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Key indicators

Single-crystal X-ray study
 $T = 298\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.005\text{ \AA}$
 R factor = 0.037
 wR factor = 0.110
Data-to-parameter ratio = 18.5For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Ethyl (4-bromo-1*H*-pyrrole-2-carboxamido)acetate

The title compound, $\text{C}_9\text{H}_{11}\text{BrN}_2\text{O}_3$, was synthesized by condensation of glycine ethyl ester with 4-bromo-2-(trichloroacetyl)pyrrole at room temperature in 78.4% yield. In the crystal structure, intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen-bond interactions link the molecules into two-dimensional sheets.

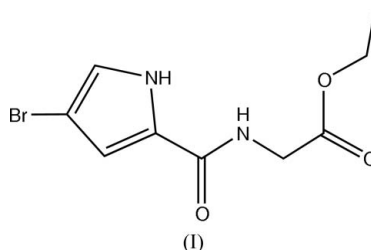
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Comment

Pyrrole derivatives are well known in many marine organisms (Faulkner, 2001), and some of them are bioactive substances (Tasdemir *et al.*, 2002). In our search for bioactive compounds, a series of brominated pyrrole-2-carboxamido acid esters, including the title compound, (I), has been synthesized by reaction of amino acid esters with brominated 2-(trichloroacetyl)pyrrole, or brominated 1-methyl-2-(trichloroacetyl)pyrrole.



We report here the structure of (I) which has been shown to inhibit *Streptococcus faecalis* and *Micrococcus luteus* moderately in pharmacological studies (Zeng *et al.*, 2004). Bond lengths and angles are unexceptional and are in good agreement with the corresponding values in [(4,5-dibromo-1-methyl-1*H*-pyrrole-2-carbonyl)amino]acetic acid methyl ester (Zeng *et al.*, 2004). In the crystal structure, there are two kinds of intermolecular hydrogen bonds. The $\text{N1}-\text{H}\cdots\text{O1}$ hydrogen bonds (Table 1) form centrosymmetric dimers (Fig. 2), which graph-set analysis describes as an $R_2^2(10)$ motif (Bernstein *et al.*, 1995). At the same time, the dimeric units are further assembled into a herring-bone pattern in the overall

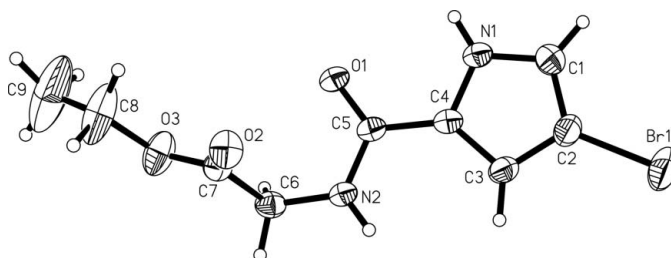


Figure 1

The molecular structure of the title compound, with the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.

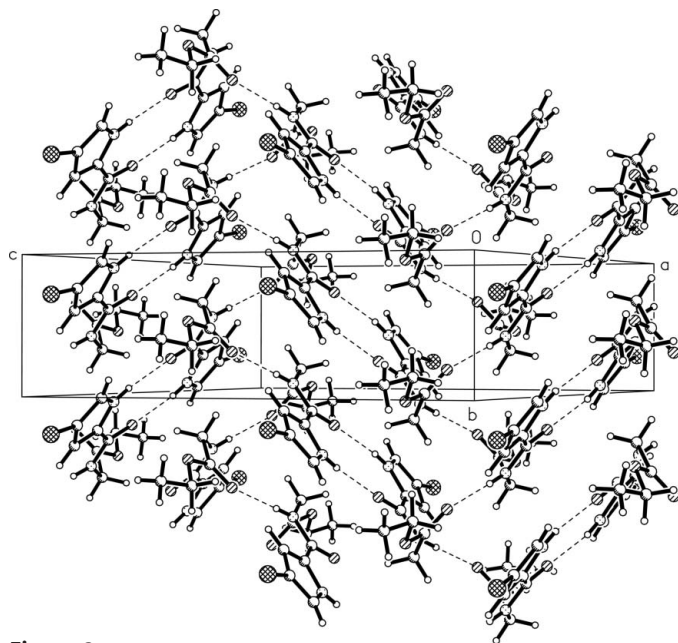


Figure 2
The crystal packing of (I), showing the two-dimensional sheet formed by hydrogen bonds (dashed lines).

