

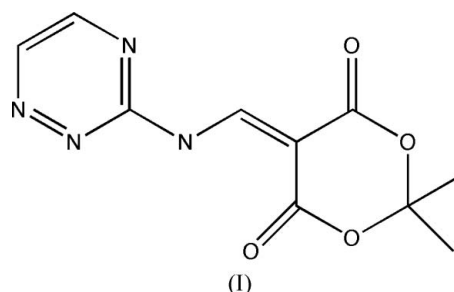
2,2-Dimethyl-5-[(1,2,4-triazin-3-ylamino)-  
methylene]-1,3-dioxane-4,6-dioneAntonio C. Joussef,<sup>a</sup>  
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## Key indicators

Single-crystal X-ray study  
 $T = 193$  K  
Mean  $\sigma(\theta) = 0.000$  Å  
Disorder in main residue  
 $R$  factor = 0.055  
 $wR$  factor = 0.161  
Data-to-parameter ratio = 8.4For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.In the title compound,  $C_{10}H_{10}N_4O_4$ , the triazine ring is nearly planar. The 1,3-dioxane-4,6-dione ring exhibits a half-chair conformation. The molecule is disordered and the structure has been refined using a split model.Received 14 June 2005  
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## Comment

Various derivatives of Meldrum's acid have been investigated by X-ray methods (Gould *et al.*, 1998; Blake *et al.*, 1997, Blake, Gould *et al.*, 1994; Blake & McNab, 1995; Blake, McNab & Morrow, 1994). As part of a continuing study of the 5-methoxymethylene Meldrum's acid derivatives as potential antiviral, antitrypanosomal and leishmanicidal, we report the structure of the title compound, (I).

In (I), the 1,3-dioxane-4,6-dione ring exhibits a half-chair conformation and the triazine ring is nearly planar. The molecule is disordered and the structure has therefore been refined using a split model. Because of the disorder some of the bond lengths and angles are unusual. The disorder components with displacement ellipsoids are shown separately in Figs. 1 and 2. The total disordered molecular structure is shown in Fig. 3. Details of the intra- and intermolecular hydrogen bonds are given in Table 1.

## Experimental

The title compound was prepared according to the literature procedure of Cassis *et al.* (1985) and recrystallized from methanol.

## Crystal data

 $C_{10}H_{10}N_4O_4$   
 $M_r = 250.22$   
Monoclinic,  $P2_1/n$   
 $a = 6.379$  (5) Å  
 $b = 18.239$  (5) Å  
 $c = 10.101$  (5) Å  
 $\beta = 107.331$  (5)°  
 $V = 1121.9$  (11) Å<sup>3</sup>  
 $Z = 4$   
 $D_x = 1.481$  Mg m<sup>-3</sup> $D_m = ?$  Mg m<sup>-3</sup>  
Mo  $K\alpha$  radiation  
Cell parameters from 25  
reflections  
 $\theta = 4.9$ – $15.4$ °  
 $\mu = 0.12$  mm<sup>-1</sup>  
 $T = 193$  (2) K  
Prism, colorless  
 $0.50 \times 0.20 \times 0.16$  mm

Data collection

Enraf–Nonius CAD-4  
diffractometer  
 $\omega/2\theta$  scans  
Absorption correction: none  
2028 measured reflections  
1924 independent reflections  
1191 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.033$

$\theta_{\text{max}} = 25.1^\circ$   
 $h = -7 \rightarrow 7$   
 $k = -21 \rightarrow 0$   
 $l = -12 \rightarrow 0$   
3 standard reflections  
every 200 reflections  
intensity decay: 1%

Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.055$   
 $wR(F^2) = 0.161$   
 $S = 1.05$   
1924 reflections  
230 parameters  
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0592P)^2 + 0.8549P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} < 0.001$   
 $\Delta\rho_{\text{max}} = 0.40 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.27 \text{ e \AA}^{-3}$

