

SUPPLEMENTARY MATERIAL

Actinobacteria associated with stingless bee biosynthesize bioactive polyketides against bacterial pathogen

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Table S1. 16S rRNA identification of actinobacteria isolated from *Mellipona scutellaris* colony.

Isolates	Isolation source	NCBI accession number	Closest relative (accession number)
ICBG1307 - <i>Streptomyces</i> sp.	<i>Mellipona scutellaris</i> nurse bee	MK608316	<i>Streptomyces kunmingensis</i> (AB184597)
ICBG1313 - <i>Streptomyces</i> sp.	<i>M. scutellaris</i> nurse bee	MK608317	<i>Streptomyces drozdowiczi</i> (AB249957)
ICBG1318 - <i>Streptomyces</i> sp.	<i>M. scutellaris</i> nurse bee	MK608318	<i>Streptomyces albiaxialis</i> (AY999901)
ICBG1319 - <i>Streptomyces</i> sp.	<i>M. scutellaris</i> nurse bee	MK608319	<i>Streptomyces xylanilyticus</i> (LC128341)
ICBG1321 - <i>Micromonospora</i> sp.	<i>M. scutellaris</i> nurse bee	MK608320	<i>Micromonospora tulbaghiae</i> (Jgi.1058868)
ICBG1323 - <i>Streptomyces</i> sp.	<i>M. scutellaris</i> foraging bee	MK608321	<i>Streptomyces olivaceus</i> (JOFH01000101)
ICBG1324 - <i>Streptomyces</i> sp.	<i>M. scutellaris</i> foraging bee	MK608322	<i>Streptomyces rhizosphaerhabitans</i> (HQ267983)

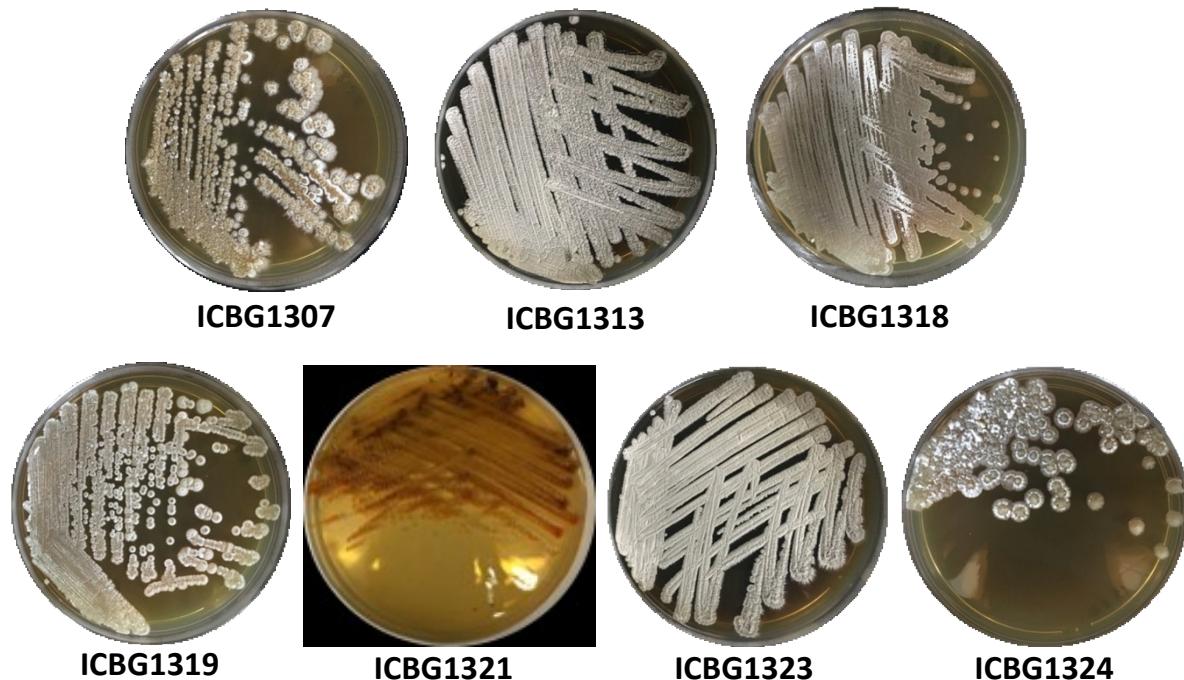


Fig. S1 Macroscopic characteristics of actinobacterial strains isolated from *M. scutellaris*, cultured in ISP-2 agar during 10 days at 30 °C (*Streptomyces* sp. ICBG1307, *Streptomyces* sp. ICBG1313, *Streptomyces* sp. ICBG1318, *Streptomyces* sp. ICBG1319, *Streptomyces* sp. ICBG1323, *Streptomyces* sp. ICBG1324, and *Micromonospora* sp. ICBG1321).

