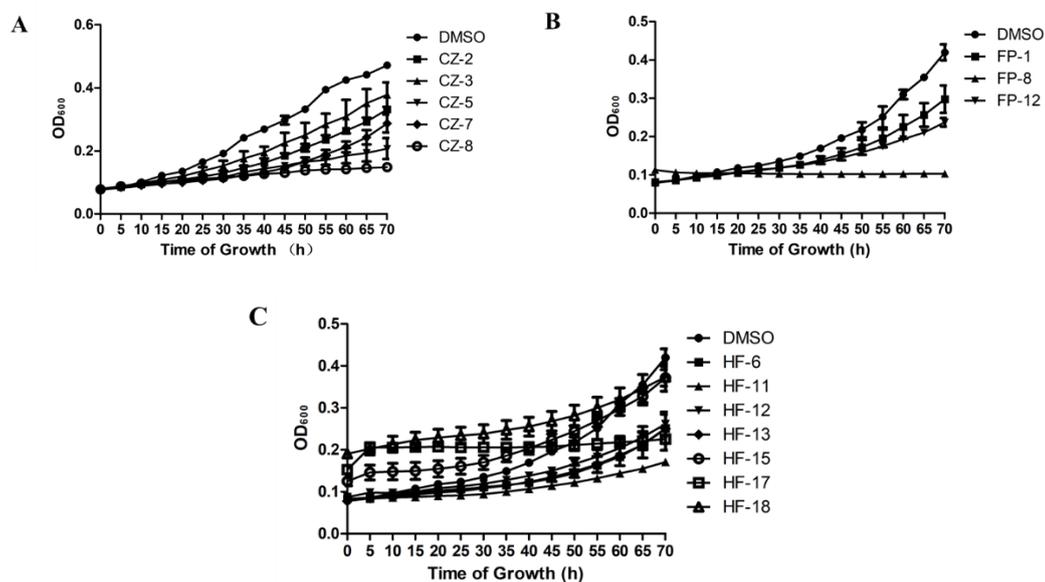




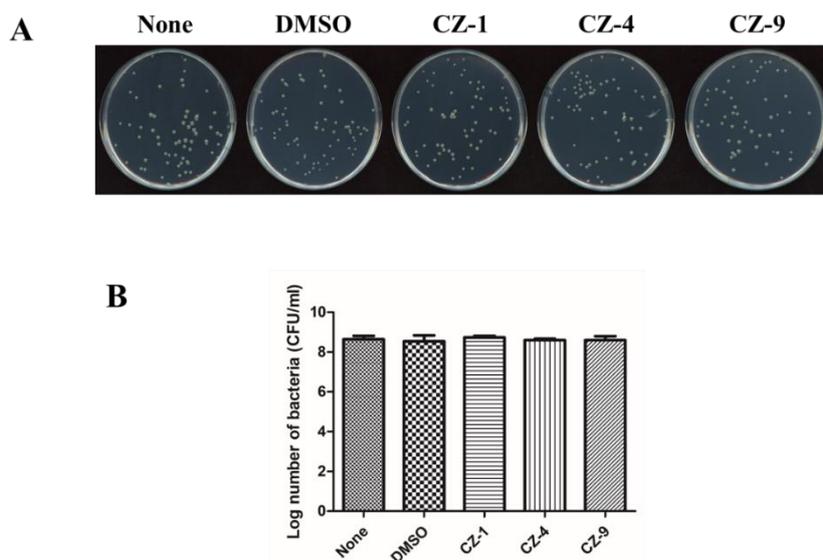
## Supplementary Materials

Phpa1-F  
**CCACTTAACGGGCAAG**CGAAAAAAGCTTTTCTCAACAACGCCCGCGGATTTGTTATCGA 60  
TTCTAAAACATTTTTTCACTTGCCCT**TTTCGCGCGTACAAGCGCAATTTTCGCAAAATTT**C 120  
TGGCGATGATGGGCTTCCGCTTCTACTGTTTGATCGGGGCGCAAAACGCGCCTCGCAGCC 180  
ACCGTGTCTAGGCACGGCTGTTGATACTAAAGACACATACCATTTA**ATCAGAGAGGAAT** 240  
Phpa1-R  
Start codon  
**CGTACGATGAAT**TCTTTGAACACACAATTCGGCGGCAGCACGTCCAACCTTCAGGTTGG 300  
CCCAAGCCAGGACACAACGTTTCGGTTCGAACCAGGGCGCAACCAGGGCATCTCGGAAAA 360  
GCAACTGGACCAGTTGCTGTGCCAGCTCATCTCGGCCCTGCTTCAGTCGAGCAAAAAATGC 420  
TGAGGAGGGTAAGGGTCAGGGTGGCGATAATGGCGGTGGCCAGGGCGCAATTCGCAGCA 480  
GGCTGGGCAGCAGAATGGCCCTCGCCATTCACCCAGATGCTGATGCATATCGTCGGAGA 540  
GATTCTCCAGGCGCAGAATGGTGGTGGTGGTGGCGGGTTCGGCGGGGGTTTCGG 600  
CGGTGACTTTAGTGGCGACCTCGGCCTCGGCACCAACCTCTCGAGCGACAGCGCATCGAT 660  
Stop codon  
**GCAGTAA** 667

**Figure S1.** DNA sequence of *hpa1* gene and its promoter region. Sequences for a PIP box, start codon, stop codon and the primers used for amplification of the promoter region are indicated as bold fonts.



