Acta Crystallographica Section E

## Structure Reports

Online
ISSN 1600-5368

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## Key indicators

Single-crystal X-ray study
$T=293 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.046$
$w R$ factor $=0.116$
Data-to-parameter ratio $=14.8$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

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## 2-(1H-1,2,3-Benzotriazol-1-ylmethyl)-1-benzoylethyl 4-chlorobenzoate

In the title compound, $\mathrm{C}_{22} \mathrm{H}_{16} \mathrm{ClN}_{3} \mathrm{O}_{3}$, molecules are linked into chains along the $c$ axis by $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonds, while other $\mathrm{C}-\mathrm{H} \cdots \mathrm{N}$ hydrogen bonds connect the chains into two-dimensional layers. The packing is further stabilized by $\pi-$ $\pi$ interactions.

## Comment

Benzotriazole derivatives have been found to exhibit a broad spectrum of pharmacological activities such as anti-inflammatory, antifungal, antitumor, antineoplastic and antidepressant activities (Al-Soud et al., 2003; Katarzyna et al., 2004). As part of a search for new benzotriazole compounds with high bioactivity, the title compound, (I), was synthesized.


The bond lengths and angles in (I) show normal values (Allen et al., 1987) and are comparable to those in the related compound 2-(1H-1,2,3-benzotriazol-1-ylmethyl)-1-benzoylethyl 4-ethylbenzoate (Wan et al., 2005). The benzotriazole system is essentially planar, with a dihedral angle of $1.9(1)^{\circ}$ between the triazole ring ( $A$, atoms $\mathrm{N} 1-\mathrm{N} 3 / \mathrm{C} 10 / \mathrm{C} 15$ ) and the benzene ring ( $B$, atoms $\mathrm{C} 10-\mathrm{C} 15$ ). The dihedral angles between the mean planes of the benzotriazole system and rings $C$ (atoms C1-C6) and $D$ (atoms C17-C22) are 38.1 (1) and $53.7(1)^{\circ}$, respectively. The dihedral angle between rings $C$ and $D$ is $88.9(1)^{\circ}$.

In the crystal structure, molecules of (I) are linked into chains along the $c$ axis by $\mathrm{C} 9-\mathrm{H} 9 A \cdots \mathrm{O} 1^{\mathrm{ii}}$ and $\mathrm{C} 11-$ $\mathrm{H} 11 A \cdots \mathrm{O} 1^{\text {ii }}$ hydrogen bonds (see Table 2 for geometry values and symmetry codes). In addition, $\mathrm{C} 3-\mathrm{H} 3 B \cdots \mathrm{~N} 3^{i}$ interactions connect the chains into two-dimensional layers in the ac plane (Fig. 2). The packing is further stabilized by $\pi-\pi$ interactions involving the benzotriazole system and benzene ring $D$, the distances being $C g 1 \cdots C g 4^{4 i i}=3.570 \AA$ and $C g 3 \cdots C g 4^{\mathrm{iii}}=3.654 \AA[C g 1, C g 3$ and $C g 4$ denote the centroids of the rings $A, B$ and $D$, respectively; symmetry code: (iii) $x, \frac{3}{2}$ $\left.-y, z-\frac{1}{2}\right]$.


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

## Experimental

Bromine ( $3.2 \mathrm{~g}, 0.02 \mathrm{~mol}$ ) was added dropwise to a solution of 3-(benzotriazol-1-yl)-1-phenylpropan-1-one $\quad(5.0 \mathrm{~g}, \quad 0.02 \mathrm{~mol})$ and sodium acetate $(1.6 \mathrm{~g}, 0.02 \mathrm{~mol})$ in acetic acid $(50 \mathrm{ml})$. The reaction was allowed to proceed for 13 h . Water $(50 \mathrm{ml})$ and chloroform ( 20 ml ) were then added. The organic layer was washed successively with saturated sodium bicarbonate solution and brine, dried over anhydrous magnesium sulfate and the chloroform solution filtered. It was cooled with ice-water and then an acetone solution $(10 \mathrm{ml})$ of 4 chlorobenzoic acid $(3.1 \mathrm{~g}, 0.02 \mathrm{~mol})$ and triethylamine $(2.8 \mathrm{ml})$ were added. The mixture was stirred at room temperature for about 2 h . The solution was then filtered and concentrated (yield 62\%). Crystals of (I) were obtained by slow evaporation of an acetone-water (1:1 $\mathrm{v} /$ $v)$ solution at room temperature over a period of one week.

