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Key indicators

Single-crystal X-ray study T = 299 KMean σ (C–C) = 0.003 Å R factor = 0.036 wR factor = 0.083 Data-to-parameter ratio = 9.0

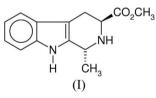
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. The title compound, $C_{14}H_{16}N_2O_2$, was obtained from the reaction between *S*-tryptophan methyl ester hydrochloride and acetaldehyde. The molecule adopts a *trans* configuration, with the methyl and methoxycarbonyl groups located on opposite sides of the central tetrahydro- β -carboline unit. Bifurcated intermolecular N-H···O/N hydrogen bonds link the molecules into chains.

hydro-β-carboline-3-carboxylate

trans-(1R,3S)-Methyl 1-methyl-1,2,3,4-tetra-

Comment

Alkaloids comprise a large and complex group of naturally occurring organic compounds, many of which contain indole or indoline groups. A large number of these molecules are biologically active. For example, vincristine and vinblastine from *Catharanthus roseus* (Johnson *et al.*, 1963) have long been established as anti-tumour alkaloids of clinical significance. Others, such as reserpine (Bein, 1953) and ajmaline (Petter & Engelmann, 1974), exhibit important cardiovascular effects. Tetrahydro- β -carbolines have been isolated from *Vinca theiodora* and other plants of South American origin and employed by Indian tribes as botanical source of intoxicating snuffs (Agurell *et al.*, 1969). Recently, tetrahydro- β -carbolines have been found in chocolate and cocoa (Herraiz, 2000). We report here the crystal structure of the title tetrahydro- β -carboline compound, (I).



The molecule adopts a *trans* configuration (Fig. 1), with the methyl and methoxycarbonyl groups located on opposite sides of the central tetrahydro- β -carboline unit. The geometry of the molecule is unexceptional, and closely comparable to that observed for the 1-(2-methylpropenyl) derivative (Bailey *et al.*, 2001). Hydrogen bonds are formed between molecules (Table 1), with N1-H1N acting as bifurcated donor to two acceptors, N2ⁱⁱⁱ and O1ⁱⁱⁱ (symmetry code in Table 1). These interactions generate chains running along the *a* axis.

Experimental

S-Tryptophan methyl ester hydrochloride (1.0 g, 0.004 mol) and acetaldehyde (2.2 ml, 0.004 mol) were dissolved in a methanol/water solution (23 ml, 75/25%, ν/ν). The mixture was refluxed for 12 h, after which time thin-layer chromatography indicated the presence of two new components [$R_{\rm F}$ = 0.36, 0.26, chloroform/methanol (9:1)] in the

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organic papers

reaction medium. The reaction mixture was cooled and the solvent evaporated under vacuum. The residue was dissolved in 14% ammonium hydroxide, extracted with chloroform and dried over anhydrous sodium sulfate. The solvent was removed under reduced pressure to leave an oil (yield 0.69 g, 72%). The diastereomeric *cis:trans* ratio was found to be 40:60 by ¹H NMR. The oil separated into ether-soluble and ether-insoluble parts. On crystallization from chloroform, the ether-soluble part gave the *cis* isomer as a light yellow solid [yield 0.23 g, 29%; m.p. 347–349 K; $R_{\rm F}$ 0.36, chloroform-methanol (9:1)]. Crystallization of the ether-insoluble part from methanol gave colourless crystals of the *trans* isomer, (I) [yield 0.46 g, 43%; m.p. 466–468 K; $R_{\rm F}$ 0.26, chloroform/methanol (9:1)].

