

## 5-[(Cyclohexylamino)methylene]-2,2-dimethyl-1,3-dioxane-4,6-dione

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## Key indicators

Single-crystal X-ray study

T = 297 K

Mean  $\sigma(\text{C}-\text{C}) = 0.006 \text{ \AA}$ 

R factor = 0.057

wR factor = 0.204

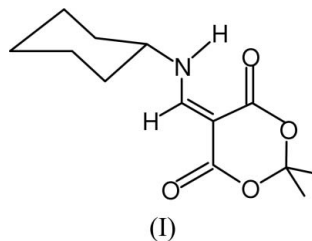
Data-to-parameter ratio = 7.7

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

In the title enaminone,  $\text{C}_{13}\text{H}_{19}\text{N}_1\text{O}_4$ , the Meldrum's acid group has a distorted boat conformation. The two independent molecules of the asymmetric unit are dimerized about a pseudo inversion center through two intermolecular N—H···O hydrogen bonds, with N···O distances of 3.049 (4) and 3.093 (4) Å. The dimers are stacked along the [010] direction.

## Comment

The synthesis of functionalized enaminones is a theme of ongoing interest owing to their potential in the synthesis of heterocyclic compounds (Negri *et al.*, 2004; Lue & Greenhill, 1997; Kuckländer, 1994; Ferraz & Pereira, 2004). A particular class of such compounds is derived from Meldrum's acid and the solid-state structural study of such derivatives has been described (Vencato *et al.*, 2004; Cunha *et al.*, 2003; Silva *et al.*, 2006; Silva *et al.*, 2005a,b, Joussef *et al.*, 2005a,b). Here we report our structural study of one enaminone, (I), derived from Meldrum's acid.



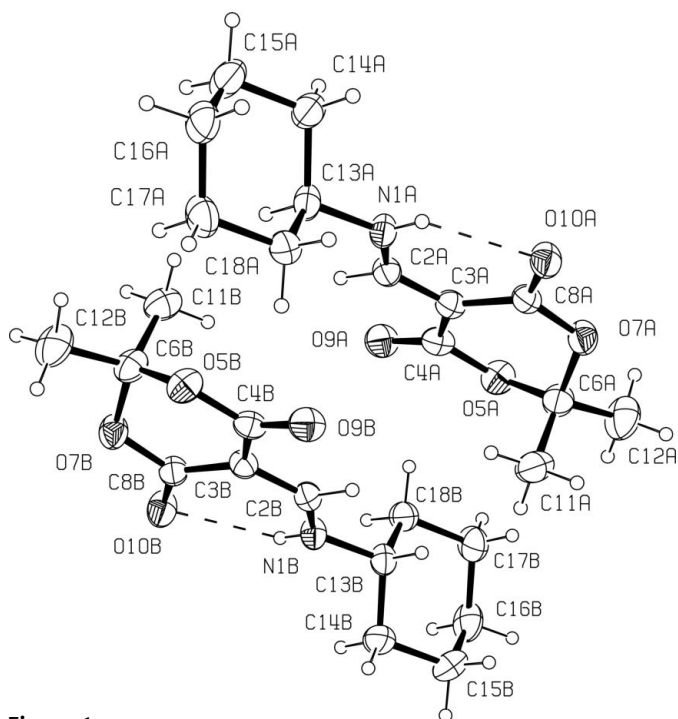
The structure has two independent molecules in the asymmetric unit (Fig. 1). Selected bond distances, angles and torsion angles are given in Table 1. In both molecules, the Meldrum's acid group has a distorted boat conformation, indicated by the Cremer & Pople (1975) parameters:  $\text{O5A} \rightarrow \text{C4A} \rightarrow \dots \rightarrow \text{C6A}$  [ $Q = 0.429$  (4) Å,  $\theta = 61.6$  (5)° and  $\varphi = 297.7$  (6)°] and  $\text{O5B} \rightarrow \text{C4B} \rightarrow \dots \rightarrow \text{C6B}$  [ $Q = 0.436$  (4) Å,  $\theta = 63.3$  (5)° and  $\varphi = 298.4$  (5)°].

There are small differences between the two molecules, as can be seen in the bond angles  $\text{O7}-\text{C6}-\text{C12}$  of 106.7 (4) and 104.5 (3)° for molecules *A* and *B*, respectively. The bond distance  $\text{O10A}-\text{C8A}$  of 1.219 (4) Å is marginally longer than the corresponding  $\text{O10B}-\text{C8B}$  distance of 1.204 (4) Å, presumably due to the fact that atom  $\text{O10A}$  is involved in a hydrogen bond as acceptor.

