

An analysis of C—H···O and C—H··· π interactions in three substituted 4-ketotetrahydroindoles

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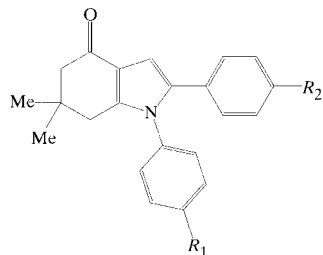
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The crystal and molecular structures of the three 4-ketotetrahydroindoles 2-(4-chlorophenyl)-1-(4-fluorophenyl)-6,6-dimethyl-4,5,6,7-tetrahydro-1*H*-indol-4-one (C₂₂H₁₉ClFNO), (I), 1-(4-fluorophenyl)-2-(4-methoxyphenyl)-6,6-dimethyl-4,5,6,7-tetrahydro-1*H*-indol-4-one (C₂₃H₂₂FNO₂), (II), and 6,6-dimethyl-1,2-diphenyl-4,5,6,7-tetrahydro-1*H*-indol-4-one (C₂₂H₂₁NO), (III), have been determined *via* single-crystal X-ray diffraction in order to study the intermolecular interactions therein. All three structures are stabilized *via* intermolecular C—H···O and C—H··· π interactions, generating different molecular motifs.

Comment

The generation of different types of packing motifs in crystalline lattices has been illustrated in detail in the literature. Different types of intermolecular interactions, such as hydrogen bonding (Desiraju, 1989, 2000, 2002; Aakeröy, 1997; Guru Row, 1999; Hunter *et al.*, 2001), C—H··· π interactions



(I) $R_1 = \text{F}$, $R_2 = \text{Cl}$

(II) $R_1 = \text{F}$, $R_2 = \text{OMe}$

(III) $R_1 = R_2 = \text{H}$

(Nishio *et al.*, 1995; Umezawa *et al.*, 1999; Takahashi *et al.*, 2000), π - π stacking (Hunter, 1993, 1994) and C—H···O hydrogen bonding (Desiraju & Steiner, 1999; Steiner, 2002) have been reviewed extensively. Recently, it has been shown that 'organic fluorine' provides similar interactions, generating

packing motifs in the presence or absence of other intermolecular forces (Prasanna & Guru Row, 2000*a,b,c*, 2001). We have been interested in the study of the role of organic fluorine in the packing of organic molecules which show biological activity (Choudhury, Urs, Guru Row & Nagarajan, 2002; Choudhury, Urs, Smith *et al.*, 2002). Against this background, we report here the crystal and molecular structures of 2-(4-chlorophenyl)-1-(4-fluorophenyl)-, (I), and 1-(4-fluorophenyl)-2-(4-methoxyphenyl)-6,6-dimethyl-4,5,6,7-tetrahydro-1*H*-indol-4-one, (II), and compare these with the corresponding unsubstituted compound 6,6-dimethyl-1,2-diphenyl-4,5,6,7-tetrahydro-1*H*-indol-4-one, (III) (Varghese *et al.*, 1986), the structure of which we have redetermined.

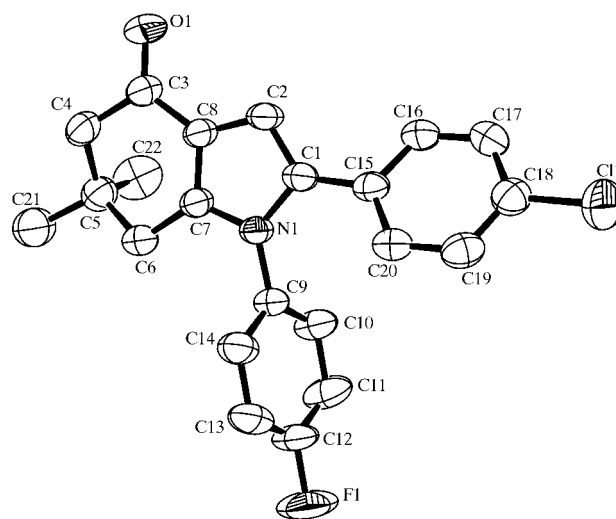


Figure 1

A view of the molecule of (I), showing the atom-numbering scheme and displacement ellipsoids drawn at the 50% probability level. H atoms have been omitted for clarity.

