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Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(C-C) = 0.004 \text{ Å}$ R factor = 0.046 wR factor = 0.126 Data-to-parameter ratio = 19.2

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Ethyl 4-[1-(4-bromophenyl)-3-methyl-5oxo-4,5-dihydro-1*H*-1,2,4-triazol-4-yliminomethyl]phenoxyacetate

In the crystal structure of the title compound, $C_{20}H_{19}BrN_4O_4$, the asymmetric unit consists of two crystallographically independent molecules, which differ in the relative orientations of the triazole and phenyl rings. In the crystal structure, both molecules form approximately centrosymmetric dimers through weak $C-H \cdots O$ intermolecular interactions. Received 21 July 2003 Accepted 31 July 2003 Online 8 August 2003

Comment

The triazole moiety possesses many pharmacological properties, such as acting as an antimicrobial (Habib *et al.*, 1997), antiviral (Ergen *et al.*, 1996), anti-HIV-1 (Invidiata *et al.*, 1996), antifungal, antimycobacterial and anticonvulsant agent (Gülerman *et al.*, 1997). It is also a highly potent eosinophilia inhibitor (Naito *et al.*, 1996) and is used as a fungicide (Crofton, 1996) and herbicide (Tada *et al.*, 1995). Some triazole derivatives have been evaluated for their antibacterial activity against both Gram-positive and Gram-negative bacteria (Bs *et al.*, 1996). In view of these important attributes, the structure determination of the title compound, (I), was performed.



A perspective view of (I), including the atomic numbering scheme, is shown in Fig. 1. The asymmetric unit consists of two crystallographically independent molecules. The overall r.m.s fit of these molecules is 0.233 Å. The corresponding bond distances and angles in the independent molecules agree with each other, but the two molecules differ in the relative orientations of the mean planes of the A/B and B/C rings, with dihedral angles of 3.83(11) [10.24(11)°] and 3.70(9)[14.14 (11)°], respectively. The dihedral angle between the mean planes of rings A and C is 5.32 (11)° [3.91 (10)°] (values in brackets are for molecule B). The bond lengths and angles of the triazole moiety in (I) are unexceptional, and comparable with those reported for related structures (Chen et al., 1998; Wang et al., 1998; Thamotharan, Parthasarathi, Sunagar et al., 2003; Thamotharan, Parthasarathi, Hunnur et al., 2003; Thamotharan, Parthasarathi, Kavali et al., 2003). In (I), the N4A - N12A - C13A - C14A torsion angle of $-178.3 (2)^{\circ}$ $[-177.3 (2)^{\circ}$ for molecule B] indicates that the imine group has an E configuration. The ethoxycarbonylmethoxy group is oriented at an angle of 29.42 $(15)^{\circ}$ [31.33 $(17)^{\circ}$] with respect to the mean plane of ring C.

10124 independent reflections 6462 reflections with $I > 2\sigma(I)$

 $w = 1/[\sigma^2(F_o^2) + (0.0552P)^2]$

Extinction correction: SHELXL97

Extinction coefficient: 0.0045 (6)

+ 0.6771P] where $P = (F_0^2 + 2F_c^2)/3$

 $(\Delta/\sigma)_{\rm max} = 0.002$ $\Delta \rho_{\rm max} = 0.52 \ {\rm e} \ {\rm \AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.42 \text{ e} \text{ Å}^{-3}$

 $R_{\rm int} = 0.026$

 $\theta_{\text{max}} = 29.5^{\circ}$ $h = -11 \rightarrow 10$

 $k = -20 \rightarrow 21$

 $l = -23 \rightarrow 23$



Figure 1

View of the two independent molecules of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are represented by circles of arbitrary radii.



Figure 2

The C-H···O contacts in two dimers of (I) [symmetry code: (#) -x, 2 - y, 1 - z, i.e. (ii) in Table 2].

The substituted phenyl ring at N2 of the triazole ring is almost coplanar [3.83 (11)°; 10.24 (10)° for molecule B] with the triazole ring, while the corresponding angles in the chloro (Thamotharan, Parthasarathi, Sunagar et al., 2003; Thamotharan, Parthasarathi, Kavali et al., 2003) and bromo (Thamotharan, Parthasarathi, Hunnur et al., 2003) derivatives of triazole are 30.63 (9)/4.86 (5) and $8.93 (14)^{\circ}$, respectively.

The exocyclic angles C16A-C17A-O29A, O22A-C22A-O23A and O22A-C22A-C21A deviate significantly from the normal value of 120° (Table 1); this may be due to steric repulsion $[H16A \cdots H21A = 2.16 \text{ Å} (H16B \cdots H21D =$ 2.13 Å in molecule B) and H24B···O22A = 2.41 Å $(H24D \cdots O22B = 2.37 \text{ Å in molecule } B)].$

In the crystal structure, atom C8B of molecule B acts as donor for a weak intermolecular C-H···O interaction with the carbonyl atom O22B of the centrosymmetrically related molecule at (-x, 2 - y, -z), which leads to an $R_2^2(34)$ motif. Atoms C10A and C10B are involved in weak intermolecular $C-H \cdots O$ interactions with atoms O3B and O3A of the centrosymmetrically related molecules at (-x, 2 - y, 1 - z) (Fig. 2) and each interaction has a graph-set motif of $R_2^2(14)$ (Bernstein et al., 1995).

