



Cytotoxic polyacetylenes related to petroformyne-1 from the marine sponge *Petrosia* sp.

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ABSTRACT

Four polyacetylenes related to petroformyne-1 were isolated from the marine sponge *Petrosia* sp. Their structures were determined on the basis of spectroscopic data and the modified Mosher analysis. They exhibit cytotoxic activity against P388 murine leukemia cells.

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1. Introduction

Long-chain polyacetylenes are characteristic metabolites in the marine sponge of the genus *Petrosia*.¹ They exhibit potent cytotoxic activity in general. Petroformynes, which represent this class of metabolites, were isolated from the Mediterranean *Petrosia fici-formis*.² Recently, closely related polyacetylenes with the same lengths but different arrangements of the same sets of functional groups were isolated from the Korean *Petrosia* sp.³

In the course of our search for cytotoxic constituents from marine invertebrates, we isolated four new polyacetylenes from the marine sponge *Petrosia* sp. collected at Kurose Hole, 30 km north of Hachijo Island. Their structures were elucidated on the basis of NMR and MS/MS data. This paper describes the isolation, structure elucidation, and biological activities of these compounds, and suggests the necessity to reexamine the structure of petroformyne-1.

2. Results and discussion

The organic layer of the extract of the sponge was dried and subjected to the modified Kupchan procedure⁴ to yield 60% MeOH, CHCl₃, and *n*-hexane layers. The CHCl₃ layer was separated by ODS flash chromatography, silica gel open column chromatography, and reversed-phase HPLC to give neopetroformynes A–D (**1**–**4**).

Neopetroformyne A (**1**) had a molecular formula of C₄₆H₆₈O₃, which was suggested by HRFABMS [*m/z* 691.5070, (M+Na)⁺, Δ +0.3 mmu]. Analysis of the ¹H NMR data in conjunction with the HSQC spectrum revealed the presence of two acetylenic protons, ten sp² methines, three oxygenated methines, and a number of methylenes. The ¹³C NMR spectrum further showed the presence of six acetylenic carbons without hydrogen. Partial structures **a–c** with unit **a** duplicating were deduced from the COSY data, and confirmed by the HMBC data (Fig. 1). The locations of units **b** and **c** in the alkyl chain were determined by an analysis of the FABMS/MS data for **1** (Fig. 2a). An intense product ion at *m/z* 306 was ascribed to the C-1 to C-19 fragment with a proton shift.⁵ The [M+Na]⁺ ion was selected as the precursor ion. Analysis of the product ions indicated that unit **a** and unit **b** were connected via one methylene, unit **b** and unit **c** were connected via one methylene, and unit **c** and unit **a** were connected via ten methylenes. Connection between unit **a** and unit **b** was supported by the TOCSY correlations between H-6 and H-10 and between H-6 and H-11.

The geometries of the Δ⁴-, Δ¹⁷- and Δ⁴²-olefines were determined as *E* on the basis of a coupling constant of 15.4 Hz, 15.7 Hz, and 15.4 Hz between the olefinic protons, respectively. The *Z*-geometries of the Δ²¹- and Δ²⁷-olefines were assigned on the basis of the chemical shifts of allylic carbons.⁶

The absolute stereochemistry of **1** was determined by the modified Mosher method⁷ applied to the three hydroxyl groups (Fig. 3a). Treatment of **1** with *R*-(–)- or *S*-(+)-MTPACI yielded (*S*)- or (*R*)-MTPA esters **5** and **6**, respectively. The Δδ values indicated the 3S,14S,44S configuration.

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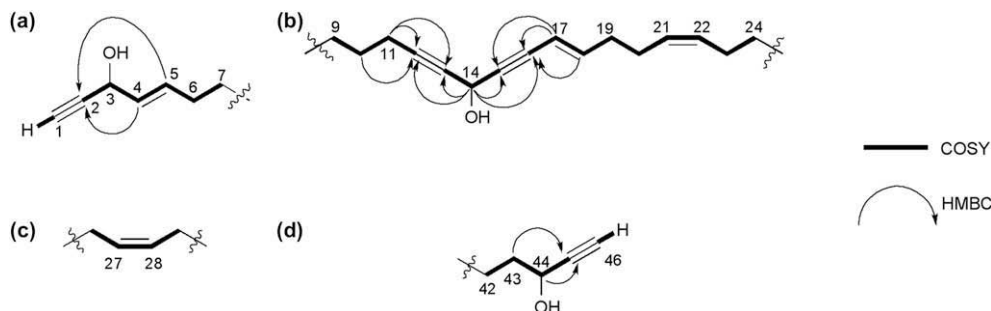


Figure 1. Partial structures **a–c** for **1** and partial structure **d** for **3**.

