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#### Key indicators

Single-crystal X-ray study  
 $T = 299$  K  
Mean  $\sigma(C-C) = 0.003$  Å  
 $R$  factor = 0.035  
 $wR$  factor = 0.094  
Data-to-parameter ratio = 10.5

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

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

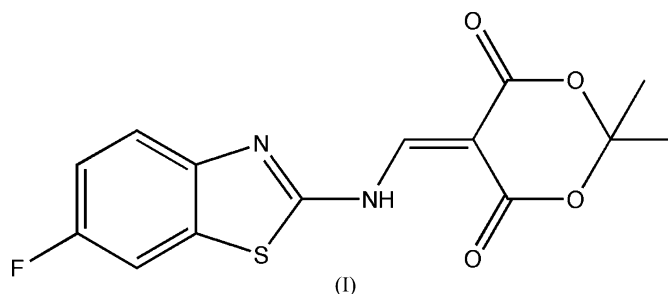
In the title compound,  $C_{20}H_{16}N_2O_4S$ , the 1,3-dioxane-4,6-dione ring exhibits a half-chair conformation. The amino H atom forms an intramolecular contact to a carbonyl O atom, forming a six-membered ring, and an intermolecular contact to a carbonyl O atom. Additional intermolecular hydrogen bonds of the types  $C-H \cdots O$  and  $C-H \cdots F$  are also observed.

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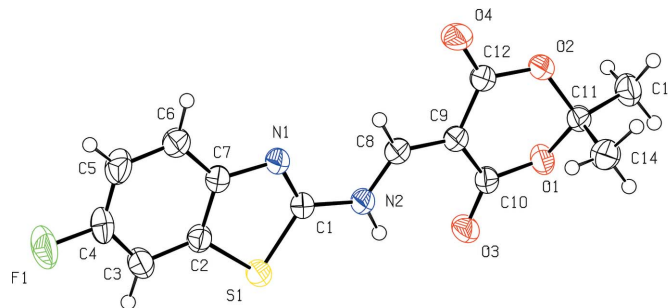
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#### Comment

We have focused on arylaminemethylene derivatives of Meldrum's acid in our search for biologically active molecules (Joussef *et al.*, 2005*a,b*; da Silva *et al.*, 2005*a,b*). As an extension of this approach, we report here an X-ray crystallographic study of the title compound, (I).



In (I), the 1,3-dioxane-4,6-dione ring exhibits a half-chair conformation. The  $C1-N2-C8-C9$  torsion angle is  $-177.1(2)^\circ$  and the  $C1-N2$  [ $1.400(2)$  Å] and  $C8-C9$  [ $1.371(3)$  Å] distances indicate delocalization. The delocalization of the N-atom lone pair into the ring of Meldrum's acid may be favoured in the direction of one of the two carbonyl groups (Blake *et al.*, 2003).



**Figure 1**  
The molecular structure of (I), with displacement ellipsoids drawn at the 50% probability level.

One intramolecular hydrogen bond of the type N—H···O was observed, together with three intermolecular hydrogen bonds of types N—H···O, C—H···O and C—H···F. These connect molecules to form a three-dimensional network, as represented in Fig. 2. Details of the hydrogen-bonding parameters are given in Table 1.

## Experimental

The title compound was prepared according to the literature procedure of Cannon *et al.* (2001) and was recrystallized from ethyl acetate (m.p. 495–596 K; yield 78%).

### Crystal data

|                                |   |
|--------------------------------|---|
| $C_{14}H_{11}FN_2O_4S$         | $D_x = 1.556 \text{ Mg m}^{-3}$           |
| $M_r = 322.31$                 | Cu $K\alpha$ radiation                    |
| Monoclinic, $P2_1/c$           | Cell parameters from 25 reflections       |
| $a = 6.433 (1) \text{ \AA}$    | $\theta = 4.6\text{--}18.6^\circ$         |
| $b = 19.532 (2) \text{ \AA}$   | $\mu = 2.41 \text{ mm}^{-1}$              |
| $c = 11.218 (2) \text{ \AA}$   | $T = 299 (2) \text{ K}$                   |
| $\beta = 102.52 (1)^\circ$     | Rod, light brown                          |
| $V = 1376.0 (4) \text{ \AA}^3$ | $0.20 \times 0.13 \times 0.10 \text{ mm}$ |
| $Z = 4$                        |   |