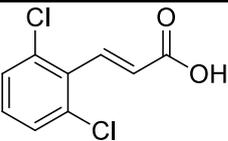
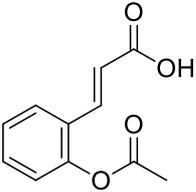
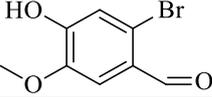
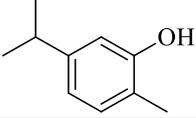
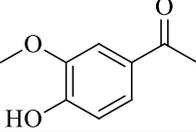
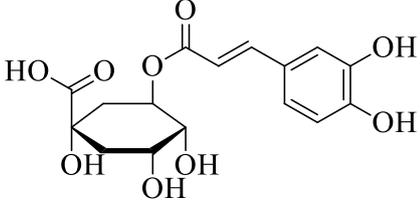
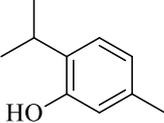
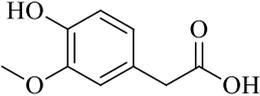
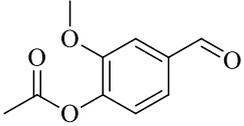
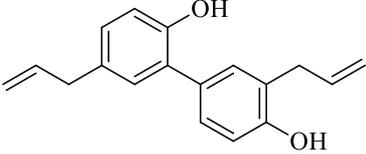
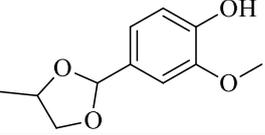
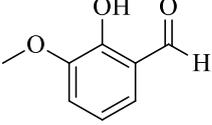
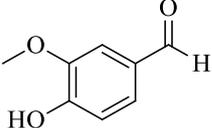
**Figure S2.** Effects of various compounds on bacterial survival, Growth rates of PXO99<sup>A</sup> in plant-mimicking medium XOM2 supplemented with 200 μmol/L compounds. The optical density at 600 nm (OD<sub>600</sub>) of the culture suspensions was measured every 1 h during the 72 h period. Three independent tests were performed with similar results.

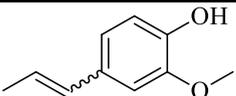
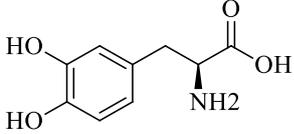
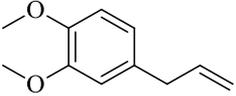
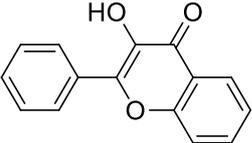
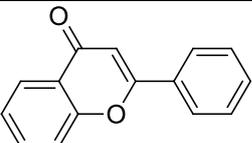
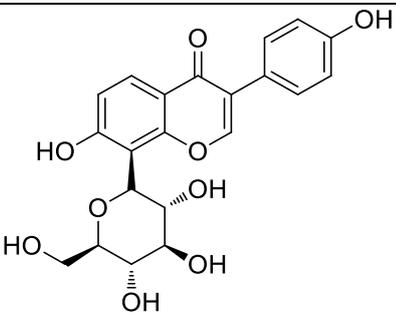
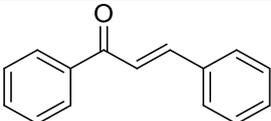
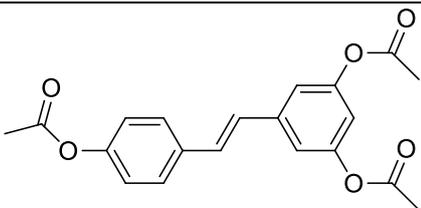
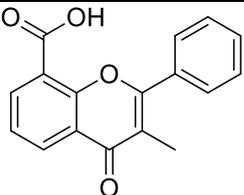
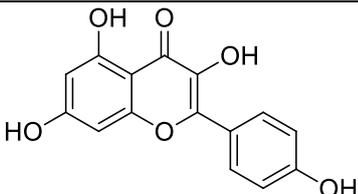


**Figure S3.** Effects of various compounds on bacterial survival. Bacterial cells were incubated with DMSO or various compounds at a concentration of 200  $\mu$ M for 2 h at 28  $^{\circ}$ C before being plating on PSA plate after serial dilutions. Three independent tests were performed with similar results.

