

**Supplementary Information for**

**Functional characterization of Vib-PT, an aromatic prenyltransferase involved in the biosynthesis of vibralactone from *Stereum vibrans***

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Na Bai and Guo-Hong Li contributed equally to this work. Author order was determined both alphabetically and in order of increasing seniority.

**Table S1. Primer sequences used for PCR and qPCR**

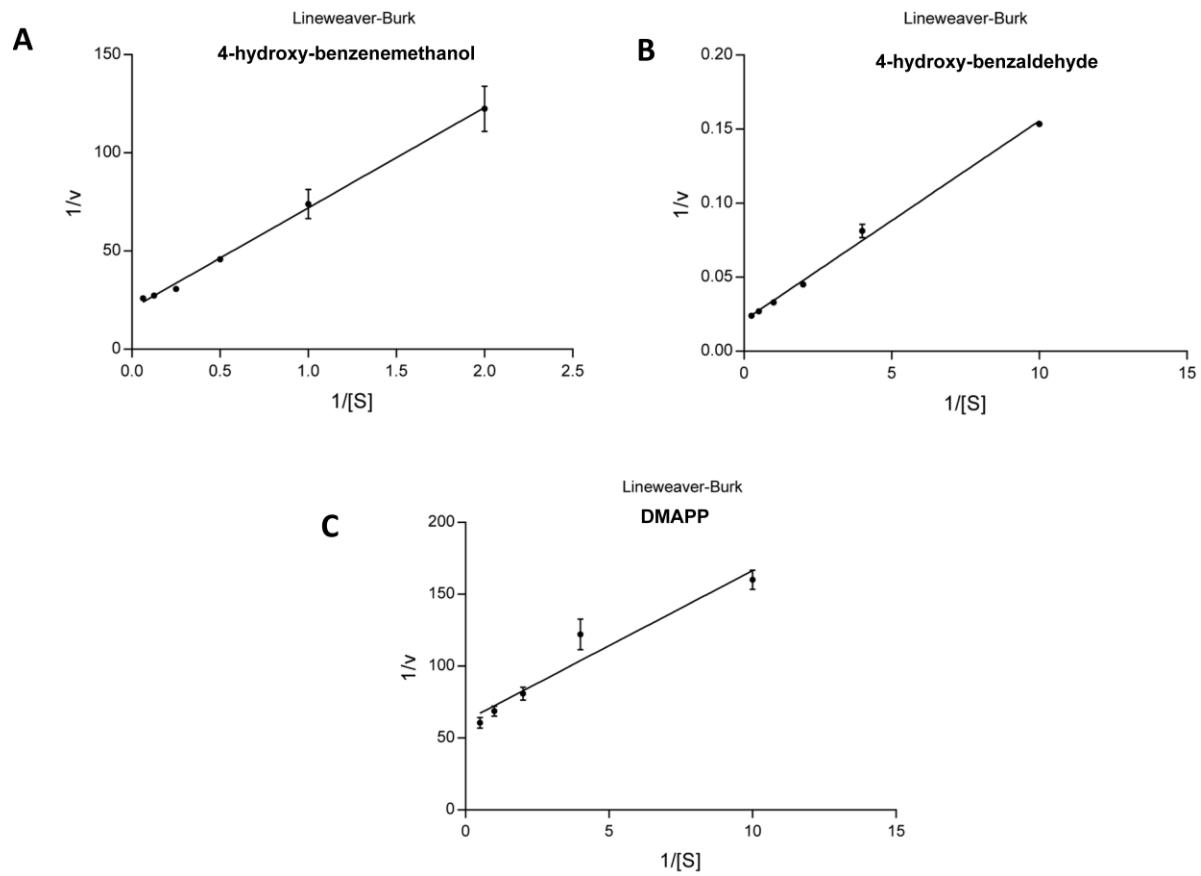
Name	Primer sequence (5'-3')
Alc_A-F	CCAAGCTTGAAAAGCTGATTGTGAT
Alc_A-R	CGGGATCCGCTAATTAAC TGAGTAGA
PT-2UP-BamH1	ATGGATCCGCTTCTCGGTCACTGTTATGGTC
PT-2DN-Hpal	ATGTTAAC TTGCGAACTCGTAGTGGTAAGGTA
PT-2iv-UP-sac1	AT GAGCTCTTGC GAAC TCGTAGTGGTAAGGTA
PT-2iv-DN-xho1-Not1	ATCTCGAGGC GGCGCCGCTTCTCGGTCACTGTTAT GGTC
PT-FW F	CCACCATGGTAGATCTTGC GAAC TCGTAGTGGTAAGG
PT-FW R	AACACTAGTCAGATCTTGAAAAGCTGATTGTGATAGTT
PT-IV F	ATCCTCTAGAGTCGACCGCTTCTCGGTCACTGTT
PT-IV R	ATGCCTGCAGGTCGACAACAAGAGAGGGTCCACAATGA
IN F	AATTCTGATGTGTAATTCTG
ALC R	GTTAACACTAGTCAGATCTT
1HF	ATTCGGCTCCAACAATGTC
1IR	GTCCACAATGATAAAGAGGG
β-tubulin F	ATGCGTGAGATTGTCCATCT
β-tubulin R	CCATGGTTCCGGGCTCGAGA
PT F	CTATCCCTCCTGGTCAACT
PT R	TTAGGAAGCTGCCTCTGC