Table 1

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N7A-H7A\cdots O14A$	0.86	2.08	2.724 (5)	131
$N7B-H7B\cdots O10B$	0.86	2.13	2.777 (19)	132
$C4B-H4B\cdots O10A^i$	0.93	2.04	2.842 (14)	143
$C4B-H4B\cdots O10B^i$	0.93	1.90	2.802 (19)	162

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

The structure is disordered and was refined with a split model. Refinement was performed with restraints for both components. The corresponding site-occupation factors were refined but later fixed to 0.80 and 0.20. All non-H atoms of the minor occupied component were refined isotropically and a common variable was used for the isotropic displacement factor of the ring N1B–C6B. The triazine ring is additionally disordered, such that atoms N2A and C4A, as well as N2B and C4B, occupy the same position. Therefore, these atoms were refined with site-occupation factors 0.5:0.5. All H atoms were positioned with idealized geometry and were refined with isotropic displacement parameters (set to 1.2 times  $U_{\text{eq}}$  of the parent atom) using a riding model with  $C-H = 0.93 \text{ \AA}$  ( $0.96 \text{ \AA}$  for methyl groups) and  $N-H = 0.86 \text{ \AA}$ .

Data collection: *CAD-4/PC Software* (Enraf–Nonius, 1993); cell refinement: *CAD-4/PC Software*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

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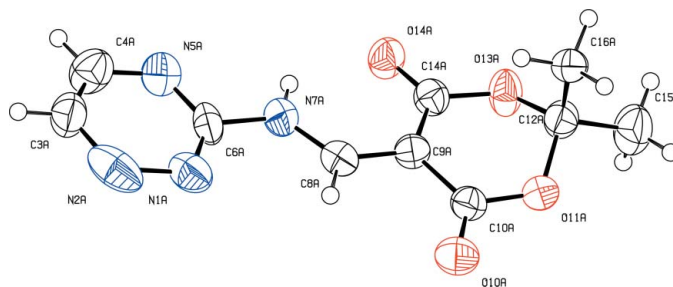


Figure 1

The molecular structure of the major component (site-occupation factor = 0.80), with the atom labelling, and with displacement ellipsoids drawn at the 50% probability level.

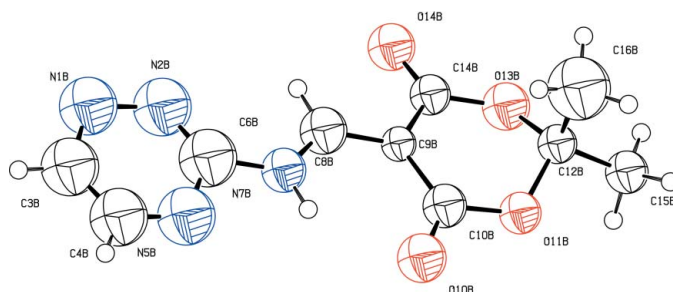


Figure 2

The molecular structure of the minor component (site-occupation factor = 0.20), with displacement spheres drawn at the 50% probability level. (Please note: this fragment was refined isotropically.)

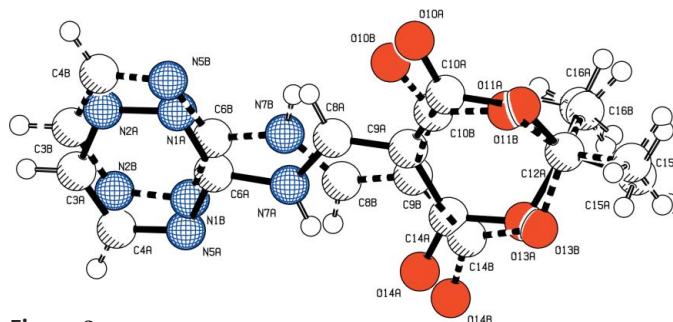


Figure 3

The disordered molecular structure of the title compound, with the atom labelling (the bonds of the minor component are drawn with dashed lines).

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