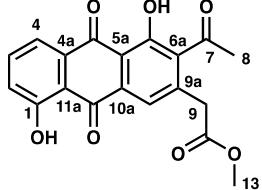
Table S2. Visual depiction of antagonist assay between isolated actinobacteria against *P. larvae* ATCC 9545, *M. anisopliae* and *B. bassiana*.

Strains	<i>P. larvae</i>	<i>M. anisopliae</i>	<i>B. bassiana</i>	Inhibition	
<i>Streptomyces</i> sp. ICBG1307	0	0	0		
<i>Streptomyces</i> sp. ICBG1313	0	0	0	3	
<i>Streptomyces</i> sp. ICBG1319	0	0	0	2	
<i>Streptomyces</i> sp. ICBG1324	0	0	0	1	
<i>Streptomyces</i> sp. ICBG1318	3	0	0	0	High
<i>Streptomyces</i> sp. ICBG1323	3	0	0		Low
<i>Micromonospora</i> sp. ICBG1321	3	0	0		

Table S3. ^1H (500 MHz) and ^{13}C (125 MHz) NMR data for compound **8** (CD_3OD).

	Compound 8	
#	^{13}C NMR	^1H NMR ($m, J = \text{Hz}$)
1	163.8 (C)	-
2	125.8 (CH)	7.34 (d, 8.1)
3	137.9 (CH)	7.76 (dd, 7.5; 8.1)
4	120.12 (CH)	7.82 (d, 7.4)
4a	134.67 (C)	-
5	188.8 (C)	-
5a	115.1 (C)	-
6	159.8 (C)	-
6a	134.14 (C)	-
7	44.18 (CH)	4.25 (d, 4.0)
8A	36.01 (CH ₂)	2.76 (d, 12.7)
8B		2.47 (dd, 12.7; 4.0)
9	85.1 (C)	-
10	58.14 (CH)	4.07 (s)
10a	142.9 (C)	-
11	120.03 (CH)	7.66 (s)
11a	134.67 (C)	
12	188.27 (C)	-
12a	116.7 (C)	-
13	23.63 (CH ₃)	1.58 (s)
14	172.1 (C)	-
15	53.05 (CH ₃)	3.82 (s)
2'	116.8 (C)	-
3'	134.8	5.72 (t, 2.1)
3a'	141.4 (C)	-
4'	21.67 (CH ₂)	2.85 (m)
5'	54.16 (CH ₂)	4.17 (m)
5a'	168.8 (C)	-

Table S4. ^1H (500 MHz) and ^{13}C (125 MHz) NMR data for compound **13** and their isomer reported by KUNNARI et al. (1999) (CDCl_3).

 Compound 13		Kunnari et al., 1999		
#	^{13}C NMR	^1H NMR ($m, J = \text{Hz}$)	^{13}C NMR	^1H NMR ($m, J = \text{Hz}$)
1	162.8 (C)	-	119.8 (CH)	7.85 (dd, 7.4; 1.2)
1-OH	-	11.93 (s)	-	-
2	124.7 (CH)	7.34 (d, 8.4)	132.7 (CH)	7.73 (dd, 8.5; 7.4)
3	137.4 (CH)	7.73 (dd, 8.4; 7.5)	124.6 (CH)	7.34 (dd, 8.5; 1.2)
4	120.2 (CH)	7.85 (d, 7.5)	162.1 (C)	-
4-OH	-	-	-	11.94 (s)
4a	133.3 (C)	-	114.8 (C)	-
5	180.9 (C)	-	180.4 (C)	-
5a	115.6 (C)	-	115.1 (C)	-
6	160.1 (C)	-	159.5 (C)	-
6-OH	-	12.53 (s)	-	12.54 (s)
6a	135.9 (C)	-	135.5 (C)	-
7	202.8 (C)	-	202.2 (C)	-
8	31.5 (CH ₃)	2.68 (s)	31.2 (CH ₃)	2.96 (s)
9	38.4 5 (CH ₂)	3.87 (s)	38.3 (CH ₂)	3.88 (s)
9a	142.5 (C)	-	141.1 (C)	-
10	122.3 (CH)	7.70 (s)	121.9 (CH)	7.71 (s)
10a	Not observed	-	132.7 (C)	-
11	181.2 (C)	-	192.1 (C)	-
11a	116.0 (C)	-	133.0 (C)	-
12	170.9 (C)	-	169.7 (C)	-
13	52.7 (CH ₃)	3.71 (s)	51.9 CH ₃)	3.72 (s)

