

Supporting Information

Identification of the Anti-infective Aborycin Biosynthetic Gene Cluster from Deep-Sea-Derived *Streptomyces* sp. SCSIO ZS0098 Enables Production in a Heterologous Host

Mingwei Shao^{1,2} Juying Ma,^{1,2} Qinglian Li,^{1,2} and Jianhua Ju^{1,2,*}

¹ CAS Key Laboratory of Tropical Marine Bio-resources and Ecology, Guangdong Key Laboratory of Marine Materia Medica, RNAM Center for Marine Microbiology, South China Sea Institute of Oceanology, Chinese Academy of Sciences, 164 West Xingang Road, Guangzhou 510301, China; jianting880720@126.com (M.S.); majunying@scsio.ac.cn (J. M.); liql@scsio.ac.cn (Q. L.)

² University of Chinese Academy of Sciences, 19 Yuquan Road, Beijing 110039, China

* Correspondence: jjju@scsio.ac.cn (J.J.); Tel./Fax: +86-20-3406-6449 (H.H.); Tel./Fax: +86-20-8902-3028 (J.J.)

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Table S1. Bacteria used in this study.

Strains	Description	Source/[Ref]
<i>E. coli</i>		
DH5α	Host strain for general clone	
BW25113/pIJ790	K-12 derivative: <i>araBAD</i> , <i>rhaBAD</i> ; host strain for Red/ET-mediated recombination	[1]
ET12567/pUZ8002	<i>dam</i> , <i>dcm</i> , <i>hsdM</i> , <i>hsdS</i> , <i>hsdR</i> , <i>catR</i> , <i>tetR</i> ; donor strain for conjugation between <i>E. coli</i> and <i>Streptomyces</i>	[2]
XL 1-Blue MR	Host strain for construction of genomic cosmid library	Stratagene
<i>Streptomyces</i>		
<i>Streptomyces coelicolor</i> M1152	Host strain for heterologous expression of the <i>abm</i> biosynthetic gene cluster	
<i>Streptomyces coelicolor</i> TK64	Host strain for heterologous expression of the <i>abm</i> biosynthetic gene cluster	
<i>Streptomyces</i> sp. SCSIO ZS0098	Wild-type producer of aborycin	This work

Table S2. Plasmids used in this study.

Plasmids	Description	Source/[Ref]
pIJ773	P1-FRT-oriT-aac(3)IV-FRT-P2	[3]
pIJ790	λ-RED (<i>gam bet exo</i>) CmlR <i>araCrep10lts</i>	[4]
pSET152AB	<i>aac(3)IV-oriT-intφC31</i>	
SuperCos1	Used for construction of genomic cosmid library	Stratagene
cosmid 15-12H	A cosmid which contains partial <i>abo</i> biosynthetic cluster	This work

Table S3. Primers used in this study.

Primer Name	Sequence (5'→3')	purpose
screen-orf(-3)-F	TGGAAGCCCGTCACCGCTA	For the screening of the genomic library
screen-orf(-3)-R	CGAACGCACCCACCAGAA	
screen-orf(+3)-F	TGGAAGCCCGTCACCGCTA	For the screening of the genomic library
screen-orf(+3)-R	CGAACGCACCCACCAGAA	
screen-aboB1-F	CGACTTCGGCTGGGTGTTCG	For the screening of the genomic library
screen-aboB1-R	TCGGCTGGAGGTGATGGGAC	