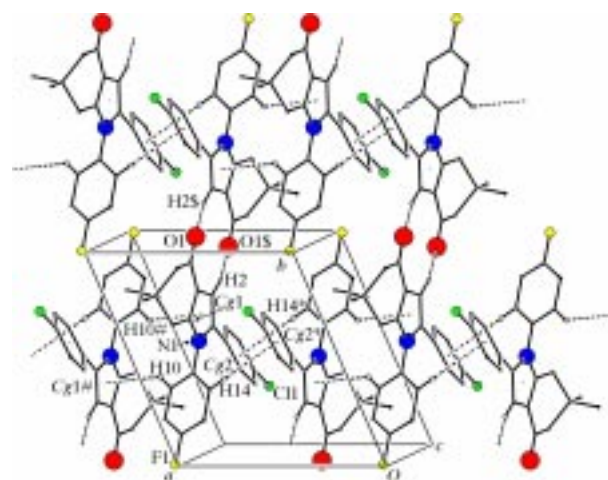


Figure 2

A packing diagram for (I), showing the C—H···O hydrogen-bonded dimer and the intermolecular C—H··· π interaction. Cg1 is the centroid of the five-membered N1/C1/C2/C8/C7 ring and Cg2 is the centroid of the C9—C14 phenyl ring. Atoms labelled with a dollar sign (\$) or hash (#) are at symmetry positions $(1 - x, 2 - y, 1 - z)$, $(2 - x, 1 - y, 1 - z)$ and $(1 - x, 1 - y, 1 - z)$, respectively.

Most unsubstituted 4-ketotetrahydroindoles have been found to be biologically inactive (Remers *et al.*, 1971) and this inactivity was attributed to the extended conjugation of the C=O group with the indole N atom (Weiss *et al.*, 1968). In order to delocalize the lone pair of electrons on N and to facilitate the activity of the keto function, a series of 1,2-diaryl derivatives was synthesized (Nagarajan *et al.*, 1985). It was found that differently substituted phenyl groups on the 1- or 2-positions facilitate biological activity. Compound (I) of the present study was inactive, whereas (II) showed considerable activity, as reported by Nagarajan *et al.* (1985). We have redetermined the structure of the unsubstituted compound, (III), since the intermolecular interactions were not discussed in detail in the earlier report of Varghese *et al.* (1986).

In compound (I), the six-membered ring of the tetrahydroindole system (Fig. 1) has an envelope conformation, with atom C5 a distance of $-0.641(3)$ Å from the C3/C4/C6/C7/C8 plane. Selected torsion angles are given in Table 1. The

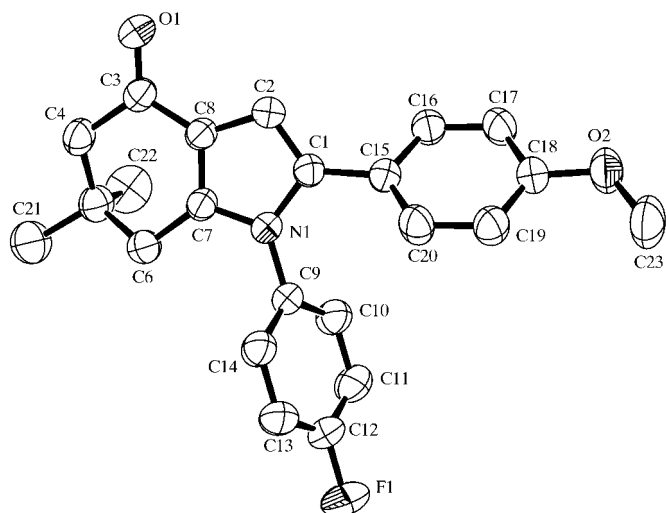


Figure 3
A view of the molecule of (II), showing the atom-numbering scheme and displacement ellipsoids drawn at the 50% probability level. H atoms have been omitted for clarity.

