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# Conformations of three heterocyclic perhydropyrrolobenzofurans and polymeric assembly via co-operative intermolecular  $C$ —H $\cdots$ O hydrogen bonds

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In 1-cyclohexyl-6,6,8a-trimethyl-3a,6,7,8a-tetrahydro-1H-1 benzofuro $[2,3-b]$ pyrrole-2,4(3H,5H)-dione, C<sub>19</sub>H<sub>27</sub>NO<sub>3</sub>, (I), and the isomorphous compounds 6,6,8a-trimethyl-1-phenyl- $3a, 6, 7, 8a$ -tetrahydro-1H-1-benzofuro[2,3-b]pyrrole-2,4(3H,5H)dione,  $C_{19}H_{21}NO_3$ , (II), and 6,6,8a-trimethyl-1-(3-pyridyl)- $3a, 6, 7, 8a$ -tetrahydro-1H-1-benzofuro[2,3-b]pyrrole-2,4(3H,5H)dione,  $C_{18}H_{20}N_2O_3$ , (III), the tetrahydrobenzo-dihydrofuropyrrolidine ring systems are folded at the cis junction of the five-membered rings, giving rise to a non-planar shape of the tricyclic cores. The dihydrofuran and pyrrolidine rings in (I) are puckered and adopt an envelope conformation. The cyclohexene rings adopt a half-chair conformation in all the molecules, while the substituent N-cyclohexyl ring in (I) assumes a chair form. Short intramolecular  $C-H\cdots O$ contacts form  $S(5)$  and  $S(6)$  motifs. The isomorphous compounds (II) and (III) are effectively isostructural, and aggregate into chains via intermolecular  $C-H\cdots O$  hydrogen bonds.

#### Comment

Co-operativity is an important property of intermolecular interactions and, thereby, molecules assemble into polymers with distinct patterns of interactions. By the process of mutual polarization within a polymeric assembly, weaker interactions acquire greater strength than they otherwise possess. A dimer of two molecules intermolecularly connected by symmetrical hydrogen bonds is more stable than if they are connected by an isolated hydrogen bond, the stability of a tetramer is greater than that of a pair of dimers, and so on (Desiraju & Steiner, 1999; Sharma & Desiraju, 1994). Polymeric assemblies formed by co-operative weak interactions are the subject of the present discussion. In this report, we present examples of a new class of tricyclic benzofuran derivatives with a benzofuropyrrolidine ring skeleton. The structural and conformational analyses of three compounds have been undertaken, namely 1-cyclohexyl-6,6,8a-trimethyl-3a,6,7,8a-tetrahydro-1H-1-benzofuro[2,3-b]pyrrole-2,4(3H,5H)-dione, (I), 1-phenyl- $6,6,8a$ -trimethyl-3a, $6,7,8a$ -tetrahydro-1H-1-benzofuro[2,3-b]pyrrole-2,4(3H,5H)-dione, (II), and 1-pyridin-3-yl-6,6,8a-trimethyl-3a,6,7,8a-tetrahydro-1H-1-benzofuro[2,3-b]pyrrole- $2,4(3H,5H)$ -dione, (III). These chiral molecules are formally derived from a perhydrofuro(or -pyrrolo)benzofuran system, and they have previously been shown to exhibit hypoglycaemic activity (Nagarajan et al., 1988).



The stereogenic centres C3a and C8a in compounds  $(I)$ -(III) (Figs. 1–3) adopt  $RS$  and  $SR$  configurations, respectively (Cahn et al., 1966). The core of each molecule consists of fused tetrahydrobenzo  $(A)$ , dihydrofuro  $(B)$  and pyrrolidine  $(C)$ rings. The C3b= $C7a$  double bond at the AB ring junction is in the range 1.341 (2)–1.344 (2) Å. In each compound, the B and  $C$  rings are folded at the  $BC$  ring junction, which is *cis*-fused (Bucourt, 1974). The angles between the best planes through ring  $B$  (atoms O8/C8a/C3a/C3b/C7a) and ring  $C$  (atoms N1/ C2/C3/C3a/C8a) are 66.4 (1), 63.8 (1) and 63.2 (1)°, respectively, while the crossed torsion angles at the junction, i.e.





 $N1 - C8a - C3a - C3b$  and  $O8 - C8a - C3a - C3$ , are 103.2 (1) and  $-130.2$  (1), 114.3 (1) and  $-121.2$  (1), and 113.4 (1) and  $-122.3$  (1)°, respectively, in the molecules of (I)–(III).

The folding at the  $BC$  ring junction gives rise to the nonplanarity of the tricyclic ring system. The structures of two analogous molecules based on a chiral tricyclic tetrahydrobenzo-dihydrofuro-tetrahydrofuran (or pyrrolidine) core, namely 1-isopropyl-6,6,8a-trimethyl-1,3a,5,6,7,8a-hexahydro-3H-1-benzofuro[2,3-b]pyrrole-2,4-dione (Narasegowda et al.,



The molecular structure of (II), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii. Numerical data refer to the internal torsion angles  $(°)$  of the individual rings (s.u. values vary between 0.1 and 0.3°). The intramolecular  $C-H\cdots O$  short contact is shown as a dashed line.



