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Supplementary Information

***Bacillus volatiles* adversely affect the physiology and ultra-structure of *Ralstonia solanacearum* and induce systemic resistance in tobacco against bacterial wilt**

Hafiz Abdul Samad Tahir, Huijun Wu, Yuedi Niu, Qin Gu, Rong Hu, Xuewen Gao *

Department of Plant Pathology, College of Plant Protection, Nanjing Agricultural University, Key Laboratory of Integrated Management of Crop Diseases and Pests, Ministry of Education, Nanjing 210095, PR China

* Corresponding author: Xuewen Gao

Address: Nanjing Agricultural University, Weigang No.1, Nanjing 210095, PR China

Email: gaoxw@njau.edu.cn

Telephone/Fax: 86-25-8439526

13 **Table S1: Bacteria used in this study**

No	Strain	Antagonistic strain or pathogen	Accession NO
1	<i>Bacillus amyloliquefacians</i> FZB42	Antagonist	NR075005.1
2	<i>Bacillus cereus</i> NMSL88	Antagonist	GU568190.1
3	<i>Bacillus amyloliquefacians</i> NMSX4	Antagonist	GU568185.1
4	<i>Bacillus subtilis</i> FA26	Antagonist	KY003098
5	<i>Bacillus artrophaeus</i> LSSC22	Antagonist	GU568193.1
6	<i>Bacillus pumulis</i> GBSW19	Antagonist	GU568202.1
7	<i>Ralstonia solanacearum</i> TBBS1	Pathogen	KY003096

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15 **Table S2: Primers used in the study**

No	Oligo Name	Sequence 5' to 3'
1	<i>16sr RNA Rsc</i> TBBS1 (F)	AGTCCACGCCCTAAACGA
2	<i>16sr RNA Rsc</i> TBBS1 (R)	CGAAGGCACCAATCCATC
3	<i>Mot A</i> (F)	GTTTTCCATCATCCAAGC
4	<i>Mot A</i> (R)	GTGCCGAACAGACCCATC
5	<i>FliT</i> (F)	GACCAGCGCCCGGACACG
6	<i>FliT</i> (R)	CGCCGCAAGGCAACGACA
7	<i>Chew</i> (F)	CGGCGACCGTCATTGGCT
8	<i>Chew</i> (R)	GCGCTGCGGATCCTTGGA
9	<i>PhcA</i> (F)	GCAACGTCTGCCTTTTCACT
10	<i>PhcA</i> (R)	CGCTGTCATGTGCATCTTCT
11	<i>HrpB</i> (F)	AGACCAAGGTGGAAGTCGTG
12	<i>HrpB</i> (R)	CGTCTTGCATGTAGCTGGTG
13	<i>EpsA</i> (F)	TTCCTCTGACCCAAGGAATG
14	<i>EpsA</i> (R)	ATCAAAGGTGTAGCCGTTGG
15	<i>EpsB</i> (F)	GGCGTTGTCCTAGGTGTCAT
16	<i>EpsB</i> (R)	CGCTTCGATAGATGGGTCAT
17	<i>EpsC</i> (F)	ACGGTGACACTTCAACCACA
18	<i>EpsC</i> (R)	GCCATGGGCTGTACAAGTTT

