Supplementary Materials: Eremophilane Sesquiterpenes from a Deep Marine-Derived Fungus, *Aspergillus* sp. SCSIOW2, Cultivated in the Presence of Epigenetic Modifying Agents

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 Table S1. Conformer distribution of 1 in solvated models (methanol) calculation at the B3LYP/aug-cc-PVDZ level.

Conformers	Contribution %			
1	26.05			
2	19.17			
3	19.17			
4	14.77			
5	6.69			
6	5.01			
7	4.03			
8	2.25			
9	2.25			
10	0.61			





Figure S1. Cont.



Figure S1. Cont.



Figure S1. Effects of SBHA and 5-AZA on secondary metabolites (SMs) production by *Aspergillus* sp. SCSIOW2. SMs were extracted from one week cultures with (+) or without (-) 1 mM SBHA and 1 mM 5-AZA. (**A**) HPLC analyses [(a) = +; (b) = -; (c-f) = standards of 1–4]; Effects of SBHA and 5-AZA on secondary metabolites (SMs) production by *Aspergillus* sp. SCSIOW2. SMs were extracted from one week cultures with (+) or without (-) 1 mM SBHA and 1 mM 5-AZA. (**B**) LCMS analyses of samples with epigenetic modifiers. (a) = HPLC of +; (b–e) = SIM287, 301, 533, 285; Effects of SBHA and 5-AZA on secondary metabolites (SMs) production by *Aspergillus* sp. SCSIOW2. SMs were extracted from one week cultures with (+) or without (-) 1 mM SBHA and 1 mM 5-AZA. (**C**) LCMS analyses of samples without epigenetic modifiers. (a) = HPLC of -; (b–e) = SIM287, 301, 533, 285; Effects of SBHA and 5-AZA on secondary metabolites (SMs) production by *Aspergillus* sp. SCSIOW2. SMs were extracted from one week cultures with (+) or without (-) 1 mM SBHA and 1mM 5-AZA. (**C**) LCMS analyses of samples without epigenetic modifiers. (a) = HPLC of -; (b–e) = SIM287, 301, 533, 285; Effects of SBHA and 5-AZA on secondary metabolites (SMs) production by *Aspergillus* sp. SCSIOW2. SMs were extracted from one week cultures with (+) or without (-) 1 mM SBHA and 1mM 5-AZA. (**C**) LCMS analyses of samples without epigenetic modifiers. (a) = HPLC of -; (b–e) = SIM287, 301, 533, 285; Effects of SBHA and 5-AZA on secondary metabolites (SMs) production by *Aspergillus* sp. SCSIOW2. SMs were extracted from one week cultures with (+) or without (-) 1 mM SBHA and 1mM 5-AZA. (**D**) TLC analysis.



Figure S2. HPLC analysis of **1** (**a**), **2** (**b**) and **3** (**c**), and profile of **1** changed to **2** and **3** after storage for 1 week in MeOH:H₂O (9:1) solution (**d**); HPLC was performed with a YMC Pack pro ODS C18 column ($4.6 \times 250 \text{ mm I.D. 5 } \mu$) eluted with MeOH-H₂O (0:100–50:50 from 0 to 10 min; 50:50–80:20 from 10 to 20 min; 1.0 mL/min) with detection at 254 nm.



Figure S3. Cont.





Figure S3. Cont.









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Figure S5. ¹H-¹H COSY spectrum of **1** in DMSO-*d*₆.







Figure S7. HMBC spectrum of 1 in DMSO-d₆.



Figure S8. NOESY spectrum of 1 in DMSO-d6.



Figure S9. Cont.



Figure S9. Cont.



Figure S9. ¹H-NMR spectrum of 2 in DMSO-*d*₆.











Figure S10. Cont.





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Figure S11. ¹H-¹H COSY spectrum of 2 in DMSO-*d*₆.







Figure S13. HMBC spectrum of 2 in DMSO-*d*₆.



Figure S14. NOESY spectrum of 2 in DMSO-d₆.



Figure S15. Cont.





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Figure S15. ¹H-NMR spectrum of 3 in DMSO-*d*₆.



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138.80	132.10 127.57 127.13 127.13	117.44	104.26	666.28 65.83 61.17	42.74 41.54 39.66 39.52 39.28 39.24	22.37	10.51
	VV			V I		V	Y



Figure S16. Cont.

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Figure S17. ¹H-¹H COSY spectrum of 3 in DMSO-*d*₆.



Figure S18. HSQC spectrum of 3 in DMSO-d₆.



Figure S19. HMBC spectrum of 3 in DMSO-*d*₆.





7.5 7.0 6.5 6.0 5.5 5.0 4.5 4.0 3.5 3.0 2.5 2.0 1.5 1.0 0.5 ppm

ppm 0.0 1.0 2.0 2.5 3.0

3.5 4.0 4.5 5.0

6.0 6.5 7.0 7.5



Figure S21. Cont.



Figure S21. Cont.

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Figure S22. Cont.





Figure S22. Cont.



Figure S22. Cont.









Figure S23. ¹H-¹H COSY spectrum of 4 in DMSO-*d*₆.



Figure S24. HSQC spectrum of 4 in DMSO-d₆.



Figure S25. HMBC spectrum of 4 in DMSO-*d*₆.



Figure S26. NOESY spectrum of 4 in DMSO-d6.







Figure S27. Cont.





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