**Table S1.** The compounds in this study.

No.	Compounds	Structure	Name
1	CZ-1		2-methoxybenzene propanoic acid
2	CZ-2		3-[2-(trifluoromethyl)phenyl]-2-propenoic acid
3	CZ-3		2-methylbenzene propanoic acid
4	CZ-4		3-(2-fluorophenyl)-2-propenoic acid
5	CZ-5		3-(2-chlorophenyl)-2-propenoic acid
6	CZ-6		3-(2-methoxyphenyl)-2-propenal
7	CZ-7		[1,1'-biphenyl]-4-carboxylic acid

8	CZ-8		3-(2,6-dichlorophenyl)-2-propenoic acid
9	CZ-9		3-[2-(acetyloxy)phenyl]-2-propenoic acid
10	FP-1		6-bromo vanillin
11	FP-2		carvacrol
12	FP-3		aceto vanillone
13	FP-4		chlorogenic acid
14	FP-5		thymol
15	FP-6		homovanillic acid
16	FP-7		vanillin acetate
17	FP-8		honokiol
18	FP-9		vanillin propylene glycol acetal
19	FP-10		2-hydroxy-3-methoxy-benzaldehyde
20	FP-11		4-hydroxy-3-methoxy-benzaldehyde

21	FP-12		isoeugenol, mixture of <i>cis</i> and <i>trans</i>
22	FP-13		3,4-dihydroxy-L-phenylalanine
23	FP-15		methyl eugenol
24	HF-1		flavonol
25	HF-6		flavone
26	HF-8		8-( $\beta$ -D-glucopyranosyl-7-hydroxy-3-(4-hydroxyphenyl)-4H-1-benzopyran-4-one
27	HF-11		<i>trans</i> -chalcone
28	HF-12		triacetyl resveratrol
29	HF-13		3-methylflavone-8-carboxylic acid
30	HF-15		3,4',5,7-tetrahydroxyflavone 3,5,7-trihydroxy-2-(4-hydroxyphenyl)-4H-1-benzopyran-4-one robigenin

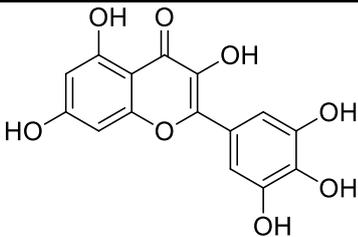
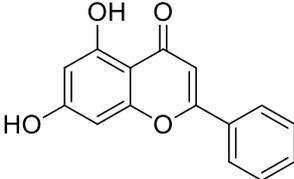
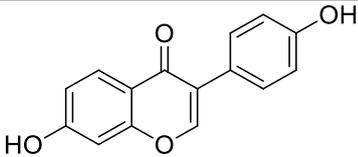
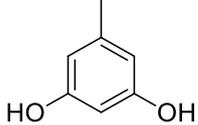
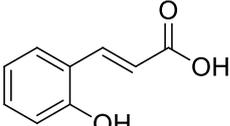
31	HF-17		3,3',4',5,5',7-hexahydroxyflavone cannabiscetin
32	HF-18		5,7-dihydroxyflavone
33	HF-19		4',7-dihydroxy-iso-flavone
34	FP-31		1,3-dihydroxy-5-methylbenzene
35	TS006		O-coumaric acid (OCA)

Table S2. Primers used in this study.

Primer name	Sequences (5'-3', restriction sites underscored)	Description of amplified product
PhrpB-F	ACGCGT <u>TCGAC</u> ATTTAAATGCGCCCAACTTC	234-bp promoter of <i>hrpB</i> <sub>1</sub>
PhrpB-R	<u>GGAATTC</u> TGTGAATCTTCTCCACACTG	
hrpG-F	TGTCCACCTGATGAACGACCCT	<i>hrpG</i> gene fragment, 165bp
hrpG-R	GGCGAATGCCGCAACGAA	
hrpX-F	AGGCACTGACCCACTTTC	<i>hrpX</i> gene fragment, 101bp
hrpX-R	ATCGGAAGCACCCTCTC	
hpa1-F	AAGCCAGGACACAACGTTTCG	<i>hpa1</i> gene fragment, 101bp
hpa1-R	GAAGCAGGGCCGAGATGAG	
hrcC-F	GCGCTTCCGGTGCGTTAC	<i>hrcC</i> gene fragment, 105bp
hrcC-R	CCCTGCTCGACCTGCTTGG	
hrcT-F	AGGGCGTGTCTGCTGTTGACC	<i>hrcT</i> gene fragment, 104bp
hrcT-R	CGCGATTCCGGGAAGACTGT	
hrcU-F	CAGGCACGCAGCCAGGAA	<i>hrcU</i> gene fragment, 86bp
hrcU-R	GCGATGCAACGGCGATAA	
hrpE-F	CGTTGTCGCCCGCCTT	<i>hrpE</i> gene fragment, 134bp
hrpE-R	GGTTCGTTGCTCGGCG	
hrpF-F	AGTCCGGCGTGCTCATCG	<i>hrpF</i> gene fragment, 102 bp
hrpF-R	AGTGCCACCGCAGTTGA	
gyrB-F	GGCGAGCACAATGGCATT	<i>gyrB</i> gene fragment, 101bp
gyrB-R	CCATCCTTCTGCGGGATGT	