Neopetroformyne B (**2**) had a molecular formula of $C_{45}H_{66}O_3$, which was established by HRFABMS [m/z 677.4914, $(M+Na)^+$, $\Delta +0.4$ mmu]. The 1H and ^{13}C NMR spectra were almost identical with those of **1**, indicating that **2** is a lower homologue of **1**: the same partial structures (**a** $\times 2$, **b**, and **c**) were deduced from the 2D NMR data. The planar structure of **2** was determined by FABMS/MS analysis (Fig. 2b). The alkyl chain between unit **c** and unit **a** was shorter than that of **1** by one methylene unit. The geometries of olefins were determined by analysis of 1H – 1H coupling constants and ^{13}C chemical shifts of allylic carbons as described above. The absolute stereochemistry was assigned as 3*S*,14*S*,43*S*, because the 1H NMR spectrum of the (*S*)-MTPA ester **7** of **2** was almost superimposable on that of **5**.

Neopetroformyne C (**3**) had a molecular formula of $C_{46}H_{70}O_3$, which was established by HRFABMS [m/z 693.5228, $(M+Na)^+$, $\Delta +0.5$ mmu], suggesting one less unsaturation than **1**. The 1H NMR data suggested the replacement of one of units **a** in **1** by unit **d** (Fig. 1). The planar structure of **3** was assigned by the FABMS/MS data (Fig. 2c). The intense product ion at m/z 306 suggested that the C-1 to C-19 portion was conserved in **3**. The tandem MS data

demonstrated that the Δ^{42} -olefin in **1** was saturated in **3**. The geometries of the olefins were determined as described for **1**. The absolute stereochemistry was assigned as 3*S*,14*S*,44*S* by applying the modified Mosher method to the (*S*)- and (*R*)-MTPA esters (**8** and **9**, respectively) (Fig. 3b).

Neopetroformyne D (**4**) had a molecular formula of $C_{45}H_{66}O_4$, which was established by HRFABMS [m/z 693.4862, $(M+Na)^+$, $\Delta +0.3$ mmu]. The 1H NMR spectrum displayed one additional oxygenated methine signal instead of a propargylic methylene in **2**. Partial structures **a** ($\times 2$) and **c** were deduced from the 2D NMR data. C-11 in partial structure **b** was oxidized to a secondary alcohol. Tandem FABMS data afforded an intense ion at m/z 322, in agreement with oxidation at C-11 (Fig. 2d). All the other product ions were larger than the corresponding product ions of **2** by 16 u, indicating that **4** differed from **2** only in the oxidation state of C-11. The geometries of the olefins were determined as described for **1**. The absolute stereochemistry was assigned as 3*S*,43*S* by comparison of the 1H NMR spectrum of the (*S*)-MTPA ester **10** with that of **5**. However, the absolute stereochemistry of C-11 and C-14 was not assigned due to the paucity of the material.

