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Shi-Hai Xu,^a* Xiao-Jian Liao,^a Xiang-Chao Zeng,^a Kai Yang,^a Dong-Hong He^a and Kai-Bei Yu^b

^aDepartment of Chemistry, Jinan University, Guangzhou, Guangdong, 510632, People's Republic of China, and ^bChengdu Branch, The Chinese Academy of Sciences, Chengdu 610041, People's Republic of China

Correspondence e-mail: txush@jnu.edu.cn

Key indicators

Single-crystal X-ray study T = 296 KMean σ (C–C) = 0.012 Å Disorder in solvent or counterion R factor = 0.037 wR factor = 0.067 Data-to-parameter ratio = 15.1

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

Dibromophakellin methanol hemisolvate

The title compound, $C_{11}H_{11}Br_2N_5O \cdot 0.5CH_3OH$, was isolated from algae (*Laurencia majuscula Lucas*) collected from the South China Sea and its crystal structure was determined. It exhibits mild antibacterial and antineoplastic activity. The crystal structure has supramolecular layers with $N-H \cdots O$ and $N-H \cdots N$ hydrogen bonds. The methanol solvent molecule is threefold disordered.

Comment

Dibromophakellin (m.p. 515 K), (I), a brominated pyrrole alkaloid, was first isolated by Sharma & Burkholder (1971) from the marine Phakellia flabellate off the coast of the Great Barrier Reef in Australia. It has also been prepared by a biomimetic synthesis (Foley & Büchi, 1982). Its structure has been confirmed by X-ray analysis of its acetate (Fedoreyev *et al.*, 1986). Despite the presence of both aminoacetal and diaminoketal functionalities dibromophakellin exhibits considerable stability toward hydrolytic reagents.

HN HN Br (I)

X-ray diffraction analysis reveals that the title compound, (I), as its methanol hemisolvate, has a compact tetracyclic core that includes a pyrrole and a guanidine unit in a fivemembered ring. There are two essentially identical molecules in the asymmetric unit, together with a molecule of methanol disordered over three orientations. Supramolecular layers are stabilized by $N-H\cdots O$ and $N-H\cdots N$ hydrogen bonds (Table 1 and Fig. 2).

Experimental

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The chopped algae were extracted with ethanol at room temperature. The extract was subjected to silica-gel column chromatography, eluting with petroleum ether containing an increasing amount of ethyl acetate and then chloroform containing an increasing amount of methanol. The fraction eluted with chloroform–methanol (3:97) gave the title compound after evaporation of the solvent.

| Crystal data | |
|---------------------------------|-------------------------------------------|
| $C_{11}H_{11}Br_2N_5O.0.5CH_4O$ | $D_x = 1.864 \text{ Mg m}^{-3}$ |
| $M_r = 405.09$ | Mo $K\alpha$ radiation |
| Monoclinic, P2 ₁ | Cell parameters from 38 |
| a = 7.646 (1) Å | reflections |
| b = 28.050 (6) Å | $\theta = 3.015.3^{\circ}$ |
| c = 7.670(1) Å | $\mu = 5.62 \text{ mm}^{-1}$ |
| $\beta = 118.65 \ (1)^{\circ}$ | T = 296 (2) K |
| $V = 1443.5 (5) \text{ Å}^3$ | Irregular, colourless |
| Z = 4 | $0.48 \times 0.30 \times 0.22 \text{ mm}$ |

