Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.002 Å R factor = 0.042 wR factor = 0.105 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.

N,N,N'N'-Tetraphenyl-2,2'-(*o*-phenylenedioxy)diacetamide

In the title compound, $C_{34}H_{28}N_2O_4$, the molecule lies on a crystallographic twofold axis. In the crystal structure, the molecules are linked by $C-H\cdots O$ hydrogen bonds into double chains along the *c* axis. These chains are interlinked *via* $C-H\cdots \pi$ interactions.

Received 14 September 2005 Accepted 19 September 2005 Online 21 September 2005

Comment

In our ongoing study of amide-type polyethers, we have synthesized the title compound, (I). Here, we report its crystal structure.



The asymmetric unit of (I) contains one half-molecule, the other half being related by a crystallographic twofold axis (Fig. 1). Bond lengths in (I) (Table 1) are within normal ranges (Allen *et al.*, 1987). The sum of the angles around atom N1 is 359.9°, implying a planar configuration. The dihedral angle between the two benzene rings attached to atom N1 is



Figure 1

The structure of (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme. Unlabelled atoms are related to labelled atoms by the symmetry operator $(-x, y, \frac{1}{2} - z)$.

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 $83.91 (5)^{\circ}$, while these two rings (C1–C6 and C7–C12) make angles of 62.68 (5) and 54.89 (6)°, respectively, with the central benzene ring.

In the crystal structure, molecules of (I) are linked into double chains along the *c* axis (Fig. 2) *via* C11-H11A···O1ⁱ hydrogen bonds (symmetry code as in Table 2). The packing is further stabilized by C-H··· π interactions involving the C1-C6 benzene ring (centroid Cg1).

Experimental

Compound (I) was prepared according to the literature method of Wen *et al.* (2005). Yellow single crystals suitable for an X-ray diffraction study were obtained by slow evaporation of a petroleum ether–ethyl acetate solution $(1:3 \nu/\nu)$ over a period of 8 d.

 $D_r = 1.290 \text{ Mg m}^{-3}$

Cell parameters from 2861

 $0.32 \times 0.19 \times 0.08 \ \text{mm}$

Mo $K\alpha$ radiation

reflections

 $\begin{array}{l} \theta = 2.4 - 26.0^{\circ} \\ \mu = 0.09 \ \mathrm{mm}^{-1} \end{array}$

T = 293 (2) K

Plate, yellow

Crystal data

 $\begin{array}{l} C_{34}H_{28}N_2O_4\\ M_r = 528.58\\ \text{Monoclinic, } C2/c\\ a = 16.0863 (10) \text{ Å}\\ b = 11.0050 (7) \text{ Å}\\ c = 16.5816 (11) \text{ Å}\\ \beta = 111.981 (1)^\circ\\ V = 2722.1 (3) \text{ Å}^3\\ Z = 4 \end{array}$

Data collection

Siemens SMART 1000 CCD area-	2674 independent reflections
detector diffractometer	2284 reflections with $I > 2\sigma(I)$
ω scans	$R_{\rm int} = 0.016$
Absorption correction: multi-scan	$\theta_{\rm max} = 26.0^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -19 \rightarrow 19$
$T_{\min} = 0.973, T_{\max} = 0.993$	$k = -13 \rightarrow 10$
7490 measured reflections	$l = -20 \rightarrow 20$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0486P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.042$	+ 1.2205P]
$wR(F^2) = 0.106$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.03	$(\Delta/\sigma)_{\rm max} = 0.001$
2674 reflections	$\Delta \rho_{\rm max} = 0.15 \ {\rm e} \ {\rm \AA}^{-3}$
181 parameters	$\Delta \rho_{\rm min} = -0.19 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table 1

Selected	geometric	parameters	(Å,	°).
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O1-C13	1.2157 (16)	N1-C13	1.3635 (17)
O2-C15	1.3740 (15)	N1-C12	1.4404 (17)
O2-C14	1.4228 (16)		
C13-N1-C12	124.88 (11)	C12-N1-C1	116.82 (10)
C13-N1-C1	118.21 (11)		



Figure 2

The crystal packing of (I), showing a hydrogen-bonded (dashed lines) double chain.

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C11 - H11A \cdots O1^{i}$ $C17 - H17A \cdots Cg1^{ii}$	0.93 0.93	2.41 2.70	3.308 (2) 3.563 (2)	162 154

Symmetry codes: (i) -x, -y + 2, -z + 1; (ii) -x, -y + 1, -z + 1. Cg1 is the centroid of ring C1–C6.

All H atoms were positioned geometrically and constrained to ride on their parent atoms, with C-H distances in the range 0.93–0.97 Å, and with $U_{iso}(H) = 1.2U_{eq}(C)$.

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 project was supported by the Programme for New Century Excellent Talents in Universities (grant No. NCET-04–0649) and the Project of Educational Administration of Shandong Province (grant No. J04B12).

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Wen, Y.-H., Zhang, S.-S., Liang, J. & Li, X.-M. (2005). Acta Cryst. E61, 096–097.