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Key indicators

Single-crystal X-ray study T = 273 K Mean σ (C–C) = 0.004 Å R factor = 0.061 wR factor = 0.139 Data-to-parameter ratio = 13.1

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. The title compound, 5-(4-nitrophenyl)-3,4,4a,5,6,10b-hexahydro-2*H*-pyrano[3,2-*c*]quinoline, $C_{18}H_{18}N_2O_3$, was obtained from the Diels–Alder reaction of *N*-benylideneaniline and dihydropyran, catalyzed by zirconium tetrachloride. This molecule is an *exo*-cycloadduct isomer. The pyran ring adopts a chair conformation, while the *N*-heterocyclic ring prefers a half-chair conformation. Molecules are associated into centrosymmetric dimers by N–H···O hydrogen bonds.

A diastereoisomer of pyrano[3,2-c]quinoline

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Imino Diels–Alder adducts. Part V

Comment

Quinoline derivatives, an important class of pharmaceuticals, exhibit psychotropic, anti-allergenic and anti-inflammatory activities (Nesterova *et al.*, 1995; Yamada *et al.*, 1992; Faber *et al.*, 1984). In addition, the pyranoquinoline moiety is present in many alkaloids. For the synthesis of pyranoquinolines, the imino Diels–Alder reaction between *N*-arylimines and dienophile dihydropyran is probably one of the most successful synthetic tools (Mahesh *et al.*, 2004). We report here the structure of a diastereoisomer, namely 5-(4-nitrophenyl)-3,4,4a,5,6,10b-hexahydro-2*H*-pyrano[3,2-*c*]quinoline, (I), as part of our ongoing structure elucidation studies.



The molecular dimensions of (I) are normal and compare well with similar structures reported in the literature (Ravikumar *et al.*, 2004; Wang *et al.*, 2004). The orientation of atoms H4 and H5, the characteristic differentiating the diastereoisomers, results in a *trans* (*exo*) configuration, the H4–C4– C5–H5 torsion angle [–169.8 (2)°] being in accordance with the coupling constant J = 10.7 Hz. Furthermore, atoms H4 and H12 are oriented in a *cis* configuration [H4–C4–C12–H12 = 49.8 (3)°].

The strain exerted during the cycloaddition process, in forming the *N*-heterocyclic ring, may be seen from the torsion angles C6-C11-C12-C4 $[-15.0 (3)^{\circ}]$ and C6-N1-C5-C4 $[53.5 (3)^{\circ}]$. The N-heterocyclic ring is in a half-chair conformation, with atoms C6 and C11 displaced by 0.057 (3) and -0.088 (2) Å, respectively, from the plane formed by the other four atoms.

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Figure 1

A view of the molecule, with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii.



Figure 2

Packing diagram viewed down the *b* axis. Dashed lines indicate $N-H\cdots O$ hydrogen bonds.

Interestingly, the pyran ring is oriented perpendicular to the quinoline ring, with the torsion angle $C6-C11-C12-O1 = 108.3 (2)^{\circ}$. The pyran ring adopts a chair conformation, as expected, with asymmetry parameter (Nardelli, 1983) $\Delta C_2(C4-C2) = 0.034$ (3). Atoms C3 and O1 are displaced by 0.602 (3) Å above and 0.647 (2) Å below the plane defined by atoms C1/C2/C4/C12 of the pyran ring.

The sum of the angles around atom N2 (359°) shows a planar configuration, with an average N–O bond length of 1.216 (3) Å. The absence of steric effects between the NO₂ group and the quinoline moiety accounts for the small dihedral angle of 4.4 (1)° between the nitro group and its attached ring. The benzene ring C13–C18 is rotated with respect to the plane of the quinoline ring by 74.5 (1)° about the C5–C13 bond.

Intermolecular hydrogen bonds between the NO₂ and quinoline groups are responsible for the formation of dimers between molecules related by an inversion centre. A possible weak interaction of the type $C-H\cdots O$ is present (Table 2). Edge-to-edge stacking of the benzene ring is also observed, as

indicated by the distance of 3.766 (2) Å between the C16···C18ⁱ atoms [symmetry code: (i) 2 - x, 2 - y, 2 - z].

Experimental

To a solution of the appropriate *N*-benzylideneaniline (5.5 mmol) in dichloromethane (5 ml) at room temperature were added 2,3-dihydrofuran (5.5 mmol) and $ZrCl_4$ (10 mol%) and the resulting mixture was stirred for 90 min. The completed reaction was quenched with water and the crude product was purified by column chromatography using 2% ethyl acetate and hexane to yield the title compound. Crystals for X-ray study were obtained by recrystallization from a methanol–water (3:1) solution.

Crystal data

$C_{18}H_{18}N_2O_3$	Z = 2
$M_r = 310.34$	$D_x = 1.309 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 9.5001 (10) Å	Cell parameters from 2569
b = 9.8823 (10) Å	reflections
c = 9.9403 (10) Å	$\theta = 2.2-26.7^{\circ}$
$\alpha = 98.039 \ (2)^{\circ}$	$\mu = 0.09 \text{ mm}^{-1}$
$\beta = 115.157 \ (2)^{\circ}$	T = 273 (2) K
$\gamma = 104.092 \ (2)^{\circ}$	Block, colorless
$V = 787.18 (14) \text{ Å}^3$	$0.20\times0.15\times0.10~\mathrm{mm}$

2344 reflections with $I > 2\sigma(I)$

 $\begin{aligned} R_{\rm int} &= 0.021 \\ \theta_{\rm max} &= 25.0^\circ \end{aligned}$

 $h = -11 \rightarrow 11$

 $k=-11\rightarrow 11$

 $l = -11 \rightarrow 11$

Data collection

Bruker SMART APEX CCD areadetector diffractometer ω scans Absorption correction: none 7650 measured reflections 2770 independent reflections

Refinement

Table 1

Selected geometric parameters (Å, °).

N1-C6	1.385 (3)	N2-O3	1.217 (3)
N1-C5	1.457 (3)	N2-C16	1.466 (3)
N2-O2	1.214 (3)		
C6-N1-C5	118.0 (2)	O2-N2-C16	118.6 (2)
O2-N2-O3	122.9 (2)	O3-N2-C16	118.5 (2)

Table 2Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	<i>D</i> -H	$H \cdots A$	$D \cdots A$	$D - H \cdots A$
$N1 - H1N \cdots O3^{i}$ $C1 - H1B \cdots O3^{ii}$	0.84 (3) 0.97	2.38 (3) 2.59	3.200 (3) 3.535 (3)	165 (2) 166
Symmetry codes: (i) 2	x - x, 2 - y, 2 - y	z; (ii) $x - 1, y - 1$	1. <i>z</i> .	

The H atom attached to the N atom was located in a difference density map and refined isotropically. All other H atoms were positioned geometrically and refined as riding, with C-H distances in the range 0.93–0.98 Å and with $U_{\rm iso} = 1.5U_{\rm eq}(C)$ for methyl H and $1.2U_{\rm eq}(C)$ for other H atoms.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL/PC* (Sheldrick, 1990); software used to prepare material for publication: *SHELXL97* and *PARST* (Nardelli, 1995).

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