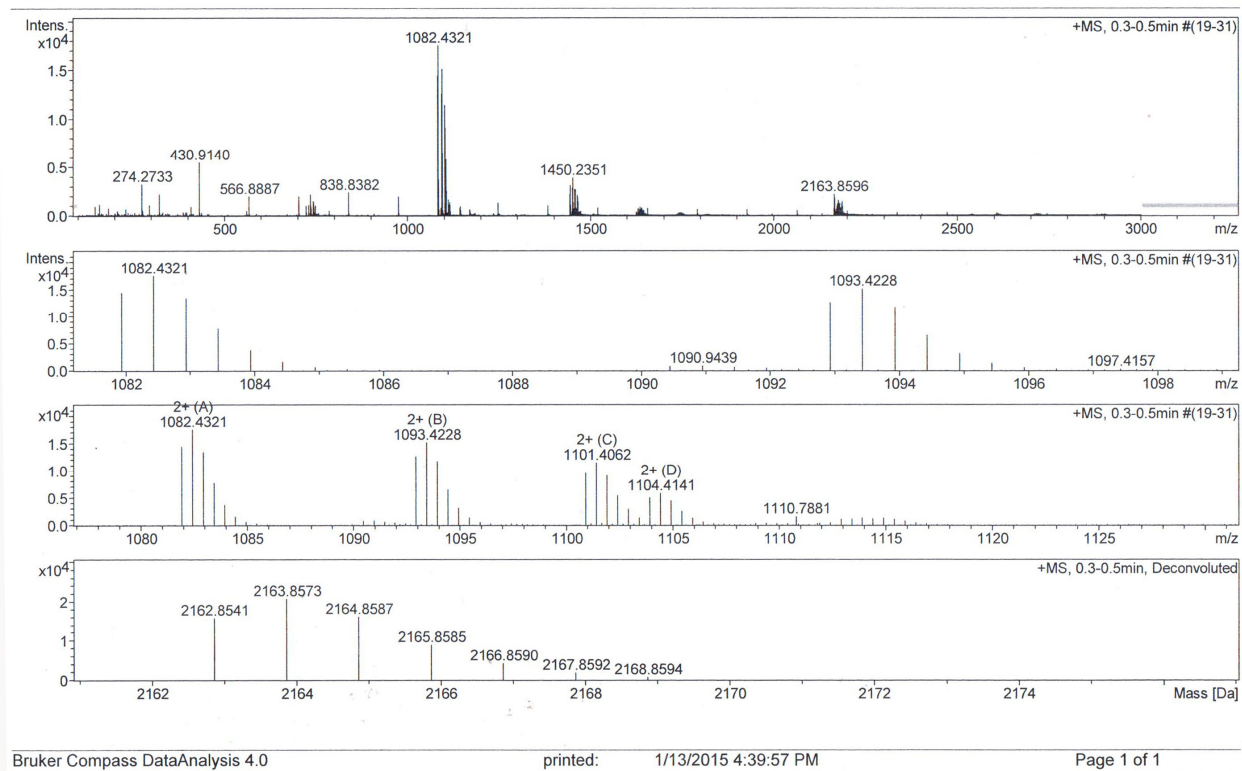


Figure S1. Positive ion peak of mass spectra of aborycin.

