.7047	0.2525	-2.791	0.00525**

Table S5. Statistical summary of GLM analysis with binomial distribution of the total mortality in planta experiment. Mortality includes mycosed larvae, pupae, and adults that developed mycosis after emerging. RCD: root collar diameter, included as covariant.

Factor	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	2.9015	0.9581	3.028	0.00246**
Cb15III	-0.8301	0.3658	-2.269	0.02324*
Cb17B	-0.2827	0.3721	-0.76	0.44746
CC5	-0.4439	0.368	-1.206	0.22774
Gc1I	-0.853	0.3887	-2.195	0.02818*
Gd12	-1.0107	0.3561	-2.839	0.00453**
RCD	-0.1567	0.111	-1.412	0.15794

Table S6. Statistical summary of beta regression analysis with of percentage of damage of root collar in planta experiment.

Factor	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	1.1920	0.1789	6.662	2.71e-11 ***
TreatCb15III	-0.7290	0.2491	-2.926	0.003432 **
TreatCb17B	-0.8382	0.2379	-3.523	0.000427 ***
TreatCC5	-0.6718	0.2394	-2.807	0.005005 **
TreatGc1I	-0.8858	0.2623	-3.377	0.000732 ***
TreatGd12	-1.1614	0.2415	-4.808	1.52e-06 ***

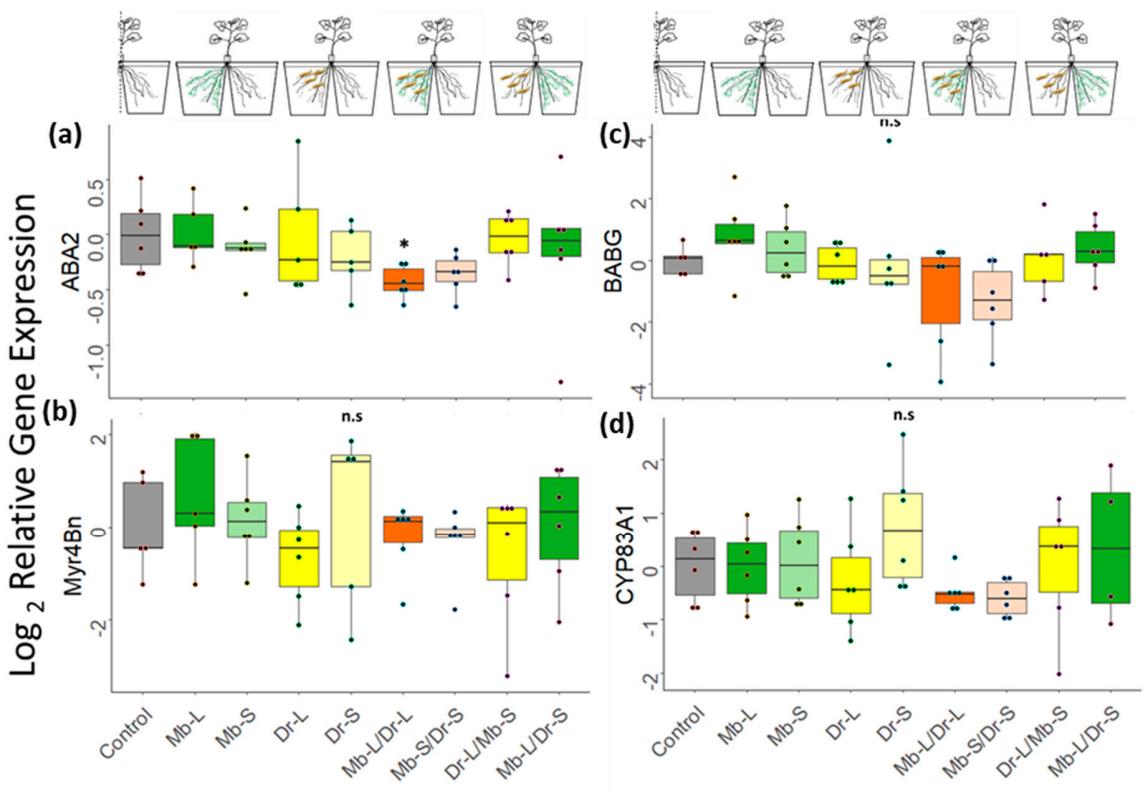


Figure S1. Normalized expression of representative genes of **(a)** abscisic acid biosynthesis *ABA2*, **(b)** myrosinase synthesis *Myr4Bn*, **(c)** beta-glucosidase biosynthesis *BABG* and **(d)** aliphatic GSL synthesis *CYP83A1*. Plants grew in a split root setup in which each compartment had either *M. brunneum* inoculation (Mb) in the local (L) or adjacent (S) compartment, *D. radicum* egg infestation (Dr) in the local (L) or adjacent (S) compartment, both treatments in same compartment (Mb-L/Dr-L), or each in adjacent compartments of the same plant (Dr-L/Mb-S; Mb-L/Dr-S). Eggs were placed 4 weeks after Mb inoculation. Plants were harvested 7 days after egg infestation. Gene expression was normalized to the housekeeping gene *AUXIN*. The boxplot shows all data points from at least 4 independent biological replicates ($n \leq 6$) in which the horizontal line represents the median, surrounded by the upper (25th) and lower (75th) percentiles.