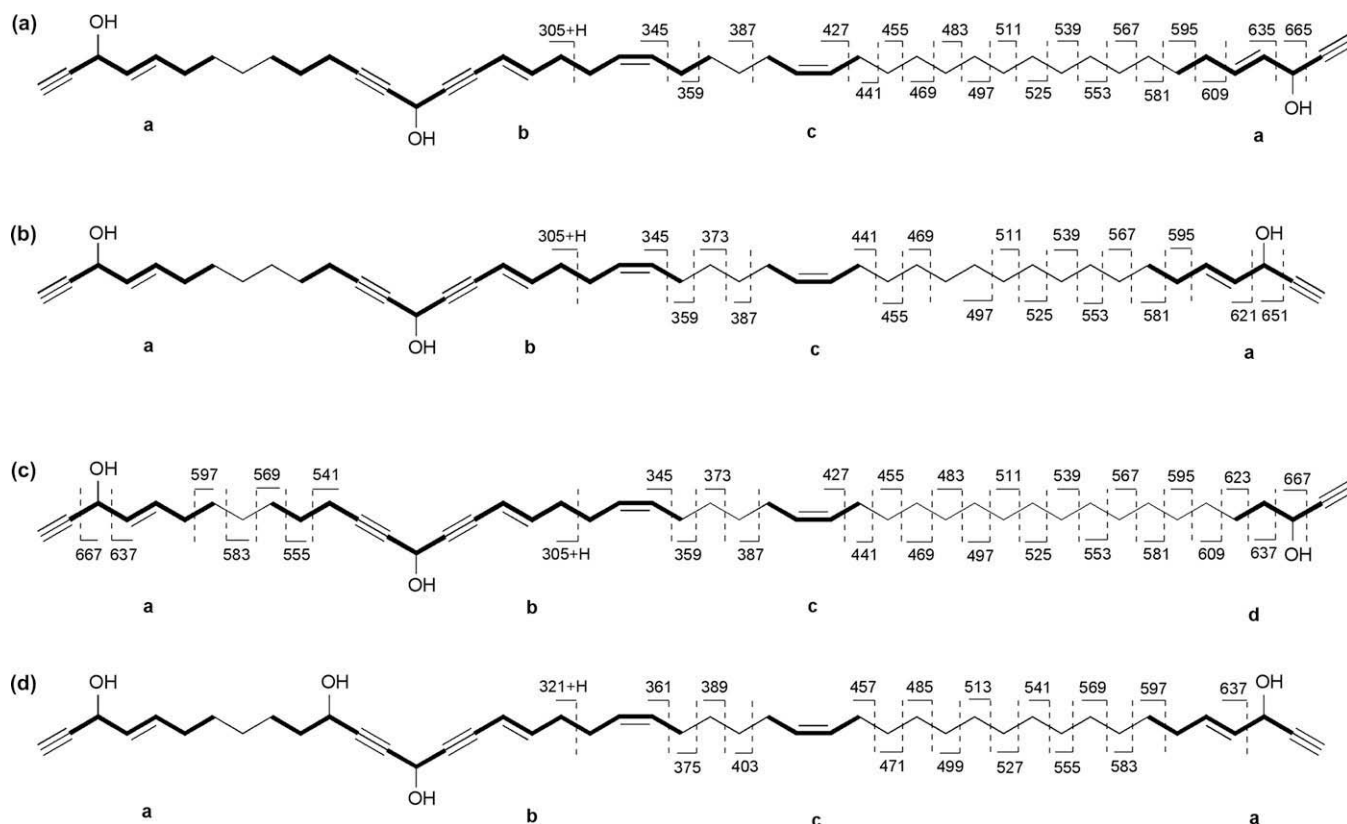


Figure 2. a)–d) The structures and FABMS/MS data of **1–4** from the $[M+Na]^+$ ion, respectively.

