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Key indicators

Single-crystal X-ray study
 $T = 273$ K
Mean $\sigma(\text{C}-\text{C}) = 0.004$ Å
 R factor = 0.039
 wR factor = 0.113
Data-to-parameter ratio = 15.2For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.***N*-[2-(Aminosulfonyl)-5,6-dihydro-6-methyl-7,7-dioxo-4*H*-thieno[2,3-*b*]-thiopyran-4-yl]acetamide**

The title compound, $\text{C}_{10}\text{H}_{14}\text{N}_2\text{O}_5\text{S}_3$, has been synthesized as a key intermediate in the synthesis of topically active carbonic anhydrase inhibitor MK-0507. $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds are responsible for the formation of centrosymmetric dimers and zigzag molecular chains. The intermolecular non-bonded $\text{S}\cdots\text{O}$ separation in the crystal structure is 3.415 (2) Å, indicating a strong intermolecular interaction between the aminosulfonyl and *S*-dioxide groups.

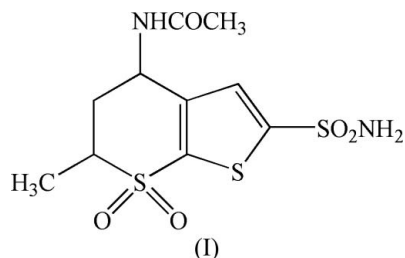
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Comment

The title compound, (I), is a key intermediate in the total synthesis of topically active carbonic anhydrase inhibitor MK-0507 (Blacklock *et al.*, 1993; Thomas *et al.*, 1997).



The molecular structure and atom-labelling scheme are shown in Fig. 1. The shortest intermolecular $\text{S}\cdots\text{O}$ contact, between S1 and O3^i , is 3.415 (2) Å [symmetry code: (i) $-x + \frac{3}{2}$, $y + \frac{1}{2}$, z], indicating a strong intermolecular interaction between these two atoms. In addition, there are also weak $\text{S}\cdots\text{C}$ interactions [non-bonded separation = 3.858 (3) Å], as well as $\text{S}\cdots\text{S}$ short contacts [3.949 > (4) Å] (Lan *et al.*, 2005).

Further examination of the crystal structure of (I) reveals possible $\text{N}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{O}$ interactions (Table 1). Intermolecular $\text{N2}-\text{H2B}\cdots\text{O3}^i$ hydrogen bonds [symmetry code: (i) $-x + \frac{3}{2}$, $y + \frac{1}{2}$, z] form zigzag molecular chains propagating in the *b*-axis direction, while $\text{N2}-\text{H2A}\cdots\text{O2}^{ii}$ hydrogen bonds [symmetry code: (ii) $-x + 2$, $-y + 2$, $-z + 1$] form centrosymmetric dimers, as shown in Fig. 2.

Experimental

The title compound was synthesized from methyl 3-(*p*-toluenesulfonyloxy)butyrate according to a literature method (Blacklock *et al.*, 1993; Ponticello *et al.*, 1987, 1988). Single crystals of (I) were grown by slow evaporation, in air, of a THF solution. Selected analytical data: pale-yellow solid, yield 89%; m.p. 524–525 K; ¹H NMR (DMSO-*d*₆, 500 MHz): δ 8.62 (*d*, 1H), 8.05 (*s*, 2H), 7.42 (*s*, 1H), 5.19 (*m*, 1H), 3.89 (*m*, 1H), 2.45 (*m*, 1H), 2.29 (*m*, 1H), 1.86 (*s*, 3H),

1.33 (d, 3H); IR (KBr) ν : 3337, 3240, 3080, 2939, 1654, 1523, 1449, 1334, 1303, 1170, 1138, 1035, 926, 873, 694 cm^{-1} ;

Crystal data

$\text{C}_{10}\text{H}_{14}\text{N}_2\text{O}_5\text{S}_3$
 $M_r = 338.41$
 Orthorhombic, *Pbca*
 $a = 9.842$ (3) Å
 $b = 15.290$ (5) Å
 $c = 19.434$ (6) Å
 $V = 2924.5$ (15) Å³
 $Z = 8$
 $D_x = 1.537$ Mg m⁻³

Mo $K\alpha$ radiation
 Cell parameters from 1543 reflections
 $\theta = 2.7$ – 21.2°
 $\mu = 0.53$ mm⁻¹
 $T = 273$ (2) K
 Prism, yellow
 $0.20 \times 0.10 \times 0.10$ mm

Data collection

Bruker SMART CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.902$, $T_{\max} = 0.949$
 13578 measured reflections

2865 independent reflections
 1657 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.058$
 $\theta_{\text{max}} = 26.0^\circ$
 $h = -12 \rightarrow 12$
 $k = -18 \rightarrow 18$
 $l = -21 \rightarrow 23$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.039$
 $wR(F^2) = 0.114$
 $S = 0.96$
 2865 reflections
 189 parameters

