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2-(4-Bromophenyl)-5-methyl-2,3-dihydro-4 <i>H</i> -1,2,4-triazol-3-one	
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Authory Emden	
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Key indicators

Single-crystal X-ray study T = 160 KMean $\sigma(\text{C-C}) = 0.003 \text{ Å}$ R factor = 0.038 wR factor = 0.092Data-to-parameter ratio = 20.4

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

2-(4-Bromophenyl)-5-methyl-2,3-dihydro-4*H*-1,2,4-triazol-3-one

In the title compound, $C_9H_8BrN_3O$, the dihedral angle between the phenyl and triazole rings is 8.93 (14)°. In the crystal, the molecules exist as centrosymmetrically related $N-H\cdots O$ hydrogen-bonded dimers.

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Comment

1,2,4-Triazole derivatives are found to be associated with diverse pharmacological activities, such as analgesic, antiasthmatic, diuretic, antifungal, antibacterial, pesticidal and anti-inflammatory activities (Thamotharan *et al.*, 2003, and references therein). Triazole derivatives have been synthesized as possible anticonvulsants, antidepressants, tranquilizers and plant-growth regulators (Bradbury & Rivett, 1991; Hirota *et al.*, 1991; Walser *et al.*, 1991). In view of this, the crystal structure determination of the title triazole derivative, (I), has been carried out in order to elucidate the molecular conformation.

$$Br$$
 NH
 CH_3

The bond lengths and angles in (I) are unexceptional and comparable with those reported for related structures (Chen et al., 1998; Wang et al., 1998; Thamotharan et al., 2003). The phenyl ring is oriented at an angle of 8.93 (14)° with respect to the plane of the triazole ring, compared with a value of $30.63~(9)^{\circ}$ reported for our previous chloro derivative of triazole (Thamotharan et al., 2003). In the crystal, centrosymmetrically related molecules form dimeric pairs through $N-H\cdots O$ intermolecular hydrogen bonds and have a graph set motif of $R_2^2(8)$ (Bernstein et al., 1995).

Experimental

The title compound was prepared by refluxing 2-(4-bromophenyl)-5-methyl-1,3,4-oxadiazolin-2-one with formamide. The solid obtained, (I), was crystallized from benzene (m.p. 503 K).

Crystal data

 $C_9H_8BrN_3O$ $M_r = 254.09$ Monoclinic, $P2_1/c$ a = 4.0333 (1) Å b = 10.8622 (2) Å c = 21.1608 (4) Å $\beta = 93.9045$ (7)° V = 924.91 (3) Å³ Z = 4 $D_x = 1.825 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation Cell parameters from 26 468 reflections $\theta = 2.0-30.0^{\circ}$ $\mu = 4.41 \text{ mm}^{-1}$ T = 160 (2) KNeedle, colourless $0.25 \times 0.08 \times 0.05 \text{ mm}$

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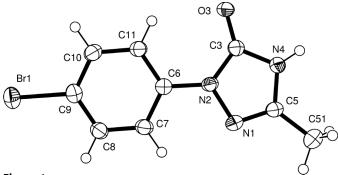


Figure 1

View of the asymmetric unit of the title compound, showing the atomlabelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are represented by circles of arbitrary radii.

Data collection

Nonius KappaCCD diffractometer φ and ω scans with κ offsets Absorption correction: multi-scan (SORTAV; Blessing, 1995) $h = -5 \rightarrow 5$ $T_{\min} = 0.550, \, T_{\max} = 0.818$ $k = -15 \rightarrow 15$ 22 923 measured reflections $l = -29 \rightarrow 29$

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0309P)^2]$ $R[F^2 > 2\sigma(F^2)] = 0.038$ + 0.9447*P*] $wR(F^2) = 0.092$ where $P = (F_o^2 + 2F_c^2)/3$ S = 1.07 $(\Delta/\sigma)_{\text{max}} = 0.001$ $\Delta \rho_{\rm max} = 0.35$ e Å 2710 reflections $\Delta \rho_{\min} = -0.75 \text{ e Å}^{-3}$ 133 parameters H atoms treated by a mixture of Extinction correction: SHELXL97 Extinction coefficient: 0.0047 (11) independent and constrained refinement

 Table 1

 Selected geometric parameters (\mathring{A} , °).

N1-C5	1.297 (3)	N4-C5	1.363 (3)
N1-N2	1.400(3)	N4-C3	1.372 (3)
N2-C3	1.372 (3)		
C5-N1-N2	103.8 (2)	N2-C3-N4	103.5 (2)
C3-N2-N1	111.8 (2)	N1-C5-N4	112.5 (2)
C5-N4-C3	108.4 (2)		

Table 2 Hydrogen-bonding geometry (Å, °).

$D-H\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D-\mathrm{H}\cdots A$
N4—H4···O3 ⁱ	0.82 (3)	1.94 (3)	2.756 (3)	171 (3)

Symmetry code: (i) -1 - x, 2 - y, -z.

The position of the amine H atom was determined from a difference Fourier map and refined freely along with its isotropic displacement parameter. The methyl H atoms were constrained to an ideal geometry (C—H = 0.98 Å), with $U_{\rm iso}({\rm H})=1.5U_{\rm eq}({\rm C})$, but were allowed to rotate freely about the C—C bond. All remaining H atoms were placed in geometrically idealized positions (C—H = 0.95–0.99 Å) and constrained to ride on their parent atoms, with $U_{\rm iso}({\rm H})=1.2U_{\rm eq}({\rm C})$.

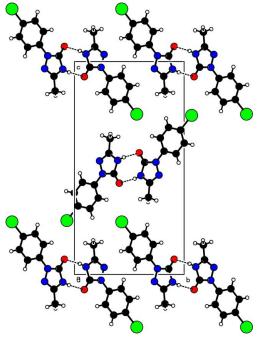


Figure 2 Packing diagram, viewed along the *a* axis..

Data collection: *COLLECT* (Nonius, 2000); cell refinement: *DENZO-SMN* (Otwinowski & Minor, 1997); data reduction: *DENZO-SMN* and *SCALEPACK* (Otwinowski & Minor, 1997); program(s) used to solve structure: *SIR*92 (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Version 1.07; Farrugia, 1997); software used to prepare material for publication: *SHELXL*97 and *PLATON* (Spek, 2002).

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