Miharadienes A–D with Unique Cyclic Skeletons from a Marine-Derived *Streptomyces miharaensis*

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Figure S1. HRESIMS data of 1.



Figure S2. ¹H NMR spectrum of miharadiene (1) in CD₃OD.



Figure S3. ¹³C NMR spectrum of miharadiene A (1) in CD₃OD.



Figure S4. ¹H-¹H COSY spectrum of miharadiene A (1) in CD₃OD.



Figure S5. HSQC spectrum of miharadiene A (1) in CD₃OD.



Figure S6. HMBC spectrum of miharadiene A (1) in CD₃OD.



Figure S7. NOESY spectrum of miharadiene A (1) in CD₃OD.



Figure S8. HRESIMS spectrum of miharadiene B (2).



Figure S9. ¹H NMR spectrum of miharadiene B (2) in CDCl₃.



Figure S10. ¹³C NMR spectrum of miharadiene B (2) in CDCl₃.



Figure S11. ¹H-¹H COSY spectrum of miharadiene B (**2**) in CDCl₃.



Figure S12. HSQC spectrum of miharadiene B (2) in CDCl₃.



Figure S13. HMBC spectrum of miharadiene B (2) in CDCl₃.

Scheme S1. Determination of absolute configuration for 2.





Figure S14. ¹H NMR spectrum of compound 2a in CD₃OD.



Figure S15. ¹³C NMR spectrum of compound 2a in CD₃OD.



Figure S16. HSQC spectrum of compound 2a in CD₃OD.



Figure S17. Selected regions of ¹H spectrum of Mosher's ester analysis of hydrolysate of 2a.



Figure S18. NOESY spectrum of compound 2a in CD₃OD.

Mass	Calc. Mass	mDa	PPM	DBE	Formula
368.1837	368.1838	-0.1	-0.3	7.5	C20 H27 N O4 Na



Figure S19. HRESIMS spectrum of miharadiene C (3).



Figure S20. ¹H NMR spectrum of miharadiene C (3) in CDCl₃.



Figure S21. ¹³C NMR spectrum of miharadiene C (3) in CDCl₃.



Figure S22. ¹H-¹H COSY spectrum of miharadiene C (3) in CDCl₃.



Figure S23. HSQC spectrum of miharadiene C (3) in CDCl₃.



Figure S24. HMBC spectrum of miharadiene C (3) in CDCl₃.



Figure S25. ¹H NMR spectrum of miharadiene C (3) in DMSO-*d*₆.



Figure S26. ¹H-¹H COSY spectrum of miharadiene C (3) in DMSO- d_6 .



Figure S27. HMBC spectrum of miharadiene C (3) in DMSO- d_6 .



Figure S28. HRESIMS spectrum of miharadiene D (4).



Figure S29. ¹H NMR spectrum of miharadiene D (4) in CD₃OD.



Figure S30. ¹³C NMR spectrum of miharadiene D (4) in CD₃OD.



Figure S31. ¹H-¹H COSY spectrum of miharadiene D (4) in CD₃OD.



Figure S32. HSQC spectrum of miharadiene D (4) in CD₃OD.



Figure S33. HMBC spectrum of miharadiene D (4) in CD₃OD.

X-ray Crystallographic analysis. Single-crystal X-ray diffraction data were collected using an Bruker SMART APEX2 ULTRA and a APEX II CCD area detector with a multilayer-monochromated Mo K α radiation ($\lambda = 0.71073$ Å) generated by a rotating anode. Data collection, data reduction, and semiempirical absorption correction were carried out using the software package APEX2.(1) All of the calculations for the structure determination were carried out using the SHELXTL package. (2) All non-H atoms were refined anisotropically. All hydrogen atoms were included in calculated positions with isotropic thermal parameters 1.2 times those of attached atoms.

(1) APEX2 (Version 2009.1–0) Data Collection and Processing Software; Bruker AXS Inc.: Madison, Wisconsin, U.S.A., 2008.
(2) SHELXTL-PC (Version 6.22) Program for Solution and Refinement of Crystal Structures; Bruker AXS Inc.: Madison, Wisconsin, U.S.A., 2001.



Crystal data for **3.** Formula= $C_{22}H_{39}CuN_5O_6$, crystal size = $0.30 \times 0.24 \times 0.13$ mm, Monoclinic, space group $P2_1/n$, a = 16.5612(2) Å, b = 9.2100(1) Å, c = 16.7191(2) Å, $\beta = 115.463(1)^\circ$, V = 2303.4(1) Å³, Z = 4, $D_c = 1.538$ g cm⁻³, $\mu = 0.669$ mm⁻¹, T = 173(2) K, F(000) = 1132, $T_{max}/T_{min} = 0.89/0.80$, $2\theta_{max} = 27.54^\circ$; 5297 reflections collected, 307 independent reflections, $R_{int} = 0.0317$, R1 = 0.03 and R2 = 0.08 for I $\ge 2\sigma(I)$, R1 = 0.39 and R2 = 0.08 for all data, GoF = 1.044.

	3
Formula	$C_{13}H_{12}BrCuN_4S_2$
M	431.84
T/K	173(2)
Crystal system	Triclinic
Space group	<i>P</i> -1
a/Å	8.6303(13)
b/Å	9.8063(12)
$c/\text{\AA}$	9.9415(14)
$lpha/^{\circ}$	69.229(8)
$\beta/^{\circ}$	71.497(6)
$\gamma/^{\circ}$	81.910(8)
$V/Å^3$	745.6(2)
Ζ	2
μ (Mo-K α)/mm ⁻¹	4.420
Crystal size/mm	$0.08 \times 0.05 \times 0.02$
Absorption correction	Multi-scan SADABS
Reflections collected	11391
Independent reflections	2898
Goodness-of-fit on F^2	1.064
Final R1, wR2 $[I > 2\sigma(I)]$	0.1134, 0.3054
(all data)	0.1456, 0.3367

Table S1. (Crystal	data	and	structural	refinement
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Figure S34. DFT optimized conformers and populations of miharadiene A (3*S*, 5*S*) above 5% population.



Figure S35. DFT optimized conformers and populations of miharadiene D (3*R*) above 2% population.

B3LYP/6-311+G(d,p) Gibbs free energy (298.15K)				
	G (Hartree)	Population (%)		
Conformer 1	-1152.483640	20.77		
Conformer 2	-1152.486823	13.53		
Conformer 3	-1152.487408	13.17		
Conformer 4	-1152.487148	9.44		
Conformer 5	-1152.485161	6.21		

Table S2. Gibbs free energies and Boltzmann distribution of conformers of 1

Table S3. Gibbs free energies and Boltzmann distribution of conformers of 4

B3LYP/6-311+G(d,p) Gibbs free energy (298.15K)				
	G (Hartree)	Population (%)		
Conformer 1	-808.849283	8.17		
Conformer 2	-808.850206	5.50		
Conformer 3	-808.848945	4.01		
Conformer 4	-808.849516	3.85		
Conformer 5	-809.274794	3.17		