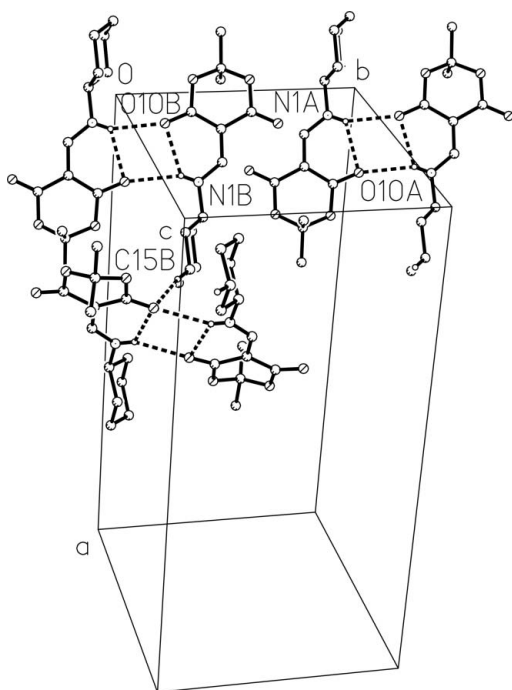
There are two intramolecular hydrogen bonds (Table 2), providing two flattened  $\text{H1}-\text{N1}-\text{C2}-\text{C3}-\text{C8}-\text{O10}$  rings for molecules *A* and *B*. The two independent molecules of the asymmetric unit are dimerized about a pseudo-inversion center through two intermolecular hydrogen bonds (Table 2). It is worth noting that atoms  $\text{H1A}$  and  $\text{H1B}$  belong to bifur-

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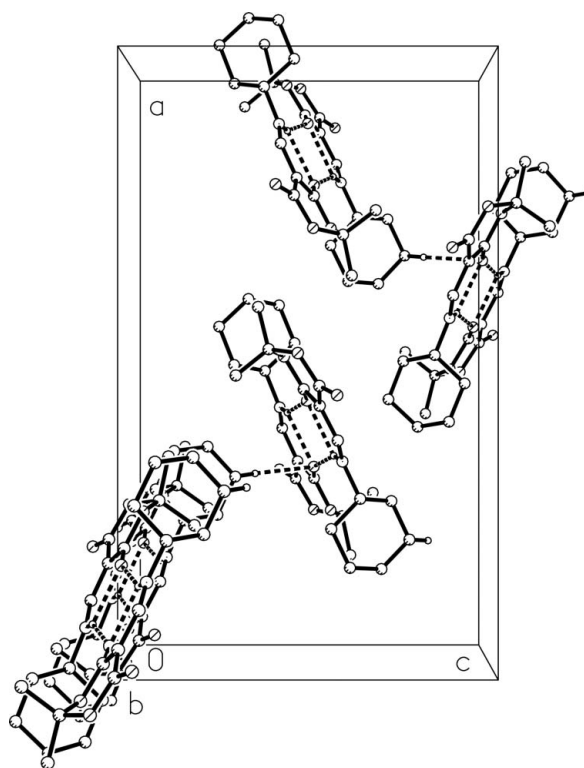


**Figure 1**  
The asymmetric unit of (I), with the atom-numbering schemes for molecules *A* and *B*. Displacement ellipsoids are drawn at the 30% probability level. Dashed lines indicate hydrogen bonds.



**Figure 2**  
Hydrogen bonding of (I). Intra- and intermolecular hydrogen bonds are shown as dashed lines. Only the H atoms involved in hydrogen bonds are shown.

cated intra- and intermolecular hydrogen bonds. The hydrogen bonding is shown in Fig. 2. The molecular packing of (I) is shown in Fig. 3, with the geometrical parameters in Table 2. The non-classical intermolecular C—H...O hydrogen



**Figure 3**  
Partial packing diagram of (I), viewed down the *b* axis. Dashed lines indicate hydrogen bonds.

bond completes the packing, with the dimers stacked parallel to the [010] direction.

## Experimental

A solution of 288 mg (2.0 mmol) of Meldrum's acid in 2 ml (26.1 mmol) of trimethyl orthoformate was heated at reflux for 3 h after which time the solvent was evaporated. The yellow solid that formed was added to 2.0 mmol of cyclohexylamine in 5 ml of CH<sub>2</sub>Cl<sub>2</sub> and the solution was allowed to stand at room temperature with stirring for 72 h. The solvent was evaporated and the crude residue was purified by silica gel 60 (70–230 mesh) column chromatography (hexane/ethyl acetate 50%), afforded (I) as a yellow solid (yield 116.7 mg, 23%; m.p. 421–422 K). Single crystals were obtained by slow evaporation of a solution of (I) in CH<sub>3</sub>Cl at room temperature.

