DINOFLAGELLATE NEUROTOXINS RELATED TO SAXITOXIN: STRUCTURES OF TOXINS C3 AND C4, AND CONFIRMATION OF THE STRUCTURE OF NEOSAXITOXIN¹

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Abstract: By x-ray crystallography of the 11 β epimer, toxins C3 and C4 are shown to be 21-sulfo-N-1hydroxysaxitoxin-11 α - and 11 β -hydroxysulfate, confirming the position and identity of the 3 substituents which, with the parent compound, form the array of 12 saxitoxins found in <u>Protogonyaulax</u>.

In the analysis of toxins from cultured <u>Protogonyaulax</u> of the northeast Pacific (2) the expected compounds <u>1</u>, <u>3</u>, <u>5</u>, <u>7</u>, <u>9</u>, and <u>11</u> were generally found to accompany somewhat larger amounts of the 21-sulfo derivatives <u>2</u>, <u>4</u>, <u>6</u>, and <u>8</u>. Although composition varied greatly among isolates, the above substances were all eventually found in relatively high concentrations. Compounds <u>10</u> (C3) and <u>12</u> (C4), the corresponding derivatives of <u>9</u> (gonyautoxin 1, GTX 1) and <u>11</u> (gonyautoxin 4, GTX 4) were at first notable for their absence but were finally detected in the mother liquors from crystallizations of <u>4</u> and <u>6</u> (3), and as trace constituents in analyses of <u>Protogonyaulax</u> clone PI07 (4).

The behavior of <u>10</u> and <u>12</u> (5,7) is largely analogous to that of <u>4</u> and <u>6</u> (2,3). Compounds <u>10</u> and <u>12</u> epimerize (TLC, NMR), the conversion of <u>12</u> to <u>10</u> predominating. Hydrolysis (0.1M HCl, 100 °C, 5 minutes; 6) converts <u>10</u> to <u>9</u> and <u>12</u> to <u>11</u> (TLC). Preliminary assays indicate that the mouse intraperitoneal potencies of crude 10 and 12 increase by factors of 40x and 7x with these conversions.

1 2 3 4 5 6 7 8 9 10 11 12	к інннн н н н н н н н н н н н н н н н н н	R2 H H H OSO; OSO; H H H OSO; OSO;	R3 H H OSO ⁻ ₃ OSO ⁻ ₃ H H H OSO ⁻ ₃ OSO ⁻ ₃ H H	R4 H SO', H SO', H SO', H SO', H SO', SO',	STX B1 GTX 2 C1 GTX 3 C2 NEO B2 GTX 1 C3 GTX 4 C4	$\begin{array}{c c} & Figure 1. \\ C4 (12) \\ \hline \\ R4 & 2 \\ 200 \\ R1 \\ H_2 N + \\ H_2 N + \\ R2 \\ R3 \\ \hline \\ R2 \\ R3 \\ \hline \\ \\ R3 \\ \hline \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ $	Table 2. 13C-NMR data for 10 carbon shift, ppm 2b 159.4 4 81.8 5 57.7 6 62.1 8b 158.8 10 52.0 11 78.3 12 98.0 13 64.6 19 154.2			
Table 1. ¹ H-NMR data ^a .										
H-5 <u>10</u> 4.50,s <u>12</u> 4.53,s a) Chemical parenthe b) Assignme		H-6 3.76 (6,6) 3.79 (7,7) shift ses ar	b ,dd ,dd s in e co opro	H-10 ^b 3.77,d (12) 3.78,dd (7,10) ppm from upling co ximate d	H-10 3.63,dd (4.6,11.9) 3.25,dd (7.0,10.7) n internal chl onstants in Hz ue to overlap	H-11 4.46,d (4.5) 4.58,dd (7,8) oroform =	H-13 4.12,dd (6.7,11.6) 4.14,dd (7.0,11.3) 7.27. Data	H-13 3.87,dd (6.4,11.3) 3.92,dd (6.7,11.6) in	 a) Chemical shifts in ppm from internal dioxane = 67.6. b) Assignments may be reversed. 	