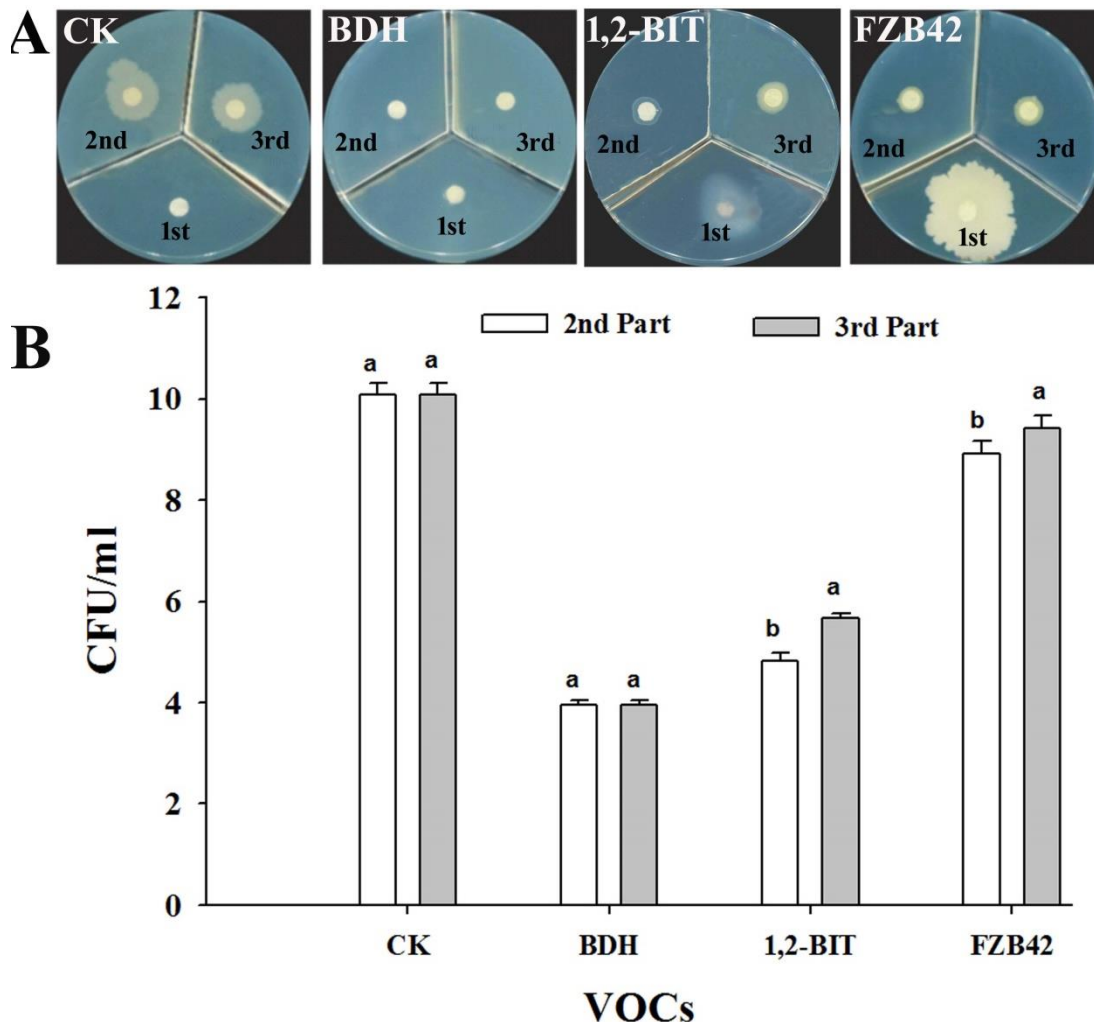
19	<i>EpsD</i> (F)	CAGCCGAGATGTGCAAATA
20	<i>EpsD</i> (R)	GGGGCAGCATCTACGATAAA
21	<i>EpsE</i> (F)	TACCGGCCACACTAGGAAAC
22	<i>EpSE</i> (R)	TTTCGCCTGCTAGCAAAAAT
23	<i>EpsF</i> (F)	GTTTCTCATCACGGCGTTTT
24	<i>EpsF</i> (R)	GGCGAAGGCTATGCTAGATG
25	<i>EpsP</i> (F)	CGTGAGATACAGGCGAGACA
26	<i>EpsP</i> (R)	GTCTTCGAACGCCATCATT
27	<i>Awr1</i> (F)	ACGTTTCCACGCATAACTCC
28	<i>Awr1</i> (R)	GTCTGGGTGGCAAAAAGTGT
29	<i>Awr3</i> (F)	CTCACGCATTCCTACAAGCA
30	<i>Awr3</i> (R)	CCAGTCTGCTCAGGTGACAA
31	<i>Awr5</i> (F)	CACATGGCGGAGAGATTTTT
32	<i>Awr5</i> (R)	TCGTAGACGTACGCCTGTTG
33	<i>PilQ</i> (F)	CAGGGACAAAACCTGGTCGT
34	<i>PilQ</i> (R)	CGGAGTCGGTAGCTCTCATC
35	<i>RRS1</i> (F)	ATGAGAAAGAGGCTCGTCAA
36	<i>RRS1</i> (R)	ACCACAACCCTCAAGCAGTT
37	<i>NPRI</i> (F)	CTGGAGCAAGCAGAAAG
38	<i>NPRI</i> (R)	TCATACGCAAATCATCG
39	<i>EDS1</i> (F)	GAG TAT CAG ACC AAG TGT GAT ATC CG
40	<i>EDS1</i> (R)	GCT GAG GTG GGA GTG TTT TCC ACC

17 **Table S3: VOC profile of *Bacillus amyloliquefacians* FZB42 and *Bacillus***
 18 ***artrophaeus* LSSC22**

