

4-(4-Nitrophenyl)-2,3,3a,4,5,9b-hexahydro-furo[3,2-c]quinoline

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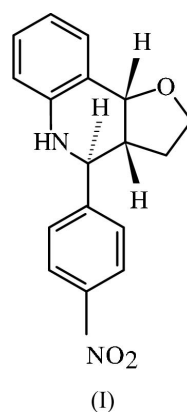
Key indicators

Single-crystal X-ray study
 $T = 273$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.056
 wR factor = 0.163
Data-to-parameter ratio = 16.3For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title compound, $\text{C}_{17}\text{H}_{16}\text{N}_2\text{O}_3$, is one of the diastereoisomers formed as the result of the Diels–Alder reaction of 1-(4-nitrophenyl)methylidene]-*N*-phenylamine and dihydrofuran catalyzed by zirconium tetrachloride. The furan ring adopts an envelope conformation, while the *N*-heterocyclic ring prefers a sofa conformation. Molecules are linked into centrosymmetric dimers by $\text{N}-\text{H}\cdots\text{O}$ hydrogen-bond interactions.

Comment

Furoquinoline derivatives have been studied extensively for a range of biological applications, such as psychotropic (Nesterova *et al.*, 1995), anti-allergic (Yamada *et al.*, 1992) and anti-inflammatory (Faber *et al.*, 1984) applications. We have recently synthesized and purified pyrano- and furoquinolines using the imino Diels–Alder reaction with ZrCl_4 as a green catalyst (Mahesh *et al.*, 2004). The structure of the title compound, (I), is reported here as part of our ongoing structural studies on this series of compounds (Ravikumar *et al.*, 2004, 2005).



A view of (I) is shown in Fig. 1. In all essential details, the molecular geometry (Table 1) is in good agreement with relevant structures found in the literature (Ma *et al.*, 1999; Batey *et al.*, 2001; Hoemann *et al.*, 2002). Generally, these diastereomers differ according to the stereochemistry (*cis/trans*) of the ring junction H atoms and the orientation (*endo/exo*) of the furan ring. The $\text{H}7-\text{C}7-\text{C}8-\text{H}8$ torsion angle of -174.4° indicates the *trans* (*exo*) configuration and is in accordance with the coupling constant $J = 13.4$ Hz. Similarly, the $\text{H}4-\text{C}4-\text{C}7-\text{H}7$ torsion angle of 35.5° indicates the *cis* arrangement of the H atoms at the furan ring junction.

The values of the $\text{C}8-\text{C}7-\text{C}4-\text{C}3$ [34.6 (2°)] and $\text{C}8-\text{N}1-\text{C}9-\text{C}3$ [-26.8 (3°)] torsion angles are due to the strain exerted during the cycloaddition process in forming the *N*-

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Imino Diels–Alder adducts.
VI.

