Further Studies on the Chemistry of the Flustra Alkaloids from the Bryozoan Flustra foliacea

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Since 1980, over a dozen novel brominated alkaloids, named flustramines, have been isolated from Scandinavian and Canadian collections of the marine bryozoan *Flustra foliacea*. This paper describes the reisolation of the known compound dihydroflustramine C (1) and the isolation of 11 new flustramines (2-4, 6-13), including two dimers (12, 13) that may be isolation artifacts. Together these compounds, some with an unexpected aryl substitution pattern, reveal an intricate network of metabolites present in the extracts of the bryozoan. The structures of these metabolites were solved using a variety of spectroscopic techniques and chemical derivatization and modification. This work also led to the recognition of an unusual rearrangement reaction that occurred slowly over a number of years.

Early chemical studies of the marine bryozoan F. foliacea were notable in several aspects. First, these studies were among the first carried out on a phylum of animals that have since yielded structurally and pharmacologically interesting compounds such as the bryostatins.¹ Second, the family of compounds identified by these studies, the flustramines, were early examples of alkaloids identified from marine sources.²⁻⁴ Moreover, the flustramines were among the first marine natural products found to have a strong structural affinity with terrestrial natural products, in this case the physostigmine alkaloids from legumes and other sources. The marine structures still have distinctive features however, such as bromination of the indole ring, prenylation of the indole ring nitrogen of flustramines A and B, and the presence of a "reverse" prenyl substitution in flustramines A and C, the latter of which lacks prenylation of the indole nitrogen. Subsequent studies led to the identification of dihydroflustramine C⁵ and also flustramine D and isoflustramine D,6 the latter two containing prenylated aromatic rings. Flustramine E,7 like flustramine B, exhibits "normal" prenylation, but like dihydroflustramine C, lacks prenylation of the indole nitrogen. Variations seen in other flustramine congeners include incorporation of oxygen atoms into the terminal pyrrolidine moiety via ring expansion as in flustrarine B,8 as an exocyclic *N*-oxide,^{6,9} or via lactamization.^{8,10}

The flustramines have attracted attention for their biological activities and as synthetic targets. These alkaloids and related compounds have been reported to have antibacterial,^{5–7} muscle relaxant,¹¹ nonspecific voltage-sensitive potassium channel blocking,¹² and subtype-specific nicotinic acetylcholine receptor blocking activities.¹³ The first synthetic studies on the flustramines led to the production of racemic flustramine B¹⁴ and debromoflustramine B.¹⁵ The later synthesis of *ent*-debromoflustramine B established the configuration of natural flustramine B as being 3a*S*, 8a*R*.¹⁶ More recently, Austin et al. reported the synthesis of chiral flustramine B identical to the natural product using a cascade addition–cyclization approach.¹⁷ Other studies have targeted racemic debromodihydroflustramine C,^{18,19} racemic flustramine

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C,²⁰ various racemic flustramines,²¹ racemic dihydroflustramine C and flustramine E,²² racemic flustramines A and C,²³ and scalemic flustramines A and B.²⁴

We report here studies building on our previous work with F. *foliacea* collected from Canadian waters. In this latest report, we have been able to identify 11 new flustramines, which differ from those previously reported in bromination patterns and by a new hydroxylation pattern on the aromatic ring. We also observed flustramines that appear to be derived by further cyclization, by isomerization to an alternative ring system, or by dimerization mediated by a single carbon block.

Results and Discussion

Following a large collection of *F. foliacea*, the cells were freshly extracted and the CH₂Cl₂-soluble fraction of the aqueous MeOH extract showed significant antibacterial and antifungal activity. Large-scale, medium-pressure reversed-phase chromatography of the extract produced several fractions containing UV-active compounds. Compounds containing the flustramine skeleton were identified on the basis of their IR and UV spectra and by characteristic fragmentation pathways by EIMS. The most useful EIMS fragmentations were the loss of one or two isoprene subunits followed by the loss of 43 mass units, corresponding to ejection of a CH₃–N=CH₂ fragment from the pyrrole moiety.^{2–7} Further purification of selected fractions by C18 reversed-phase HPLC yielded the known dihydroflustramine C (1) and 11 new flustramine derivatives (2–4, 6–13).

The major metabolite 1, first reported in this laboratory,⁵ was identified by comparison of its spectroscopic data with that originally reported. A new product, designated flustramine F (2), possessed a molecular formula of C₁₈H₂₃BrN₂O ([M]⁺ 362.0987) requiring eight degrees of unsaturation. The spectroscopic data for 2 were very similar to those of 1, and the difference of 42 mass units between 1 and 2 corresponds to the addition of C₂H₂O. In CDCl₃ the ¹H NMR resonances were broad, while in DMSO- d_6 at elevated temperature (367 K) the resonances became sharp. Despite this effect, the NMR data obtained for 2 in CDCl₃ (Tables 1 and 2) were very similar to those for 1, the only significant differences being the appearance of acetate resonances in 2 ($\delta_{\rm H}$ 2.31 ppm, br s; δ_C 23.8 ppm, CH₃; 169.0 ppm, qC) and the absence of the resonance assigned to the benzylic amine proton. Collectively, these data suggested that 2 was the N-acetate of 1. The broad signals observed in the ¹H NMR spectra of 2 at standard temperature were ascribed to the presence of rotational isomers of the N-acetate moiety. This was confirmed by acetylation of 1 to yield a product that was spectroscopically identical to 2.

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The isotope pattern and accurate mass for the parent ion observed in the EIMS of flustramine G (3) indicated a molecular formula ($C_{16}H_{20}Br_2N_2$; [M]⁺ 397.9990) related to that of **1** by addition of a bromine atom. The most important differences between the ¹H NMR spectroscopic data for **3** and **2** were that the aromatic region of the spectrum of **3** contained only two aromatic singlets (δ_H 7.30 and 6.82 ppm), suggesting a *para* relationship between the protons. Complete assignment of the proton and carbon resonances was possible by detailed 2D NMR studies (Tables 1 and 2) and led to structure **3** for flustramine G.

The molecular formula $C_{21}H_{29}BrN_2O$ ([M]⁺ 404.1460) obtained for flustramine H (**4**) suggested that **4** contained additional C_5H_8 (isoprene) and OH moieties compared to **1**. Indeed, the EIMS fragmentation pattern confirmed the presence of a geranyl moiety, from which two isoprene subunits are lost sequentially from the molecular ion to give fragment ions at m/z 335/337 and 267/269. The presence of a geranyl moiety was supported by analysis of the NMR data (Tables 1 and 2), which showed three olefinic methyls (δ_H 1.59, 1.57, and 1.66 ppm) and two olefinic methines (δ_H 5.12 and 5.03 ppm). In addition, 2D COSY and HMBC analysis enabled complete assignment of the geranyl moiety. HMBC correlations were observed from the two olefinic methyls, H₃-16 and H₃-17, to C-14, which in turn was connected by successive COSY correlations to C-13, then C-12. The HMBC correlation between the olefinic methyl resonance H_3 -18 and C-12 confirmed the link between the two C_5 isoprene units. The point of attachment of the C_{10} side chain to the aromatic nucleus was established by the HMBC correlation between the H_2 -9 protons and C-3b.