<i>Bacillus</i> strain	RT (min)	Relative peak area (%)	Compound	Abbreviation used	Degree of inhibition
<i>Bacillus amyloliquefacians</i> FZB42	4.567	3.829	Silanediol, dimethyl	SDD	–
	5.618	3.574	1,2-Benz isothiazol-3(2H)-one	1,2-BIT	+++
	6.963	1.865	Benzeneacetamide	BAM	++
	7.631	2.573	Oxime,methoxy-phenyl	OMP	NT
	7.787	3.494	(1R)-2,6,6-Trimethylbicyclo[3.1.1]hept-2-ene	TMB	+
	8.207	1.557	Benzoic acid, 2-formyl-4,6-dimethoxy-,8,8-dimethoxyoct-2-yl	BA	+
	8.949	1.695	Benzaldehyde	BDH	+++
	9.543	2.277	Sulfurous acid, cyclohexylmethyl isobutyl ester	SCE	–
	10.148	1.406	6-Tridecen, 2,2,4,10,12,12-hexamethyl-7-(3,5,5-trimethylhexyl)-	6 -THT	NT
	10.304	2.116	2-Undecanethiol, 2-methyl	2-UT,2-M	–
	10.734	6.490	Dodecane , 1-fluoro	DCF	++
	12.262	1.695	Dodecane	DCN	++
	14.421	2.380	Phenol, 2-(1,1-dimethyl)-5-methyl-	PH	–
<i>Bacillus artrophaeus</i> LSSC22	1.301	1.073	1,3 –Butadiene	1,3-BDN	++
	5.631	5.995	1,2-Benz isothiazol-3(2H)-one	1,2-BIT	+++
	7.729	2.245	(1R)-2,6,6-Trimethylbicyclo[3.1.1]hept-2-ene	TMB	+
	8.764	3.483	Benzoic acid	BA	+
	10.487	2.204	1-octyn-3ol, 4- ethyl-	1,OTN	++
	10.712	8.354	Dodecane , 1-fluoro	DCF	++
	11.575	1.491	Undecanal, 2-methyl	UDM	++
	12.264	2.768	Dodecane	DCN	++
	14.413	2.694	Phenol, 2-(1,1-dimethylethyl)-6-methyl	PH	–
	16.669	1.213	Cyclohexene, 3-(1,5-dimethyl-4-hexenyl)-6-methylene-	CHN	NT

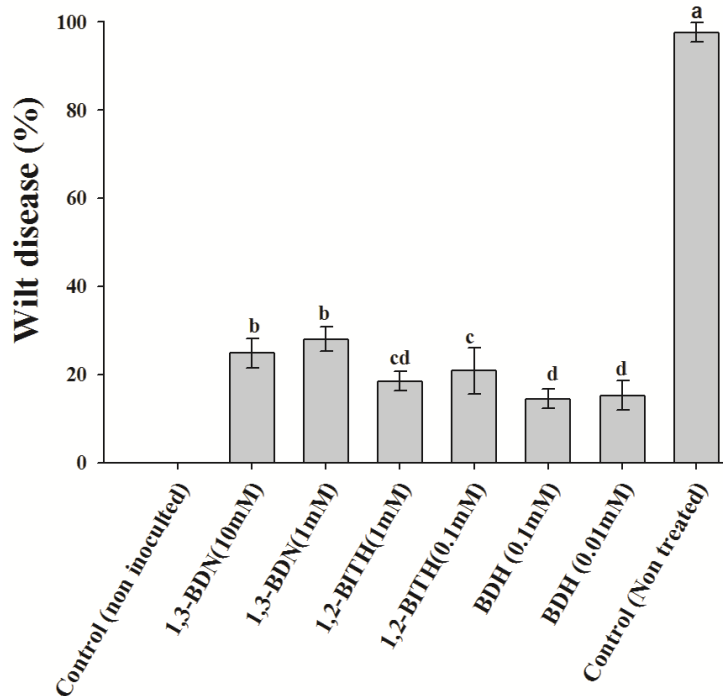
19 Minor air-peaks were excluded from the total analysis representing $\leq 1\%$ of the total

20 area. Similar compounds found in both inoculated and non-inoculated MS medium
 21 were also not included. **RT**, retention time, - no inhibition, + $\leq 10\%$ inhibition rate,
 22 ++ 10-30 % inhibition rate, +++ More than 30 % inhibition, **NT** Not tested



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 24 **Fig S1 Effect of the concentration of VOCs** The experiment was conducted in three
 25 partition plates with completely sealed portions without any air movement except holes
 26 in two walls of the partitions. An arrangement was made so that VOCs could move
 27 from the first partition, encompassing *Bacillus* or a synthetic chemical, to the second
 28 partition and then from the second partition to the third partition. The second and third
 29 partition both were inoculated with 10 μ l of *Rsc* (18-24h) culture. The whole
 30 experiment was repeated three times, including three replicates per experiment. Letters

31 above error bars represent significant differences according to Duncan's multiple-range
32 test (P=0.05) using SPSS software (SPSS, Chicago, IL).



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34 **Fig S-2 BDH, 1,2-BITH and 1,3-BDN reduced the virulence of *Ralstonia***
35 ***solanacearum* and induced systemic resistance** I-plates prepared with
36 one-half-strength Murashige and Skoog solid medium, and 5-6-day-old emerging
37 tobacco seedlings (seven seedlings/plate) were dipped in the suspension of *Rsc* (10^7
38 CFU/mL) cells and transplanted into one compartment. In the non-inoculated control,
39 the roots were dipped in sterile water. Chemicals (10mM and 1mM 1,3-BDN, 1mM and
40 0.1mM 1,2-BITH while 0.1mM and 0.01mM BDH) were used in the other
41 compartment. Wilt symptoms were observed after 7 days of inoculation, and the data
42 were recorded. Error bars indicate standard deviations of the means. Different letters
43 above error bars represent significant differences according to Duncan's multiple-range
44 test (P=0.05) using SPSS software (SPSS, Chicago, IL).