+MS2(887.3451), 45.2963-67.9444eV, 16.9min #994

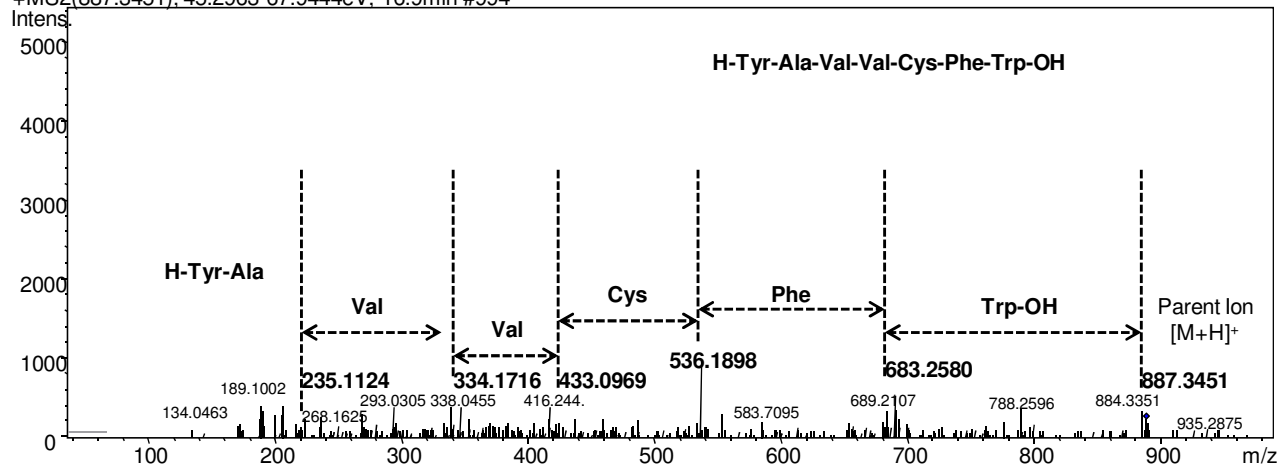


Figure S2. Tandem mass analysis and derived structure of a peptide (parent ion $[M + H]^+$ at $m/z = 887.3451$) yielded from acid hydrolysate of aborycin.