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Data collection

Siemens P4 diffractometer ω scans Absorption correction: multi-scan (SHELXTL; Siemens, 1994) $T_{min} = 0.104, T_{max} = 0.290$ 6923 measured reflections 5939 independent reflections 3078 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.037$ $wR(F^2) = 0.067$ S = 0.805939 reflections 393 parameters H atoms treated by a mixture of independent and constrained refinement $\begin{aligned} R_{\text{int}} &= 0.037\\ \theta_{\text{max}} &= 26.5^{\circ}\\ h &= -9 \rightarrow 9\\ k &= -35 \rightarrow 35\\ l &= -9 \rightarrow 8\\ 3 \text{ standard reflections}\\ \text{every 97 reflections}\\ \text{intensity decay: 7.6\%} \end{aligned}$

$$\begin{split} &w = 1/[\sigma^2(F_o^2) + (0.023P)^2] \\ &where \ P = (F_o^2 + 2F_c^2)/3 \\ &(\Delta/\sigma)_{\rm max} = 0.001 \\ &\Delta\rho_{\rm max} = 0.29 \ {\rm e} \ {\rm \AA}^{-3} \\ &\Delta\rho_{\rm min} = -0.43 \ {\rm e} \ {\rm \AA}^{-3} \\ &{\rm Absolute \ structure: \ Flack \ (1983);} \\ &2877 \ {\rm Friedel \ pairs} \\ &{\rm Flack \ parameter} = 0.000 \ (11) \end{split}$$

Table 1

Hydrogen-bonding geometry (Å, °).

| $D - \mathbf{H} \cdot \cdot \cdot A$ | D-H | $H \cdot \cdot \cdot A$ | $D \cdot \cdot \cdot A$ | $D - \mathbf{H} \cdot \cdot \cdot A$ |
|----------------------------------------|------------|-------------------------|-------------------------|--------------------------------------|
| N3-H3N···O1 ⁱ | 0.856 (10) | 2.14 (3) | 2.921 (7) | 151 (4) |
| $N5-H5BN\cdotsO1^{i}$ | 0.86 | 2.36 | 3.096 (7) | 143 |
| N5−H5AN···N4′ ⁱⁱ | 0.86 | 2.13 | 2.981 (8) | 171 |
| $N3' - H3N' \cdots O1'^{iii}$ | 0.855 (10) | 2.13 (3) | 2.833 (7) | 139 (5) |
| $N5' - H5'D \cdots O1'^{iii}$ | 0.86 | 2.34 | 3.062 (7) | 142 |
| $N5' - H5'C \cdot \cdot \cdot N4^{iv}$ | 0.86 | 2.12 | 2.961 (7) | 167 |
| | | | | |

Symmetry codes: (i) x, y, 1+z; (ii) $1-x, \frac{1}{2}+y, 1-z$; (iii) x-1, y, z; (iv) $1-x, y-\frac{1}{2}, 1-z$.

The methanol solvent molecule is disordered over three orientations on approximately the same site. These orientations were refined with a C–O bond-length restraint of 1.480 (4) Å and riding H atoms (C–H = 0.96 Å and O–H = 0.82 Å). In the molecule of dibromophakellin, H atoms were positioned geometrically and refined as riding, with C–H = 0.93–0.98 Å and N–H = 0.86 Å, except that the H atoms attached to ring N atoms were refined freely. For all constrained H atoms, $U_{iso}(H) = 1.2U_{eq}$ (parent atom).

Data collection: *XSCANS* (Siemens, 1994); cell refinement: *XSCANS*; data reduction: *SHELXTL* (Siemens, 1994); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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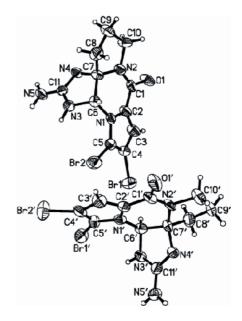


Figure 1

The two molecules in the asymmetric unit of the title compound, shown with 50% probability ellipsoids. The disordered solvent molecule is not shown.

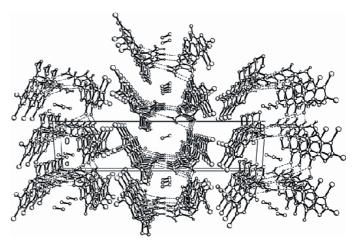


Figure 2

The packing of the title compound, with hydrogen bonds shown as dashed lines. H atoms of the disordered solvent molecule have been omitted, and only one methanol orientation is shown on each site.

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