The remaining differences between 4 and 1 centered on the nature and substitution pattern of the aromatic nucleus. The ¹H NMR spectrum of 4 revealed the presence of two rather than three aromatic protons, suggesting the aromatic ring to be the site of hydroxylation. This was supported by ¹³C NMR resonances at $\delta_{\rm C}$ 152.2 and 93.4 ppm, consistent with hydroxylated and brominated aromatic carbons, respectively. However, the location of these groups on the aromatic ring was ambiguous. In the ¹H NMR spectra, the two aromatic resonances showed a large *ortho* coupling (J =8.5 Hz), but one proton was significantly more shielded than the other ($\delta_{\rm H}$ 6.02 cf. 6.99 ppm), suggesting that it is *ortho* to either the phenolic moiety or the nitrogen of the indole system. Regardless of chemical shift correlations, six possible substitution patterns can accommodate *ortho* coupling of the aromatic protons, as illustrated in Figure 1 (substructures A–F).

To determine the correct substitution pattern, two derivatives were made and characterized. The first derivative, debromoflustramine H (**5**), was formed by reaction of flustramine H with LiAlH₄. EIMS analysis of the product showed a single molecular ion at m/z 326 with no isotope peaks, confirming the replacement of bromine by hydrogen. Inspection of the ¹H NMR spectrum confirmed the presence of three aromatic protons ($\delta_{\rm H}$ 6.14 ppm, d (J = 7.9 Hz); 6.19 ppm, d (J = 7.9 Hz); 6.90 ppm, dd (J = 7.9, 7.9 Hz)). Each aromatic proton showed a large *ortho* coupling, eliminating substructures E and F, as these would have resulted in one *meta*-coupled proton.

It was considered that the methoxy derivative would be useful in identifying which of the four remaining substructures were correct. Treatment of 4 with K₂CO₃ and MeI yielded several quite polar compounds. Surprisingly, the major product was shown to be the rearranged permethylated ammonium cation 19 (Scheme 1). HRFABMS revealed the molecular formula of 19 as C25H38BrN2O $([M]^+ 461/463)$, suggesting the addition of four methyl groups rather than the expected three. The presence of four additional methyl groups in 19 was supported by the ¹H NMR spectra, which showed the presence of three downfield singlet methyl resonances, one of which integrated for nine protons and was more deshielded than the others. The ^{13}C and ^{1}H NMR data (δ_{H} 3.45 ppm, 9H; δ_{C} 53.5 ppm) were consistent with the three degenerate methyl groups being attached to a positively charged nitrogen atom. The ¹H chemical shift of the remaining N-methyl resonance was somewhat deshielded ($\delta_{\rm H}$ 3.99 ppm), whereas the associated carbon resonance was relatively shielded ($\delta_{\rm C}$ 32.7 ppm), values typical of an *N*-methyl indole group ($\delta_{\rm H}$ 3.40–4.20 ppm; $\delta_{\rm C}$ 28.0–34.0 ppm).^{25–27} The remaining 3H resonance was assigned as a methoxy signal on the basis of the chemical shift data ($\delta_{\rm H}$ 3.89 ppm, 3H; $\delta_{\rm C}$ 55.6 ppm).

Better resolution of the methyl resonances was achieved by collecting ¹H NMR spectroscopic data of the product in a 60% benzene- d_6 /CDCl₃ solution. Signals for the methyl groups were now observed at $\delta_{\rm H}$ 3.38 ppm (3H), $\delta_{\rm C}$ 55.5 ppm (OMe); $\delta_{\rm H}$ 2.60 ppm (9H), $\delta_{\rm C}$ 52.5 ppm (NMe₃); and $\delta_{\rm H}$ 3.68 ppm (3H), $\delta_{\rm C}$ 33.9 ppm (indole N-Me). HMBC correlations helped to derive the new indole substructure and the attachment points of the side chains. The H₂-8 protons showed a long-range correlation to a carbon at 128.5 ppm (C-3). The H_2 -10 protons showed a long-range correlation to a carbon at 139.0 ppm (C-2) as well as correlations into the geranyl side chain. This established that the two side chains are attached to the indole at different positions. The indole N-Me resonance at $\delta_{\rm H}$ 3.68 ppm showed HMBC correlations to two deshielded quaternary carbon resonances at $\delta_{\rm C}$ 135.0 ppm (C-7a) and $\delta_{\rm C}$ 139.0 ppm (C-2), confirming the point of attachment of the geranyl side chain on C-2. This was supported by correlations in the NOESY spectrum from the indole N-Me resonance to the geranyl side-chain

Table 1. ¹³C NMR Assignments for Flustramines 2–4 and 6–12 in CDCl₃ ($\delta_{\rm C}$, mult)^{*a*}

pos.	2	3	4	6	7	8	9	10	11	12
2	54.0, CH ₂	52.9, CH ₂	52.6, CH ₂	52.6, CH ₂	52.5, CH ₂	52.4, CH ₂	52.6, CH ₂	52.5, CH ₂	51.9, CH ₂	54.0, CH ₂
3	32.6, CH ₂	34.5, CH ₂	36.5, CH ₂	36.2, CH ₂	36.7, CH ₂	36.8, CH ₂	36.2, CH ₂	37.2, CH ₂	35.3, CH ₂	33.9, CH ₂
3a	64.0, qC	64.3, qC	59.3, qC	59.4, qC	59.7, qC	59.6, qC	59.2, qC	59.1, qC	57.7, qC	64.9, qC
3b	127.6, qC	134.9, qC	119.9, qC	119.6, qC	120.2, qC	120.1, qC	119.3, qC	122.8, qC	115.8, qC	132.1, qC
4	126.0, CH	129.5, CH	152.2, qC	152.0, qC	147.6, qC	147.5, qC	150.5, qC	150.3, qC	137.8, qC	126.2, CH
5	126.7, CH	123.3, qC	108.3, CH	108.8, CH	99.1, qC	99.1, qC	120.8, qC	117.0, qC	100.6, qC	119.8, CH
6	119.2, qC	111.6, qC	130.9, CH	131.2, CH	132.0, CH	132.0, CH	130.9, CH	121.9, CH	131.3, CH	121.8, qC
7	121.2, CH	113.0, CH	93.4, qC	93.7, qC	93.7, qC	93.7, qC	93.5, qC	99.0, qC	114.9, qC	109.4, CH
7a	144.4, qC	150.9, qC	149.5, qC	149.3, qC	149.6, qC	149.6, qC	148.0, qC	146.0, qC	158.3, qC	152.1, qC
8a	87.3, CH	84.8, CH	86.1, CH	86.3, CH	86.3, CH	86.1, CH	86.8, CH	86.3, CH	106.2, CH	90.5, CH
9	40.8, qC	41.2, qC	35.7, CH ₂	35.9, CH ₂	35.2, CH ₂	35.3, CH ₂	35.9, CH ₂	36.5, CH ₂	34.8, CH ₂	41.8, qC
10	22.4, CH ₃	22.4, CH ₃	120.6, CH	120.3, CH	120.4, CH	120.3, CH	120.5, CH	120.2, CH	119.7, CH	23.8, CH ₃
11	23.3, CH ₃	23.1, CH ₃	138.3, qC	135.5, qC	137.9, qC	134.3, qC	135.1, qC	134.1, qC	134.7, qC	23.1, CH ₃
12	144.2, CH	143.9, CH	39.9, CH ₂	18.2, CH ₃	39.9, CH ₂	18.1, CH ₃	18.2, CH ₃	18.0, CH ₃	18.2, CH ₃	144.9, CH
13	112.7, CH ₂	113.9, CH ₂	26.7, CH ₂	26.0, CH ₃	26.7, CH ₂	26.0, CH ₃	26.0, CH ₃	25.6, CH ₃	25.9, CH ₃	113.5, CH ₂
14	38.9, CH ₃	36.7, CH ₃	124.2, CH	36.9, CH ₃	124.2, CH	37.0, CH ₃	29.4, CH ₂	100.0, CH	27.4, CH ₂	41.8, CH ₃
15			131.4, qC		131.3, qC		122.2, CH	161.6, qC	122.2, CH	58.2, CH ₂
16			17.7, CH ₃		17.7, CH ₃		134.8, qC	69.1, qC	132.4, qC	
17			25.7, CH ₃		25.7, CH ₃		17.9, CH ₃	28.7, CH ₃	25.8, CH ₃	
18			16.4, CH ₃		16.3, CH ₃		25.8, CH ₃	28.7, CH ₃	17.7, CH ₃	
19			37.0, CH ₃		37.1, CH ₃		37.0, CH ₃	36.9, CH ₃	35.1, CH ₃	
\underline{COCH}_3	169.0, qC									
COCH ₃	23.8. ĈH ₃									

^{*a*} Recorded at 125 MHz; referenced to solvent at δ 77.0 ppm.