### Data collection

|  |                                    |
|--|------------------------------------|
| Enraf–Nonius CAD-4 diffractometer      | $\theta_{\text{max}} = 67.0^\circ$ |
| $\omega/2\theta$ scans                 | $h = -7 \rightarrow 1$             |
| Absorption correction: none            | $k = -23 \rightarrow 0$            |
| 2707 measured reflections              | $l = -13 \rightarrow 13$           |
| 2454 independent reflections           | 3 standard reflections             |
| 1949 reflections with $I > 2\sigma(I)$ | frequency: 120 min                 |
| $R_{\text{int}} = 0.024$               | intensity decay: 2.0%              |

### Refinement

|                                 |  |
|---------------------------------|--|
| Refinement on $F^2$             | $w = 1/[\sigma^2(F_o^2) + (0.047P)^2 + 0.3778P]$     |
| $R[F^2 > 2\sigma(F^2)] = 0.035$ | where $P = (F_o^2 + 2F_c^2)/3$                       |
| $wR(F^2) = 0.094$               | $(\Delta/\sigma)_{\text{max}} = 0.004$               |
| $S = 1.01$                      | $\Delta\rho_{\text{max}} = 0.26 \text{ e \AA}^{-3}$  |
| 2454 reflections                | $\Delta\rho_{\text{min}} = -0.21 \text{ e \AA}^{-3}$ |
| 233 parameters                  | Extinction correction: <i>SHELXL97</i>               |
| Only H-atom coordinates refined | Extinction coefficient: 0.0043 (3)                   |

**Table 1**

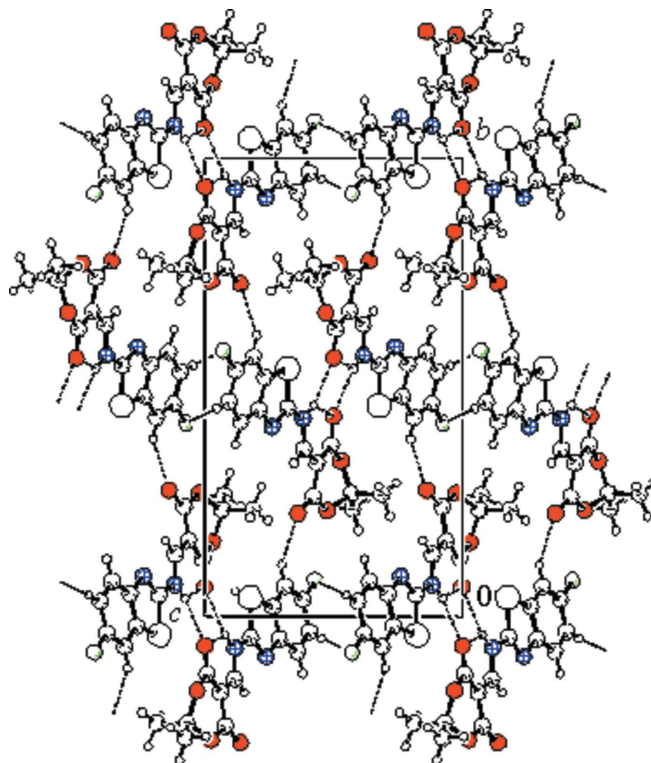
Hydrogen-bond geometry ( $\text{\AA}, ^\circ$ ).

| $D\text{—}H\cdots A$      | $D\text{—}H$ | $H\cdots A$ | $D\cdots A$ | $D\text{—}H\cdots A$ |
|---------------------------|--------------|-------------|-------------|----------------------|
| N2—H2N···O3               | 0.89 (2)     | 2.19 (2)    | 2.801 (2)   | 125.5 (19)           |
| N2—H2N···O3 <sup>i</sup>  | 0.89 (2)     | 2.23 (2)    | 3.045 (2)   | 151 (2)              |
| C3—H3···O4 <sup>ii</sup>  | 0.94 (3)     | 2.58 (3)    | 3.394 (3)   | 145 (2)              |
| C5—H5···F1 <sup>iii</sup> | 0.90 (3)     | 2.46 (3)    | 3.356 (3)   | 174 (2)              |

Symmetry codes: (i)  $-x + 1, -y, -z$ ; (ii)  $-x + 2, y - \frac{1}{2}, -z + \frac{1}{2}$ ; (iii)  $-x + 4, -y, -z + 1$ .

All H atoms were located in a difference map, refined [ $0.90 (3)\text{--}1.03 (3) \text{ \AA}$ ] and assigned isotropic displacement parameters of  $1.2U_{\text{eq}}$  (parent atom).

Data collection: *CAD-4-PC* (Enraf–Nonius, 1993); cell refinement: *CAD-4-PC*; data reduction: *REDU4* (Stoe & Cie, 1987); program(s)



**Figure 2**

The molecular packing (arbitrary spheres) of (I), viewed down the  $a$  axis, with hydrogen bonds shown as dashed lines.

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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