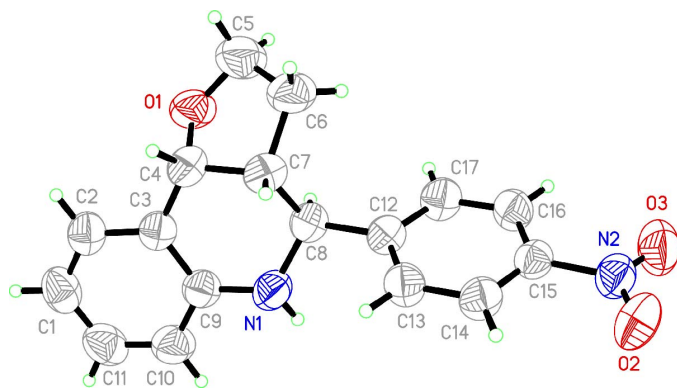


Figure 1
A view of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

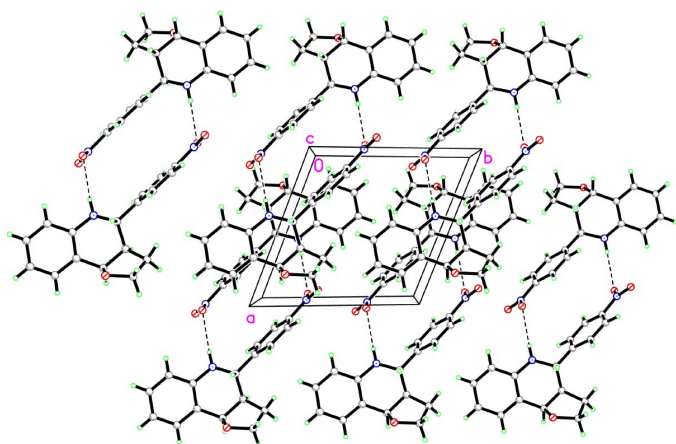


Figure 2
Packing diagram, viewed down the *c* axis, showing the dimers. Dashed lines indicate N—H...O hydrogen bonds.

heterocyclic ring. The *N*-heterocyclic ring of the quinoline moiety adopts a sofa conformation [asymmetry parameter $\Delta C_2(C3-C9) = 0.064(1)$; Nardelli, 1983], with atom C8 displaced by $0.626(2)$ Å from the mean plane defined by the atoms N1, C9, C3, C4 and C7.

The furan ring is oriented perpendicular to the quinoline ring, with atom O1 displaced by $-1.374(1)$ Å from the N1/C9/C3/C4/C7 least-squares plane. The furan ring adopts an envelope conformation [asymmetry parameter $\Delta C_s(C4) = 0.008(1)$].

The absence of steric effects between the NO₂ group and the quinoline moiety accounts for the small dihedral angle of $6.9(2)^\circ$ between the nitro group and its attached ring. The phenyl substituent at C8 is rotated about the C8—C12 bond by $74.2(1)^\circ$.

Intermolecular hydrogen bonds between the NO₂ and quinoline groups lead to the formation of dimers between molecules related by an inversion center (Table 2), generating a characteristic $R_2^2(18)$ -type motif (Bernstein *et al.*, 1995).

Experimental

To a solution of 1-(4-nitrophenyl)methylidene-*N*-phenylamine (5.5 mmol) in dichloromethane (5 ml) at room temperature were

added 2,3-dihydrofuran (5.5 mmol) and ZrCl₄ (10 mol%) with stirring for a period of 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 suitable for X-ray study were obtained by recrystallization from a methanol–water (3:1) solution.

Crystal data

C₁₇H₁₆N₂O₃
 $M_r = 296.32$
 Triclinic, $P\bar{1}$
 $a = 9.1090(10)$ Å
 $b = 9.2626(10)$ Å
 $c = 9.7553(11)$ Å
 $\alpha = 103.411(2)^\circ$
 $\beta = 102.791(2)^\circ$
 $\gamma = 107.081(2)^\circ$
 $V = 727.28(14)$ Å³

$Z = 2$
 $D_x = 1.353$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 2986 reflections
 $\theta = 2.4$ – 27.8°
 $\mu = 0.09$ mm⁻¹
 $T = 273(2)$ K
 Block, yellow
 $0.20 \times 0.15 \times 0.10$ mm

Data collection

Bruker SMART APEX CCD area-detector diffractometer
 ω scans
 Absorption correction: none
 8354 measured reflections
 3308 independent reflections

2549 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.017$
 $\theta_{max} = 28.0^\circ$
 $h = -11 \rightarrow 12$
 $k = -12 \rightarrow 12$
 $l = -12 \rightarrow 12$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.056$
 $wR(F^2) = 0.163$
 $S = 1.03$
 3308 reflections
 203 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0819P)^2 + 0.1431P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} < 0.001$
 $\Delta\rho_{max} = 0.39$ e Å⁻³
 $\Delta\rho_{min} = -0.16$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

O2—N2	1.221(2)	N1—C8	1.453(2)
O3—N2	1.224(2)	N2—C15	1.472(2)
N1—C9	1.389(2)		
C9—N1—C8	118.39(14)	O2—N2—C15	118.52(16)
O2—N2—O3	123.25(17)	O3—N2—C15	118.21(16)

Table 2

Hydrogen-bond geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1N...O2 ⁱ	0.84(2)	2.34(2)	3.164(2)	171(2)

Symmetry code: (i) $-x, -y, -z + 1$.

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

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