The molecular structure of (III), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii. Numerical data refer to the internal torsion angles  $(°)$  of the individual rings (s.u. values vary between 0.1 and 0.3°). The intramolecular  $C-H\cdots O$  short contact is shown as a dashed line.

2006), (IV), and 6,6,8a-trimethyl-3a,6,7,8a-tetrahydrobenzo[b]furo[3,2-d]furan-2,4(3H,5H)-dione (Nagaraj et al., 2005), (V), also possess a similar non-planar shape of the tricyclic core, and the equivalent crossed torsion angles at the junctions of the five-membered rings are  $99.7(1)$  and  $-132.8$  (1) (values correspond to an inverted image of the reported structure), and  $104.0$  (1) and  $-127.2$  (1)°, respectively.

The superposition of all five available structures of perhydrofuro(or -pyrrolo)benzofuran derivatives is shown in supplementary Fig. 5. The conformations of the substituent rings, *i.e.* N-cyclohexyl in  $(I)$ , N-phenyl in  $(II)$  and N-pyridyl in (III), are described by torsion angles  $C8a-N1-C12-C13$  of  $-112.4$  (1),  $-117.2$  (2) and  $-120.2$  (1)°, respectively.

The isomorphous compounds (II) and (III), with a difference of one atom [C14 in (II) and N14 in (III)] are isostructural. The degree of isostructurality has been quantitatively described by two descriptors, *i.e.* the unit-cell similarity index,  $\Pi$ , which is the difference between orthogonalized lattice parameters, and the isostructurality index,  $I_i(n)$ , where *n* is the number of distance differences between identical non-H atoms (Kálmán et al., 1993). The calculated values of  $\Pi = 0.019$ and  $I_i(24) = 99.7\%$  indicate the structures are close to the ideal case of isostructrality.

The internal torsion angles of the heterocyclic rings are listed in Figs.  $1-3$ . Ring A (cyclohexene) adopts a half-chair  $(C_2)$  conformation in all three molecules. However, its conformation, i.e. a half-chair  $(C_2)$  versus a sofa  $(C_s)$ , is hardly distinguishable in the present structures (Bucourt, 1974). The N-cyclohexyl ring in (I) assumes a chair form, with the larger tricyclic system in the equatorial position. The puckering (Cremer & Pople, 1975) and asymmetric (Duax et al., 1976) parameters of individual rings are provided in supplementary Table 4. The five-membered B and C rings in  $(I)$  are puckered and adopt envelope  $(C_s)$  conformations (Fuchs, 1978), with atoms C8a and C3a at the flaps of the envelopes. Atoms C8a and C3a are  $0.18(1)$  and  $0.27(1)$  Å, respectively, out of the best planes formed by the other four atoms of the ring. Rings  $B$  and  $C$  are planar in (II) and (III). The least-squares planes





The crystal packing, showing the co-operative  $C-H\cdots O$  hydrogenbonded polymeric association of isostructural compounds (II) and (III), viewed along the a axis. The representative diagram corresponds to the packing in (II). Atoms labelled with an asterisk (\*) or a hash (#) are at the symmetry positions  $(-x + 1, -y + 1, -z + 1)$  and  $(x + 1, y, z)$ , respectively. Intermolecular  $C-H\cdot\cdot\cdot O$  hydrogen bonds are shown as dashed lines.

formed by atoms of rings A and B (C3a/C3b/C4/C5/C7/C7a/ O8/C8a) and ring C (N1/C2/O2/C3/C3a/C8a) intercept at an angle of 65.8 (1) $^{\circ}$  in (II) and 64.2 (1) $^{\circ}$  in (III).

The parameters for intramolecular short contacts and intermolecular hydrogen bonds are given in Tables 1-3. The conserved intramolecular C $-H$  $\cdots$ O short contacts in (I)-(III) were observed between the donors of the N-substituents and atom O8 of ring B (Figs. 1–3). The short contact  $C12$ H12 $\cdots$ O8 forms an S(5) motif (Bernstein *et al.*, 1995) in (I), while  $S(6)$  motifs are formed by C17–H17 $\cdots$ O8 in (II) and (III). Additionally, an intramolecular  $C17 - H171 \cdots O2$ contact forms an  $S(6)$  motif in (I). The crystal packing in (I) is entirely due to van der Waals interactions. The crystal structures of isostructural compounds (II) and (III) are held together primarily by intermolecular  $C-H\cdots O$  hydrogen bonds (Tables 2 and 3), forming chains of rings along [100]  $(Fig. 4)$ . The significance of the co-operativity of weak intermolecular interactions for molecular self-assembly is elucidated in the present examples.

## Experimental

The synthetic procedures used for the preparation of compounds  $(I)$ -(III) are as described in the literature (Nagarajan et al., 1988). Single crystals suitable for X-ray diffraction were grown by slow evaporation of solutions containing the following solvents (in a 1:1 ratio): for (I), ethanol and water; for (II), dichloromethane and hexane; for (III), benzene and hexane.