Table 2. ¹H NMR Assignments for Flustramines 2–4 and 6–7 in CDCl₃ ($\delta_{\rm H}$, mult, J in Hz)^a

pos.	2	3	4	6	7
2	2.61, m	2.67, m	2.66, m	2.60, m	2.63, m
		2.55, td (8.0, 6.0)		2.80, m	
3	1.89, m	1.82, dt (12.0, 6.0)	2.05, dt (12.4, 7.3)	2.13, m	2.05, dt (12.5, 7.2)
	2.35, m	2.32, dt (12.0, 8.0)	2.20, dt (12.4, 5.4)	2.23, m	2.20, dt (12.5, 5.7)
4	7.08, br d (8)	7.30, s			
5	7.19, dd (8.1, 1.4)		6.02, d (8.5)	6.09, d (8.6)	
6			6.99, d (8.5)	7.00, d (8.6)	7.38, s
7	8.23, br s	6.82, s			
8a	4.77, br s	4.51, br s	4.33, s	4.44, s	4.35, s
9			2.50, dd (14.5, 8.4)	2.60, m	2.44, m
			2.66, m		2.71, dd (14.7, 6.6)
10	1.08, s	1.04, s	5.12, t (7.3)	5.05, br t (7)	4.97, br t (7)
11	0.99, s	1.00, s			
12	5.31, dd (17.4, 10.8)	5.93, dd (17.4, 10.9)	2.00, m	1.61, s	1.96, m
13	5.06, br m	<i>trans</i> 5.03, d (17.6) <i>cis</i> 5.12, d (10.9)	2.00, m	1.67, s	1.96, m
14	2.49, s	2.41, s	5.03, br t (7.3)	2.50, s	5.01, m
16			1.66, s		1.67, s
17			1.57, s		1.58, s
18			1.59, s		1.59, s
19			2.45, s		2.43, s
NH		4.51, br s	4.39, br s	4.49, br s	4.37, br s
OH					5.30, br s
\overline{OOCH}_3	2.31, br s				

^{*a*} Recorded at 500 MHz; referenced to residual solvent at δ 7.24 ppm.



Figure 1. Possible substitution patterns for flustramine H (4).

protons H_2 -10 and H-11 (weak). NOESY correlations observed between protons in the two side chains (H_2 -8 and H_2 -10) indicated that the two chains were adjacent on the indole. Structure **19** was fully consistent with the FABMS/MS data, which showed a loss of the N-Me₃ group, followed by sequential losses of side-chain fragments and the bromine atom.

A plausible mechanism by which **19** may be formed through ring-opening and Wagner–Meerwein rearrangement is depicted in Scheme 1. Expected methylation of the phenol group and indole nitrogen, yielding **14**, is followed by further methylation to yield the quaternary ammonium compound **15**. Ring-opening to produce **16** followed by a 1,2-alkyl shift of the geranyl chain and subsequent loss of a proton quenches the positive charge and forms the indole nucleus. Further methylation yields the charged ammonium salt **19**. This variant of the Wagner–Meerwein rearrangement is well known, and it has been demonstrated that the more electron rich R group will generally migrate preferentially,²⁸ though stability of the product and bond alignment can also affect which R group migrates.²⁹ Molecular modeling showed that in this case the two possible products had very similar energy minima (approximately

Scheme 1. Methylation and Rearrangement of Flustramine H



50 kJ/mol; Sculpt modeling program).³⁰ The two R groups were also similarly disposed toward the vacant p-orbital of the carbocation center, implying that the electron-donating ability of the two R groups is the dominant factor in deciding the product of Wagner–Meerwein rearrangement.

Fortunately the unexpected methylation product **19** was able to supply an important piece of information in determining the structure of flustramine H. A strong NOESY correlation from the OMe resonance ($\delta_{\rm H}$ 3.38 ppm) in **19** to the shielded aromatic methine at $\delta_{\rm H}$ 6.05 ppm was observed. This indicated that flustramine H must contain an aromatic methine adjacent to the hydroxyl group, thus eliminating substructures B and C. Of the two remaining substructures, A and D, the latter was disfavored by the absence of a NOESY correlation between the methoxy protons and the indole methyl. Thus, our NMR coupling data and chemical derivatization results were most consistent with substructure A as incorporated in **4**.

To further distinguish between substructures A and D in Figure 1, a series of deuterium exchange experiments were carried out with 4 and four model phenolic compounds, 20–23. In every case, exchange of the phenolic proton with deuterium in the model compounds resulted in an upfield shift of the resonance frequencies of the phenolic carbon and the two *ortho* carbon atoms. The isotope shifts for the four model compounds are summarized in Figure 2. The largest chemical shift changes for all the model compounds occur at the phenolic carbon and its *ortho* carbons, with smaller shifts occurring at the more distant carbons. In 4, the largest shift occurred at the phenolic carbon C-4 ($\Delta \delta$ –0.19 ppm) and the adjacent *ortho* carbons C-5 ($\Delta \delta$ –0.17 ppm) and C-3b ($\Delta \delta$ + 0.11 ppm). The chemical shift changes for the remaining indole carbon resonances were all less than 0.06 ppm. In the alternative substructure D, significant changes in the chemical shifts of the



Figure 2. Model compounds for deuterium shift experiments with observed isotope shift ($\Delta\delta$) shown for each carbon.

two most deshielded quaternary carbons (C-7 and C-7a), as well as the shielded aromatic carbon (C-6), would be predicted, but these were not observed. Thus the results of the isotope-shift experiments were consistent with substructure A, and this result in combination with the other spectroscopic and chemical data described above led to the structural assignment of flustramine H as **4**.

Substructure A, however, required bromination at a site different from that in previously isolated flustramines, which would be more consistent with substructure C. Chemical shift predictions were calculated using ACD/CNMR Predictor and ACD/HNMR Predictor (both from Advanced Chemistry Development, Inc.) for model compounds (4, 24-26) based on substructures A and C to further test our assigned structure (Figure 3). The results of the calculations further confirmed substructure A over substructure C. Specifically, the chemical shift differences for the ¹H and ¹³C resonances predicted for positions 4 and 5 clearly favored a para relationship between the Br and OH groups as in substructure A. This was further supported by ab initio chemical shift calculations using two model compounds and comparing the shift data as illustrated in Figure 3. Additionally, the model compound with an ortho relationship between the Br and OH groups (as in substructure C, Figure 1) cannot account for the shielded aromatic proton ($\delta_{\rm H}$ 6.02 ppm; $\delta_{\rm C}$ 108.3 ppm) observed in the NMR spectra of flustramine H.

