

## 3-(4-Methoxyphenyl)-1',6,6-trimethyl-4,5,6,7-tetrahydrospiro[1-benzofuran-2,3'(3H,2'H)-1H-indole]-2',4-dione

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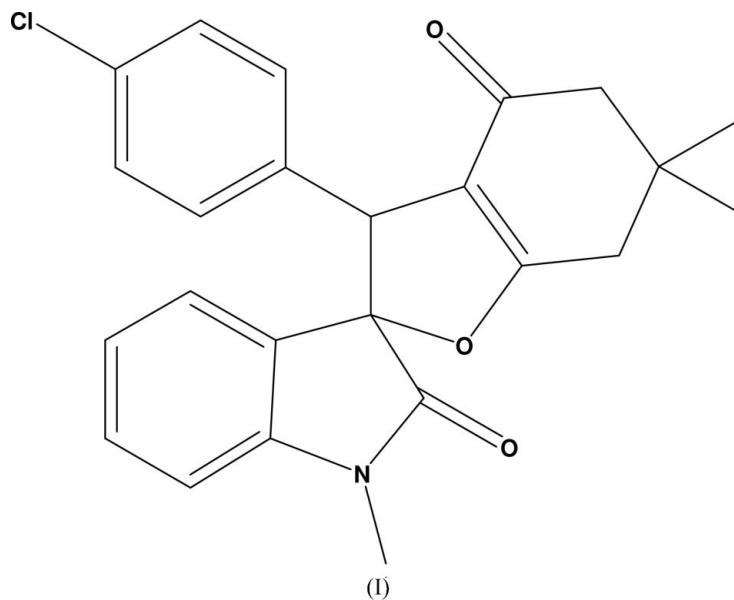
## Key indicators

Single-crystal X-ray study  
 $T = 293$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å  
 $R$  factor = 0.047  
 $wR$  factor = 0.138  
Data-to-parameter ratio = 18.7For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title compound,  $\text{C}_{24}\text{H}_{22}\text{ClNO}_3$ , contains two independent molecules in the asymmetric unit. In one of the molecules, the furan ring is in a twist conformation, whereas in the other it adopts an envelope conformation. In both molecules, the cyclohexenone rings adopt envelope conformations. The crystal structure is stabilized by intermolecular  $\text{C}-\text{H}\cdots\text{O}$  interactions.

## Comment

Spiro compounds represent an important class of naturally occurring substances characterized by highly pronounced biological properties (Kobayashi *et al.*, 1991). The spiro-oxindole ring system is found in a number of alkaloids, such as horsifiline, spiro-tryprostatin and (+)-elacomine (Hilton *et al.*, 2000). Spiro-oxindole derivatives find wide biological application as antimicrobial and antitumour agents and as inhibitors of the human NK1 receptor (Kornet & Thio, 1976). We report here the crystal structure of the title compound, (I).

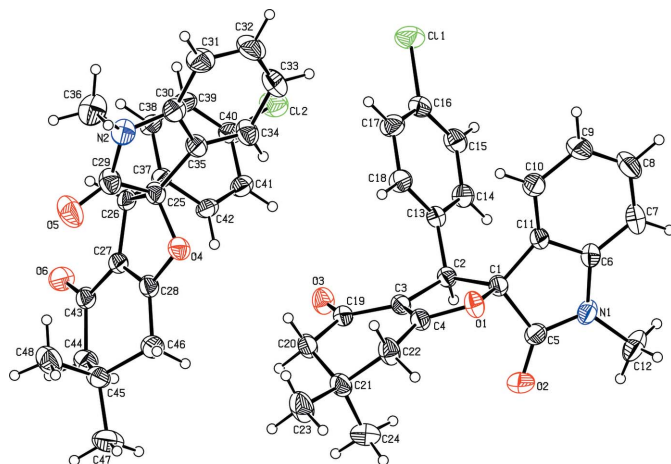


The asymmetric unit of (I) contains two molecules, *A* (C11/N1/O1–O3/C1–C24) and *B* (C12/N2/O4–O6/C25–C48). The corresponding bond lengths and angles of these two molecules agree with each other and show normal values (Allen *et al.*, 1987), except at the spiro junctions. In both molecules *A* and *B*, the N atom exhibits  $sp^2$  hybridization.

