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Key indicators

Single-crystal X-ray study T = 150 K Mean σ (C–C) = 0.003 Å R factor = 0.041 wR factor = 0.104 Data-to-parameter ratio = 17.4

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

(7*S*)-*N*-(4-Phenyltricyclo[4.2.1.0^{2,5}]non-3-ene-3-carbonyl)-2,10-camphorsultam

The regio- and absolute stereochemistry of the title compound, $C_{26}H_{31}NO_3S$, has been established by X-ray analysis. The molecular dimensions are in the normal range. In the crystal structure, weak $C-H\cdots O$ interactions link molecules, related by 2_1 screw axes, into extended tapes in the *b*-axis direction.

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Comment

Recently, we reported the first examples of asymmetric ruthenium-catalyzed [2 + 2]-cycloadditions between bicyclic alkenes and a chiral acetylenic acyl sultam (Villeneuve & Tam, 2004). Two different diastereoisomers could be formed in the cycloadditions. When norbornene was used as the bicyclic alkene component and phenyl–acetylene–acyl camphorsultam was used as the alkyne component, two diastereoisomers were obtained in a 131:1 ratio. These diastereoisomers were separated by fractional recystallization. The absolute stereo-chemistry of the major isomer, (I), was established by the single-crystal X-ray diffraction analysis described here.



A view of the title compound is shown in Fig. 1. The structure determination establishes the absolute configuration of the following atoms in Fig. 1: C2-S, C5-R, C8-R, C14-S, C15-R, C18-S and C19-R. The molecular dimensions, a selection of which are listed in Table 1, are in the normal range. Weak $C-H \cdots O$ interactions, the geometric details of which are given in Table 2, link molecules, related by 2_1 screw axes, into extended tapes in the b direction (Fig. 2). The hydrogenbonding motif consists of two main graph-set descriptors (Bernstein et al., 1995); C(6) chains are formed via the C1-H1B···O3ⁱ [symmetry code: (i) $1 - x, \frac{1}{2} + y, \frac{3}{2} - z$] interactions and $S_2^1(6)$ rings are fromed through the C3-H3B···O1ⁱⁱ and C8-H8A···O1ⁱⁱ [symmetry code: (ii) 1 - x, $y - \frac{1}{2}, \frac{3}{2} - z$] interactions. Both of these interactions propagate along the same 2_1 screw axes, but in opposite directions with respect to the polarity of the $C1 - H1B \cdot \cdot \cdot O3^{1}$ hydrogen bond.

Experimental

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Mo $K\alpha$ radiation

reflections

 $\mu = 0.18~\mathrm{mm}^{-1}$

T = 150 (1) K

 $\begin{aligned} R_{\rm int} &= 0.071\\ \theta_{\rm max} &= 27.5^\circ \end{aligned}$

 $h = -9 \rightarrow 9$

 $k = -15 \rightarrow 15$

 $l = -34 \rightarrow 34$

 $(\Delta/\sigma)_{\rm max} = 0.001$

 $\Delta \rho_{\rm max} = 0.23 \ {\rm e} \ {\rm \AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.29 \ {\rm e} \ {\rm \AA}^{-3}$

1990 Friedel pairs

Flack parameter = -0.02 (7)

Extinction correction: SHELXL97

Extinction coefficient: 0.0106 (16) Absolute structure: Flack (1983);

Cut needle, colourless $0.26 \times 0.20 \times 0.20$ mm

 $\theta = 2.6-27.5^{\circ}$

Cell parameters from 31 147

3962 reflections with $I > 2\sigma(I)$

Crystal data

 $C_{26}H_{31}NO_{3}S$ $M_{r} = 437.58$ Orthorhombic, $P2_{1}2_{1}2_{1}$ a = 7.2089 (1) Å b = 11.6317 (2) Å c = 26.3219 (5) Å $V = 2207.14 (6) \text{ Å}^{3}$ Z = 4 $D_{x} = 1.317 \text{ Mg m}^{-3}$

Data collection

Nonius KappaCCD diffractometer φ scans and ω scans with κ offsets Absorption correction: multi-scan (*SORTAV*; Blessing, 1995) $T_{min} = 0.921, T_{max} = 0.963$ 18 496 measured reflections 4876 independent reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.042$ $wR(F^2) = 0.104$ S = 1.044876 reflections 281 parameters H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.0524P)^2 + 0.0897P]$ $where P = (F_o^2 + 2F_c^2)/3$

Table 1

Selected geometric parameters (Å, °).

\$1-N1	1.6921 (17)	C12-C13	1.364 (3)
S1-C1	1.7885 (19)	C12-C19	1.517 (3)
N1-C11	1.407 (2)	C13-C14	1.516 (3)
N1-C8	1.486 (2)	C14-C19	1.573 (3)
02-\$1-01	117.76 (12)	C12-C13-C14	93.26 (15)
N1-S1-C1	95.61 (9)	C13-C14-C19	86.69 (13)
C11-N1-C8	116.39 (15)	C12-C19-C14	85.40 (13)
C13-C12-C19	94.63 (14)		

Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$C1-H1B\cdots O3^{i}$	0.99	2.41	3.361 (3)	161
$C3-H3B\cdots O1^{ii}$	0.99	2.51	3.377 (3)	146
$C8-H8A\cdotsO1^{ii}$	1.00	2.50	3.425 (2)	153
Symmetry codes: (i) 1	$-x, \frac{1}{2}+y, \frac{3}{2}-z$	z; (ii) $1 - x, y - x$	$\frac{1}{2}, \frac{3}{2} - z.$	

All H atoms were placed in calculated positions, with C–H distances ranging from 0.95 to 1.00 Å, and included in the refinement in a riding-model approximation, with $U_{\rm iso} = 1.2U_{\rm eq}(\rm C)$ (1.5 $U_{\rm eq}$ for methyl).

Data collection: *COLLECT* (Nonius, 1997–2002); cell refinement: *DENZO–SMN* (Otwinowski & Minor, 1997); data reduction: *DENZO–SMN*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2001); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.



Figure 1

View of the molecule of (I), showing the crystallographic labeling scheme. Displacement ellipsoids are drawn at the 30% probability level.



Figure 2

Packing diagram (Spek, 2004) showing weak intermolecular $C-H\cdots O$ interactions as dashed lines. S atoms are shown in green, O atoms in red, and N atoms in blue. All H atoms except H1*B*, H3*B* and H8*A* have been omitted for clarity

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