It should be mentioned that compound **25**, an isomer of **4**, has been reported along with other novel compounds as constituents of *F. foliacea*.¹² In this study, **25** corresponds to substructure C in



Figure 3. Predicted ¹³C and ¹H NMR shifts for model compounds **4** and **24–26**. (A) Structures of the model compounds. (B) Experimental and predicted shifts: (a) calculated using ACD/ Laboratories software; (b) calculated *ab initio* using GIAO-HF within Gaussian 03 and the 6-31G** basis set.

Table 3. ¹H NMR Assignments for Flustramines 8–12 in CDCl₃ ($\delta_{\rm H}$, mult)^{*a*}

pos.	8	9	10	11	12
2	2.62, m	2.66, m	2.70, br m	2.63, ddd (10.4, 8.8, 5.6) 2.85, ddd (8.8, 7.0, 1.8)	2.70, m
3	2.05, dt (12.7, 6.7)	2.05, m	1.80, m	1.97, ddd (11.8, 10.4, 7.0)	1.62, m
	2.19, dt (12.7, 5.9)	2.19, m	2.20, m	2.09, ddd (11.8, 5.6, 1.8)	2.39, m
4					7.00, d (8.0)
5					6.80, dd (8.0, 1.6)
6	7.25, s	6.89, s	7.34, s	6.98, s	
7					6.75, d (1.6)
8a	4.39, br s	4.28, s	4.49, br s	5.21, br s	4.56, br s
9	2.47, dd (15.0, 8.1)	2.52, m	2.70, br m	2.59, m	
	2.69, dd (15.0, 6.6)	2.58, m			
10	4.96, br t (7)	5.05, br t (6.8)	5.06, br t (7)	5.01, br t (7)	0.99, s
11					1.10, s
12	1.61, s	1.60, s	1.63, s	1.67, s	6.03, dd (17.6, 10.5)
13	1.65, s	1.68, s	1.55, s	1.64, s	trans 5.10, d (17.6)
					<i>cis</i> 5.14, d (10.5)
14	2.44, s	3.22, br d (7)	6.41, s	3.10, dd (15.9, 7.5)	2.52, s
				3.19, dd (15.9, 7.5)	
15		5.24, br t (7)		5.23, t (7.5)	4.89, s
17		1.76, s	1.64, s	1.73, s	
18		1.76, s	1.64, s	1.69, s	
19		2.44, s	2.47, s	2.57, s	
OH	5.30, br s				
N <u>H</u>	4.39, br s	4.24, br s	4.41, br s		
NH_2				3.94, br s	

^{*a*} Recorded at 500 MHz; referenced to residual solvent at δ 7.24 ppm.

Figure 1, which is the substitution pattern that one might expect given precedent in the flustramine family. However, the reported spectroscopic data are nearly identical to those of **4**. In light of our studies, the reported structure should be reconsidered.

The detailed structural characterization of **4** revealed an additional substitution pattern in these alkaloids that, once established, permitted straightforward identification of the related metabolite flustramine I (**6**). The molecular formula for **6** was established as $C_{16}H_{21}BrN_2O$ ([M]⁺ 336.0853), indicating the loss of an isoprene unit (C_5H_8) when compared with the molecular formula of **4**. The ¹H NMR data (Table 2) for **6** confirmed this relationship by displaying resonances for only one olefinic methine and two olefinic methyls, consistent with a regular C_5 isoprene unit and unlike the reversed isoprene side chain found in **1**, **2**, and **3**. The ¹H and ¹³C NMR data for the indoline ring system in **4** and **6** were virtually identical, indicating the functional groups and substitution pattern are the same in both molecules. Extensive 2D NMR analysis including HMQC, HMBC, COSY, and NOESY confirmed the structure.

A characteristic triplet pattern for the molecular ion in the mass spectra of two additional compounds, **7** and **8**, indicated they were both dibromo derivatives. The first, flustramine J (**7**), possessed a molecular formula of $C_{21}H_{28}Br_2N_2O$ ($[M + H]^+$ 483.0644), indicating the molecule contained an additional bromine atom compared with **4**. Inspection of the ¹H NMR data (Table 2) revealed only one aromatic proton (δ_H 7.38 ppm), hence locating the additional bromine on the aromatic ring. The chemical shift of the aromatic methine suggested that it was *meta* to the phenolic moiety, implying that the two bromines were at C-5 and C-7. The ¹H and ¹³C NMR data for the isoprenoid side chain were very similar to those of **4**, indicating that the only difference between the compounds was in the aromatic portion of the molecule. The structure was fully supported by 2D NMR analysis including HMQC, HMBC, COSY, and NOESY.

The molecular formula $C_{16}H_{20}Br_2N_2O$ ([M]⁺ 413.9944) for the second dibromo derivative, flustramine K (8), suggested the compound was the dibromo derivative of 6. Indeed, comparison of the ¹H NMR data for 4, 6, 7, and 8 (Tables 2 and 3) confirmed that 8 was the 5,7-dibromo analogue of 6 in the same manner that 7 was the 5,7-dibromo analogue of 4. The ¹H and ¹³C NMR data for the isoprenoid side chain in 8 were also similar to that of 6,

establishing the same C_5 side chain. 2D NMR analysis supported this structure.

Accurate mass measurements on the molecular ion of flustramine L (9) ($C_{21}H_{29}BrN_2O$; [M]⁺ 404.1464) suggested it was an isomer of 4, although preliminary inspection of the ¹H NMR data (Table 3) indicated that 9 was quite different from 4. The most obvious difference in the ¹H NMR data was the presence of only one aromatic methine ($\delta_{\rm H}$ 6.89 ppm) and two separate C5 isoprene units. For example, the ¹H NMR spectrum of **9** displayed two olefinic methines ($\delta_{\rm H}$ 5.05 and 5.24 ppm) but four olefinic methyls at $\delta_{\rm H}$ 1.60 (3H), 1.68 (3H), and 1.76 ppm (6H), indicative of two C₅ isoprene units as opposed to a single C₁₀ chain. This information, together with the presence of only one aromatic methine, was consistent with one isoprene subunit attached to the aromatic ring, and this was confirmed by HMBC correlations from the benzylic methylene protons ($\delta_{\rm H}$ 3.22 ppm) to C-6 and C-3b. Furthermore, comparison of the ¹³C NMR data (Table 1) for the physostigmine carbon skeleton of 9 and 6 showed they were very similar (within 1.5 ppm), with the exception of the chemical shift for C-5, suggesting this as the point of attachment. The interconversion of flustramines L and N, described shortly, confirmed this.

