

## 2,2-Dimethyl-5-[(4-methylthiazol-2-ylamino)-methylene]-1,3-dioxane-4,6-dione

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In the title compound,  $C_{11}H_{12}N_2O_4S$ , the thiazole ring is nearly planar. The 1,3-dioxane-4,6-dione ring exhibits a half-chair conformation. The NH group forms an intramolecular contact to a carbonyl O atom, forming a six-membered ring, and an intermolecular N—H···O hydrogen bond, with an H···O distance of 2.16 Å, is also observed.

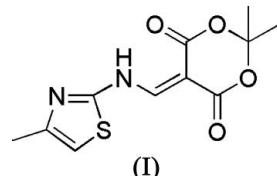
### Comment

Thiazoles constitute an important class of heterocyclic compounds for which pharmacological properties have been reported (Krasovsky *et al.*, 2002). 5-Arylaminemethylene Meldrum's acid analogs play an important role in heterocyclic chemistry as pivotal intermediates in the formation of cyclic products (Chen, 1991). As part of a continuing study of the conformation in the solid state of 5-aminomethylene Meldrum's acid derivatives (Joussef *et al.*, 2005*a,b*; da Silva *et al.*, 2005*a,b*, 2006), we report an X-ray crystallographic study of the title compound, (I).

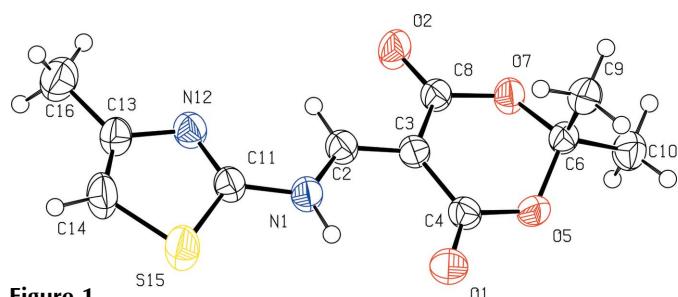
### Key indicators

Single-crystal X-ray study  
 T = 293 K  
 Mean  $\sigma(C-C)$  = 0.003 Å  
 R factor = 0.042  
 wR factor = 0.125  
 Data-to-parameter ratio = 13.2

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



In (I), the 1,3-dioxane-4,6-dione ring exhibits a half-chair conformation. The dihedral angle C11—N1—C2—C3 is 174.5 (2)° and the distances N1—C11 and C2—C3 indicate delocalization of the conjugated system. The amino H atom has an intramolecular contact to O1, with an H···O distance of 2.24 Å, forming a six-membered ring. The delocalization of the N-atom lone pair into the Meldrum's acid ring may be favoured in the direction of one of the two carbonyl groups (Blake *et al.*, 2003). Details of the hydrogen bonding are given in Table 1.



**Figure 1**

The molecular structure of (I), with the atom labeling and displacement ellipsoids drawn at the 50% probability level.

## Experimental

The title compound was prepared according to a literature procedure (Cassis *et al.*, 1985) and was recrystallized from methanol (m.p.: 468–470 K).

### Crystal data

$C_{11}H_{12}N_2O_4S$	$D_x = 1.467 \text{ Mg m}^{-3}$
$M_r = 268.29$	$\text{Mo } K\alpha \text{ radiation}$
Monoclinic, $P2_1/c$	Cell parameters from 25 reflections
$a = 5.6384 (6) \text{ \AA}$	$\theta = 9.0\text{--}18.1^\circ$
$b = 18.562 (4) \text{ \AA}$	$\mu = 0.28 \text{ mm}^{-1}$
$c = 11.920 (2) \text{ \AA}$	$T = 293 (2) \text{ K}$
$\beta = 103.14 (1)^\circ$	Irregular block, yellow
$V = 1214.9 (3) \text{ \AA}^3$	$0.36 \times 0.26 \times 0.23 \text{ mm}$
$Z = 4$	

### Data collection

Enraf–Nonius CAD-4 diffractometer  
 $\omega$ -2 $\theta$  scans  
Absorption correction: none  
2249 measured reflections  
2144 independent reflections  
1590 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.024$

$\theta_{\text{max}} = 25.1^\circ$   
 $h = -6 \rightarrow 6$   
 $k = -22 \rightarrow 0$   
 $l = -14 \rightarrow 0$   
3 standard reflections every 200 reflections  
intensity decay: 1%

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.042$   
 $wR(F^2) = 0.125$   
 $S = 1.04$   
2144 reflections  
163 parameters  
H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0636P)^2 + 0.5695P]$$

where  $P = (F_o^2 + 2F_c^2)/3$

 $(\Delta/\sigma)_{\text{max}} < 0.001$ 
 $\Delta\rho_{\text{max}} = 0.33 \text{ e } \text{\AA}^{-3}$ 
 $\Delta\rho_{\text{min}} = -0.30 \text{ e } \text{\AA}^{-3}$

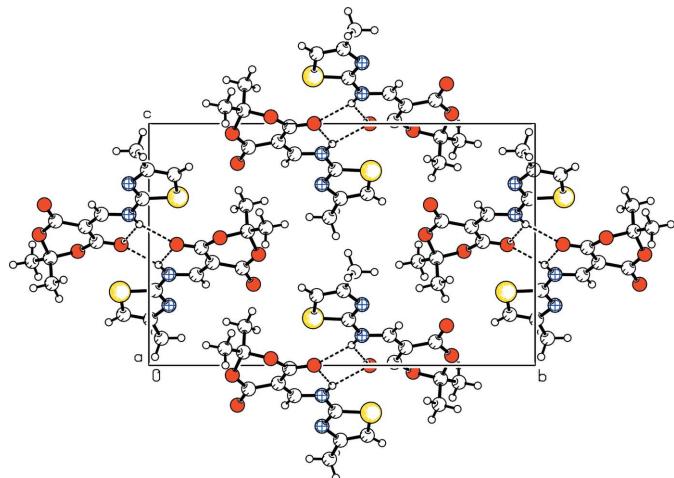
**Table 1**  
Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1—H1 $\cdots$ O1	0.91	2.24	2.836 (3)	123
N1—H1 $\cdots$ O1 <sup>i</sup>	0.91	2.16	2.996 (3)	154

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

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, 1.5 for methyl groups) using a riding model with N—H = 0.91  $\text{\AA}$ , C—H = 0.93  $\text{\AA}$  (aromatic), 0.96  $\text{\AA}$  (methyl).

Data collection: CAD-4 EXPRESS (Enraf–Nonius, 1993); cell refinement: CAD-4 EXPRESS; data reduction: XCAD4 (Harms &



**Figure 2**

Molecular packing of (I), with hydrogen bonds shown as dashed lines.

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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