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Key indicators

Single-crystal X-ray study T = 295 K Mean σ (C–C) = 0.002 Å R factor = 0.046 wR factor = 0.133 Data-to-parameter ratio = 18.5

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

11-Phenyl-7,8,9,10,10a,11,12,13,14,15decahydro-5*H*-3,1-benzoxazino[2,1-e]acridine

2-Hydroxy-8-phenyltricyclo $[7.3.1.0^{2,7}]$ tridecan-13-one reacts with 2-aminobenzyl alcohol to form the title compound, C₂₆H₂₉NO, which crystallizes with two independent molecules in the asymmetric unit. The N atom shows a nearly planar coordination owing to the strain imposed on the fused-ring compound.

Comment

2-Hydroxy-8-phenyltricyclo[7.3.1.0^{2,7}]tridecan-13-one (trivial name: phenyl ketol) condenses through its carbonyl group with primary amines to form Schiff bases that undergo cyclization owing to the proximity of the hydroxy group, as noted with, for example, aniline and substituted anilines, which afford 9-phenyl-10-organyldecahydroacridines (Alekseev et al., 1976; Kruglyakova et al., 1977). The reaction of phenyl ketol with o-aminobenzyl alcohol probably yields as the initial product 10-(2-hydroxymethylphenyl)-1,2,3,4,5,-6,7,8,9,10-decahydro-9-phenylacridine; the -CH₂OH group then adds across one of the two double bonds, giving rise to the title compound, (I). The compound crystallizes with two independent molecules in the asymmetric unit; the bond dimensions in both are normal, and there are no important intermolecular interactions in the crystal structure. The compound is expected to undergo oxidative coupling with methylene-active compounds in the presence of manganese dioxide (Slabko et al., 1990).



Experimental

Commercially available phenyl ketol (2.0 g, 7.0 mmol) was dissolved in *p*-xylene (50 ml) and to the solution was added *o*-aminobenzyl alcohol (1.0 g, 8.1 mmol). The reaction was catalysed by *p*-toluenesulfonic acid (5 mg); the mixture was heated for 35 h in a Dean–Stark apparatus. On completion of the reaction, the solvent was removed under reduced pressure. The oily residue was triturated with ethanol (10 ml) to yield a solid compound that was purified by column chromatography on alumina, with *n*-hexane as eluant, in about 30% yield (0.81 g; m.p. 435–437 K). The molecular weight of the compound was established by mass spectrometry.

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Figure 1

View of one of the two independent molecules of (I). Displacement ellipsoids are drawn at the 50% probability level and H atoms are drawn as spheres of arbitrary radii.

Crystal data

 $\begin{array}{l} C_{26}H_{29}NO\\ M_r = 371.50\\ Triclinic, P\overline{1}\\ a = 10.653 \ (1) \ \text{\AA}\\ b = 11.427 \ (1) \ \text{\AA}\\ c = 17.337 \ (2) \ \text{\AA}\\ a \approx 91.124 \ (2)^{\circ}\\ \beta = 100.810 \ (2)^{\circ}\\ \gamma = 100.191 \ (2)^{\circ}\\ V = 2037.4 \ (4) \ \text{\AA}^{3} \end{array}$

Z = 4 $D_x = 1.211 \text{ Mg m}^{-3}$ Mo K\alpha radiation Cell parameters from 956 reflections $\theta = 2.4-27.6^{\circ}$ $\mu = 0.07 \text{ mm}^{-1}$ T = 295 (2) K Prism, colorless $0.40 \times 0.35 \times 0.25 \text{ mm}$ Data collection

H-atom parameters constrained

Bruker SMART 1000 CCD area- detector diffractometer φ and ω scans Absorption correction: None 25549 measured reflections 9331 independent reflections	6699 reflections with $I > 2\sigma(I)$ $R_{int} = 0.033$ $\theta_{max} = 27.5^{\circ}$ $h = -13 \rightarrow 13$ $k = -14 \rightarrow 14$ $l = -22 \rightarrow 22$
Refinement	
Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.046$ $wR(F^2) = 0.133$ S = 1.01 9331 reflections	$w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0534P)^{2} + 0.6554P]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3 (\Delta/\sigma)_{max} = 0.001 \Delta\rho_{max} = 0.49 \text{ e} \text{ Å}_{a}^{-3}$
505 parameters	$\Delta \rho_{\rm min} = -0.28 \ {\rm e} \ {\rm A}^{-3}$

H atoms were placed at calculated positions [C-H = 0.93-0.96 Å]and $U_{iso}(H) = 1.2U_{eq}(C)]$ and were included in the refinement in the riding-model approximation.

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1998); software used to prepare material for publication: *SHELXL97*.

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