

Ethyl 3-[[[(2-chlorophenyl)sulfonyl]imino]-2-[[[(3,5-dimethoxyphenyl)amino]carbonyl]-3-(methylsulfanyl)propanoate

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Key indicators

Single-crystal X-ray study

$T = 293$ K

Mean $\sigma(\text{C}-\text{C}) = 0.010$ Å

Disorder in main residue

R factor = 0.043

wR factor = 0.152

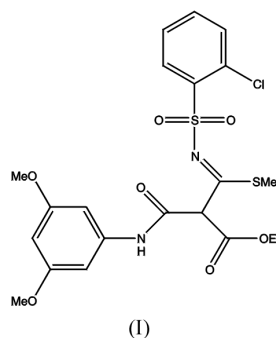
Data-to-parameter ratio = 14.9

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

The title compound, $\text{C}_{21}\text{H}_{23}\text{ClN}_2\text{O}_7\text{S}_2$, is a representative of moderately active hydroxyacid synthase inhibitors. The structure has a sulfonylimine bond [$-\text{N}=\text{C}(-\text{SMe})-\text{CH}$] and not sulfonylamino [$-\text{NH}-\text{C}(-\text{SMe})=\text{C}$], contrary to expectations. There is a strong intramolecular hydrogen bond between the imine H and the acid carbonyl group, and a weaker one between the imine carbonyl group and a benzene H atom.

Comment

This report is part of a general study on acetohydroxyacid synthase inhibitors (McFadden *et al.*, 1993). The title compound, (I), was reported in that paper as a molecule with a double bond between atoms C2 and C3. However, a crystal structure determination (Fig. 1 and Table 1) shows that the double bond occurs between atoms N1 and C2 [1.339 (6) Å], indicating a minimum energy structure with a sulfonylimine link. There are intramolecular $\text{N1}-\text{H1}\cdots\text{O32}$ and $\text{C12}-\text{H12}\cdots\text{O21}$ hydrogen bonds in (I). Compared with the inactive compound 3-[(2-chlorophenyl)sulfonylamino]-2-cyano-*N*-(3,5-dimethoxyphenyl)-3-methylsulfanyl-2-propenamide (Kennard *et al.*, 2003), compound (I) has different torsion angles [$\text{N1}-\text{C2}-\text{C3}-\text{C4} = 155.5$ (5) and -179.67 (17)°, $\text{C2}-\text{C3}-\text{C4}-\text{N5} = -69.2$ (7) and -3.9 (3)°, and $\text{C3}-\text{C4}-\text{N5}-\text{S6} = 0.7$ (9) and 170.64 (15)°].



Experimental

The synthesis of (I) has been reported by McFadden *et al.* (1993).

Crystal data

$\text{C}_{21}\text{H}_{23}\text{ClN}_2\text{O}_7\text{S}_2$

$M_r = 514.98$

Monoclinic, $P2_1/n$

$a = 8.546$ (4) Å

$b = 24.166$ (3) Å

$c = 12.057$ (8) Å

$\beta = 104.67$ (2)°

$V = 2409$ (2) Å³

$Z = 4$

$D_x = 1.420$ Mg m⁻³

Mo $K\alpha$ radiation

Cell parameters from 25

reflections

$\theta = 10-12^\circ$

$\mu = 0.38$ mm⁻¹

$T = 293$ (2) K

Plate, colourless

$0.30 \times 0.05 \times 0.05$ mm

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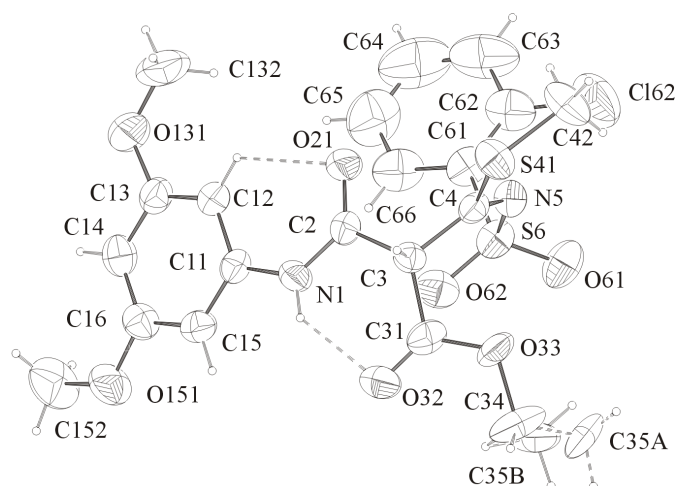


Figure 1

The molecular structure of (I), showing displacement ellipsoids at the 50% probability level. The occupation factors of atoms C35A and C35B are 40 and 60%, respectively.

Data collection

Enraf–Nonius CAD-4
diffractometer
 ω - 2θ scans
Absorption correction: ψ scan
(North *et al.*, 1968)
 $T_{\min} = 0.933$, $T_{\max} = 0.979$
4570 measured reflections
4233 independent reflections
1419 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.152$
 $S = 0.91$
4233 reflections
307 parameters

$R_{\text{int}} = 0.037$
 $\theta_{\text{max}} = 25.0^\circ$
 $h = -4 \rightarrow 10$
 $k = -11 \rightarrow 28$
 $l = -14 \rightarrow 13$
3 standard reflections
frequency: 120 min
intensity decay: none

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0626P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.011$
 $\Delta\rho_{\text{max}} = 0.24 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.25 \text{ e } \text{\AA}^{-3}$

Table 1

Selected geometric parameters (\AA , $^\circ$).

C2–C3	1.546 (6)	C3–C4	1.540 (7)
C12–C11–N1–C2	–11.3 (8)	C3–C4–N5–S6	1.0 (8)
C11–N1–C2–C3	–172.8 (5)	C4–N5–S6–C61	112.7 (5)
N1–C2–C3–C4	155.6 (5)	N5–S6–C61–C62	65.8 (5)
C2–C3–C4–N5	–69.4 (7)		

The terminal atom C35 of the ethyl group is disordered over two sites, C35A and C35B. Their occupation factors were assigned, from refinement, to be 40 and 60%, respectively. The ratios of atomic displacement parameters for the various directions for C35A are large and indicate serious disorder.

Data collection: *SDP* (Frenz, 1985); cell refinement: *SDP*; data reduction: *WinGX* (Farrugia, 1999); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97*; molecular graphics: *PLATON98* (Spek, 1988); software used to prepare material for publication: *SHELXL97*.

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