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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.002 Å R factor = 0.045 wR factor = 0.133 Data-to-parameter ratio = 15.3

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

N,N'-Bis(4-methylphenyl)-3-oxapentanediamide

The title compound, $C_{18}H_{20}N_2O_3$, has twofold rotation symmetry. Molecules are linked into ribbons along the *c* axis by $N-H\cdots O$ and $C-H\cdots O$ hydrogen bonds.

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Comment

We have recently reported the structure of N,N'-bis(4methoxyphenyl)-3-oxapentanediamide, (II), (Wen *et al.*, 2006). In our ongoing studies of amide-type acyclic polyethers, the title compound, (I), has been prepared.



In the molecule of (I), a crystallographic twofold rotation axis passes through O2. All bond lengths and angles are within normal ranges (Allen *et al.*, 1987), and comparable to those in the compound (II). There exists one intramolecular hydrogen bond, C6–H6A···O1 (Table 2), forming a six-membered ring. In the crystal structure, molecules are linked into ribbons along the *c* axis (Fig. 2) by N1–H1A···O1ⁱ and C4– H4A···O1ⁱ intermolecular hydrogen bonds (symmetry code as in Table 2).

Experimental

 $SOCl_2$ (5.0 ml, 0.08 mol) was slowly added to a solution of oxydiacetic acid (2.68 g, 0.02 mol) in benzene (20 ml). After stirring for 3 h at



ellipsoids and the atom numbering scheme. The suffix A corresponds to

Figure 1 The molecular structure of (I), showing 50% probability displacement

symmetry code $(1 - x, y, \frac{1}{2} - z)$

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Figure 2

A view down the b axis, showing the ribbons. Hydrogen bonds are indicated by dashed lines.

343 K, the mixture turned clear and was stirred for a further 2 h. Benzene and excess SOCl₂ were then removed under reduced pressure to give oxydiacetic acid dichloride. This compound (1.71 g, 0.01 mol) in benzene (20 ml) was added dropwise to a solution of *p*-methylaniline (2.14 g, 0.02 mol) and pyridine (2 ml) in benzene (40 ml), and the mixture was stirred at 343 K for 12 h. After cooling to room temperature, the mixture was washed three times with water and then filtered. The title compound was recrystallized from benzene as a light-brown powder. Dark-brown single crystals suitable for X-ray diffraction study were obtained by slow evaporation of an ethanol–ethyl acetate (1:16, v/v) solution over a period of one month.

Crystal data

 $\begin{array}{l} C_{18}H_{20}N_2O_3\\ M_r=312.36\\ Monoclinic, C2/c\\ a=29.059 (3) Å\\ b=6.2798 (7) Å\\ c=8.9209 (10) Å\\ \beta=90.299 (2)^\circ\\ V=1627.9 (3) Å^3 \end{array}$

Data collection

Siemens SMART 1000 CCD areadetector diffractometer ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{\min} = 0.972, T_{\max} = 0.990$ Z = 4 D_x = 1.274 Mg m⁻³ Mo K α radiation μ = 0.09 mm⁻¹ T = 293 (2) K Column, dark-brown 0.33 × 0.24 × 0.11 mm

4509 measured reflections 1606 independent reflections 1278 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.018$ $\theta_{\text{max}} = 26.0^{\circ}$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0709P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.045$	+ 0.5191P]
$vR(F^2) = 0.133$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.04	$(\Delta/\sigma)_{\rm max} < 0.001$
606 reflections	$\Delta \rho_{\rm max} = 0.17 \text{ e } \text{\AA}^{-3}$
05 parameters	$\Delta \rho_{\rm min} = -0.15 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table 1

Selected	bond	lengths	(A)).

01-C8	1.2216 (18)	N1-C8	1.346 (2)
O2-C9	1.4059 (16)	N1-C5	1.4166 (19)

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

, e .

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N1 - H1A \cdots O1^{i}$ $C4 - H4A \cdots O1^{i}$	0.86	2.32	3.131 (2) 3.298 (2)	158 138
$C6-H6A\cdots O1$	0.93	2.40	2.920 (2)	115
	-			

Symmetry code: (i) $x, -y, z - \frac{1}{2}$.

All H atoms were located in a difference Fourier map and constrained to ride on their parent atoms, with C-H = 0.93-0.97 Å and N-H = 0.86 Å, and with $U_{iso}(H) = 1.2U_{eq}(C,N)$ [$U_{iso}(H) = 1.5U_{eq}(C)$ for methyl H atoms].

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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