dihedral angle between the 4-fluorophenyl and 4-chlorophenyl rings is $69.84(7)^\circ$. The packing of the molecules in the crystal lattice (Fig. 2) is stabilized *via* C—H...O and C—H... π interactions (Table 2). There are two independent C—H... π interactions, which link the molecules in the *b* direction. Thus, molecules packed around the inversion centres at $(\frac{1}{2}, \frac{1}{2}, \frac{1}{2})$ and $(1, \frac{1}{2}, \frac{1}{2})$ are linked by pairs of independent C—H... π interactions. C—H...O interactions about the inversion centre at $(\frac{1}{2}, 1, \frac{1}{2})$ then serve to link these chains to form sheets of molecules in the *bc* plane. It is of interest to note that there are no interactions involving the halogen atoms, though the structure contains Cl as well as F.

In compound (II), the six-membered ring of the tetrahydroindole system (Fig. 3) also has an envelope conformation, with atom C5 a distance of $-0.618(2)$ Å from the C3/C4/C6/C7/C8 plane, similar to what was observed for (I). Selected torsion angles are given in Table 3. The dihedral angle between the 4-fluorophenyl and 4-methoxyphenyl rings is $60.94(4)^\circ$. The molecules pack in the unit cell (Fig. 4) *via* two different C—H...O interactions (Table 4) in the $(10\bar{1})$ plane,

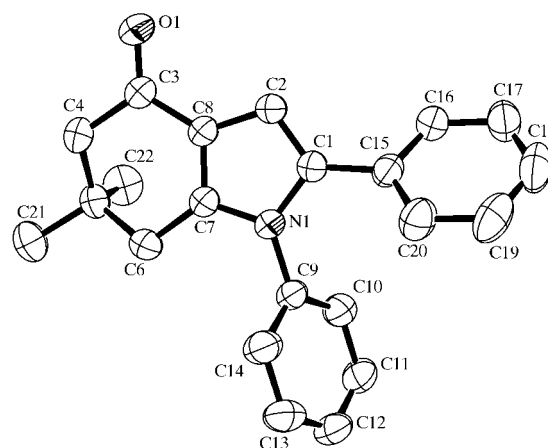


Figure 5
A view of the molecule of (III), showing the atom-numbering scheme and displacement ellipsoids drawn at the 50% probability level. H atoms have been omitted for clarity.

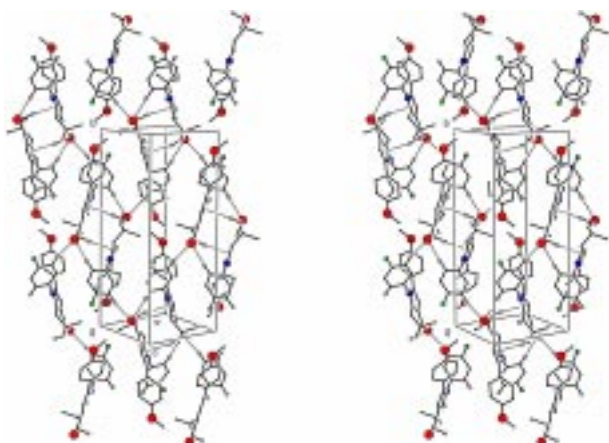


Figure 4
A stereoview of the packing of (II), showing the C—H...O hydrogen-bonded sheet motif and C—H... π interactions.