## Crystal data

## $\mathrm{C}_{22} \mathrm{H}_{16} \mathrm{ClN}_{3} \mathrm{O}_{3}$

$M_{r}=405.83$
Monoclinic, $P 2_{1} / c$
$a=9.712$ (3) $\AA$
$b=20.379$ (5) $\AA$
$c=10.009(3) \AA$
$\beta=90.392$ (4) ${ }^{\circ}$
$V=1980.9(9) \AA^{3}$
$Z=4$
$D_{x}=1.361 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
Cell parameters from 2380
$\quad$ reflections
$\theta=2.3-22.3^{\circ}$
$\mu=0.22 \mathrm{~mm}^{-1}$
$T=293(2) \mathrm{K}$
Column, colorless
$0.28 \times 0.11 \times 0.09 \mathrm{~mm}$

## Data collection

Siemens SMART 1000 CCD area-
detector diffractometer

## $\omega$ scans

Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
$T_{\text {min }}=0.941, T_{\text {max }}=0.967$
11133 measured reflections

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.046$
$w R\left(F^{2}\right)=0.116$
$S=1.02$
3885 reflections
262 parameters
H -atom parameters constrained


Figure 2
The two-dimensional layers of (I), viewed down the $b$ axis. Dashed lines indicate $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}, \mathrm{N}$ interactions.

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

| $\mathrm{Cl} 1-\mathrm{C} 20$ | $1.743(2)$ | $\mathrm{O} 3-\mathrm{C} 16$ | $1.202(2)$ |
| :--- | :--- | :--- | :--- |
| $\mathrm{O} 1-\mathrm{C} 7$ | $1.218(2)$ | $\mathrm{C} 7-\mathrm{C} 8$ | $1.527(3)$ |
| $\mathrm{O} 2-\mathrm{C} 16$ | $1.357(2)$ | $\mathrm{C} 8-\mathrm{C} 9$ | $1.514(3)$ |
| $\mathrm{O} 2-\mathrm{C} 8$ | $1.436(2)$ |  |  |

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

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{C} 3-\mathrm{H} 3 B \cdots \mathrm{~N}^{\mathrm{i}}$ | 0.93 | 2.58 | $3.506(4)$ | 175 |
| $\mathrm{C}^{\mathrm{ii}}-\mathrm{H} 9 A \cdots \mathrm{O}^{\mathrm{ii}}$ | 0.97 | 2.49 | $3.453(3)$ | 173 |
| ${\mathrm{C} 11-\mathrm{H} 11 A \cdots \mathrm{O}^{1 i}}^{\mathrm{ii}}$ | 0.93 | 2.52 | $3.316(3)$ | 143 |

Symmetry codes: (i) $x+1,-y+\frac{3}{2}, z-\frac{1}{2}$; (ii) $x,-y+\frac{3}{2}, z-\frac{1}{2}$.

All H atoms were located in difference Fourier maps and constrained to ride on their parent atoms, with $\mathrm{C}-\mathrm{H}$ distances in the range $0.93-0.98 \AA$ and with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$.

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).

This study was supported by the Project of Educational Administration of Shandong Province (No. J04B12) and the Outstanding Adult-Young Scientific Research Encouraging Foundation of Shandong Province (No. 2005BS04007).

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