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Key indicators

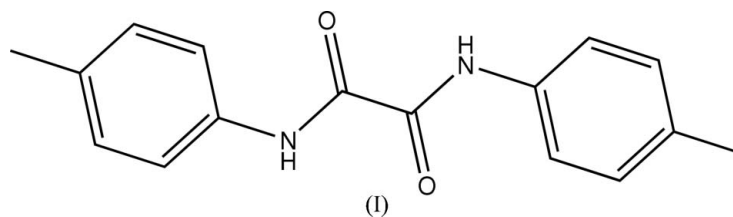
Single-crystal X-ray study
 $T = 293\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$
 R factor = 0.053
 wR factor = 0.155
Data-to-parameter ratio = 14.3For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.*N,N'*-Bis(2-methylphenyl)oxamide

The molecule of the title compound, $\text{C}_{16}\text{H}_{16}\text{N}_2\text{O}_2$, lies on a crystallographically imposed center of symmetry at the midpoint of the C—C bond of the oxamide unit. An intramolecular C—H \cdots O hydrogen bond forms a six-membered ring, and molecules are linked into ribbons along the a axis by N—H \cdots O hydrogen bonds.

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Comment

We have recently reported the synthesis and crystal structure of *N,N'*-diphenyloxalamide, (II) (Wen *et al.*, 2006). In our ongoing studies of oxalamide derivatives, the title compound, (I), was obtained.



The title compound has a crystallographically imposed center of symmetry at the mid-point of the C—C bond of the oxamide unit. Bond lengths and angles in (I) are within normal ranges (Allen *et al.*, 1987) and comparable to those found in (II). The dihedral angle between the benzene ring and the $\text{N1}/\text{C7}/\text{O1}/\text{N1}'/\text{C7}'/\text{O1}'$ plane [symmetry code: (i) $-x, 1 - y, -z$] is $24.5(1)^\circ$. An intramolecular C5—H5A \cdots O1 hydrogen bond forms a six-membered ring, and molecules are linked into ribbons along the a axis (Fig. 2) by N—H \cdots O hydrogen bonds (Table 1).

Experimental

To a solution of 2-ethoxyaniline (21.8 g, 0.2 mol) in benzene (70 ml) was added dropwise a solution of oxalyl chloride (6.4 g, 0.05 mol) in benzene (30 ml), and the mixture was stirred at 343 K for 7 h. After

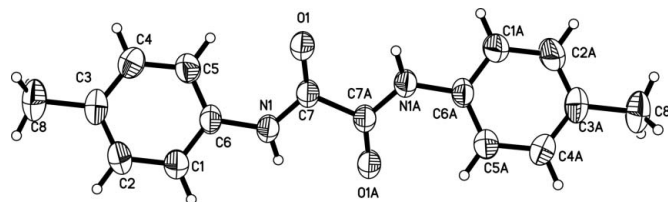


Figure 1

The structure of compound (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme. [Symmetry code: (A) $-x, 1 - y, -z$.]

cooling to room temperature, water (50 ml) was added to the reaction and the organic phase was washed 3 times with water to give (I) as a solid. The title compound was obtained after drying at room temperature for 3 d. Colorless single crystals suitable for X-ray diffraction study were obtained by slow evaporation of an *N,N*-dimethylformamide (DMF) solution over a period of 9 h.

Crystal data

$C_{16}H_{16}N_2O_2$	$V = 343.83 (15) \text{ \AA}^3$
$M_r = 268.31$	$Z = 1$
Triclinic, $P\bar{1}$	$D_x = 1.296 \text{ Mg m}^{-3}$
$a = 5.3504 (13) \text{ \AA}$	Mo $K\alpha$ radiation
$b = 6.7371 (17) \text{ \AA}$	$\mu = 0.09 \text{ mm}^{-1}$
$c = 10.228 (3) \text{ \AA}$	$T = 293 (2) \text{ K}$
$\alpha = 99.582 (4)^\circ$	Plate, colorless
$\beta = 96.152 (4)^\circ$	$0.26 \times 0.18 \times 0.07 \text{ mm}$
$\gamma = 106.527 (4)^\circ$	

Data collection

Siemens SMART 1000 CCD area-detector diffractometer	1934 measured reflections
ω scans	1311 independent reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	1111 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.978$, $T_{\max} = 0.994$	$R_{\text{int}} = 0.014$
	$\theta_{\max} = 26.0^\circ$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0866P)^2 + 0.0923P]$
$R[F^2 > 2\sigma(F^2)] = 0.053$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.156$	$(\Delta/\sigma)_{\max} < 0.001$
$S = 1.03$	$\Delta\rho_{\max} = 0.23 \text{ e \AA}^{-3}$
1311 reflections	$\Delta\rho_{\min} = -0.17 \text{ e \AA}^{-3}$
92 parameters	Extinction correction: <i>SHELXL97</i>
H-atom parameters constrained	Extinction coefficient: 0.14 (3)

Table 1

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N1-H1B\cdots O1^{ii}$	0.86	2.39	3.171 (3)	151
$N1-H1B\cdots O1^i$	0.86	2.26	2.676 (2)	110
$C5-H5A\cdots O1$	0.93	2.42	2.953 (3)	116

Symmetry codes: (i) $-x, -y + 1, -z$; (ii) $x + 1, y, z$.

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

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

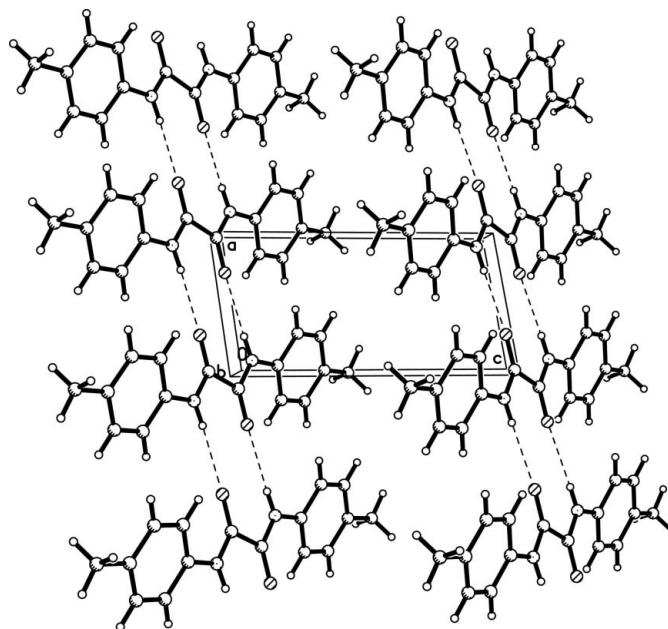


Figure 2

A view down the b axis, showing the ribbons generated by hydrogen bonds (indicated by dashed lines).

structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*, *PARST* (Nardelli, 1995) and *PLATON* (Spek, 2003).

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