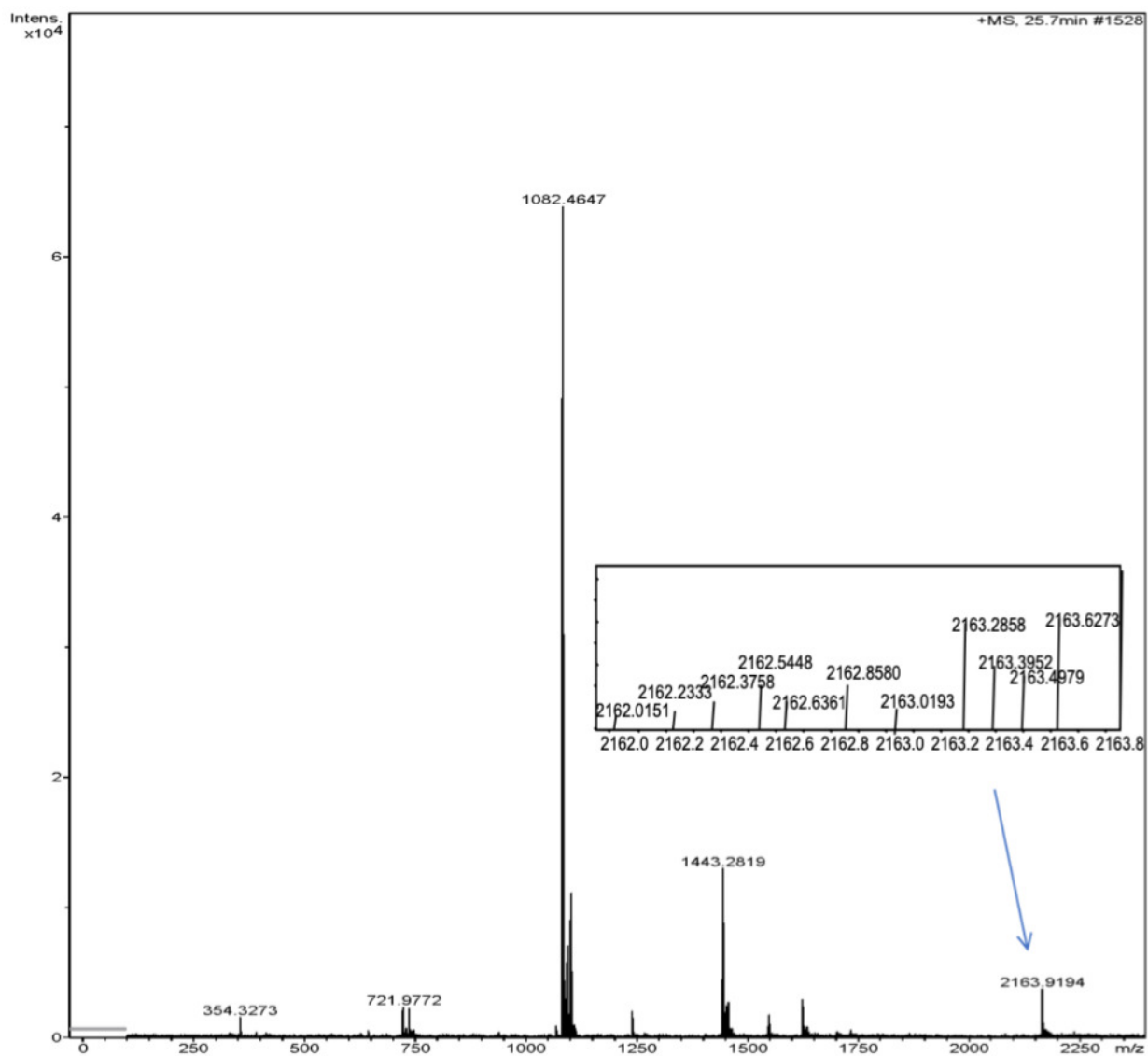


Figure S3. LC-MS analysis of aborycin accumulated in the cultures of *Streptomyces coelicolor* M1152/1512H (Positive mode).

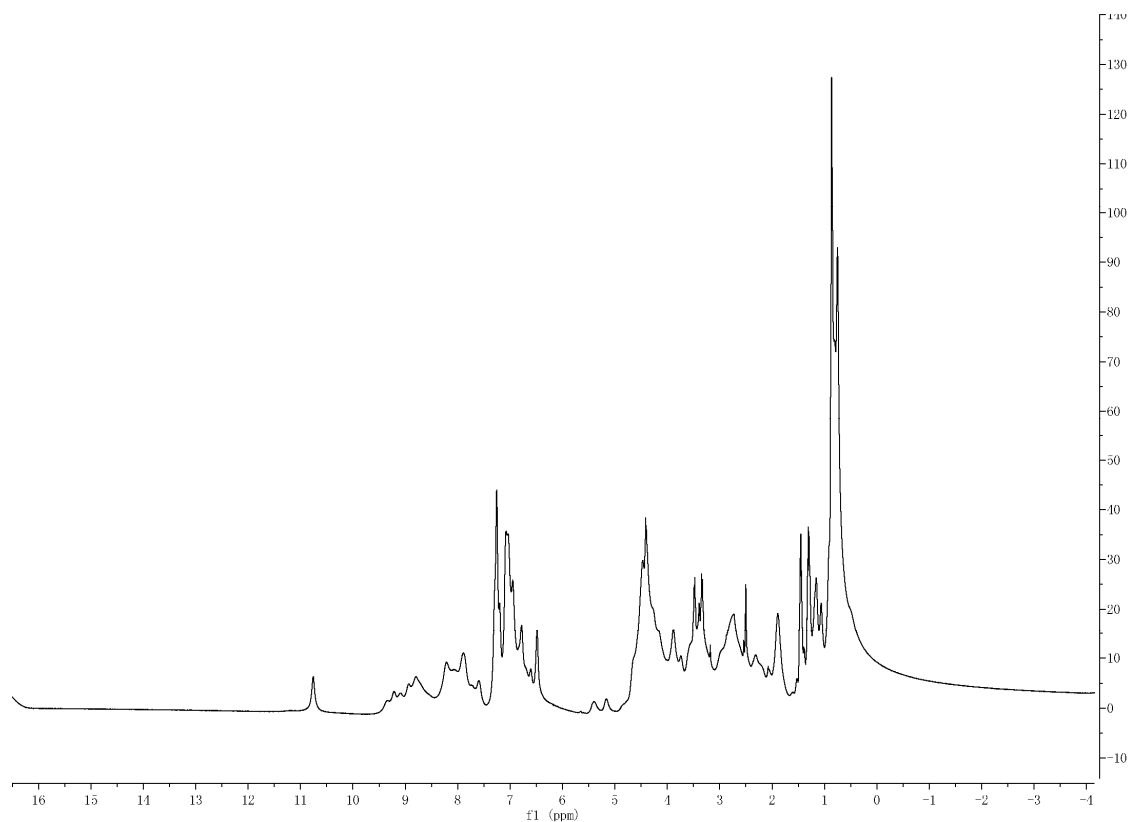


Figure S4. ^1H NMR (500 MHz) spectrum of aborycin in $\text{DMSO-}d_6$.