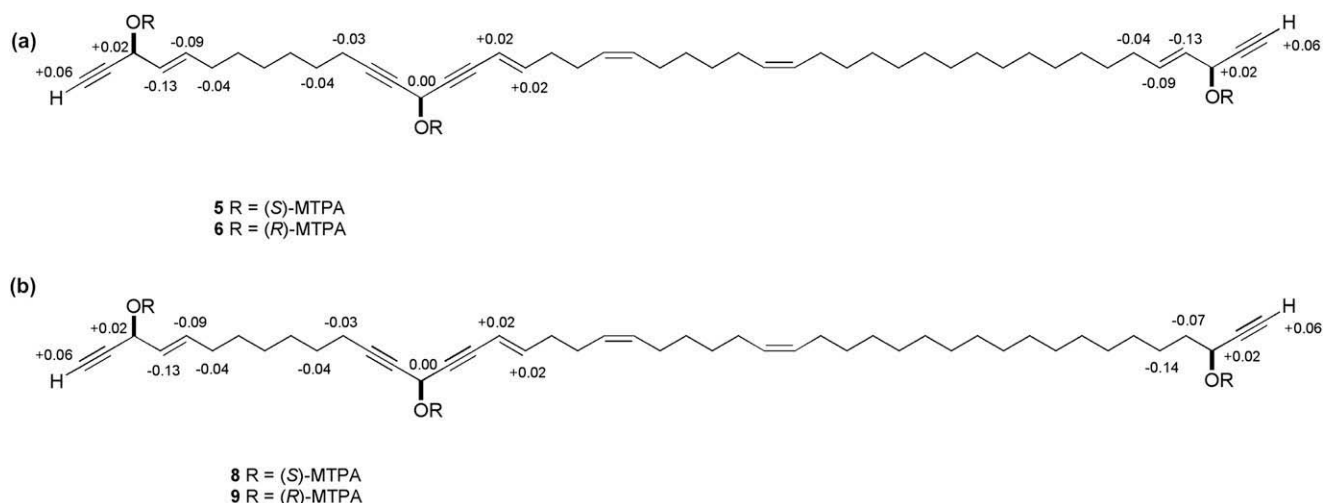


Figure 3. $\Delta\delta_{S-R}$ values for the MTPA esters of **1** and **3**.

Neopetroformyne A–D (**1**–**4**) exhibit cytotoxic activity against P388 murine leukemia cells with IC_{50} values of 0.089 $\mu\text{g/mL}$, 0.2 $\mu\text{g/mL}$, 0.45 $\mu\text{g/mL}$, and 0.45 $\mu\text{g/mL}$, respectively.

3. Conclusion

Neopetroformyne A has the same molecular formula as petroformyne-1 as well as the four sets of partial structures (Fig. 4). The structure of petroformyne-1 was assigned by analysis of ozonolysis products and EIMS data of the tri-TMS ether. The NMR data of neopetroformyne A were indistinguishable from those of petroformyne-1 and the same products are expected to be formed by ozonolysis of the two compounds. We consider it necessary to procure the FABMS/MS data of petroformyne-1 in order to confirm the proposed structure.⁸

4. Experimental section

4.1. General procedures

Optical rotations were measured on a JASCO DIP-1000 digital polarimeter in MeOH. NMR spectra were recorded on a JEOL delta 600 NMR spectrometer at 600 MHz for ^1H and 150 MHz for ^{13}C . ^1H and ^{13}C chemical shifts were referenced to the solvent peaks at δ_{H} 3.31 and δ_{C} 49.15 for CD_3OD , and at δ_{H} 7.27 and δ_{C} 77.23 for CDCl_3 . FAB mass spectra were measured on a JMS-700 T mass spectrometer.

4.2. Animal material

The sponge *Petrosia* sp. was collected by dredging at a depth of 150 m at Kurose Hole, 30 km north of Hachijo island (33°21'N;

139°40'E), in 2007, immediately frozen, and kept at -20°C until used. The voucher specimen was deposited at the Misaki Marine Biological Station, The University of Tokyo.

4.3. Extraction and isolation

The sample (600 g) was extracted with MeOH (2×3 L) and EtOH (1×3 L), and the extracts were combined and concentrated in vacuo. The residue was suspended in H_2O (500 mL) and extracted with CHCl_3 (3×500 mL) and *n*-BuOH (2×500 mL). The CHCl_3 extract was partitioned between 90% MeOH and *n*-hexane. The 90% MeOH layer was diluted with H_2O to yield a 60% MeOH solution and then extracted with CHCl_3 . The CHCl_3 layer was concentrated and separated by ODS flash chromatography to give six fractions (A–F). The fraction E (100% MeOH fraction) was separated by silica gel open column chromatography to give 14 fractions (A'–N'). The active fraction C' (*n*-hexane/EtOAc (7:3) fraction) was further separated by reversed-phase HPLC (COSMOSIL 5C₁₈-AR-II, 20×250 mm) with 60% 1-PrOH to give 20.1 mg of neopetroformyne A (**1**), 1.2 mg of neopetroformyne B (**2**), and the active fraction A''. The active fraction A'' was purified by reversed-phase HPLC (Phenomenex 5-Phenylhexyl, 10×250 mm) with 60% 1-PrOH to give 0.3 mg of neopetroformyne C (**3**). The active fraction E' (*n*-hexane/EtOAc (6:4) fraction) was separated by reversed-phase HPLC (COSMOSIL 5C₁₈-AR-II, 10×250 mm) with 60% 1-PrOH to give 0.2 mg of neopetroformyne D (**4**).

