Supporting Information: Isolation of Isotrichophycin

C and Trichophycins G-I from a Collection of

Trichodesmium thiebautii

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Figure S32. Comparison of ¹H NMR spectra of *S*-MTPA and *R*-MTPA esters of trichophycin A with partial $\Delta \delta S - \delta R$ values noted (500 MHz, CDCl₃) as well as HRESIMS values.

Figure S33. Comparison of ¹H NMR spectra of *S*-MTPA and *R*-MTPA esters of **3** with partial $\Delta \delta S - \delta R$ values noted (500 MHz, CDCl₃).

Figure S34. EC₅₀ curves of doxorubicin, **1-3**, trichophycin A triacetate and isotrichophycin C diacetate.

Figure S35. 16S rRNA phylogenetic tree aligning *Trichodesmium* species in this study with *T. thiebautii* strains.

pos	$\delta_{\rm H}$ (<i>J</i> in Hz)	TOCSY	
1a	5.17 m	2	
1u 1b	5.17, m 5.14 m	2	
2	5 84 m	- 1 3b	
- 3a	2.32 m	2, 3b, 4	
3h	2.10 m	2,33,4	
4	$3.47 \text{ ovl}\text{p}^{a}$	$\frac{2}{3a}$ 3b 5	
5	1 63 m	4 6a 6b 22	
6a	142 ddd (135 86 46)	5 6b	
6b	1.00. ddd (14.2, 9.2, 5.6)	5. 6a. 7	
7	1.54. ovlp	21	
8a	1.35. ovlp	7. 8b	
8b	1.19. m	8a, 9b	
9a	1.50. ovlp		
9b	1.45, m		
10	3.79, m	9a, 9b, 11a, 11b	
11a	2.39, dd (13.7, 8.8)	10, 11b	
11b	2.28, dd (13.5, 4.2)	10, 11a	
12			
13	3.47, ovlp	20	
14	· •		
15	7.18, d (7.4)	16	
16	7.31, t (7.4)	17	
17	7.24, m		
18	7.31, t (7.4)		
19	7.18, d (7.4)	18	
20	6.00, s	13	
21	0.91, ovlp	7	
22	0.91, ovlp	5	

 Table S1. NMR Data for Trichophycin G (2) (500 MHz for ¹H NMR, CDCl₃).

^aoverlapping signals

pos	δc , type	$\delta_{\rm H}$ (<i>J</i> in Hz)	HMBC	COSY	$\Delta \delta^{13} C^a$
19	1179 CH	5.15 m		2	0.0
16 1h	117.9, CH2	5.13, m		2	0.0
2	135.6 CH	5.13, m 5.87 m		1 3a 3h	0.0
2 3a	39.0 CH2	2.07, m		1, 5 u , 50 4	0.0
3h	59.0, 0112	2.20, m 2.17 m		4	0.1
4	739 CH	3.56 dt (8.5, 4.0)		3a 3h 5	0.0
5	37.8 CH	$1.54 \text{ ovln}^{\text{b}}$		4 27	0.0
5 6a	33.0 CH ₂	1.5 i, ovip		6h	0.0
6h	55.0, 0112	1.10, m 1.26 ovln		60 6a	0.0
00 7a	27.4 CH2	1.20, 0Vlp		7h	0.0
7h	27.1, 0112	1 35 ovlp		78 7a	0.0
8a	267	1 42 ovlp		, a	0.0
8b		1 32 m			
9	34.6 CH2	1 42 ovlp	10 11	8b	0.0
10	74.5 CH	3 50 dt (7 6 4 0)	10, 11	9 11	0.0
11	35.2. CH	1.61. m		10.26	0.0
12a	40.8, CH ₂	1.41, ovlp	10, 25, 26	11, 13	0.0
12b	, _	1.02, dt (14.1, 7.4)	10, 25, 26	11, 12a	
13	29.8, CH	1.57, ovlp	, ,	14a, 25	0.0
14a	32.6, CH ₂	1.42, ovlp		14b	0.1
14b	,	1.26, ovlp		14a, 15a	
15a	34.8, CH ₂	1.55, ovlp		15b	0.0
15b		1.44, ovlp		15a	
16	70.6, CH	3.82, m		15a, 17a, 17b	0.1
17a	38.2, CH ₂	2.46, dd (13.6, 8.7)	16, 18, 19, 24	16, 17b	0.0
17b		2.36, dd (13.6, 4.4)	16, 18, 24	16, 17a	
18	139.0, C				
19	34.2, CH ₂	2.28, m	18, 20, 21, 24	20	
20	26.3, CH ₂	1.68, m	18, 19, 21, 22	19a, 19b	
21	17.8, CH ₂	2.20	19, 22	20	
22	83.5, C				
23	68.8, CH	1.98, t (2.7)		21	
24	114.9, CH	5.99, s	18	19a	
25	20.4, CH ₃	0.90, ovlp	12, 13	13	0.0
26	14.0, CH ₃	0.86, d (6.8)	10, 11, 12	11	0.1
27	14.0, CH ₃	0.91, ovlp	4, 5, 6	5	0.1

 Table S2. NMR Data for Trichophycin H (3) (500 MHz for ¹H NMR, CDCl₃).