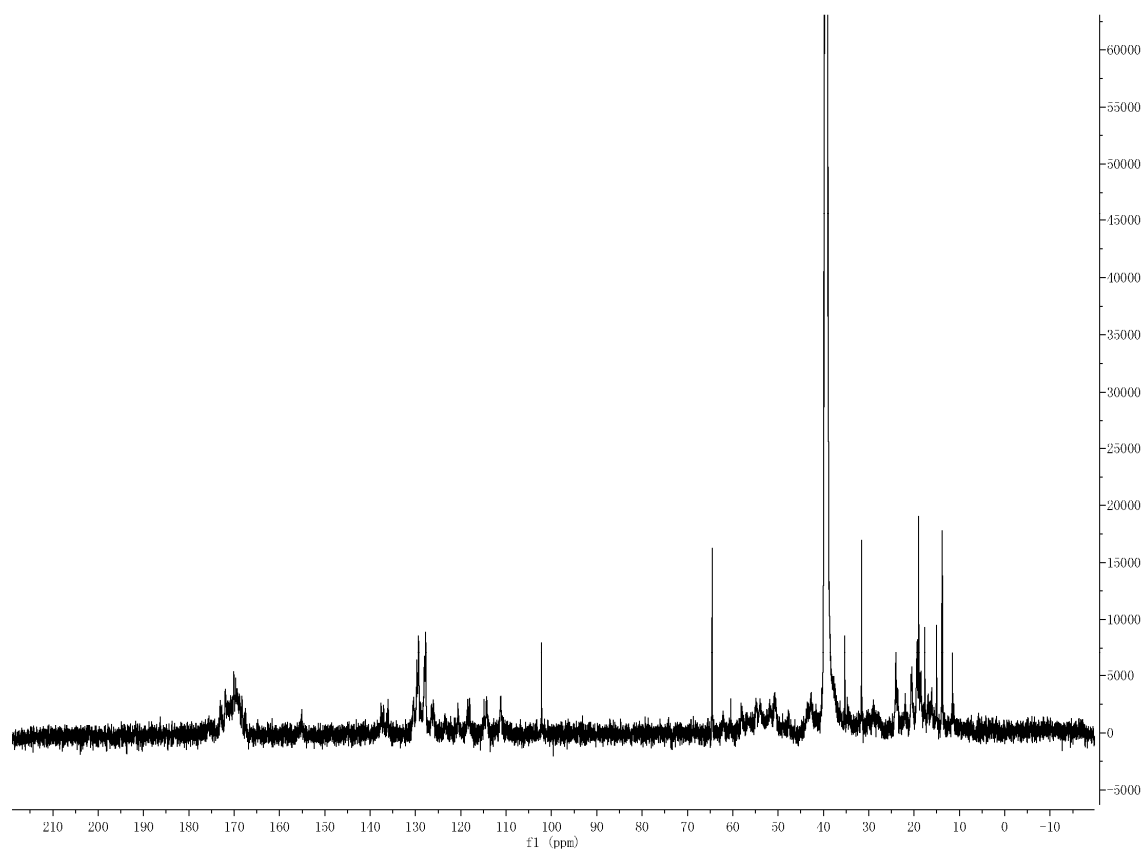


Figure S5. ^{13}C NMR (125 MHz) spectrum of aborycin in $\text{DMSO}-d_6$.

References

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