The molecular formula $(C_{21}H_{27}BrN_2O_2; [M]^+ 418.1265)$ of flustramine M (10) corresponded to the molecular formula of 9, plus the addition of H₂O. The ¹H and ¹³C NMR data for 10 revealed the chemical shift resonances for C-2, C-3, C-3a, C-8a, and C-9-C-13 were very similar to the corresponding positions in 9, suggesting that this portion of both molecules was identical. However, the resonances assigned to the isoprene aromatic substituent in 9 were replaced in 10 by two new olefinic carbon resonances ($\delta_{\rm C}$ 161.6 ppm, qC; and $\delta_{\rm C}$ 100.0 ppm, CH; $\delta_{\rm H}$ 6.41 ppm) and a new deshielded singlet ($\delta_{\rm C}$ 69.1 ppm) in addition to two aliphatic methyl groups. These shifts suggest a five-membered heteroaromatic ring system and a tertiary alcohol. Two coincident deshielded methyl groups ($\delta_{\rm H}$ 1.64 ppm, $\delta_{\rm C}$ 28.7 ppm) showed HMBC correlations with the alcohol carbon and also with the furano singlet C-15 ($\delta_{\rm C}$ 161.6 ppm), suggesting an allylic carbon bearing a hydroxy group and the two methyls. A new aromatic methine ($\delta_{\rm H}$ 6.41 ppm) also showed HMBC correlations with C-15 and C-4 ($\delta_{\rm C}$ 150.3 ppm), which supported the presence of the furanobenzene moiety in 10. NOESY correlations from H-14 to H₃-17, H₃-18, and H-6 confirmed the substitution pattern as shown. One plausible origin for 10 could

Scheme 2. Rearrangement of 4 $(R_1 = C_{10}H_{17}, R_2 = H)$ and 9 $(R_1 = C_5H_9, R_2 = C_5H_9)$



involve intramolecular cyclization of 9. This raises the question of whether or not 10 is an isolation artifact produced from 9. No conversion of 9 into 10 was observed, even after several years of storage or after exposure to mild acid (pTSOH in EtOH), making it unlikely that 10 was an artifact. However, 9 does slowly interconvert with its isomer flustramine N (11) (see below).

The mass spectrometric data for flustramine N (11) $(C_{21}H_{29}BrN_2O; [M]^+ 404.1468)$ established that it was isomeric with 4 and 9. The ¹H NMR and ¹³C NMR data for positions C-2, C-9-C-13, and C-14–C-19 in 11 were similar to those for 9, suggesting that these portions of the molecule were the same. However, by the same comparison, the resonances associated with the methine at position 8a were quite different. For all of the other phenolic flustramines described to date, the H-8a methine resonates between $\delta_{\rm H}$ 4.20 and 4.50 ppm, whereas in **11**, H-8a was considerably deshielded at $\delta_{\rm H}$ 5.21 ppm. Similarly, the ¹³C resonance for C-8a was also deshielded and now appeared at $\delta_{\rm C}$ 106.2 ppm, consistent with a carbon bonded to at least one oxygen. While the imine proton of the other flustramines usually resonates in the region $\delta_{\rm H} 4.2-4.5$ ppm, the NMR data of flustramine N showed two protons not bonded to carbon resonating at $\delta_{\rm H}$ 3.94 ppm. This denoted the presence of a primary rather than a secondary amine in flustramine N and, taken with the other data, suggested that the compound did not possess the physostigmine skeleton. Instead, structure 11 was further supported by additional NMR data. The amine protons showed an HMBC correlation with the shielded bromo-substituted carbon C-5 ($\delta_{\rm C}$ 100.6 ppm), as well as C-3b ($\delta_{\rm C}$ 115.8 ppm). The benzylic methylene protons H₂-14 ($\delta_{\rm H}$ 3.10 and 3.19 ppm) of the isoprene substituent showed HMBC correlations with C-7 ($\delta_{\rm C}$ 114.9 ppm), C-6 (δ_{C} 131.3 ppm), and C-7a (δ_{C} 158.3 ppm), as well as the olefinic carbons C-15 ($\delta_{\rm C}$ 122.2 ppm) and C-16 ($\delta_{\rm C}$ 132.4 ppm). An additional correlation between H-8a and C-7a helped to further establish the substitution pattern as shown, which was also supported by NOESY correlations between H2-14 and H-6 and between NH₂ and H₂-9.

Following prolonged storage (\sim 5 y) of **11**, the ¹H NMR spectra displayed new resonances, which upon closer inspection matched those for 9. This prompted investigation of 9, which had been stored for a similar period of time, and the ¹H NMR data revealed an approximate 1:1 ratio of 9 and 11, confirming interconversion of the two compounds. A possible mechanism is presented in Scheme 2. For such a mechanism to operate, it is essential to have a phenolic group located at position C-4, and so the presence of 9 and 11 and their interconversion adds further indirect support for the proposed structure of 4. In order for this to be valid, it would be expected that the corresponding interconversion product of 4 would be observed following extended storage. Indeed, careful reinvestigation of the ¹³C NMR data of 4 suggested that such an interconversion was occurring. The characteristic resonance assigned to the deshielded methine C-8a was particularly distinct. However, the isomer of 4 was present only in trace amounts (<5%), suggesting that the interconversion of 4 must proceed even more slowly than that of 9.

The isolation of flustramine O(12) initially posed an interesting conundrum: the mass spectrometric data showed a molecular formula of $C_{33}H_{42}Br_2N_4$ ([M]⁺ 652.1769), yet the ¹H NMR and ¹³C NMR data (Tables 1 and 3) suggested that the compound was very similar to the known metabolite 1.5 In fact, the only significant differences in the ¹H spectra of **1** and **12** were the absence of the amine proton in the spectrum of 12 and the appearance of a new singlet at $\delta_{\rm H}$ 4.89 ppm. The ¹³C NMR data were also very similar, with the exception of the carbon resonances assigned to C-8a ($\delta_{\rm C}$ 90.5 ppm), C-14 ($\delta_{\rm C}$ 41.8 ppm), and a new resonance at $\delta_{\rm C}$ 58.2 ppm. This latter resonance ($\delta_{\rm H}$ 4.89 ppm/ $\delta_{\rm C}$ 58.2 ppm) integrated for only one proton, yet the DEPT spectra showed that it was a methylene. This observation, combined with a molecular formula that corresponded to two 1 units plus an additional methylene, suggested that flustramine O was a dimer of 1 linked through a methylene group. The absence of the imine resonance in the ¹H NMR spectrum of 12 implied that the two 1 molecules were linked through their respective imine functions by a single methylene bridge.

The molecular formula of flustramine P (13) (C₃₈H₅₀Br₂N₄O; [M]⁺ 736.2370) indicated that it too was substantially larger than most flustramines. On the basis of the example of 12, it appeared that this latest compound was composed of two flustramine molecules linked in some fashion. However, in contrast to 12, the ¹H and ¹³C NMR data for 13 clearly showed that this dimer was not symmetrical, nor did it contain two identical subunits (Table 4). Instead, the data displayed resonances for a single methylene group, a unit of 1, and a unit of a derivative of 4. Furthermore, the ¹H and ¹³C NMR data for the **1** portion of the molecule were essentially identical with the data for 12, suggesting that it was similarly linked through the indoline nitrogen. This was supported by the presence of only one non-carbon-bonded proton resonance in the NMR spectra of 13. The NMR data for the flustramine H subunit established the presence of only one aromatic methine (Table 4), implying a linkage through the aromatic ring of the flustramine H moiety. The deshielded nature of this aromatic methine ($\delta_{\rm H}$ 7.10 ppm) suggested the substitution of flustramine H at C-5'. These conclusions were supported by HMBC analyses, which showed correlations from the bridging methylene protons (H₂-15; $\delta_{\rm H}$ 4.14 and 4.25 ppm) to C-7a and C-8a of the dihydroflustramine C subunit and to C-4' and C-6' of the flustramine H subunit, thus confirming the structure of flustramine P as 13. The isolation of these methylene-linked compounds was surprising. As natural products, they could arise by joining two flustramine units through a one-carbon unit derived from formate. However, while it is interesting to speculate on a biosynthetic origin, it is more likely they are artifacts of the workup process that employed CH₂Cl₂.