4.3.1. Neopetroformyne A (**1**)

Yellowish oil; $[\alpha]_{\text{D}}^{20.5} +19$ (c 0.45, MeOH); ^1H NMR (CD_3OD) and ^{13}C NMR (CD_3OD) data, see Table 1; HRFABMS m/z 691.5070 (calcd for $\text{C}_{46}\text{H}_{68}\text{O}_3\text{Na}$, 691.5067).

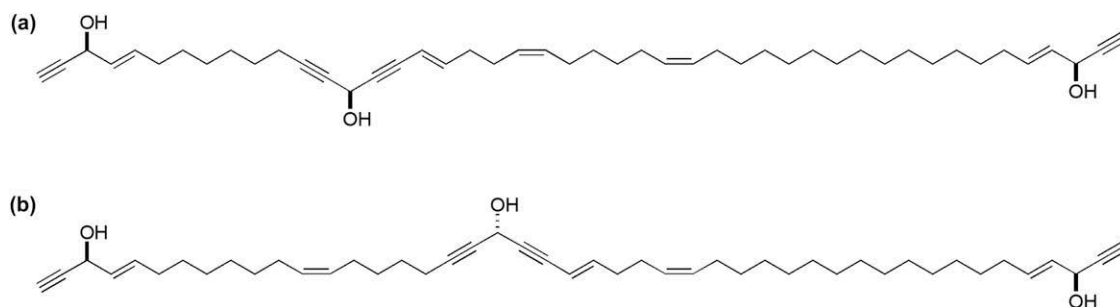


Figure 4. The structures of a) **1** and b) petroformyne-1.

