

***N,N'*-Bis(3,5-dimethoxyphenyl)oxamide****Wen Gu<sup>a\*</sup> and Qing-Gang Tang<sup>b</sup>**<sup>a</sup>Institute of Functional Biomolecules, State Key Laboratory of Pharmaceutical Biotechnology, Nanjing University, Nanjing 210093, People's Republic of China, and <sup>b</sup>Department of Applied Chemistry, College of Science, Nanjing<sup>U</sup>niversity of Technology, Xinmofan Road No. 5 Nanjing, Nanjing 210009, People's Republic of China

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**Key indicators**

Single-crystal X-ray study

 $T = 293$  KMean  $\sigma(\text{C}-\text{C}) = 0.005$  Å $R$  factor = 0.069 $wR$  factor = 0.152

Data-to-parameter ratio = 14.1

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

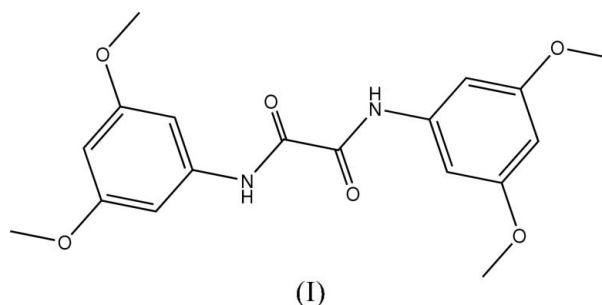
The molecule of the title compound,  $\text{C}_{18}\text{H}_{20}\text{N}_2\text{O}_6$ , lies on a crystallographically imposed center of symmetry at the mid-point of the C—C bond of the oxamide unit. The crystal structure is stabilized by  $\pi$ – $\pi$  interactions.

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**Comment**

*N,N'*-Diphenyloxamide and its derivatives have been widely applied in a number of materials as anti-oxidants, ultraviolet absorbents and metal ion passivators (Feng *et al.*, 1997). We report here the synthesis and crystal structure of *N,N'*-bis(3,5-dimethoxyphenyl)oxamide, (I).



Compound (I) has a crystallographically imposed center of symmetry at the mid-point of the C9—C9<sup>i</sup> bond [symmetry code: (i)  $2 - x, 1 - y, -z$ ]. The bond lengths and angles are within normal ranges (Allen *et al.*, 1987). However, the C9—C9<sup>i</sup> bond length [1.527 (7) Å] is longer than the typical  $\text{Csp}^2$ — $\text{Csp}^2$  single bond distance (1.460 Å). The molecule of (I) is almost planar. Weak N—H···O and C—H···O interactions (Table 1) are observed in the molecular structure. The crystal structure is stabilized by  $\pi$ – $\pi$  interactions between the benzene rings at  $(x, y, z)$  and  $(1 - x, 1 - y, 1 - z)$ , with a centroid–centroid distance of 3.670 (2) Å.

**Experimental**

To a solution of 3,5-dimethoxyaniline (15.3 g, 0.1 mol) in  $\text{CH}_2\text{Cl}_2$  (30 ml) was added dropwise a solution of oxalyl chloride (3.2 g, 0.025 mol) in  $\text{CH}_2\text{Cl}_2$  (10 ml), and the mixture was stirred at 353 K for 2 h. After cooling to room temperature, the solvent was removed *in vacuo*, and the residue was subjected to column chromatography (petroleum ether/EtOAc 3:1) to give (I) as a yellow solid (5.6 g, 62%). Single crystals of (I) suitable for an X-ray diffraction study were obtained by slow evaporation of a  $\text{CHCl}_3$  solution at room temperature over a period of 3 d.

Crystal data

C<sub>18</sub>H<sub>20</sub>N<sub>2</sub>O<sub>6</sub>  
*M<sub>r</sub>* = 360.36  
 Monoclinic, *P*2<sub>1</sub>/*c*  
*a* = 6.7851 (7) Å  
*b* = 15.6234 (15) Å  
*c* = 8.0308 (8) Å  
 β = 92.59 (3)°  
*V* = 850.45 (15) Å<sup>3</sup>

*Z* = 2  
*D<sub>x</sub>* = 1.407 Mg m<sup>-3</sup>  
 Mo *K*α radiation  
 μ = 0.11 mm<sup>-1</sup>  
*T* = 293 (2) K  
 Block, yellow  
 0.20 × 0.20 × 0.10 mm

Data collection

Bruker SMART CCD area-detector diffractometer  
 φ and ω scans  
 Absorption correction: multi-scan (*SADABS*; Bruker, 2001)  
*T<sub>min</sub>* = 0.979, *T<sub>max</sub>* = 0.989

1805 measured reflections  
 1672 independent reflections  
 747 reflections with *I* > 2σ(*I*)  
*R<sub>int</sub>* = 0.061  
 θ<sub>max</sub> = 26.0°

Refinement

Refinement on *F*<sup>2</sup>  
*R*[*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.069  
*wR*(*F*<sup>2</sup>) = 0.152  
*S* = 0.98  
 1672 reflections  
 119 parameters  
 H-atom parameters constrained

*w* = 1/[σ<sup>2</sup>(*F<sub>o</sub>*<sup>2</sup>) + (0.0455*P*)<sup>2</sup>]  
 where *P* = (*F<sub>o</sub>*<sup>2</sup> + 2*F<sub>c</sub>*<sup>2</sup>)/3  
 (Δ/σ)<sub>max</sub> = 0.001  
 Δρ<sub>max</sub> = 0.20 e Å<sup>-3</sup>  
 Δρ<sub>min</sub> = -0.20 e Å<sup>-3</sup>  
 Extinction correction: *SHELXL97*  
 Extinction coefficient: 0.023 (3)

Table 1

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—H1...O1 <sup>i</sup>	0.86	2.22	2.665 (4)	112
C2—H2...O1	0.93	2.33	2.935 (5)	123

Symmetry code: (i) -*x* + 2, -*y* + 1, -*z*.

H atoms were placed in idealized positions and refined as riding, with C—H = 0.93 or 0.96 Å, N—H = 0.86 Å and *U*<sub>iso</sub>(H) = 1.2*U*<sub>eq</sub>(C,N), or 1.5*U*<sub>eq</sub>(C) for methyl H.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve

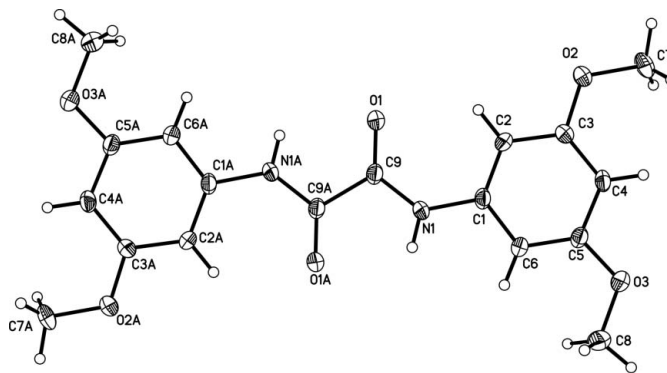


Figure 1

The molecular structure of (I), showing the labeling scheme. Displacement ellipsoids are drawn at the 30% probability level. The suffix A corresponds to symmetry code (2 - *x*, 1 - *y*, -*z*).

structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2001); software used to prepare material for publication: *SHELXL97*.

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