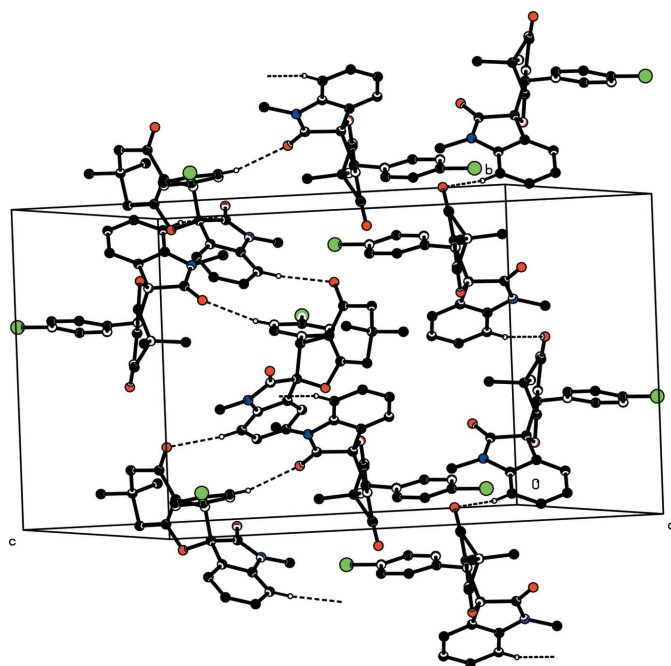
In molecule *A*, the dihydropyrrolone ring adopts an envelope conformation, with atom C1 at the flap [deviation 0.138 (1) Å], whereas in molecule *B* this ring is planar. The

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**Figure 1**  
The asymmetric unit of (I), showing 30% probability displacement ellipsoids.



**Figure 2**  
The crystal packing of (I). Dashed lines indicate hydrogen bonds and H atoms not involved in the interactions have been omitted.

puckering parameters (Cremer & Pople, 1975) and the smallest displacement asymmetry parameters (Nardelli, 1983) are  $q_2 = 0.084$  (2) Å,  $\varphi = 247.0$  (1)° and  $\Delta_2(C_3) = 3.0$  (2). In molecule *A*, the furan ring adopts a twist conformation, with a pseudo-twofold axis passing through atom C3 and the midpoint of the O1—C1 bond [ $q_2 = 0.130$  (1) Å and  $\varphi = 206.4$  (7)°]. In molecule *B*, the furan ring adopts an envelope conformation, with atom C25 deviating by 0.374 (2) Å from the mean plane of the rest of the atoms in that ring [ $q_2 = 0.234$  (2) Å,  $\varphi = 38.6$  (4)° and  $\Delta_s(C_{25}) = 1.1$  (2)]. In both molecules *A* and *B*, the cyclohexenone rings adopt envelope

conformations, with atoms C21 and C45 deviating by 0.649 (2) and 0.641 (2) Å, respectively, from the mean plane through the remaining atoms in the corresponding rings. The puckering parameters  $q_2$ ,  $q_3$ ,  $Q$  and  $\theta$ , respectively, are 0.383 (2) Å, 0.261 (2) Å, 0.463 (2) Å and 55.7 (2)° for the C3/C4/C19—C22 ring, and 0.385 (2) Å, −0.247 (2) Å, 0.458 (2) Å and 122.7 (2)° for the C27/C28/C43—C46 ring. The dihedral angles between the two benzene rings in molecules *A* and *B* are 48.5 (1)° and 53.6 (1)°, respectively.

In the crystal structure of (I), molecules *A* are linked *via* C7—H7···O3<sup>i</sup> hydrogen bonds, generating a zigzag C(9) chain (Bernstein *et al.*, 1995) running along the *b* axis. Similarly, molecules *B* are linked *via* C31—H31···O6<sup>iii</sup> hydrogen bonds to form another C(9) chain running along the *b* axis. These chains are cross-linked by C15—H15···O5<sup>ii</sup> hydrogen bonds (Fig. 2). (Symmetry codes are given in Table 2.)