 $a\Delta\delta^{13}C$ (trichophycin A-3) ^boverlapping signals

pos	δc , type	$\delta_{\rm H} (J \text{ in Hz})$	HMBC	COSY
1a	117.3, CH ₂	5.07, m	3	2, 3
1b		5.03, m	3	2, 3
2	134.1, CH	5.71, m		1a, 1b, 3
3	35.8, CH ₂	2.27, m	1, 2, 4	2, 4
4	76.6, CH	4.80, ddd (8.0, 5.9, 4.5)	2	3, 5
5	36.0, CH	1.67, m		4, 17
6a	29.2, CH ₂	1.50, m		6b, 7
6b		1.17, m		5, 6a, 7
7	27.6, CH ₂	2.14, m	6, 8, 16	6a, 6b
8	141.9, C			
9	41.2, CH ₂	3.36, m	7, 8, 10, 11/15, 16	16
10	138.2, C			
11/15	128.9, CH	7.15, d (7.4)	9,13	9, 12/14
12/14	128.5, CH	7.30, t (7.4)	10	11/15
13	126.6, CH	7.22, t (7.4)		
16	114.2, CH	5.84, s	7, 8, 9	9
17	15.2, CH ₃	0.90, d (6.8)	4, 5, 6	5
18	170.7, C			
19	21.2, CH ₃	2.02, s	18	

Table S3. NMR Data for Trichophycin I (4) (500 MHz for ¹H NMR, CDCl₃)

Molecule	Configuration	¹³ C NMR C-17 ^a	
Trichophycin A	Ζ	38.2	
Isotrichophycin C (1)	Ζ	38.2	
Trichophycin H (3)	Ζ	38.2	
Smenlactone B	Ζ	38.9	
Smenlactone C	Ζ	38.9	
Isomalyngamide K	Ζ	39.0	
Trichophycin B	E	42.7	
Trichophycin C	E	42.8	
Trichophycin D	E	43.1	
Trichophycin E	E	43.1	
Malyngamide K	E	44.0	
aC 17 in trich on hyperin	٨		

 Table S4. NMR Data for Proposed Chlorovinylidene Configuration

^aC-17 in trichophycin A



Figure S1. Previously reported polyketide metabolites from *Trichodesmium* 2014 collection and new metabolites from this report. In this report, we address the absolute configuration of the previously published trichophycin A.



Figure S3. ¹H NMR spectrum of isotrichophycin C (1) (800 MHz, CDCl₃).

Figure S5. HSQC spectrum of 1.

Figure S6. HMBC spectrum of 1.

Figure S7. DQF-COSY spectrum of 1.

Figure S8. TOCSY spectrum of 1.

Figure S9. NOESY spectrum of 1.

Figure S10. HRESIMS of 1.

Figure S11. Comparison of trichophycin C and isotrichophycin C (1) ECD spectra (0.4 mg/mL, CH₃CN).

Figure S12. ¹H NMR spectrum of trichophycin G (2) (500 MHz, CDCl₃).

Figure S13. COSY spectrum of 2.

Figure S14. HRESIMS of 2.

Figure S15. Comparison of ¹H NMR spectra: trichophycin A (800 MHz, CDCl₃) and trichophycin H (3) (500 MHz, CDCl₃).

Figure S16. ¹H NMR of trichophycin H (**3**) (500 MHz, CDCl₃).

Figure S19. COSY spectrum of 3.

Figure S20. HRESIMS of 3.

Figure S22. HSQC of 4.

Figure S23. HMBC of 4.

Figure S24. COSY of 4.

Figure S25. NOESY of 4.

Figure S26. HRESIMS of 4.

Figure S27. ¹H NMR of trichophycin A triacetate (500 MHz, CDCl₃).

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Figure S34. EC₅₀ curves of doxorubicin, 1-3, trichophycin A triacetate and isotrichophycin C diacetate.

Figure S35. 16S rRNA phylogenetic tree aligning *Trichodesmium* species in this study (PI2014, red arrow) with *T. thiebautii* strains. The tree was created using the Maximum Likelihood method and the Tamura-Nei model. The bootstrap consensus tree is inferred from 1000 replicates and the percentage of replicate trees in which the associated taxa clustered together in the bootstrap test are shown next to branches. Analysis was conducted in MEGA. Genbank accession numbers of sequences are noted in parentheses.