Compound <u>12</u> crystallizes (8) in space group $P2_12_12_1$ with cell dimensions a = 12.037(8) Å, b = 16.255(10) Å, c = 11.652(9) Å, and Z = 4. D_c for $C_{10}H_{17}O_{12}N_7S_2\cdot 4H_2O$, mw 563.40, is 1.642 g·cm⁻³. Final refinements of the diffraction data attained an R factor of 0.067 for the 30 data set. A plot of the structure is shown in Figure 1 with the calculated positions of hydrogens on carbons 5, 6, and 11 shown for clarity. Note the oxygen bonded to N-1.

¹H-NMR data (270 MHz, D_2O) for <u>10</u> and <u>12</u> (Table 1) are consistent with those for the other saxitoxins (2, 3, 9–11). It is noteworthy that the chemical shifts observed for H-5, H-6, and both H-13 in <u>10</u> and <u>12</u> are significantly downfield from the corresponding shifts for <u>4</u> and <u>6</u> (3), consistent with the presence of N-1-OH in <u>10</u> and <u>12</u>. ¹³C-NMR data (67.9 MHz, H₂O/D₂O, broad-band decoupled) for <u>10</u> (Table 2) correlate well with those for <u>1-8</u> (2,9). Of particular interest are the resonances for C-6 (cf. <u>7</u>, 61.6; <u>8</u>, 62.0), C-10 (cf. <u>3</u>, 51.1; <u>4</u>, 51.5), C-11 (cf. <u>3</u>, 77.7; <u>4</u>, 78.1), and C-19 (cf. <u>2</u>, 154.8; <u>4</u>, 154.6; <u>6</u>, 154.7; <u>8</u>, 154.3). The resonance at 154-155 ppm appears diagnostic for the sulfamate carbonyl carbon in the saxitoxins (2).

The structural relationships among compounds 1-8, 9, and 11 have been established (2, 3, 6, 10-12). The data presented here linking 10 and 12 with 9 and 11, coupled with the x-ray structure for 12, secure the structures for the entire array and in particular serve to confirm the position and identity of the N-1-hydroxyl substituent of neosaxitoxin, 7 (9,13).

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- 5. Under the previously described analytical conditions (6), toxins 10 and 12 elute from BioGel P2 overlapping but slightly later than 4 and 6 respectively. Each migrates to slightly higher Rf on TLC than 4 and 6, respectively. When sprayed with 1% hydrogen peroxide and heated 15 minutes at 120°C (L. J. Buckley, M. Ikawa, and J. J. Sasner, Jr. J. Agric. Food Chem. 24: 107 (1976)) compounds 10 and 12 tend, like 7, 8, 9, and 11, to form spots with yellowish fluorescence rather than the blue fluorescence obtained with compounds 1-6.
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- 7. Preparative chromatography of the mixed group C toxins $(\frac{4}{2}, \frac{6}{2}, \frac{10}{2}, \frac{12}{2})$ under neutral or weakly acidic conditions on a number of media failed to give useful separation. Facile resolution of $\underline{10}$ and $\underline{12}$ from $\underline{4}$ and $\underline{6}$ was finally achieved (2) by chromatography on BioGel P-2 in 0.1M pyridine, under which conditions $\underline{10}$ and $\underline{12}$ elute before and well separated from $\underline{4}$ and $\underline{6}$. Compounds $\underline{10}$ and $\underline{12}$ can be resolved from each other by chromatography on the same gel with either water or 0.1M acetic acid and then crystallized by methods similar to those used for $\underline{4}$ and $\underline{6}$ (2,3).
- 8. A crystal of <u>12</u> was mounted in a capillary with a trace of mother liquor and data collected at room temperature using graphite monochromated MoK_{α} radiation. Of the 1733 reflections collected, 1535 were classified as observed. The structure was solved by direct methods using the program MULTAN 80 and atoms verified in difference Fourier maps using the program set CRYM.
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