The relative configuration of the ring juncture joining C-3a and C-8a was found to be *cis* for the novel flustramine congeners on the basis of the presence of NOE interactions between H-8a and the isoprenyl/reversed-isoprenyl/geranyl side chains.³ The relative configuration of **11** could not be assigned unambiguously due to overlap of multiple resonances. Furthermore, all of the newly isolated flustramines described here exhibited levorotation. Although we only report the relative configuration for the novel flustramines, it is worth noting that enantioselective syntheses of derivatives of flustramine B^{16,17,24} and flustramine A²⁴ have consistently shown levorotatory species having an absolute configuration equivalent to that shown for **1–10**, **12**, and **13**, although the *R* and *S* designations vary with the types of substituents at C-3a and N-8.

The flustramines described here are a family of biologically active, biosynthetically related alkaloids. The antimicrobial potency was determined for a number of flustramines (Table 5), with most of the tested compounds showing broad-spectrum activity. The two dimers **12** and **13** did not exhibit any detectable activity. Acetylation of the indole nitrogen in **2** relative to **1** also reduced the potency

Table 4. Complete NMR Assignments for Flustramine P (13) in $CDCl_3^a$

pos.	$\delta_{\rm C}$ (mult.)	$\delta_{\mathrm{H}} \left(J \text{ in Hz} \right)$	pos.	$\delta_{\rm C}$ (mult.)	$\delta_{ m H}~(J~{ m in~Hz})$
2	52.4, CH ₂	2.65, m	2'	52.4, CH ₂	2.65, m
3	32.8, CH ₂	1.78, dd (12.5, 5.0)	3'	36.3, CH ₂	2.04, m
		2.32, m			2.19, m
3a	64.6, qC		3a'	59.4, qC	
3b	130.8, qC		3b'	116.2, qC	
4	125.5, CH	6.90, d (7.9)	4'	153.4, qC	
5	119.1, CH	6.64, dd (7.9, 1.6)	5'	122.3, qC	
6	122.1, qC		6'	132.7, CH	7.10, s
7	108.8, CH	6.47, d (1.6)	7'	91.7, qC	
7a	151.3, qC		7a′	148.2, qC	
8a	93.6, CH	4.40, s	8a'	86.1, CH	4.22, s
9	40.7, qC	,	9'	35.1, CH ₂	2.53, m, 2.70, m
10	23.8, CH ₃	0.94, s	10'	121.0, CH	5.00, m
11	23.0, CH ₃	1.07, s	11'	137.1, gC	
12	144.7, CH	5.94, dd (17.4, 11.0)	12'	40.0, CH ₂	1.97, m
13	114.0, CH ₂	trans 5.02, d (17.4)	13'	26.8, CH ₂	1.97, m
	, _	<i>cis</i> 5.13, d (11.0)	14'	124.4, CH	5.03, m
14	41.7, CH ₃	2.48, s	15'	131.2, gC	,
15	48.3, CH ₂	4.14, d (15.0)	16'	17.7. CH ₃	1.67. s
		4.25, d (15.0)	17'	25.8, CH ₃	1.58, s
			18'	16.4, CH ₃	1.60, s
			19'	37.2. CH ₃	2.43. s
			NH	,,	4.30 br s

^{*a*} Recorded at 300 MHz; referenced to residual solvent at $\delta_{\rm H}$ 7.24 ppm and $\delta_{\rm C}$ 77.0 ppm. Assignment supported by HMQC and HMBC experiments obtained at 500 MHz.

Table 5. Antimicrobial Activity of Flustramines: Zone ofInhibition (mm)

	2	5	6	9	12	13
mol/disk ($\times 10^{6}$)	1.4	1.3	1.5	1.2	0.9	0.7
E. coli		22	25	24		
M. luteus		23	29	25		
C. utilis	18	28	29	30		
S. cerevisiae		16	15	20		

against most of the tested organisms, suggesting that antimicrobial activity is hindered by substitution of the indoline nitrogen.

Experimental Section

General Experimental Procedures. Optical rotations were determined with a Perkin-Elmer 141 polarimeter. Ultraviolet and infrared spectra were recorded with a GCA-MacPherson Series 700 and a Perkin-Elmer 283B spectrophotometer. The ¹H NMR and ¹³C NMR data were obtained at either 500 and 125 MHz, respectively, or 300 and 75 MHz, as indicated, on a Bruker AMX 500 or Bruker AML 300 spectrometer, respectively. Electron impact mass spectra were obtained with a Dupont model 21-110B double-focusing spectrometer used in electrical detection mode. FAB mass spectra were obtained using a Micromass/AutoSpec oa-TOF hybrid sector/time-of-flight mass spectometer (Manchester, UK). LSIMS (liquid secondary ion mass spectrometry)/MS/MS experiments were performed using a collision energy of 800 eV and Xe collision gas. The LSIMS matrix used was a 1:1 mixture of *m*-nitrobenzyl alcohol and glycerol with 1% trifluoroacetic acid. Exact mass measurements were carried out at a mass resolution of 10 000 using voltage scanning, and the peaks were calibrated using polyethylene glycol reference peaks. TLC was carried out using glassbacked silica TLC plates (Merck) and 100% EtOAc as solvent and were visualized with a vanillin spray reagent (0.5 g vanillin/100 mL H₂SO₄/EtOH, 4:1) and gentle heat. Large-scale HPLC was performed with a Waters Associates Prep System, and semipreparative HPLC was carried out using a Waters Associates model 6000 pump, U6K injector, and a model 450 variable-wavelength detector.

Animal Material. The bryozoan *F. foliacea* was collected in the Minas Basin of the Bay of Fundy off the New Brunswick and Nova Scotian shores and identified by Prof. Sherman Blakeney of the Department of Biology, Acadia University, Wolfville, N.S. A voucher specimen has been retained at the Nova Scotia Museum, Halifax, N.S., under the catalog number 17573.

Antimicrobial Assay. Antimicrobial activity was determined using a standard disk-diffusion assay. A solution of the test compound in MeOH was applied to a 6 mm diameter filter paper and air-dried. The disk was transferred to an agar plate overgrown with the test organism, and the development of zones of growth inhibition was measured from the outer edge of the disk. The strains used for testing were *Escherichia coli* (ATCC 11775), *Micrococcus luteus* (ATCC 49732), *Candida utilis* (ATCC 9950), and *Saccharomyces cerevisiae* (ATCC 9763).

Isolation of Flustramines. The bryozoan (6 kg) was ground to a pulp and extracted with methanol at room temperature for 2 days. The methanolic extract (10 L) was concentrated under reduced pressure until most of the methanol was removed, and the residue was shaken against dichloromethane. Half of the dichloromethane-soluble fraction (15.0 g) was chromatographed on a pad of silica gel (Merck) to yield two indole-rich fractions eluting with 0-25% EtOAc/Et₂O (1.726 and 1.756 g). These fractions were further purified by preparative liquid chromatography (C18 Lobar column, 30×2 cm, 100 mL/min, 80-100%MeOH in H₂O; and silica, Merck, TLC grade, 1.5 mL/min, 0-10% MeOH in CH₂Cl₂) to yield a number of fractions including pure dihydroflustramine C (1; 455.9 mg). Other fractions contained complex mixtures of the indole alkaloids, which were eventually purified (after several preparative silica and reversed-phase steps involving the pooling of similar fractions) by semipreparative HPLC (Whatman Partisil M9 10/50 ODSII 10% H₂O/MeOH (0.2% Et₃N) or CSC S5W silica column, 25×0.9 cm, 4% MeOH/CH₂Cl₂) to yield 1 and flustramines F–P.