Table 1¹H and ¹³C NMR data for Neopetroformyne A–D (**1–4**) in CD₃OD

No.	1		2		3		4	
	δ_{H} , mult.	δ_{C}	δ_{H} , mult.	δ_{C}	δ_{H} , mult.	δ_{C}	δ_{H} , mult.	δ_{C}
1	2.86 ^b s	74.7	2.87 br	74.7	2.87 s	74.7	2.86 br	74.5
2		84.9		84.9		84.8		84.5
3	4.75 (d, 6.1)	63.3	4.75 (d, 6.0)	63.3	4.75 (d, 5.9)	63.3	4.75 (d, 5.5)	63.0
4	5.55 m	130.7 ^c	5.55 m	130.8 ^b	5.56 m	130.9 ^b	5.56 m	130.5
5	5.85 (dt, 7.2, 15.4)	134.2 ^d	5.85 (dt, 7.2, 15.6)	134.2 ^c	5.85 (dt, 7.2, 15.6)	134.2	5.85 (dt, 7.2, 15.6)	134.2
6	2.07 m	33.1 ^e	2.07 m	33.1 ^d	2.08 m	33.1	2.08 m	32.7
7	1.43 m	29.8	1.43 m	29.8	1.43 m	29.8	1.43 m	29.8
8	1.30–1.39 ^a m	29.0–32.0 ^a	1.30–1.39 ^a m	29.0–32.0 ^a	1.30–1.39 ^a m	29.0–32.0 ^a	1.30–1.39 ^a m	29.0–32.0 ^a
9	1.44 m	29.0–32.0 ^a	1.43 m	29.0–32.0 ^a	1.44 m	29.0–32.0 ^a	1.48 m	26.2
10	1.52 (quint, 7.2)	29.7	1.52 (quint, 7.3)	29.7	1.52 (quint, 7.1)	29.7	1.67 m	38.7
11	2.23 br t	19.5	2.23 br t	19.4	2.23 br t	19.4	4.34 br t	62.7
12		85.1		85.1		85.1		85.7
13		79.5		79.5		79.5		82.6
14	5.14 br s	53.0	5.14 br s	53.0	5.14 br s	53.0	5.21 br s	52.7
15		87.2		87.2		87.2		86.7
16		82.7		82.7		82.7		82.8
17	5.53 m	110.6	5.53 m	110.6	5.54 m	110.6	5.54 m	110.2
18	6.14 (dt, 6.7, 15.7)	146.1	6.14 (dt, 6.7, 16.0)	146.1	6.14 (dt, 6.4, 15.5)	146.1	6.15 (dt, 6.4, 16.0)	146.1
19	2.17 m	34.3	2.17 m	34.3	2.17 m	34.3	2.17 m	34.2
20	2.16 m	27.7	2.16 m	27.7	2.16 m	27.7	2.16 m	27.5
21	5.35 m	129.6	5.35 m	129.6	5.34 m	129.6	5.34 m	129.2
22	5.40 m	131.9	5.40 m	131.9	5.40 m	131.9	5.40 m	131.7
23	2.05 m	28.2 ^f	2.04 m	28.1 ^e	2.05 m	28.2 ^c	2.04 m	28.2
24–25	1.30–1.39 ^a m	29.0–32.0 ^a	1.30–1.39 ^a m	29.0–32.0 ^a	1.30–1.39 ^a m	29.0–32.0 ^a	1.30–1.39 ^a m	29.0–32.0 ^a
26	2.05 m	28.2 ^f	2.04 m	28.2 ^e	2.05 m	28.2 ^c	2.04 m	28.2
27	5.35 m	130.9 ^c	5.35 m	130.9 ^b	5.36 m	130.9 ^b	5.35 m	130.8
28	5.35 m	131.1 ^c	5.35 m	131.1 ^b	5.36 m	131.1 ^b	5.35 m	130.8
29	2.05 m	28.3 ^f	2.04 m	28.3 ^e	2.05 m	28.3 ^c	2.04 m	28.2
30–38	1.30–1.39 ^a m	29.0–32.0 ^a	1.30–1.39 ^a m	29.0–32.0 ^a	1.30–1.39 ^a m	29.0–32.0 ^a	1.30–1.39 ^a m	29.0–32.0 ^a
39	1.30–1.39 ^a m	29.0–32.0 ^a	1.43 m	29.8	1.30–1.39 ^a m	29.0–32.0 ^a	1.43 m	29.8
40	1.43 m	29.8	2.07 m	33.2 ^d	1.30–1.39 ^a m	29.0–32.0 ^a	2.08 m	32.7
41	2.07 m	33.2 ^e	5.85 (dt, 7.2, 15.6)	134.3 ^c	1.30–1.39 ^a m	29.0–32.0 ^a	5.85 (dt, 7.2, 15.6)	134.2
42	5.85 (dt, 7.2, 15.4)	134.3 ^d	5.55 m	130.9 ^b	1.46 m	26.5	5.56 m	130.5
43	5.55 m	130.8 ^c	4.75 (d, 6.0)	63.3	1.65 m	39.1	4.75 (d, 5.5)	63.0
44	4.75 (d, 6.1)	63.3		84.9	4.27 (t, 6.6)	62.8		84.5
45		84.9	2.87 br	74.7		86.3	2.86 br	74.5
46	2.87 ^b s	74.7			2.76 s	73.5		

^a ¹H and ¹³C chemical shifts were overlapped.^b Assignments may be interchanged.^c Assignments may be interchanged.^d Assignments may be interchanged.^e Assignments may be interchanged.^f Assignments may be interchanged.

4.3.2. Neopetroformyne B (**2**)

Yellowish oil; $[\alpha]_{\text{D}}^{20.2} +21$ (c 0.06, MeOH); ¹H NMR (CD₃OD) and ¹³C NMR (CD₃OD) data, see Table 1; HRFABMS *m/z* 677.4914 (calcd for C₄₅H₆₆O₃Na, 677.4910).

4.3.3. Neopetroformyne C (**3**)

Colorless oil; $[\alpha]_{\text{D}}^{22.0} -15$ (c 0.015, MeOH); ¹H NMR (CD₃OD) and ¹³C NMR (CD₃OD) data, see Table 1; HRFABMS *m/z* 693.5228 (calcd for C₄₆H₇₀O₃Na, 693.5223).

4.3.4. Neopetroformyne D (**4**)

Colorless oil; $[\alpha]_{\text{D}}^{21.4} +20$ (c 0.01, MeOH); ¹H NMR (CD₃OD) and ¹³C NMR (CD₃OD) data, see Table 1; HRFABMS *m/z* 693.4862 (calcd for C₄₅H₆₆O₄Na, 693.4859).