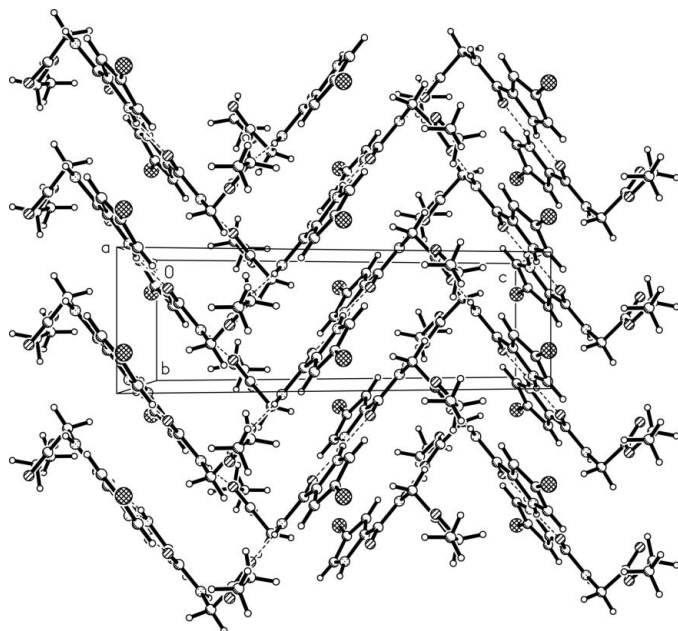


Figure 3
The crystal packing of (I). Dashed lines indicate hydrogen bonds.

crystal packing (Fig. 3). Individual dimeric units are linked by $N2-H \cdots O2$ hydrogen bonds, generating two-dimensional sheets (Fig. 2).

Experimental

A hydrochloric acid salt of glycine ethyl ester (0.70 g, 5 mmol) and 4-bromo-2-(trichloroacetyl)pyrrole (1.46 g, 5 mmol) were added to acetonitrile (12 ml), followed by the dropwise addition of triethylamine (1.4 ml). The mixture was stirred at room temperature for 10 h and then poured into water. After filtration, the precipitate was collected as a pale-yellow solid. The impure product was dissolved in

ethanol at room temperature. Colorless monoclinic crystals suitable for X-ray analysis (m.p. 454 K, 78.4% yield) grew over a period of 7 d when the solution was exposed to air. $^1\text{H NMR}$ (DMSO- d_6 , 300 Hz): 11.88 (*brs*, 1H), 8.57 (*t*, 1H), 7.01–6.99 (*m*, 1H), 6.87–6.86 (*m*, 1H), 3.96 (*d*, 2H), 3.53 (*q*, 2H), 1.21 (*t*, 3H); IR(KBr): 3375, 3217, 3140, 1732, 1644, 1566, 1524, 1335, 1224, 1130; analysis calculated for $\text{C}_9\text{H}_{11}\text{BrN}_2\text{O}_3$: C 39.29, H 4.03, N 10.18%; found: C 39.41, H 3.95, N 10.10%.

Crystal data

$\text{C}_9\text{H}_{11}\text{BrN}_2\text{O}_3$
 $M_r = 275.11$
Monoclinic, $P2_1/c$
 $a = 14.944$ (2) Å
 $b = 5.0501$ (8) Å
 $c = 16.264$ (3) Å
 $\beta = 107.843$ (3)°
 $V = 1168.4$ (3) Å³
 $Z = 4$

$D_x = 1.564$ Mg m⁻³
Mo $K\alpha$ radiation
Cell parameters from 1926 reflections
 $\theta = 2.6$ – 26.3 °
 $\mu = 3.51$ mm⁻¹
 $T = 298$ (2) K
Block, colourless
 $0.50 \times 0.38 \times 0.34$ mm

Data collection

Bruker SMART 1K CCD area-detector diffractometer
 φ and ω scans
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.202$, $T_{\max} = 0.303$
6732 measured reflections

2550 independent reflections
1562 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.020$
 $\theta_{\text{max}} = 27.0$ °
 $h = -16 \rightarrow 19$
 $k = -5 \rightarrow 6$
 $l = -20 \rightarrow 19$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.110$
 $S = 1.03$
2550 reflections
138 parameters
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0463P)^2 + 0.5034P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.46$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.34$ e Å⁻³
Extinction correction: SHELXTL
Extinction coefficient: 0.0053 (10)

Table 1
Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$N1-H1A \cdots O1^i$	0.86	1.97	2.788 (3)	159
$N2-H2 \cdots O2^{ii}$	0.86	2.08	2.935 (3)	174

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

H atoms were positioned geometrically ($C-H = 0.96$ Å for CH_3 , $C-H = 0.97$ Å for CH_2 , $C-H = 0.93$ Å for CH and $N-H = 0.86$ Å) and refined using a riding model, with $U_{\text{iso}} = 1.2U_{\text{eq}}$ ($1.5U_{\text{eq}}$ for the methyl group) of the parent atom.

Data collection: SMART (Bruker, 1999); cell refinement: SAINT-Plus (Bruker, 1999); data reduction: SAINT-Plus; program(s) used to solve structure: SHELXTL (Bruker, 1997); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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