Dihydroflustramine C (1): crystalline solid (616 mg), all spectroscopic data in good agreement with literature data.⁵

Acetylation of 1. Magnesium turnings (0.07 g, 2.9 mmol), an iodine crystal, and ethyl bromide (0.25 mL, 0.37 g, 3.3 mmol) were stirred in diethyl ether (5 mL) and gently heated for 20 min. An ethereal solution of 1 (64 mg in 1 mL, 0.2 mmol) and acetyl chloride (150 μ L, 2 mmol) was added dropwise, and the solution maintained at reflux for 20 min. After this time the reaction mixture was poured onto crushed ice, acidified (AcOH), and extracted with CH₂Cl₂ (3 × 15 mL). The organic solvent was evaporated under reduced pressure to yield a mixture (44 mg). This mixture was purified by HPLC (Whatman ODSII 10% H₂O/MeOH, 3 mL/min) to yield pure *N*-acetyl dihydroflustramine C (24.1 mg). This product was spectroscopically identical to **2**.

Flustramine F (2): oil (31 mg); $[\alpha]_D -22$ (*c* 0.20, CHCl₃); UV (MeOH) λ_{max} (log ϵ) 219 (4.41), 252 (3.98), 287 (3.39); IR (film) ν_{max} 3250, 2970, 1660, 1585, 1470, 1410, 1385, 1290, 1110 cm⁻¹; ¹³C NMR Table 1, ¹H NMR Table 2; HREIMS *m*/*z* 362.0987 (calcd for C₁₈H₂₃⁷⁹BrN₂O 362.0993); *m*/*z* 364.0972 (calcd for C₁₈H₂₃⁸¹BrN₂O 364.0974); TLC *R*_f 0.36 (gray upon spraying and heating).

Flustramine G (3): oil (7 mg); $[\alpha]_D - 80$ (*c* 0.21, CHCl₃); UV (MeOH) λ_{max} 212, 247, 310; IR (film) ν_{max} 3420, 2960, 1590, 1470, 1410, 1350, 1160, 1090 cm⁻¹; ¹³C NMR Table 1, ¹H NMR Table 2; EIMS *m*/*z* (%) 398/400/402 (15/30/15), 329/331/333 (60/100/55), 286/

288/290 (15/20/15), 250/252 (20/20), 171 (20), 128 (10), 69 (25); HREIMS m/z 397.9990 (calcd for $C_{16}H_{20}^{79}Br_2N_2$ 397.9994); m/z 401.9960 (calcd for $C_{16}H_{20}^{81}Br_2N_2$ 401.9953); TLC R_f 0.17 (yellow upon spraying then gray upon heating).

Flustramine H (4): oil (36 mg); $[\alpha]_D - 58$ (*c* 0.30, CHCl₃); UV (MeOH) λ_{max} (log ϵ) 217 (4.77), 246 (4.20), 295 (3.84); IR (film) ν_{max} 3415, 2975, 1623, 1593, 1469, 1380, 1100, 1037 cm⁻¹; ¹³C NMR Table 1, ¹H NMR Table 2; EIMS *m/z* (%) 404/406 (48/50), 335/337 (70/70), 267/269 (98/100), 224/226 (25/28), 188 (22), 117 (12), 69 (32); HREIMS *m/z* 404.1460 (calcd for C₂₁H₂₉⁷⁹BrN₂O 404.1464); TLC *R*_f 0.31 (purple upon spraying and heating).

Debromoflustramine H (5): 4 (12 mg) and LiAlH₄ (2 mg) were heated at reflux in dry diethyl ether (5 mL) for 5 h before quenching with H₂O (100 mL). The product was extracted with diethyl ether and the organic solvent removed under reduced pressure to yield a mixture, which was purified by HPLC (Whatman ODSII 10% H₂O/MeOH (0.3% Et₃N), 3 mL/min) to yield **4** (2.5 mg) and **5** (2.2 mg) as an oil: ¹H NMR (CDCl₃, 300 MHz) 6.90 (dd, 7.9 Hz, H-6), 6.19 (d, 7.9 Hz, H-5 or H-7), 6.14 (7.9 Hz, H-5 or H-7), 5.00 (br m, H₂-9, H_b-2), 2.57 (s, H₃-19), 2.43 (m, H_a-3), 2.24 (m, H_b-3), 1.94 (m, H₂-12 and H₂-13), 1.65 (H₃-18 or H₃-16), 1.60 (s, H₃-18 or H₃-16), 1.56 (s, H₃-17); EIMS *m/z* (%) 326 (35), 269 (15), 257 (98), 214 (25), 189 (98), 146 (100), 86 (99); HRFABMS *m/z* 327.2444 (calcd. for [M + H]⁺ C₂₁H₃₁N₂O 327.2436).

Flustramine I (6): oil (20 mg); $[\alpha]_D - 87$ (*c* 0.18, CHCl₃); UV (MeOH) λ_{max} (log ϵ) 217 (4.79), 297 (3.65); IR (film) ν_{max} 2980, 2940, 1675, 1640, 1520, 1430, 1395, 1200, 1170 cm⁻¹; ¹³C NMR Table 1, ¹H NMR Table 2; EIMS *m/z* (%) 336/338 (60/50), 279/281 (70/70), 267/269 (100/80), 224/226 (60/55), 188 (40), 171 (10), 159 (6), 146 (5), 84 (70); HREIMS *m/z* 336.0853 (calcd for C₁₆H₂₁⁷⁹BrN₂O 336.0838); *m/z* 338.0823 (calcd for C₁₆H₂₁⁸¹BrN₂O 338.0817); TLC *R*_f 0.16 (purple upon spraying and heating).

Flustramine J (7): oil (8 mg); $[\alpha]_D -40$ (*c* 0.15, CHCl₃); UV (MeOH) λ_{max} 217, 293; IR (film) ν_{max} 3410, 2970, 2930, 1720, 1670, 1600, 1450, 1380, 1270, 1160 cm⁻¹; ¹³C NMR Table 1, ¹H NMR Table 2; EIMS *m/z* (%) 482/484/486 (30/65/30), 413/415/417 (50/100/50), 345/347/349 (15/30/17), 302/304/306 (5/10/5), 69 (20); HRFABMS *m/z* 483.0644 (calcd for [M + H]⁺ C₂₁H₂₉⁷⁹Br₂N₂O 483.0647); *m/z* 485.0633 (calcd for [M + H]⁺ C₂₁H₂₉⁷⁹Br⁸¹BrN₂O 485.0626); TLC *R*_f 0.41 (purple upon spraying and heating).

Flustramine K (8): oil (2.7 mg); $[\alpha]_D - 29$ (*c* 0.05, CHCl₃); UV (MeOH) λ_{max} 215, 245, 303; IR (film) ν_{max} 3226, 2960, 1720, 1600, 1450, 1300, 1260, 1160 cm⁻¹; ¹³C NMR Table 1, ¹H NMR Table 3; EIMS *m*/*z* (%) 414/416/418 (35/65/30), 357/359/361 (50/100/50), 345/ 347/349 (70/100/50), 302/304/306 (30/55/30), 225 (20), 116 (20), 96 (35), 69 (75); HRFABMS *m*/*z* 413.9944 (calcd for [M + H]⁺ C₁₆H₂₀⁷⁹Br₂N₂O 413.9943); *m*/*z* 415.9926 (calcd for [M + H]⁺ C₁₆H₂₀⁷⁹Br⁸¹BrN₂O 415.9923); TLC *R*_f 0.28 (mustard upon spraying and heating).