Experimental

The title compound was prepared by heating 4-amino-2-(*p*-bromophenyl)-5-methyl-2,4-dihydro-3*H*-1,2,4-triazol-3-one with *p*-hydroxybenzaldehyde to give the corresponding hydrazone, and then further reaction with ethyl bromoacetate in the presence of dry K₂CO₃ and KI. The solid obtained, (I), was crystallized from acetone (m.p. 428–431 K). The crystal was bounded by the $\{010\}$, $\{01\overline{1}\}$ and {610} pinacoids.

Crystal data

| $C_{20}H_{19}BrN_4O_4$ | Z = 4 |
|----------------------------------|---|
| $M_r = 459.30$ | $D_x = 1.526 \text{ Mg m}^{-3}$ |
| Triclinic, P1 | Mo $K\alpha$ radiation |
| a = 8.5176 (7) Å | Cell parameters from 15908 |
| b = 15.5566 (12) Å | reflections |
| c = 17.2643 (15) Å | $\theta = 2.4-29.5^{\circ}$ |
| $\alpha = 113.407 \ (6)^{\circ}$ | $\mu = 2.09 \text{ mm}^{-1}$ |
| $\beta = 94.601 \ (7)^{\circ}$ | T = 293 (2) K |
| $\gamma = 103.763 \ (6)^{\circ}$ | Obelisk, translucent colourless |
| V = 1999.5 (3) Å ³ | $0.46 \times 0.11 \times 0.08 \text{ mm}$ |
| | |

Data collection

Stoe IPDS diffractometer φ scans Absorption correction: by integration (X-RED in IPDS; Stoe & Cie, 1997) $T_{\min} = 0.537, T_{\max} = 0.861$ 19783 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.046$ $wR(F^2) = 0.126$ S = 1.0210124 reflections 528 parameters H-atom parameters constrained

Table 1

Selected geometric parameters (Å, °).

| N1A - C5A | 1.289 (3) | N1B-C5B | 1.293 (3) |
|-----------------|-------------|-------------|-------------|
| N1A - N2A | 1.394 (3) | N1B-N2B | 1.397 (3) |
| N2A - C3A | 1.370 (3) | N2B-C3B | 1.378 (3) |
| N4A - C5A | 1.370 (3) | N4B-C5B | 1.373 (3) |
| N4A - C3A | 1.394 (3) | N4B-C3B | 1.391 (3) |
| | | | |
| C5A - N1A - N2A | 104.67 (19) | C5B-N1B-N2B | 105.1 (2) |
| C3A - N2A - N1A | 112.46 (19) | C3B-N2B-N1B | 111.83 (19) |
| C5A-N4A-C3A | 108.56 (19) | C5B-N4B-C3B | 108.7 (2) |
| N2A - C3A - N4A | 102.40 (19) | N2B-C3B-N4B | 102.8 (2) |
| N1A - C5A - N4A | 111.9 (2) | N1B-C5B-N4B | 111.6 (2) |
| | | | |

| lable | 2 | | |
|-------|------------------------------|-----|-----|
| С-Н. | $\cdot \cdot O$ interactions | (Å, | °). |

| $D - H \cdots A$ | $D-\mathrm{H}$ | $H \cdot \cdot \cdot A$ | $D \cdots A$ | $D - \mathbf{H} \cdots A$ |
|--|----------------|-------------------------|--------------|---------------------------|
| $C8B - H8B \cdot \cdot \cdot O22B^{i}$ | 0.93 | 2.54 | 3.430 (4) | 161 |
| $C10A - H10A \cdots O3B^{ii}$ | 0.93 | 2.45 | 3.371 (3) | 172 |
| $C10B - H10B \cdot \cdot \cdot O3A^{ii}$ | 0.93 | 2.39 | 3.314 (3) | 171 |

Symmetry codes: (i) -x, 2 - y, -z; (ii) -x, 2 - y, 1 - z.

Methyl H atoms were constrained to an ideal geometry (C–H = 0.96 Å), with $U_{iso}(H) = 1.5U_{eq}(C)$, but were allowed to rotate freely about the parent C–C bond. All remaining H atoms were placed in geometrically idealized positions (C–H = 0.93–0.97 Å) and constrained to ride on their parent atoms, with $U_{iso}(H) = 1.2U_{eq}(C)$.

Data collection: *IPDS* (Stoe & Cie, 1997); cell refinement: *IPDS*; data reduction: *IPDS*; program(s) used to solve structure: *DIRDIF99* (Beurskens *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP3* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97* and *PLATON* (Spek, 2003).

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