### Crystal data

C<sub>13</sub>H<sub>19</sub>NO<sub>4</sub>  
*M<sub>r</sub>* = 253.29  
 Orthorhombic, *Pca*2<sub>1</sub>  
*a* = 20.905 (2) Å  
*b* = 10.222 (1) Å  
*c* = 12.516 (6) Å  
*V* = 2674.6 (13) Å<sup>3</sup>

*Z* = 8  
*D<sub>x</sub>* = 1.258 Mg m<sup>-3</sup>  
 Cu *K*α radiation  
 μ = 0.77 mm<sup>-1</sup>  
*T* = 297 (2) K  
 Prism, yellow  
 0.35 × 0.35 × 0.30 mm

### Data collection

Enraf–Nonius CAD-4 diffractometer  
 ω/2θ scans  
 Absorption correction: none  
 2698 measured reflections  
 2558 independent reflections

2404 reflections with *I* > 2σ(*I*)  
*R*<sub>int</sub> = 0.025  
 θ<sub>max</sub> = 68.0°  
 2 standard reflections  
 frequency: 120 min  
 intensity decay: 1%

## Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.057$   
 $wR(F^2) = 0.204$   
 $S = 1.20$   
 2558 reflections  
 332 parameters  
 H atoms treated by a mixture of  
 independent and constrained  
 refinement

$$w = 1/[\sigma^2(F_o^2) + (0.1659P)^2 + 0.0648P]$$

where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.005$   
 $\Delta\rho_{\max} = 0.39 \text{ e } \text{Å}^{-3}$   
 $\Delta\rho_{\min} = -0.40 \text{ e } \text{Å}^{-3}$   
 Extinction correction: *SHELXL97*  
 Extinction coefficient: 0.054 (4)

Table 1

Selected geometric parameters (Å, °).

N1A—C2A	1.294 (4)	N1B—C2B	1.287 (5)
N1A—C13A	1.469 (4)	N1B—C13B	1.472 (4)
C4A—O9A	1.208 (5)	C4B—O9B	1.213 (4)
C4A—O5A	1.373 (5)	C4B—O5B	1.368 (5)
O5A—C6A	1.441 (5)	O5B—C6B	1.429 (5)
C6A—O7A	1.435 (4)	C6B—O7B	1.441 (5)
O7A—C8A	1.364 (4)	O7B—C8B	1.373 (4)
C8A—O10A	1.219 (4)	C8B—O10B	1.204 (4)
C2A—N1A—C13A	123.1 (3)	C2B—N1B—C13B	122.8 (3)
O7A—C6A—C12A	106.7 (4)	O7B—C6B—C12B	104.5 (4)
C18A—C13A—C14A	111.1 (3)	C18B—C13B—C14B	111.8 (3)
C15A—C16A—C17A	112.7 (4)	C15B—C16B—C17B	111.4 (4)

Table 2

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1A—H1A <sup>i</sup> ···O10A	0.80 (5)	2.27 (5)	2.791 (4)	124 (4)
N1B—H1B <sup>i</sup> ···O10B	0.77 (5)	2.27 (5)	2.796 (4)	126 (4)
N1A—H1A <sup>i</sup> ···O10B <sup>i</sup>	0.80 (5)	2.31 (5)	3.049 (4)	154 (4)
N1B—H1B <sup>i</sup> ···O10A <sup>ii</sup>	0.77 (5)	2.38 (5)	3.093 (4)	154 (4)
C15B—H15D <sup>i</sup> ···O10A <sup>iii</sup>	0.97	2.58	3.536 (6)	170

Symmetry codes: (i)  $x, y + 1, z$ ; (ii)  $x, y - 1, z$ ; (iii)  $-x + \frac{1}{2}, y - 1, z + \frac{1}{2}$ .

H atoms bonded to atoms N1A and N1B were found in a difference map and their positions were refined freely, with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$ . All other H atoms were placed in calculated positions ( $C-H = 0.96$  and  $0.97$  Å) and treated as riding atoms [ $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  or  $1.5U_{\text{eq}}(\text{methyl C})$ ]. Due to the absence of any significant anomalous scatterers in the compound, Friedel pairs were merged.

Data collection: *CAD-4-PC* (Enraf–Nonius, 1993); cell refinement: *CAD-4-PC*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXL97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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