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0614P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.35$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.32$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N2}-\text{H2B}\cdots\text{O3}^{\text{i}}$	0.81 (4)	2.14 (4)	2.876 (4)	150 (3)
$\text{N2}-\text{H2A}\cdots\text{O2}^{\text{ii}}$	0.85 (3)	2.14 (3)	2.962 (4)	161 (3)
$\text{N1}-\text{H1D}\cdots\text{O1}^{\text{iii}}$	0.86	2.07	2.861 (3)	152
$\text{C5}-\text{H5A}\cdots\text{O1}^{\text{iii}}$	0.98	2.52	3.346 (3)	142
$\text{C4}-\text{H4A}\cdots\text{O2}^{\text{iv}}$	0.97	2.66	3.623 (4)	174
$\text{C3}-\text{H3A}\cdots\text{O4}^{\text{v}}$	0.98	2.53	3.276 (3)	132

Symmetry codes: (i) $-x + \frac{3}{2}, y + \frac{1}{2}, z$; (ii) $-x + 2, -y + 2, -z + 1$; (iii) $x + \frac{1}{2}, y, -z + \frac{3}{2}$; (iv) $x - \frac{1}{2}, y, -z + \frac{3}{2}$; (v) $-x + 1, -y + 2, -z + 1$.

H atoms were included using a riding model and were constrained to have $\text{C}-\text{H} = 0.95$ Å, $\text{N}-\text{H} = 0.86$ Å and $U_{\text{iso}} = 1.2U_{\text{eq}}$ of the parent atom. H atoms attached to N2 were refined freely.

Data collection: SMART (Bruker, 1999); cell refinement: SMART; data reduction: SAINT (Bruker, 1999); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: ORTEP-3 (Farrugia, 1997); software used to prepare material for publication: SHELXTL (Sheldrick, 1997b).

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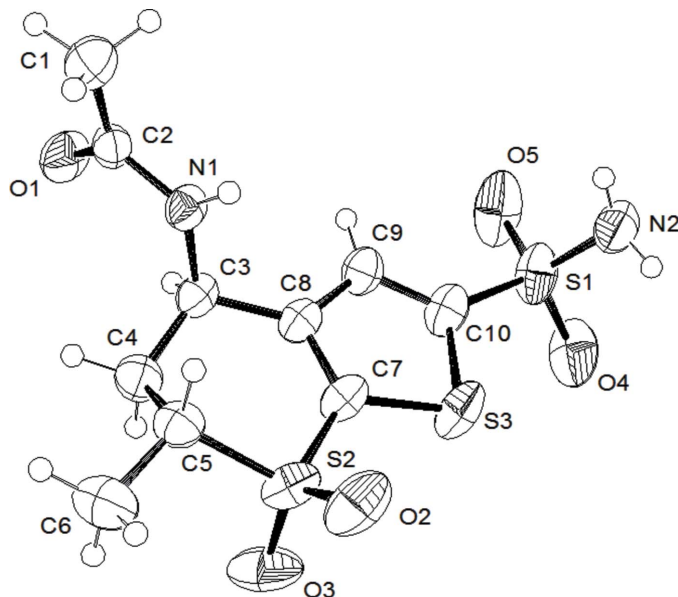


Figure 1

The molecular structure of the title compound. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii.

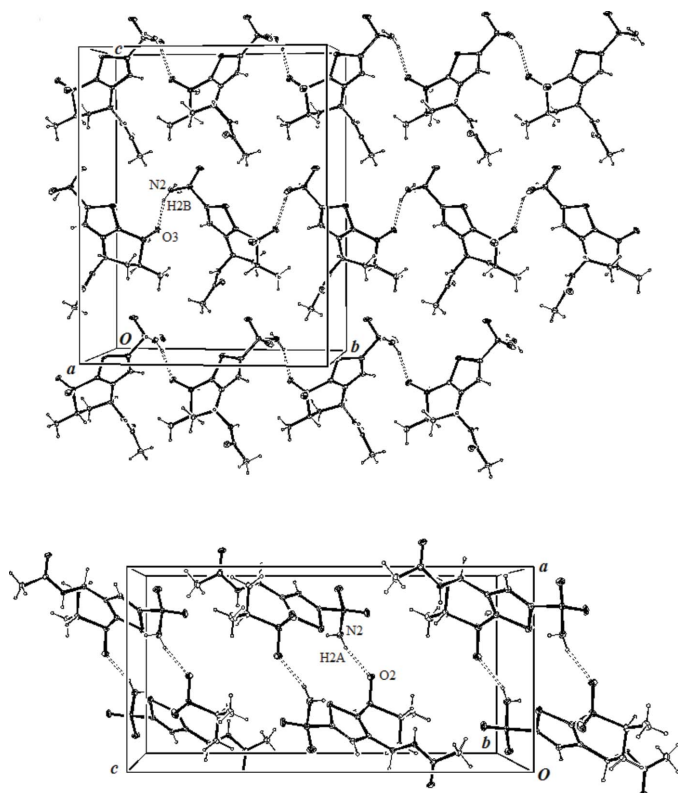


Figure 2

The crystal structure of (I), viewed in two directions. Dashed lines indicate $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds.

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