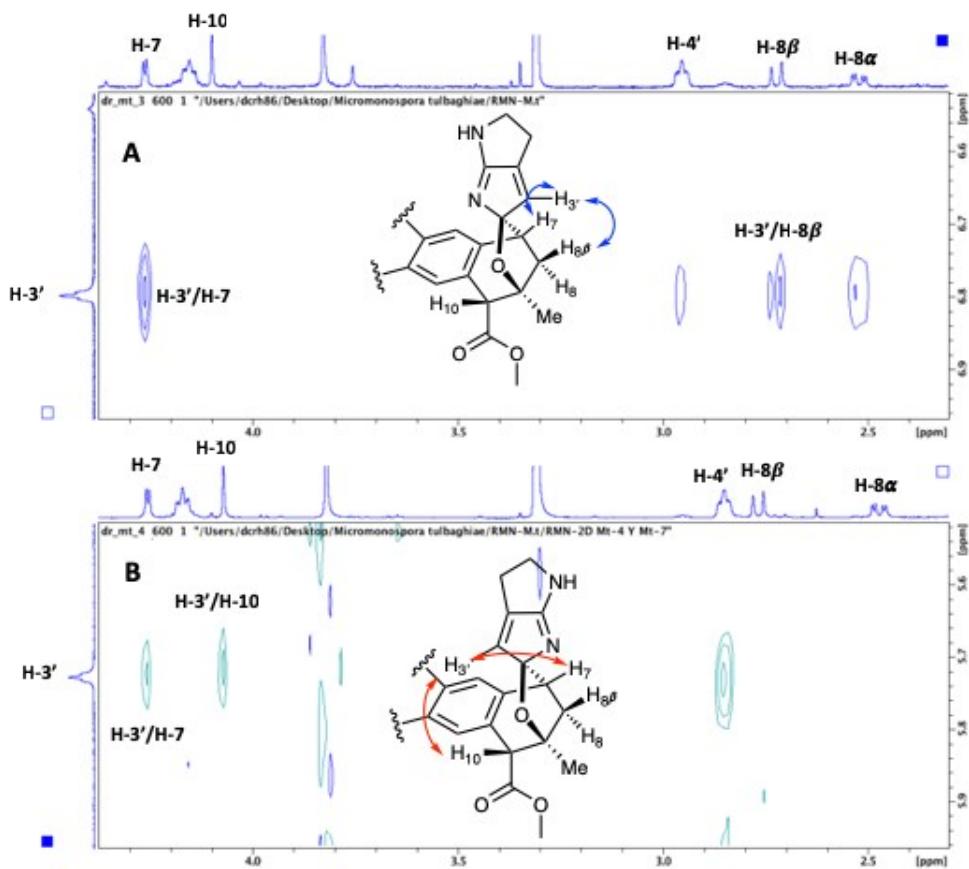


Fig. S2. NOESY correlations. A) compounds **7**; B) compound **8**.