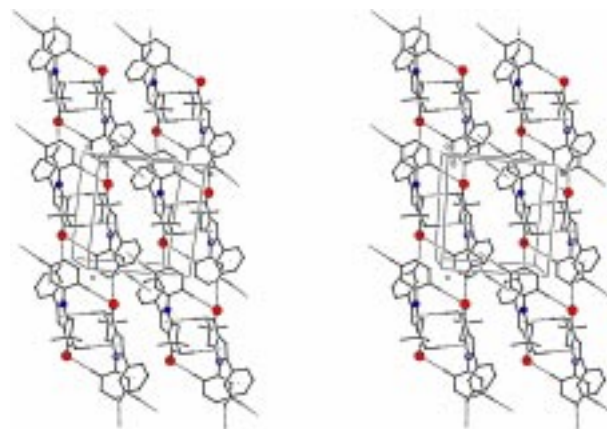


Figure 6
A stereoview of the packing of (III), showing the C—H...O hydrogen-bonded ladder motif and C—H... π interactions.

generating a sheet structure. This is further stabilized by an additional C—H... π interaction, which reinforces the two-dimensional sheet (Fig. 4). Once again, it is to be noted that the F atom does not generate any significant interaction.

In compound (III), the six-membered ring of the tetrahydroindole system (Fig. 5) again has an envelope conformation, with atom C5 a distance of -0.632 (2) Å from the C3/C4/C6/C7/C8 plane. Selected torsion angles are given in Table 5. The dihedral angle between the phenyl rings is 62.99 (6)°. The variation in dihedral angles in (I), (II) and (III) clearly shows that the conformations of the molecules are significantly different among the three compounds. The molecules of (III) pack in the unit cell *via* two different C—H...O interactions (Fig. 6 and Table 6), generating a 'ladder' motif which is further reinforced by a C—H... π interaction involving C4—H4B (Table 6). These ladders are then linked to form a two-dimensional sheet in the (101) plane *via* a further C—H... π interaction involving C11—H11 and the centroid of the C9—C14 phenyl ring (Cg2) at $(-x, 1-y, 2-z)$.

These three structures illustrate the co-existence of C—H...O and C—H... π interactions and their ability to generate different packing motifs with altered molecular conformations. Such subtle variations might have a significant impact on the biological activities of such compounds. Indeed, only compound (II) is found to be biologically active.

Experimental

Compounds (I)–(III) were synthesized according to the procedure of Nagarajan *et al.* (1985). Single crystals suitable for X-ray diffraction studies were grown from solutions in acetone (analytical reagent) by slow evaporation at 283 K.

Compound (I)

Crystal data

C₂₂H₁₉ClFNO
M_r = 367.83
 Triclinic, *P* $\bar{1}$
a = 9.246 (3) Å
b = 10.228 (4) Å
c = 11.177 (4) Å
 α = 80.529 (6)°
 β = 86.335 (7)°
 γ = 65.120 (6)°
V = 945.7 (6) Å³

Z = 2
D_x = 1.292 Mg m⁻³
 Mo *K* α radiation
 Cell parameters from 2435 reflections
 θ = 2.3–21.2°
 μ = 0.22 mm⁻¹
T = 293 (2) K
 Plate, colourless
 0.50 × 0.30 × 0.20 mm

Data collection

Bruker SMART CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: empirical (SADABS; Sheldrick, 1996)
T_{min} = 0.844, *T_{max}* = 0.957
 9987 measured reflections

3788 independent reflections
 2811 reflections with *I* > 2 σ (*I*)
R_{int} = 0.024
 θ_{\max} = 26.4°
h = -11 → 11
k = -12 → 12
l = -13 → 13

Refinement

Refinement on *F*²
R(*F*) = 0.050
wR(*F*²) = 0.118
S = 1.04
 3788 reflections
 311 parameters
 All H-atom parameters refined

$w = 1/[\sigma^2(F_o^2) + (0.0455P)^2 + 0.2584P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.20$ e Å⁻³
 $\Delta\rho_{\min} = -0.22$ e Å⁻³

Table 1
Selected torsion angles (°) for (I).