 $V = 848.50$  (3)  $\AA^3$ 

 $D_x = 1.242$  Mg m<sup>-3</sup> Mo  $K\alpha$  radiation  $\mu = 0.08$  mm<sup>-1</sup>  $T - 120$  (2) K Plate, colourless  $0.24 \times 0.16 \times 0.03$  mm

18064 measured reflections 3327 independent reflections 2990 reflections with  $I > 2\sigma(I)$ 

 $R_{\rm int} = 0.039$  $\theta_{\text{max}} = 26.0^{\circ}$ 

 $Z = 2$ 

## Compound (I)

Crystal data

 $C_{19}H_{27}NO_3$  $M_r = 317.42$ Triclinic,  $P\overline{1}$  $a = 9.4708(2)$  Å  $b = 10.2139(2)$  Å  $c = 10.7512$  (2) A  $\alpha = 105.804$  (1)<sup>o</sup>  $\beta = 99.141$  (1)<sup>o</sup>  $\gamma = 116.320(1)^{\circ}$ 

Data collection

Nonius KappaCCD area-detector diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: multi-scan (SADABS; Sheldrick, 2003)  $T_{\text{min}} = 0.891, T_{\text{max}} = 0.998$ 

#### **Refinement**



#### Table 1

Hydrogen-bond and short-contact geometry  $(\mathring{A}, \degree)$  for (I).



## Compound (II)

#### Crystal data



Nonius KappaCCD area-detector diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: multi-scan (SADABS; Sheldrick, 2003)  $T_{\text{min}} = 0.903$ ,  $T_{\text{max}} = 0.997$ 

## Refinement

Refinement on  $F^2$  $R[F^2 > 2\sigma(F^2)] = 0.044$  $wR(F^2) = 0.117$  $S = 1.07$ 3132 reflections 292 parameters All H-atom parameters refined 14845 measured reflections 3132 independent reflections 2562 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.047$  $\theta_{\text{max}} = 26.0^{\circ}$ 

 $w = 1/[\sigma^2 (F_o^2) + (0.0532P)^2]$ + 0.358P] where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\text{max}} = 0.001$  $\Delta \rho_{\text{max}} = 0.19 \text{ e A}^{-3}$  $\Delta \rho_{\rm min} = -0.27$  e  $\rm \AA^{-3}$ 

13914 measured reflections 3040 independent reflections 2769 reflections with  $I > 2\sigma(I)$ 

 $^{2} + 2F_{c}^{2})/3$ 

 $R_{\text{int}} = 0.035$  $\theta_{\text{max}} = 26.0^{\circ}$ 

#### Table 2

Hydrogen-bond and short-contact geometry  $(\mathring{A}, \circ)$  for (II).



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

## Compound (III)

Crystal data



## Data collection

Nonius KappaCCD area-detector diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: multi-scan (SADABS; Sheldrick, 2003)  $T_{\min} = 0.913, T_{\max} = 0.992$ 

## **Refinement**



#### Table 3

Hydrogen-bond and short-contact geometry  $(A, \circ)$  for (III).



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

Larger-than-expected values of residual electron density were observed in (I) and (III) [the values of  $\Delta \rho_{\text{max}}$  and  $\Delta \rho_{\text{min}}$  were 0.75 and  $-0.88 \text{ e A}^{-3}$ , respectively, in (I), and 0.82 and  $-0.87 \text{ e A}^{-3}$ , respectively, in (III)]. This was attributed to the presence of a few poorly fitting low-angle reflections  $[(011), (\overline{111}), (020), (111)$  and  $(\overline{21}2)$  in (I), and (112), (100), (111), (022), (013), (122) and (012) in (III)], which appeared to have been truncated by the beam stop. These reflections were omitted during the final cycles of refinement. The residual electron density was then featureless and the residual factor  $R$  dropped from 0.053 to 0.039 in (I), and from 0.056 to 0.039 in (III) for observed data.

H atoms were located in difference maps and were refined freely. Refined methine, methylene and methyl C-H distances are as follows: for (I), 0.99 (2), 0.97 (2)–1.01 (2) and 0.96 (2)–1.02 (2) Å; for (II), 0.97 (2), 0.96 (2)–1.01 (2) and 0.97 (2)–1.03 (2) Å, with aromatic  $C-H = 0.96$  (2)–1.00 (2) Å; for (III), 0.99 (2), 0.97 (2)–1.02 (2) and 0.98 (2)–1.02 (2) Å, with aromatic C–H = 0.97 (2)–1.01 (2) Å.

For all compounds, data collection: COLLECT (Nonius, 1998); cell refinement: DENZO (Otwinowski & Minor, 1997) and COLLECT; data reduction: DENZO and COLLECT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 (Farrugia, 1997), PLATON (Spek, 2003) and INSIGHTII (Accelrys, 2002); software used to prepare material for publication: SHELXL97 and PLATON.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: GD3014). An additional figure and table are also available. Services for accessing these data are described at the back of the journal.

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