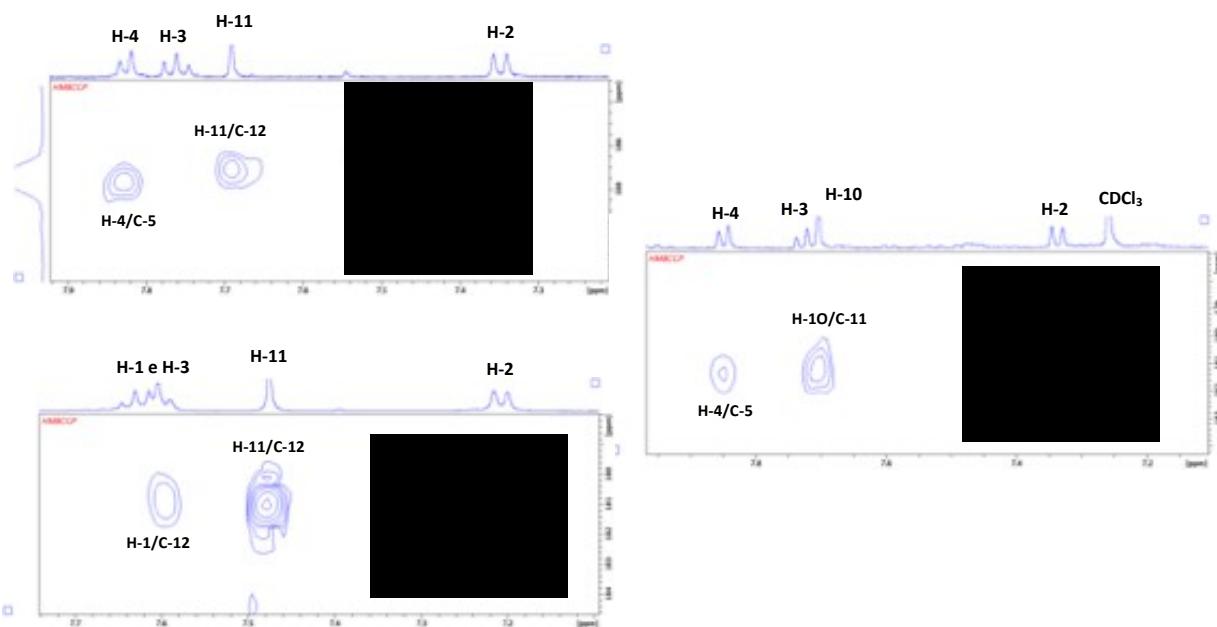


Fig. S3. gHMBC correlations of compounds **7**, **9** and **13**.