**Table S2. Strains and plasmids**

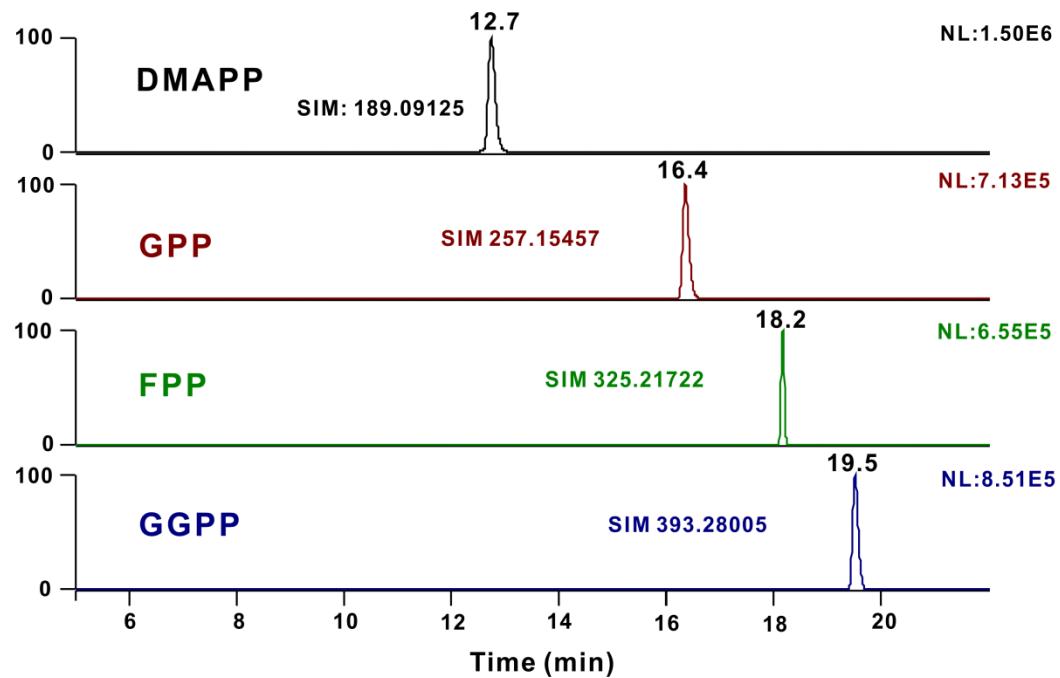
Strain or Vector	Properties	Reference or Source
<i>E. coli</i> BL21(DE3)	F-, ompT, hsdS(rBB-mB-), gal, dcm (DE3)	Novagen
<i>Agrobacterium tumefaciens</i> AGL1	Genotype C58 RecA (rif R/carB) Ti pTiBo542DT-DNA (strepR) Succinamopine.	Biomed
<i>Agrobacterium tumefaciens</i> LBA4404	Genotype Ach5 (RIF R) Ti pAL4404 (strepR) Octopine.	Biomed
<i>Agrobacterium tumefaciens</i> EHA105	Genotype C58 (rif R) Ti pEHA105 (pTiBo542DT-DNA) Succinamopine.	Biomed
<i>Agrobacterium tumefaciens</i> GV3101	Genotype C58 (rif R) Ti pMP90 (pTiC58DT-DNA) ( gentR) Nopaline.	Biomed
<i>Stereum vibrans</i>	Wild-type, vibralactone producer	Zhao et al., 2013
<i>Stereum vibrans</i> A2	vib-PT RNA silencing transformant by targeted PCR	This study
<i>Stereum vibrans</i> G7	vib-PT RNA silencing transformant by targeted PCR	This study
pUC18	pUC cloning vector	Addgene
pC-HYG-YR	Yeast recombinational cloning compatible <i>Agrobacterium tumefaciens</i> ternary vector containing a hygromycin selectable marker on transfer DNA	Addgene
pUCH2-8	pUCH2-8 containing the hygromycin B phosphotransferase gene, as the selectable marker for transformation of <i>Aspergillus</i>	Alexander et al., 1998
pMCB17apx	Promoter of alcohol dehydrogenase <i>alcA</i> (p)	Efimov, 2003
pUC18-RNAi	pUC18 harboring the second intron of the <i>Arabidopsis</i> small nuclear ribonuclear protein D1 (locus At 4g02840)	Kumar et al., 2012
pUNZ101	The promoter of <i>alcA</i> (p) was cloned into the expression vector pUCH2-8	This study
pUNZ103	A 1103-bp and reverse-complement of vib-PT cloned into the two ends of the intron of the pUC18-RNAi vector	This study
pUNZ104	The 2456-bp RNA silencing cassette digested with BamHI and NotI and cloned into pUNZ101	This study
pYUZ10	A 1479-bp element containing the <i>alcA</i> promoter and vib-PT forward sequence subcloned into the pC-HYG-YR vector	This study
pYUZ11	A 1357-bp element containing the intron and Vib-PT reverse-complemented sequence cloned into pYUZ10	This study