4.4. Assay for the cytotoxicity against P388 cells

Cytotoxicity was determined as described.⁹

4.5. Preparation of MTPA esters

To a solution of the compound (**1**: 0.5 mg, **2**: 0.2 mg, **3**: 0.1 mg, **4**: 70 μ g) in CH₂Cl₂ (100 μ L) was added (*R*)-MTPACI (5 μ L) and DMAP (1 mg) and the mixture was left at rt for 5 min. The mixture was

partitioned between 0.1 M NaHCO₃ and CHCl₃, and the CHCl₃ layer was successively washed with 0.1 N HCl and H₂O. The organic layer was concentrated and separated by preparative TLC to afford the (*S*)-MTPA ester. A (*R*)-MTPA ester was prepared in the same way.

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Supplementary data

NMR spectra for compounds **1–10** are available free of charge. Supplementary data associated with this article can be found in the online version, at doi:10.1016/j.tet.2009.04.091.

References and notes

- Minto, R. E.; Blacklock, B. J. *Prog. Lipid Res.* **2008**, *47*, 233–306.
- (a) Cimino, G.; De Giulio, A.; De Rosa, S.; Di Marzo, V. *Tetrahedron Lett.* **1989**, *30*, 3563–3566; (b) Guo, Y.; Gavagnin, M.; Trivellone, E.; Cimino, G. *Tetrahedron* **1994**, *50*, 13261–13268; (c) Guo, Y.; Gavagnin, M.; Trivellone, E.; Cimino, G. *J. Nat. Prod.* **1995**, *58*, 712–722; (d) Guo, Y.; Gavagnin, M.; Salierno, C.; Cimino, G. *J. Nat. Prod.* **1998**, *61*, 333–337.

3. (a) Seo, Y.; Cho, K. W.; Rho, J.-R.; Shin, J. *Tetrahedron* **1998**, *54*, 447–462; (b) Kim, J. S.; Im, K. S.; Jung, J. H.; Kim, Y.-L.; Kim, J.; Shim, C. J.; Lee, C.-O. *Tetrahedron* **1998**, *54*, 3151–3158; (c) Shin, J.; Seo, Y.; Cho, K. W. *J. Nat. Prod.* **1998**, *61*, 1268–1273; (d) Kim, J. S.; Lim, Y. J.; Im, K. S.; Jung, J. H.; Shim, C. J.; Lee, C. O.; Hong, J.; Lee, H. J. *J. Nat. Prod.* **1999**, *62*, 554–559; (e) Lim, Y. J.; Kim, J. S.; Im, K. S.; Jung, J. H.; Lee, C.-O.; Hong, J.; Kim, D.-K. *J. Nat. Prod.* **1999**, *62*, 1215–1217; (f) Lim, Y. J.; Park, H. S.; Im, K. S.; Lee, C.-O.; Hong, J.; Lee, M.-Y.; Kim, D.-K.; Jung, J. H. *J. Nat. Prod.* **2001**, *64*, 46–53; (g) Lim, Y. J.; Lee, C.-O.; Hong, J.; Kim, D.-K.; Im, K. S.; Jung, J. H. *J. Nat. Prod.* **2001**, *64*, 1565–1567.
4. Kupchan, S. M.; Britton, R. W.; Ziegler, M. F.; Sigel, C. W. *J. Org. Chem.* **1973**, *38*, 178–179.
5. Okamoto, C.; Nakao, Y.; Fujita, T.; Iwashita, T.; van Soest, R. W. M.; Fusetani, N.; Matsunaga, S. *J. Nat. Prod.* **2007**, *70*, 1816–1819.
6. Cimino, G.; De Giulio, A.; De Rosa, S.; Di Marzo, V. *J. Nat. Prod.* **1990**, *53*, 345–353.
7. Ohtani, I.; Kusumi, T.; Kashman, Y.; Kakisawa, H. *J. Am. Chem. Soc.* **1991**, *113*, 4092–4096.
8. We were not able to compare the tandem FAB/MS data, because petroformyne-1 was no more available at Istituto per la Chimica di Molecole d'Interesse Biologico del CNR.
9. Ueoka, R.; Nakao, Y.; Fujii, S.; van Soest, R. W. M.; Matsunaga, S. *J. Nat. Prod.* **2008**, *71*, 1089–1091.