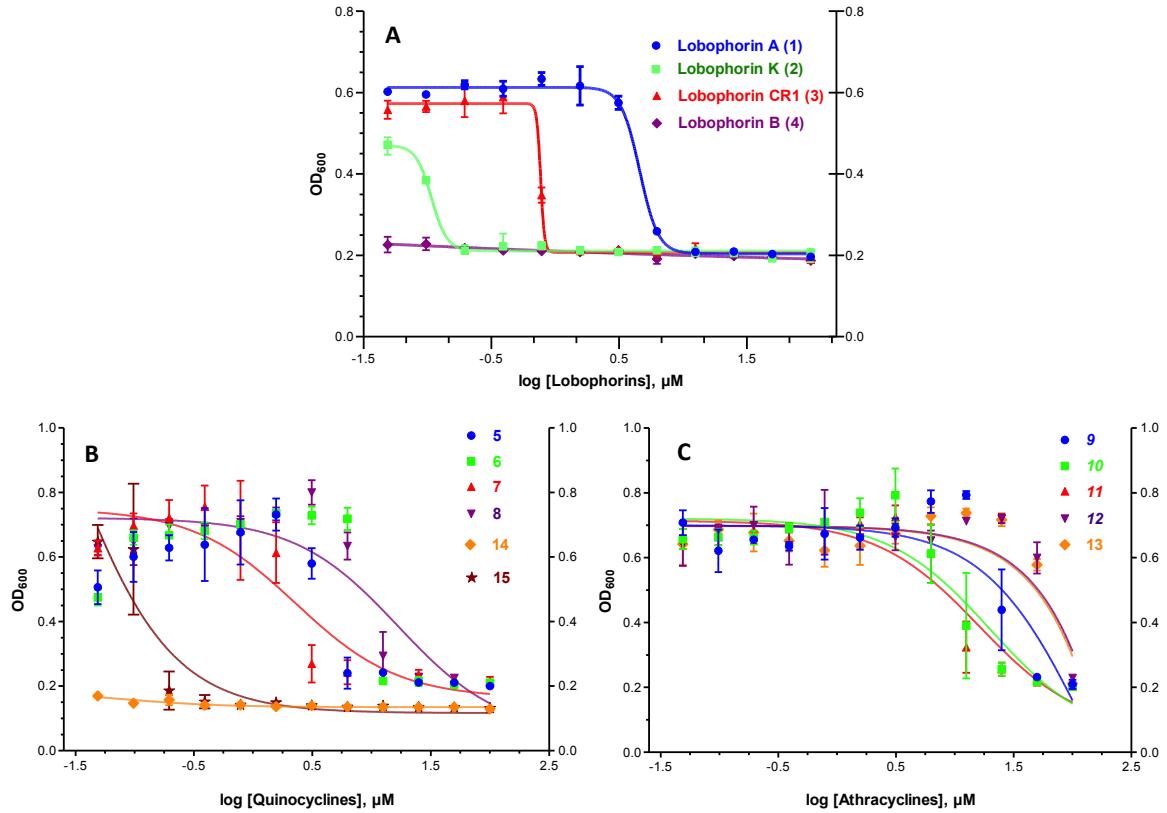


Figure S4. Dose-response curve using the isolated compounds (**1-15**) against *P. larvae* ATCC 9545. A) Lobophorins (**1-4**); B) Quinocyclines (**5-8** and **14-15**) and C) Anthracyclines (**9-13**). The dose-response curves log (Concentration of compounds **1-15**) versus OD₆₀₀ values of *P. larvae* ATCC 9545 were obtained using GraphPad Prism 5.0 software, through of nonlinear regression analysis and were performed in triplicate.

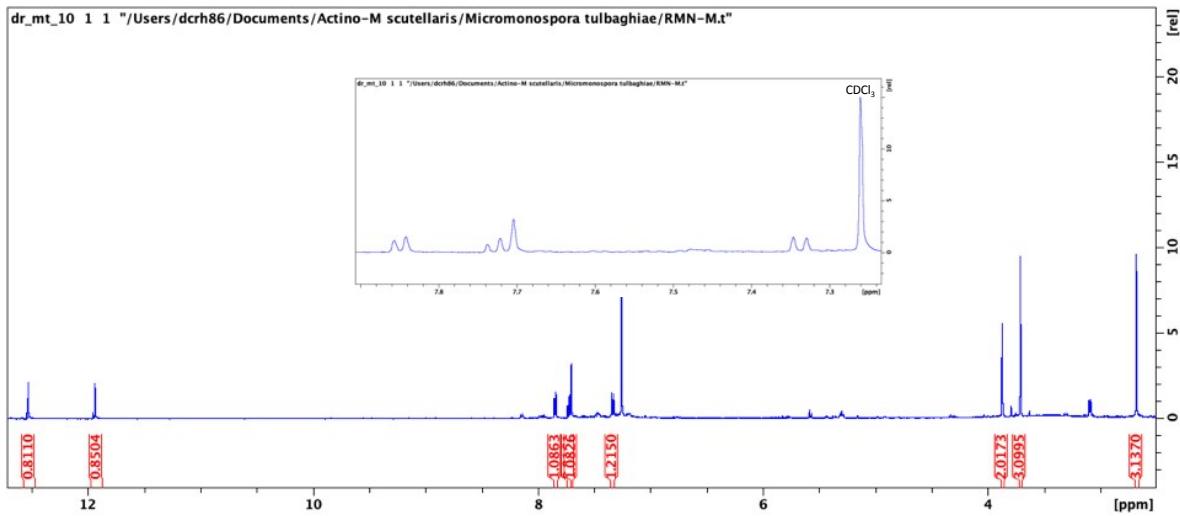


Fig. S5. ¹H NMR spectrum of **13** (500 MHz, CDCl₃).