Z = 4

 $D_x = 1.325 \text{ Mg m}^{-3}$

Mo Ka radiation

Block, colourless

 $0.60 \times 0.24 \times 0.14 \text{ mm}$

 $\mu = 0.09 \text{ mm}^{-1}$

T = 299 K

Crystal data

 $\begin{array}{l} C_{14}H_{16}N_2O_2\\ M_r = 244.29\\ \text{Orthorhombic, } P2_12_12_1\\ a = 7.9948 \ (10) \text{ \AA}\\ b = 9.6751 \ (10) \text{ \AA}\\ c = 15.8301 \ (8) \text{ \AA}\\ V = 1224.5 \ (2) \text{ \AA}^3 \end{array}$

Data collection

Bruker–Nonius KappaCCD diffractometer φ and ω scans Absorption correction: none 12651 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.036$ $wR(F^2) = 0.083$ S = 1.101468 reflections 164 parameters H-atom parameters constrained

1274 reflections with $I > 2\sigma(I)$ $R_{int} = 0.057$
$\theta_{\rm max} = 26.5^{\circ}$
$4 = \frac{2}{2} (\pi^2) + (0.022 \pi)^2$
$w = 1/[\sigma^2(F_o^2) + (0.032P)^2]$

1468 independent reflections

 $\begin{array}{l} & (1000 L^{2})^{+} (0.002L^{2})^{+} \\ & + 0.29P] \\ \text{where } P = (F_{o}^{-2} + 2F_{c}^{-2})/3 \\ (\Delta/\sigma)_{\max} < 0.001 \\ \Delta\rho_{\max} = 0.16 \text{ e } \text{\AA}^{-3} \\ \Delta\rho_{\min} = -0.14 \text{ e } \text{\AA}^{-3} \\ \text{Extinction correction: SHELXL97} \\ \text{Extinction coefficient: } 0.028 (4) \end{array}$

Table 1	I
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		0	
Hydrogen-bond	geometry	(À.	°).
	8	(,	

$D - \mathbf{H} \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$\begin{array}{c} N1 - H1 N \cdots O1^{i} \\ N1 - H1 N \cdots N2^{i} \end{array}$	0.92	2.18	2.976 (2)	145
	0.92	2.46	3.142 (2)	131

Symmetry code: (i) $x - \frac{1}{2}, -y + \frac{3}{2}, -z + 1$.

Due to the absence of significant anomalous scattering effects, Friedel pairs were merged prior to refinement. The absolute configuration was assigned on the basis of the unchanging *S* configuration at C11. Most H atoms were visible in difference Fourier maps, but those bound to C atoms were placed in calculated positions and allowed to ride during subsequent refinement with $U_{iso}(H) = 1.2U_{eq}(C)$ $[1.5U_{eq}(C)$ for methyl groups]. H atoms bound to N atoms were included in their as-found positions (N1-H1N = 0.92 and N2-H2N = 0.96 Å) and allowed to ride with $U_{iso}(H) = 1.2U_{eq}(N)$.

Data collection: *COLLECT* (Nonius, 1999); cell refinement: *DIRAX/LSQ* (Duisenberg, 1992); data reduction: *EVALCCD* (Duisenberg *et al.*, 2003); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *DIAMOND* (Brandenburg, 2006); software used to prepare material for publication: *maXus* (Mackay *et al.*, 1999).

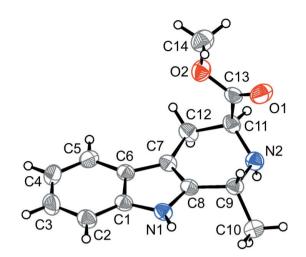


Figure 1

The molecular structure of (I), showing displacement ellipsoids drawn at the 50% probability level for non-H atoms.

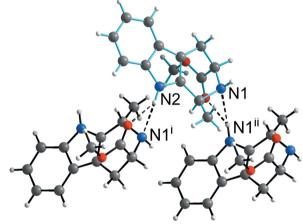


Figure 2

The hydrogen-bonding pattern in (I) (hydrogen bonds are shown as dashed lines). [Symmetry codes: (i) $x + \frac{1}{2}, -y + \frac{1}{2}, -z + 1$; (ii) $x - \frac{1}{2}, -y + \frac{1}{2}, -z + 1$.]

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