## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=293 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.004 \AA$
$R$ factor $=0.042$
$w R$ factor $=0.097$
Data-to-parameter ratio $=9.1$

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## 2-(2H-Benzotriazol-2-yl)-1-phenylethanone

The molecule of the title compound, $\mathrm{C}_{14} \mathrm{H}_{11} \mathrm{~N}_{3} \mathrm{O}$, is nonplanar, with a dihedral angle of $72.06(9)^{\circ}$ between the benzene and benzotriazole planes. Molecules are linked into two-dimensional layers via weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{N}$ and $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ interactions.

## Comment

In recent years, benzotriazoles, especially, those substituted at the 2-position of the heterocycle, have attracted special attention (Voronkov et al., 2003). A variety of benzotriazoles exhibit growth-inhibiting activities against some microorganisms and other derivatives are endowed with anti-inflammatory properties (Zelnik \& Strehlau, 1971). We report here the structure of a 2 -substituted benzotriazole compound, (I).

(I)

The bond lengths and angles are within normal ranges (Allen et al., 1987), and the bonds in the benzotriazole group show a characteristic intermediate length between single and double bonds because of the conjugated $\pi$ system (Table 1). The C9-C14 benzene and triazole rings are essentially coplanar, with a dihedral angle of $0.7(1)^{\circ}$ between these two rings, while the benzotriazole plane and the phenyl ring twist $72.06(9)^{\circ}$ from each other. In the crystal structure, molecules are linked into two-dimensional layers via $\mathrm{C}-\mathrm{H} \cdots \mathrm{N}$ and $\mathrm{C}-$ $\mathrm{H} \cdots \mathrm{O}$ interactions (Table 2 and Fig. 2).

## Experimental

To a solution of 2-(2H-bromine-2-yl)-1-phenylethanone $(20 \mathrm{~g}$, 0.1 mol ) in acetone ( 80 ml ) was added 1,2,3-benzotriazole ( 11.9 g , 0.1 mol ). The mixture was heated under reflux for 1 h , yielding a copious precipitate. Colorless single crystals suitable for X-ray diffraction were obtained by slow evaporation of an ethyl acetate and petroleum ether $(1: 1 \mathrm{v} / \mathrm{v})$ solution over a period of one week.

## Crystal data

$$
\begin{aligned}
& \mathrm{C}_{14} \mathrm{H}_{11} \mathrm{~N}_{3} \mathrm{O} \\
& M_{r}=237.26 \\
& \text { Tetragonal, } P 4_{3} 2_{1} 2 \\
& a=8.1925(3) \AA \\
& c=35.46(3) \AA \\
& V=2380.1(2) \AA \AA^{3} \\
& Z=8 \\
& D_{x}=1.324 \mathrm{Mg} \mathrm{~m}^{-3}
\end{aligned}
$$

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Figure 1
The structure of (I), showing 50\% probability displacement ellipsoids and the atom-numbering scheme.

## Data collection

Siemens SMART 1000 CCD areadetector diffractometer $\omega$ scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
$T_{\text {min }}=0.970, T_{\text {max }}=0.993$
14062 measured reflections

1477 independent reflections
1277 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.039$
$\theta_{\text {max }}=26.0^{\circ}$
$h=-10 \rightarrow 9$
$k=-10 \rightarrow 10$
$l=-43 \rightarrow 30$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.042$
$w R\left(F^{2}\right)=0.098$
$S=1.20$
1477 reflections
163 parameters
H -atom parameters constrained

$$
\begin{aligned}
& \begin{array}{c}
w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.0436 P)^{2}\right. \\
\quad+0.1331 P] \\
\text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
(\Delta / \sigma)_{\max }<0.001 \\
\Delta \rho_{\max }=0.14 \mathrm{e}^{-3} \AA^{-3} \\
\Delta \rho_{\min }=
\end{array}-0.12 \mathrm{e}^{-3}
\end{aligned}
$$

Table 1
Selected bond lengths ( $\AA$ ).

| $\mathrm{O} 1-\mathrm{C} 7$ | $1.196(3)$ | $\mathrm{N} 2-\mathrm{N} 3$ | $1.323(2)$ |
| :--- | :--- | :--- | :--- |
| $\mathrm{N} 1-\mathrm{N} 2$ | $1.319(2)$ | $\mathrm{N} 2-\mathrm{C} 8$ | $1.449(3)$ |
| $\mathrm{N} 1-\mathrm{C} 9$ | $1.348(3)$ | $\mathrm{N} 3-\mathrm{C} 14$ | $1.347(3)$ |

Table 2
Hydrogen-bond geometry $\left({ }^{\circ},{ }^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{C} 8-\mathrm{H} 8 A \cdots \mathrm{~N}^{\mathrm{i}}$ | 0.97 | 2.54 | $3.471(3)$ | 160 |
| $\mathrm{C} 11-\mathrm{H} 11 A \cdots \mathrm{O}^{\mathrm{ii}}$ | 0.93 | 2.52 | $3.439(3)$ | 170 |
| Symmetry codes: (i) $x+\frac{1}{2},-y+\frac{1}{2},-z+\frac{1}{4}$; (ii) $x+\frac{1}{2},-y+\frac{3}{2},-z+\frac{1}{4}$ |  |  |  |  |

All H atoms were located in a difference Fourier map and constrained to ride on their parent atoms, with $\mathrm{C}-\mathrm{H}=0.93$ or $0.97 \AA$, and with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$. The Friedel pairs were merged before


Figure 2
A packing view down the $c$ axis. Hydrogen bonds are indicated by dashed lines.
the final refinement because of the absence of significant anomalous scattering effects.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 1997); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL, PARST (Nardelli, 1995) and PLATON (Spek, 2003).

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