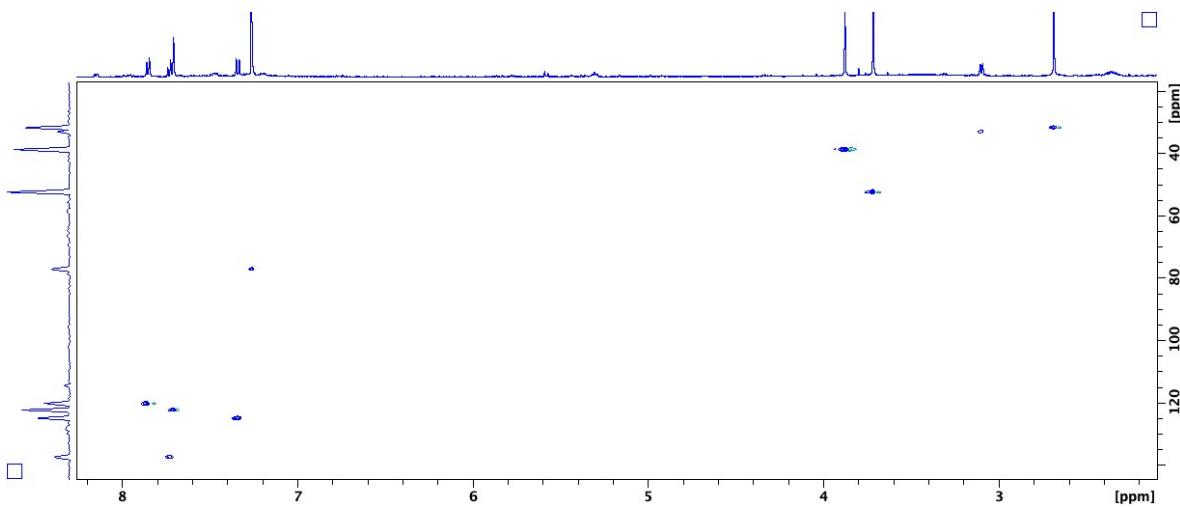


Fig. S6. gHSQC spectrum of **13** in CDCl₃.

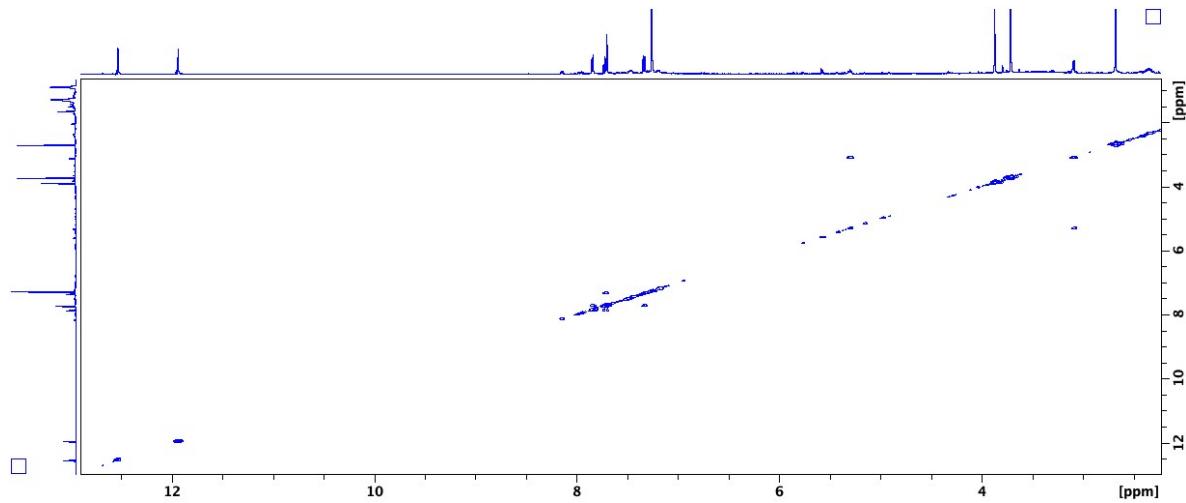


Fig. S7. *g*COSY spectrum of **13** in CDCl_3 .