C9—N1—C1—C15	-3.7 (3)	C5—C6—C7—C8	20.6 (3)
O1—C3—C4—C5	145.5 (2)	C6—C7—C8—C3	1.4 (3)
C8—C3—C4—C5	-36.3 (3)	C4—C3—C8—C7	6.1 (3)
C3—C4—C5—C6	57.3 (3)	C7—N1—C9—C10	106.0 (2)
C4—C5—C6—C7	-47.0 (2)	N1—C1—C15—C16	148.67 (18)
C9—N1—C7—C8	-177.68 (16)		

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

Cg1 is the centroid of the five-membered N1/C1/C2/C8/C7 ring and Cg2 is the centroid of the C9—C14 phenyl ring.

<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>
C2—H2...O1 ⁱ	0.90 (2)	2.50 (2)	3.358 (3)	158 (2)
C10—H10...Cg1 ⁱⁱ	0.96 (2)	2.93	3.723 (3)	142
C14—H14...Cg2 ⁱⁱⁱ	0.94 (2)	2.90	3.804 (3)	162

Symmetry codes: (i) $1-x, 2-y, 1-z$; (ii) $2-x, 1-y, 1-z$; (iii) $1-x, 1-y, 1-z$.

Compound (II)

Crystal data

C₂₃H₂₂FNO₂
M_r = 363.42
 Monoclinic, *P*_{2₁}/*n*
a = 10.112 (2) Å
b = 19.073 (4) Å
c = 11.287 (3) Å
 β = 116.348 (3)°
V = 1950.8 (8) Å³
Z = 4

D_x = 1.237 Mg m⁻³
 Mo *K* α radiation
 Cell parameters from 974 reflections
 θ = 2.7–24.3°
 μ = 0.09 mm⁻¹
T = 293 (2) K
 Prism, colourless
 0.40 × 0.30 × 0.20 mm

Data collection

Bruker SMART CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
T_{min} = 0.887, *T_{max}* = 0.983
 14 313 measured reflections

3562 independent reflections
 2818 reflections with *I* > 2 σ (*I*)
R_{int} = 0.024
 θ_{\max} = 25.4°
h = -12 → 12
k = -22 → 21
l = -13 → 13

Table 3
Selected torsion angles (°) for (II).

C9—N1—C1—C15	0.7 (2)	C5—C6—C7—C8	22.9 (2)
O1—C3—C4—C5	151.96 (15)	C6—C7—C8—C3	1.0 (2)
C8—C3—C4—C5	-29.9 (2)	C4—C3—C8—C7	1.9 (2)
C3—C4—C5—C6	53.03 (19)	C7—N1—C9—C10	112.95 (17)
C4—C5—C6—C7	-46.9 (2)	N1—C1—C15—C16	136.24 (16)
C9—N1—C7—C8	-178.73 (13)		

Table 4
Hydrogen-bonding geometry (Å, °) for (II).

Cg1 is the centroid of the five-membered N1/C1/C2/C8/C7 ring.

<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>
C11—H11...O1 ⁱ	0.97 (2)	2.41 (2)	3.366 (2)	169 (1)
C14—H14...O1 ⁱⁱ	0.96 (2)	2.48 (2)	3.428 (2)	173 (1)
C4—H4B...Cg1 ⁱⁱⁱ	1.00 (2)	2.69	3.606 (2)	166

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

Table 5
Selected torsion angles (°) for (III).

