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Key indicators

Single-crystal X-ray study T = 113 K Mean σ (C–C) = 0.002 Å R factor = 0.038 wR factor = 0.086 Data-to-parameter ratio = 19.1

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

The title compound, $C_{21}H_{21}NO_4S$, is the key intermediate in the synthesis of tazobactam. The crystal structure is stabilized by intermolecular $C-H\cdots O$ hydrogen bonds.

Diphenylmethyl 1-oxo-1-penicillanate

Comment

Tazobactam is a widely used β -lactamase inhibitor (Bai *et al.*, 2001). The title compound, (I), is the key intermediate for the synthesis of tazobactam and its structure is reported here (Fig. 1).



All bond lengths and angles in (I) are within normal ranges (Allen *et al.*, 1987). The crystal structure is stabilized by weak $C-H\cdots O$ intermolecular hydrogen bonds (Fig. 2 and Table 1).



Figure 1

The molecular structure of (I), showing the atom-numbering scheme and 30% probability displacement ellipsoids.

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organic papers

The five-membered thiazolidine ring (N1, C3, S1, C4 and C5) adopts an envelope conformation (Spek, 2003), with an r.m.s. deviation from the mean plane through atoms N1, C3, C4 and C5 of 0.019 Å and the S1 atom 0.830 (2) Å from that plane. The four-membered azetidine ring is in a distorted planar conformation, with an r.m.s. deviation from the plane through atoms N1, C1, C2 and C3 of 0.059 Å. The dihedral angle between the N1/C3/C4/C5 and N1/C1/C2/C3 planes is $42.29 (9)^{\circ}$. The two phenyl rings of the diphenylmethyl group are inclined at an angle of $77.13(5)^{\circ}$.

Experimental

The title compound was prepared according to the procedure of Bai et al. (2001), using chiral (+)-6-aminopenicillanic acid as the starting material. Colourless single crystals of (I) were grown by slow evaporation of an acetone solution.

Z = 4

Block, colorless

 $0.24 \times 0.22 \times 0.10 \ \mathrm{mm}$

Crystal data

C21H21NO4S $M_r = 383.45$ Orthorhombic, P212121 a = 7.9978 (14) Å b = 9.5959 (17) Å c = 24.020 (5) Å V = 1843.4 (6) Å³

Data collection

Rigaku Saturn diffractometer ω scans Absorption correction: numerical (NUMABS; Higashi, 1999) $T_{\min} = 0.940, T_{\max} = 0.980$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.038$ $wR(F^2) = 0.086$ S = 1.064730 reflections 248 parameters H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.0396P)^2$ + 0.2494P] where $P = (F_0^2 + 2F_c^2)/3$

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$\begin{array}{c} \hline C8-H8C\cdots O1^{i} \\ C15-H15\cdots O2^{ii} \\ C21-H21\cdots O2^{ii} \end{array}$	0.98	2.44	3.320 (2)	150
	0.95	2.40	3.160 (2)	137
	0.95	2.48	3.323 (2)	149

Symmetry codes: (i) x - 1, y, z; (ii) x, y - 1, z.



16269 measured reflections 4730 independent reflections 4400 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.044$ $\theta_{\rm max} = 28.7^{\circ}$

 $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.31 \text{ e } \text{\AA}^{-3}$ $\Delta \rho_{\rm min} = -0.27 \text{ e } \text{\AA}^{-3}$ Extinction correction: SHELXL97 Extinction coefficient: 0.0277 (16) Absolute structure: Flack (1983), 2021 Friedel pairs Flack parameter: -0.05 (6)



Figure 2 Packing diagram for (I), with $C-H\cdots O$ hydrogen bonds drawn as dashed lines

All H atoms were refined using a riding model, with C-H = 0.95 Å and $U_{iso}(H) = 1.2U_{ea}(C)$ for aromatic, C-H = 1.00 Å and $U_{iso}(H) =$ $1.2U_{eq}(C)$ for CH, \dot{C} -H = 0.99 Å and $U_{iso}(H) = 1.2U_{eq}(C)$ for CH₂ and C-H = 0.98 Å and $U_{iso}(H) = 1.5U_{eq}(C)$ for $CH_3 H$ atoms.

Data collection: CrystalClear (Rigaku/MSC, 2005); cell refinement: CrystalClear; data reduction: CrystalClear; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1997); software used to prepare material for publication: CrystalStructure (Rigaku/MSC, 2005) and PLATON (Spek, 2003).

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