

Yong-Hong Wen, Xue-Mei Li,
Long Wang and Shu-Sheng
Zhang*College of Chemistry and Molecular
Engineering, Qingdao University of Science and
Technology, 266042 Qingdao, Shandong,
People's Republic of China

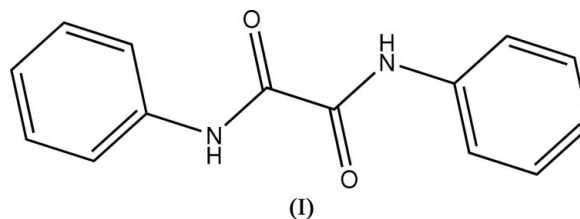
Correspondence e-mail: shushzhang@126.com

Key indicatorsSingle-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.002$ Å
 R factor = 0.037
 wR factor = 0.103
Data-to-parameter ratio = 13.7For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.***N,N'*-Diphenyloxalamide**

The molecule of the title compound, $\text{C}_{14}\text{H}_{12}\text{N}_2\text{O}_2$, lies on a crystallographically imposed center of symmetry at the mid-point of the C—C bond of the oxalamide unit. Molecules are linked into ribbons along the a axis by $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds.

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N,N'-Diphenyloxalamide and its derivatives have been widely applied in a number of materials as anti-oxidants, ultraviolet absorbents and/or metal ion passivators (Feng *et al.*, 1997). We report here the synthesis and structure of *N,N'*-diphenyloxalamide, (I) (Fig. 1 and Table 1).



The title compound has a crystallographically imposed center of symmetry at the mid-point of the C—C bond of the oxalamide unit. The bond lengths in the oxalamide unit show intermediate values due to π conjugation effects arising from the presence of the two C=O double bonds. All other bond lengths and angles are within normal ranges (Allen *et al.*, 1987). The molecule deviates somewhat from planarity, with the two O atoms twisted to opposite sides of the molecular plane, with a maximum deviation for O1 from the mean plane through C1—C7/N1 of 0.254 (4) Å. Intramolecular hydrogen bonds (C5—H5 \cdots O1) form six-membered rings between the phenyl rings and the carbonyl O atoms. In the crystal structure, molecules are linked into ribbons along the a axis (Fig. 2) by inversion-related intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds (Table 2).

Experimental

To a solution of aniline (16.6 g, 0.2 mol) in benzene (90 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 6 h. After cooling to room temperature, 50 ml water was added and the organic phase was washed three times with water to give a white solid. The title compound was obtained after drying at room temperature for 48 h. Colorless single crystals suitable for X-ray diffraction were obtained by slow evaporation of an *N,N*-dimethylformamide solution over a period of 6 h.

Crystal data

C₁₄H₁₂N₂O₂
M_r = 240.26
 Monoclinic, *P*2₁/*c*
a = 5.3207 (7) Å
b = 5.3723 (7) Å
c = 20.5227 (19) Å
 β = 101.437 (3)°
V = 574.98 (12) Å³

Z = 2
D_x = 1.388 Mg m⁻³
 Mo *K*α radiation
 μ = 0.10 mm⁻¹
T = 293 (2) K
 Block, colorless
 0.43 × 0.21 × 0.20 mm

Data collection

Siemens SMART 1000 CCD area detector diffractometer
 ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
T_{min} = 0.960, *T_{max}* = 0.981

3076 measured reflections
 1133 independent reflections
 1039 reflections with *I* > 2σ(*I*)
R_{int} = 0.019
 θ_{max} = 26.0°

Refinement

Refinement on *F*²
R [*F*² > 2σ(*F*²)] = 0.037
wR (*F*²) = 0.103
S = 1.08
 1133 reflections
 83 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0546P)^2 + 0.089P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} < 0.001$
 $\Delta\rho_{max} = 0.18 \text{ e \AA}^{-3}$
 $\Delta\rho_{min} = -0.16 \text{ e \AA}^{-3}$
 Extinction correction: SHELXL97
 Extinction coefficient: 0.203 (17)

Table 1

Selected geometric parameters (Å, °).

O1—C7	1.2209 (14)	N1—C6	1.4182 (15)
N1—C7	1.3366 (16)	C7—C7 ⁱ	1.543 (2)
C7—N1—C6	127.27 (10)	O1—C7—C7 ⁱ	121.16 (13)
O1—C7—N1	126.95 (11)	N1—C7—C7 ⁱ	111.89 (12)

Symmetry code: (i) $-x + 2, -y - 2, -z$.

Table 2

Hydrogen-bond geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1B...O1 ⁱⁱ	0.86	2.38	3.161 (1)	152
N1—H1B...O1 ⁱ	0.86	2.26	2.680 (1)	110
C5—H5A...O1	0.93	2.39	2.925 (2)	116

Symmetry codes: (i) $-x + 2, -y - 2, -z$; (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 Å, N—H = 0.86 Å and *U*_{iso}(H) = 1.2*U*_{eq}(C,N).

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).

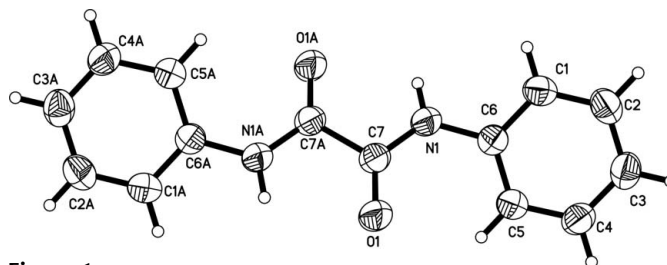


Figure 1

The structure of compound (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme. Atoms labelled with the suffix 'A' are related to the other atoms by the symmetry code (2 - *x*, -2 - *y*, -*z*), corresponding to (i) in Table 1.

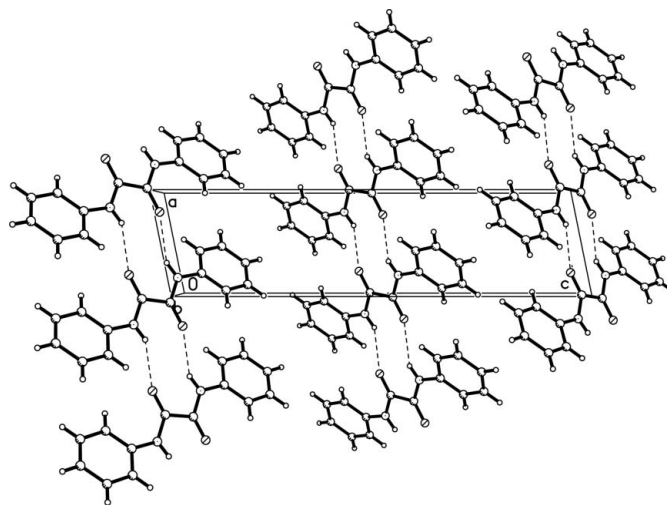


Figure 2

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

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