## References

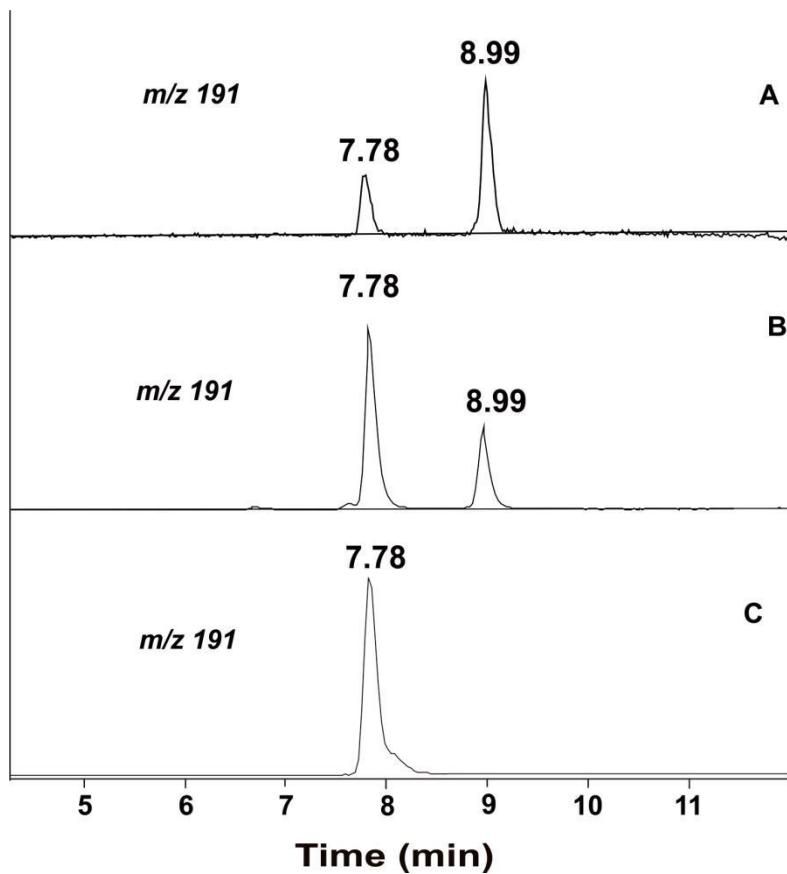
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**Figure S1. Michaelis-Menten plots of three substrates.** (A) Lineweaver-Burk of 4-hydroxy-benzenemethanol. (B) Lineweaver-Burk of 4-hydroxy-benzaldehyde. (C) Lineweaver-Burk of DMAPP.



**Figure S2. SIM chromatograms of the products by UPLC-HR-ESI-MS analysis.** The donor specificity of Vib-PT was probed using DMAPP, GPP, FPP or GGPP and 4-hydroxy-benzaldehyde as an acceptor. **DMAPP** shows the specific ion of product for DMAPP and 4-hydroxy-benzaldehyde selected ion monitoring (SIM) peak at  $m/z$  189.09125 [M - H] $^-$ ; **GPP** shows the specific ion of the product for GPP and 4-hydroxy-benzaldehyde SIM peak at  $m/z$  257.15457 [M - H] $^-$ ; **FPP** shows the specific ion of product for FPP and 4-hydroxy-benzaldehyde SIM peak at  $m/z$  325.21722 [M - H] $^-$ ; and **GGPP** shows the specific ion of the product for GGPP and 4-hydroxy-benzaldehyde SIM peak at  $m/z$  393.28005 [M - H] $^-$ . The retention times of these products were 12.7, 16.4, 18.2 and 19.5 min, consistent with their polarity.



**Figure S3. Chromatogram profiles for the products of 4-hydroxy-benzenemethanol by UPLC-MS analysis. (A)** Extracted ion chromatographs of UPLC-MS analysis of the products for DMAPP and 4-hydroxy-benzenemethanol Vib-PT for 1 h. **(B)** Extracted ion chromatographs of UPLC-MS analysis of the products for DMAPP and 4-hydroxy-benzenemethanol Vib-PT for 20 h. **(C)** 4OPBM was fast converted into 4O3PBM under acidic (such as 0.1% formic acid) and nonenzymatic conditions.