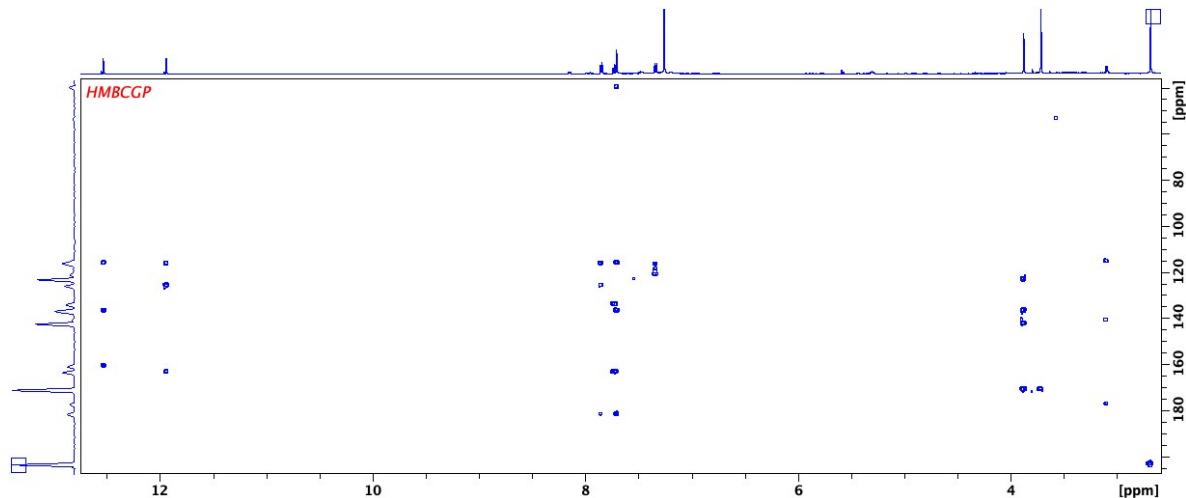


Fig. S8. *g*HMBC spectrum of **13** in CDCl_3 .

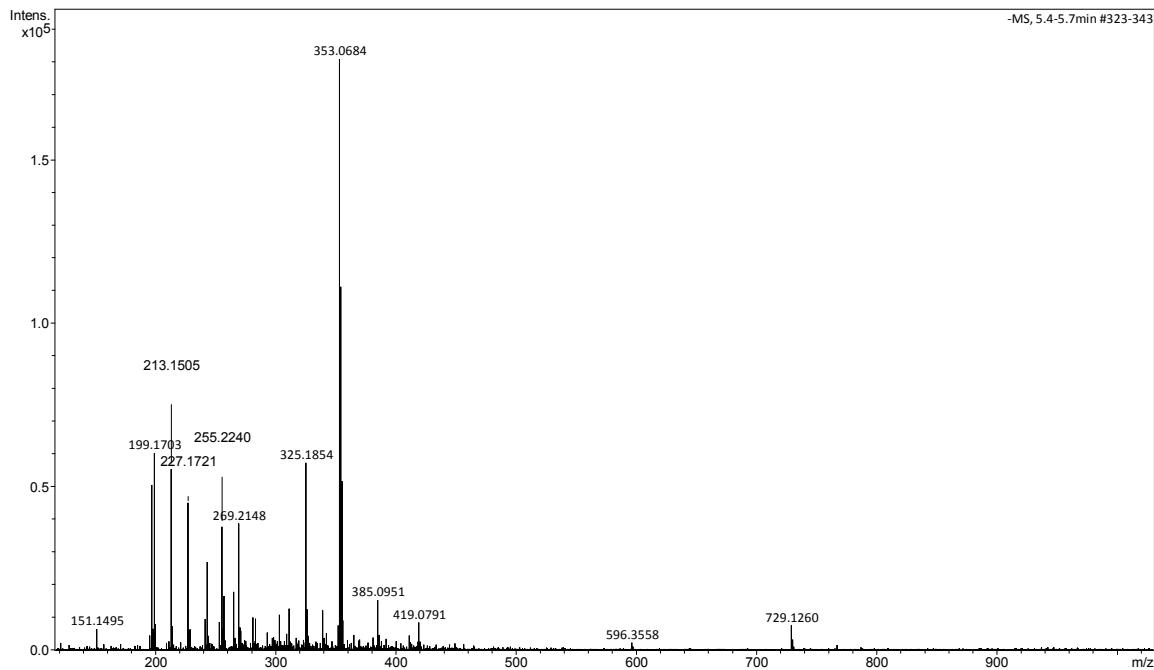


Fig. S9. The HRESIMS $[M-H]^-$ spectrum of **13**, m/z (experimental mass) = 353.0684 ($[M-H]^-$), m/z (calculated mass [calcd.]) = 353.0667 ($[M-H]^-$) consistent with the molecular formula $C_{19}H_{14}O_7$.

Notes and references

T. Kunnari, J. Kantola, K. Ylihonko, K. D. Klika, P. Mäntsälä, J. Hakala, *J. Chem. Soc. Perkin. Trans.*, 1999, 2, 1649-1652.