Flustramine L (9): oil (5.4 mg); $[\alpha]_D -53$ (*c* 0.30, CHCl₃); UV (MeOH) λ_{max} (log ϵ) 218 (4.82), 302 (3.83); IR (film) ν_{max} 3420, 2970, 2930, 2800, 1620, 1590, 1470, 1460, 1380, 1350, 1270, 1150 cm⁻¹; ¹³C NMR Table 1, ¹H NMR Table 3; EIMS *m/z* (%) 404/406 (80/100), 361/363 (15/20), 347/349 (45/60), 335/337 (70/70), 291/292 (25/20), 279/280 (35/35), 238 (35), 69 (32); HREIMS *m/z* 404.1464 (calcd for C₂₁H₂₉⁷⁹BrN₂O 404.1464); *m/z* 406.1450 (calcd for C₂₁H₂₉⁸¹BrN₂O 406.1443); TLC *R*_f 0.22 (yellow upon spraying and heating).

Flustramine M (10): oil (2.8 mg); $[\alpha]_D - 159$ (*c* 0.08, CHCl₃); UV (MeOH) λ_{max} (log ϵ) 218 (4.50), 209 (4.06), 318 (3.85); IR (film) ν_{max} 3420, 2970, 2940, 2870, 2800, 1640, 1460, 1430, 1380, 1340, 1260, 1140 cm⁻¹; ¹³C NMR Table 1, ¹H NMR Table 3; EIMS *m/z* (%) 418/ 420 (25/25), 400/402 (5/5), 349/351 (40/40), 279/281 (10/10), 243 (70), 159 (100), 69 (32); HREIMS *m/z* 418.1265 (calcd for C₂₁H₂₇⁷⁹BrN₂O₂ 418.1256); *m/z* 420.1242 (calcd for C₂₁H₂₇⁸¹BrN₂O₂ 420.1236); TLC *R*_f 0.13 (orange upon spraying and heating).

Flustramine N (11): oil (2.7 mg); $[\alpha]_D - 61$ (*c* 0.28, CHCl₃); UV (MeOH) λ_{max} (log ϵ) 217 (4.73), 299 (3.79); IR (film) ν_{max} 3490, 3390, 2975, 2860, 2815, 1630, 1480, 1430, 1360, 1260, 1150 cm⁻¹; ¹³C NMR Table 1, ¹H NMR Table 3; EIMS *m/z* (%) 404/406 (35/25), 335/337 (10/05), 279/281 (100/90), 251 (5), 157 (2), 69 (18); EIMS *m/z* 404.1468 (calcd for C₂₁H₂₉⁷⁹BrN₂O 404.1464); *m/z* 406.1454 (calcd for C₂₁H₂₉⁸¹BrN₂O 406.1443); TLC *R_f* 0.20 (yellow upon spraying and heating).

Flustramine O (12): oil (46.0 mg); $[\alpha]_D -211$ (*c* 0.63, CHCl₃); UV (MeOH) λ_{max} (log ϵ) 215 (4.82), 259 (4.32), 312 (3.89); IR (film) ν_{max} 3090, 2970, 2870, 2800, 1640, 1590, 1480, 1420, 1310, 1270, 1170 cm⁻¹; ¹³C NMR Table 1, ¹H NMR Table 3; EIMS *m/z* (%) 652/ 654/656 (20/40/20), 583/585/586 (70/100/40), 320 (80), 251/253 (30/ 30), 172 (75), 69 (50); HREIMS *m/z* 652.1769 (calcd for C₃₃H₄₂⁷⁹Br₂N₄ 652.1777); *m/z* 654.1758 (calcd for C₃₃H₄₂⁸¹Br₂N₄ 654.1758); TLC *R_f* 0.35 (yellow upon spraying, then gray upon heating).

Flustramine P (13): oil (25.0 mg); $[\alpha]_D - 5$ (*c* 0.44, CHCl₃); UV (MeOH) λ_{max} (log ϵ) 213 (5.06), 259 (4.53), 305 (4.07); IR (film) ν_{max} 3420, 2970, 2930, 2790, 1620, 1590, 1490, 1320, 1160 cm⁻¹; ¹³C NMR Table 4, ¹H NMR Table 4; EIMS *m/z* (%) 736/738/740 (1/2/1), 667/669/667 (2/4/2), 320/322 (30/35), 251/253 (95/100), 208/210 (30/35), 172 (35), 128 (25), 69 (15); EIMS *m/z* 736.2370 (calcd for C₃₈H₅₀⁷⁹Br₂N₄O 736.2352); *m/z* 738.2347 (calcd for C₃₈H₅₀⁷⁹Br⁸¹BrN₄O 738.2331); TLC *R_f* 0.22 (purple upon spraying and heating).

Permethylflustramine H (19). 4 (1.5 mg) was dissolved in acetone (1 mL). To the stirred solution were added MeI (0.1 mL) and K₂CO₃ (20 mg), and the solution was allowed to stir at room temperature for 12 h. The reaction was diluted with H₂O and extracted with CH₂Cl₂ (3 \times 5 mL) to yield, after removal of the organic solvent, a mixture of polar compounds (3 mg). Initial purification by silica flash chromatography yielded one fraction of interest (0.96 mg). Further purification by HPLC (Zorbax RX-C8, 0.46×25 cm, gradient elution: 67% ACN/ H₂O (0.1% TFA) to 90% ACN/H₂O (0.1% TFA) over 9 min, 0.5 mL/ min) yielded a major fraction, **19** (0.40 mg), as an oil: ¹H NMR (60% benzene-d₆/CDCl₃, 500 MHz) 7.15 (H-6, under solvent), 6.05 (d, 7.5 Hz, H-5), 5.05 (m, H-15), 4.89 (m, H-11), 3.63 (s, H₃-20), 3.38 (s, OMe), 3.26 (brd, 4 Hz, H_2 -10), 2.85 (m, H_2 -9), 2.72 (H_2 -8), 2.60 (s, NMe3), 2.06 (m, H2-14), 1.98 (m, H2-13), 1.68 (s, H3-19), 1.62 (s, H3-18), 1.50 (H₃-17); ¹³C NMR (60% benzene-*d*₆/CDCl₃, from HMQC and HMBC, not all carbons seen) 153.9 (C-4), 139.0 (C-2), 138.0 (C-12), 135.0 (C-7a), 132.0 (C-16), 128.5 (C-3), 124.5 (C-15), 121.0 (C-11), 120.0 (C-6 or C-3a), 102.0 (C-5), 96.1 (C-7), 67.5 (C-9), 55.5 (OMe), 52.5 (NMe₃), 40.0 (C-13), 33.9 (C-20), 27.0 (C-14), 26.2 (C-17), 23.4 (C-10), 18.2 (C-18), 16.8 (C-19); FABMS m/z (%) 461/463 (50/40), 402/404 (15/15), 338 (25), 246 (100), 226 (47); FABMS/MS ion selected 463.50 m/z (%) 404 (100), 389 (2), 349 (35), 335 (20), 334 (20), 320 (57), 306 (27), 293 (54), 281 (68), 280 (57), 266 (53), 254 (30), 225 (50); FABMS/MS ion selected 461.50 m/z (%) 402 (100), 387 (28), 347 (25), 333 (20), 332 (23), 318 (60), 304 (26), 291 (50), 279 (75), 278 (60), 264 (55), 252 (33), 225 (53); HRFABMS m/z 461.2178 (calcd for [M]⁺ C₂₅H₃₈⁷⁹BrN₂O 461.2162); *m/z* 463.2156 (calcd for C₂₅H₃₈⁸¹BrN₂O 463.2147).

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Supporting Information Available: ¹H NMR spectra for **2–13**; ¹³C NMR and HMQC spectra for **4**. This material is available free of charge via the Internet at http://pubs.acs.org.

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