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Key indicators

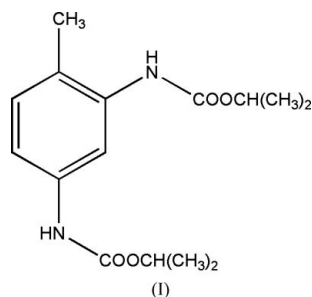
Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.040
 wR factor = 0.104
Data-to-parameter ratio = 16.6For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Diisopropyl *N,N'*-(4-methyl-*m*-phenylene)-
dicarbamateThe crystal structure of the title compound, $\text{C}_{15}\text{H}_{22}\text{N}_2\text{O}_4$, is
stabilized by two intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds.

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Comment

The title compound, (I), is a derivative of toluene-2,4-bis-
(isocyanate), prepared by the reaction of 4-methylbenzene-
1,3-diamine and isopropyl carbonochloridate according to the
method of Baskakow & Mel'nikow (1954). In the present
paper, the crystal structure of (I) (Fig. 1) is reported. The
crystal packing projected on to the bc face is shown in Fig. 2.
Selected geometric parameters are given in Table 1. In the
crystal structure, there are two intermolecular $\text{N}-\text{H}\cdots\text{O}$
hydrogen bonds (Table 2).

Experimental

Toluene-2,4-bis(isocyanate) (TDI) was provided by Hebei Cangzhou
Dahua Co. Ltd. The raw material was purified twice by a melt crys-
tallization process. Compound (I) was prepared by the reaction of
purified TDI (98%) with isopropyl alcohol (m.p. 408.7 K; differential
scanning calorimetry). Colourless block-shaped single crystals
suitable for X-ray diffraction were obtained by slow evaporation of a
methanol solution at room temperature over a period of 5 d.

Crystal data

 $\text{C}_{15}\text{H}_{22}\text{N}_2\text{O}_4$
 $M_r = 294.35$
Monoclinic, Cc
 $a = 9.0708$ (18) Å
 $b = 15.997$ (6) Å
 $c = 12.113$ (2) Å
 $\beta = 111.28$ (3)°
 $V = 1637.8$ (7) Å³
 $Z = 4$ $D_x = 1.194$ Mg m⁻³
Mo $K\alpha$ radiation
Cell parameters from 8078
reflections
 $\theta = 3.1$ – 27.5 °
 $\mu = 0.09$ mm⁻¹
 $T = 293$ (2) K
Block, colourless
 $0.31 \times 0.24 \times 0.14$ mm

Data collection

Rigaku R-AXIS RAPID IP area-
detector diffractometer
 ω scans
Absorption correction: multi-scan
(*ABSCOR*; Higashi, 1995)
 $T_{\min} = 0.973$, $T_{\max} = 0.988$
7930 measured reflections3520 independent reflections
2978 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.038$
 $\theta_{\text{max}} = 27.5$ °
 $h = -11 \rightarrow 11$
 $k = -20 \rightarrow 18$
 $l = -15 \rightarrow 15$

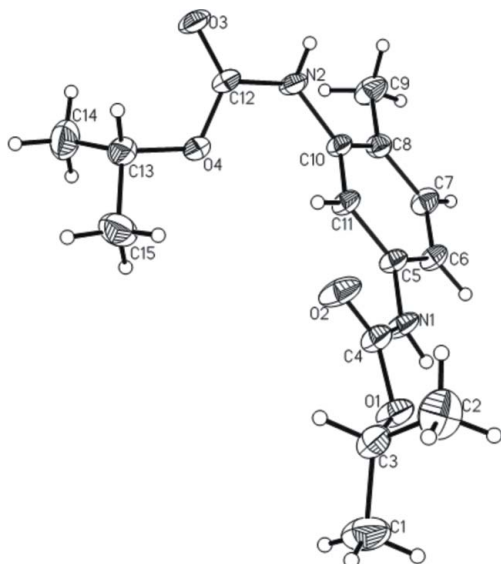


Figure 1
An *ORTEP* (Johnson, 1976) view of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level.

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.104$
 $S = 1.03$
 3520 reflections
 212 parameters
 H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0584P)^2 + 0.1074P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.008$
 $\Delta\rho_{\max} = 0.13 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.16 \text{ e } \text{\AA}^{-3}$

Table 1
Selected geometric parameters (\AA , $^\circ$).

N1—C4	1.343 (3)	N2—C12	1.348 (2)
N1—C5	1.406 (2)	N2—C10	1.425 (2)
C4—N1—C5	127.87 (15)	C12—N2—C10	128.96 (16)
C3—O1—C4—O2	−3.6 (3)	C13—O4—C12—O3	−4.5 (3)
C3—O1—C4—N1	177.11 (17)	C13—O4—C12—N2	176.84 (16)
C8—C10—N2—C12	145.9 (2)	C6—C5—N1—C4	168.6 (2)

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1A \cdots O3 ⁱ	0.86	2.03	2.873 (2)	166
N2—H2A \cdots O2 ⁱⁱ	0.86	2.13	2.975 (2)	168

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

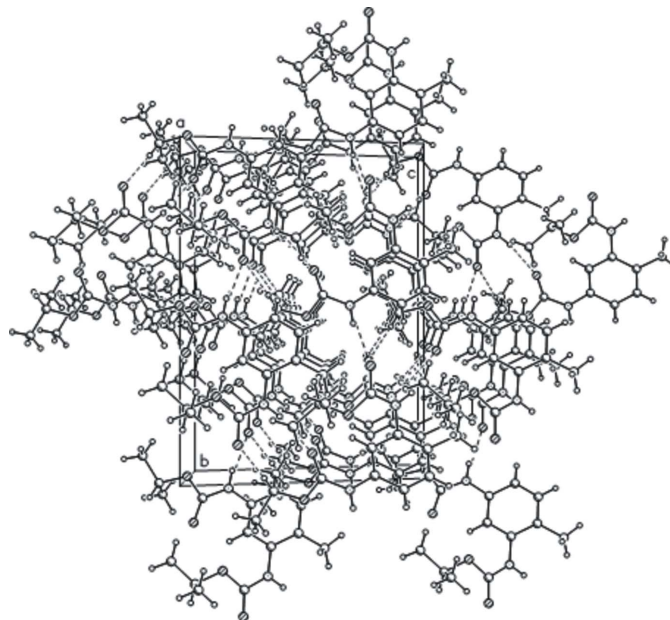


Figure 2
The molecular packing of compound (I), viewed along the a axis. Dashed lines indicate hydrogen bonds.

H atoms were placed in calculated positions and constrained to ride on their parent atoms, with C—H = 0.93–0.96 \AA and N—H = 0.86 \AA , and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$ and $1.5U_{\text{eq}}(\text{methyl C})$.

Data collection: *RAPID-AUTO* (Rigaku, 2004); cell refinement: *RAPID-AUTO*; data reduction: *RAPID-AUTO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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