C9–N1–C1–C15	–5.4 (2)	C5–C6–C7–C8	20.6 (2)
O1–C3–C4–C5	152.69 (14)	C6–C7–C8–C3	4.3 (2)
C8–C3–C4–C5	–29.93 (19)	C4–C3–C8–C7	0.08 (19)
C3–C4–C5–C6	53.75 (17)	C7–N1–C9–C10	115.85 (15)
C4–C5–C6–C7	–46.84 (17)	N1–C1–C15–C16	137.79 (15)
C9–N1–C7–C8	–176.85 (11)		

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0502P)^2 + 0.3423P]$
$R(F) = 0.041$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.106$	$(\Delta/\sigma)_{\max} < 0.001$
$S = 1.04$	$\Delta\rho_{\max} = 0.15 \text{ e } \text{Å}^{-3}$
3562 reflections	$\Delta\rho_{\min} = -0.19 \text{ e } \text{Å}^{-3}$
332 parameters	
All H-atom parameters refined	

Table 6
Hydrogen-bonding geometry (Å, °) for (III).

$Cg1$ is the centroid of the five-membered N1/C1/C2/C8/C7 ring and $Cg2$ is the centroid of the C9–C14 phenyl ring.

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
C14–H14 \cdots O1 ⁱ	0.97 (2)	2.57 (2)	3.533 (3)	173 (1)
C19–H19 \cdots O1 ⁱⁱ	0.93 (2)	2.58 (2)	3.389 (3)	145 (2)
C4–H4B \cdots Cg1 ⁱ	0.99 (2)	2.74	3.723 (3)	172
C11–H11 \cdots Cg2 ⁱⁱⁱ	0.97 (2)	2.75	3.620 (3)	149

Symmetry codes: (i) $1-x, -y, 1-z$; (ii) $x-1, y, 1+z$; (iii) $-x, 1-y, 2-z$.

Compound (III)

Crystal data

$C_{22}H_{21}NO$	$Z = 2$
$M_r = 315.40$	$D_x = 1.169 \text{ Mg m}^{-3}$
Triclinic, $P\bar{1}$	Mo $K\alpha$ radiation
$a = 9.315 (5) \text{ Å}$	Cell parameters from 985 reflections
$b = 9.793 (5) \text{ Å}$	$\theta = 2.4\text{--}24.2^\circ$
$c = 11.048 (6) \text{ Å}$	$\mu = 0.07 \text{ mm}^{-1}$
$\alpha = 81.362 (9)^\circ$	$T = 293 (2) \text{ K}$
$\beta = 65.691 (7)^\circ$	Block, colourless
$\gamma = 77.999 (8)^\circ$	$0.35 \times 0.30 \times 0.20 \text{ mm}$
$V = 896.0 (8) \text{ Å}^3$	

Data collection

Bruker SMART CCD area-detector diffractometer	3259 independent reflections
φ and ω scans	2604 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	$R_{\text{int}} = 0.014$
$T_{\min} = 0.946, T_{\max} = 0.986$	$\theta_{\max} = 25.4^\circ$
7395 measured reflections	$h = -11 \rightarrow 11$
	$k = -11 \rightarrow 11$
	$l = -12 \rightarrow 13$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0528P)^2 + 0.121P]$
$R(F) = 0.040$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.104$	$(\Delta/\sigma)_{\max} < 0.001$
$S = 1.04$	$\Delta\rho_{\max} = 0.13 \text{ e } \text{Å}^{-3}$
3259 reflections	$\Delta\rho_{\min} = -0.17 \text{ e } \text{Å}^{-3}$
301 parameters	
All H-atom parameters refined	

The ranges of the C–H bond lengths in (I), (II) and (III) are 0.91 (3)–1.08 (3), 0.93 (2)–1.01 (3) and 0.93 (2)–1.05 (2) Å, respectively.

For all three title compounds, data collection: *SMART* (Bruker, 2000); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 2000); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1993); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997), *CAMERON* (Watkin *et al.*, 1993) and *PLATON* (Spek, 2003); software used to prepare material for publication: *WinGX* (Farrugia, 1999) and *PLATON*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: FG1731). Services for accessing these data are described at the back of the journal.

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