## Experimental

To a stirred mixture of 3-(4-chlorobenzylidene)-1-methyloxindole (1.11 mmol, 0.3 g, 1 equivalent), dimedone (1.11 mmol, 0.154 g, 1 equivalent) and NaHCO<sub>3</sub> (3.33 mmol, 0.277 g, 3 equivalents) in acetonitrile (10 ml), ceric ammonium nitrate (2.78 mmol, 1.52 g, 2.5 equivalents) dissolved in acetonitrile (5 ml) was added dropwise at 273 K under an N<sub>2</sub> atmosphere. The reaction mixture was stirred until the reaction was complete, as monitored by thin-layer chromatography. Water was added to the mixture, and the product was extracted into ethyl acetate (2 × 20 ml) and then dried over anhydrous Na<sub>2</sub>SO<sub>4</sub>. Removal of the solvent under reduced pressure gave a crude mixture of isomers, which was purified and separated by column chromatography on silica gel (100–200 mesh), with ethyl acetate–hexane (3:9) as eluent, to afford the isomers (1:1) in an overall yield of 0.294 g (65%). One of the isomers, which was obtained as a white crystalline solid, was recrystallized from ethyl acetate and used for the X-ray diffraction study.

### Crystal data

C <sub>24</sub> H <sub>22</sub> ClNO <sub>3</sub>	$Z = 8$
$M_r = 407.88$	$D_x = 1.286$ Mg m <sup>−3</sup>
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
$a = 13.9638$ (8) Å	$\mu = 0.21$ mm <sup>−1</sup>
$b = 13.8490$ (8) Å	$T = 293$ (2) K
$c = 22.1754$ (13) Å	Block, colourless
$\beta = 100.851$ (1)°	0.26 × 0.25 × 0.23 mm
$V = 4211.7$ (4) Å <sup>3</sup>	

### Data collection

Bruker SMART APEX CCD area-detector diffractometer	9908 independent reflections
$\omega$ scans	7812 reflections with $I > 2\sigma(I)$
Absorption correction: none	$R_{int} = 0.022$
47537 measured reflections	$\theta_{max} = 28.0^\circ$

### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0754P)^2 + 0.8149P]$
$R[F^2 > 2\sigma(F^2)] = 0.047$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.138$	$(\Delta/\sigma)_{max} = 0.001$
$S = 1.02$	$\Delta\rho_{max} = 0.28$ e Å <sup>−3</sup>
9908 reflections	$\Delta\rho_{min} = -0.27$ e Å <sup>−3</sup>
529 parameters	
H-atom parameters constrained	

**Table 1**  
Selected bond lengths (Å).

C25—O4	1.461 (2)	C43—O6	1.223 (2)
C25—C35	1.503 (2)	C1—O1	1.455 (2)
C25—C29	1.550 (2)	C1—C11	1.493 (2)
C25—C26	1.571 (2)	C1—C5	1.542 (2)
C26—C37	1.510 (2)	C1—C2	1.580 (2)
C26—C27	1.511 (2)	C2—C3	1.506 (2)
C27—C28	1.335 (2)	C2—C13	1.512 (2)
C28—O4	1.357 (2)	C3—C4	1.330 (2)
C29—O5	1.212 (2)	C4—O1	1.353 (2)
C30—N2	1.402 (2)	C5—O2	1.213 (2)
C36—N2	1.455 (2)	C5—N1	1.353 (2)
C40—Cl2	1.740 (2)	C6—N1	1.407 (2)

**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>
C7—H7...O3 <sup>i</sup>	0.93	2.42	3.337 (3)	169
C15—H15...O5 <sup>ii</sup>	0.93	2.47	3.295 (2)	148
C31—H31...O6 <sup>iii</sup>	0.93	2.41	3.299 (3)	160

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

All H atoms were placed in idealized positions and refined using a riding model, with C—H = 0.93 (aromatic), 0.98 (CH), 0.97 (CH<sub>2</sub>) or 0.96 Å (methyl), and with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  or  $1.5U_{\text{eq}}(\text{C})$  for the methyl groups. A rotating-group model was used for the methyl groups.

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: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97* and *PARST* (Nardelli, 1995).

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