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## Two Cyclohexanespiro-5'-hydantoin Monohydrates

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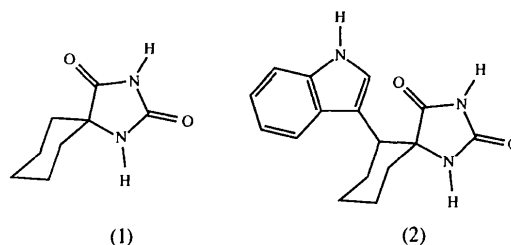
### Abstract

Cyclohexanespiro-5'-hydantoin monohydrate,  $C_8H_{12}N_2O_2 \cdot H_2O$ , has a chair-shaped cyclohexane ring with endocyclic torsion-angle magnitudes in the range  $54.4(2)$ – $56.3(2)^\circ$ . All potential hydrogen-bond donors are involved in intermolecular hydrogen bonding, with lengths in the range  $2.760(2)$ – $2.908(2)$  Å. In its indolyl adduct, 2-(3-indolyl)cyclohexanespiro-5'-hydantoin monohydrate,  $C_{16}H_{17}N_3O_2 \cdot H_2O$ , the cyclohexane moiety adopts a chair conformation with the indolyl substituent in an equatorial position. The N—H portion of the hydantoin ring is *cis* to indolyl, while the C=O of the hydantoin is *trans*. Endocyclic torsion-angle magnitudes of the cyclohexane ring are in the range  $54.2(2)$ – $56.7(2)^\circ$ . All potential hydrogen-bond donors are involved in intermolecular hydrogen bonds, with lengths  $2.828(2)$ – $3.187(2)$  Å.

### Comment

Of the aromatic amino acids, tryptophan is easiest to selectively excite due to its red-shifted absorbance. Therefore, it would be useful to use tryptophan as a fluorescent probe to study protein conformations. All of the quenching mechanisms of tryptophan in a protein environment are not fully understood. In order to study these mechanisms, it would be beneficial to limit the flexibility of the molecule. As a result, a constrained derivative of tryptophan can be used to help define how orientation of the peptide bond affects quenching (Chen, Liu, Yu & Barkley, 1996). As part of a synthetic effort to prepare these compounds, we have synthesized cyclohexanespiro-5'-hydantoin, (1), and 2-(3-indolyl)cyclohexanespiro-5'-hydantoin, (2), and have determined the crystal structures of their crystalline monohydrates as part of the complete characterization of the reaction products.  $\alpha,\alpha$ -Disubstituted amino acids are of interest because of their ability to promote specific secondary structures when incorporated into peptides. Compound (1) was synthesized as a precursor to an  $\alpha,\alpha$ -disubstituted cyclohexane amino acid, *i.e.* 1-amino-

1-cyclohexane carboxylic acid, and compound (2) as a precursor to a constrained tryptophan derivative.



The bond distances and angles within the hydantoin (2,4-imidazolidinedione) units of both molecules are normal, typical of those observed in 5,5-dimethylhydantoin (Cassady & Hawkinson, 1982) and in other spiro-5'-hydantoins (Vilches, Florencio, Smith-Verdier & Garcia-Blanco, 1981; Smith-Verdier, Florencio, & Garcia-Blanco, 1977, 1979; Florencio, Smith-Verdier & Garcia-Blanco, 1978, 1982). The hydantoin unit is very nearly planar in both molecules, with maximum deviations from planarity of  $0.014(2)$  Å (O1) in (1) and  $0.031(2)$  Å (O2 and N3') in (2). The indole ring system of (2) exhibits a maximum deviation of  $0.024(2)$  Å (for C12) from planarity and this plane forms a dihedral angle of  $58.83(4)^\circ$  with that of the hydantoin. The cyclohexane rings of both molecules are in chair conformations, with endocyclic torsion-angle magnitudes of  $54.4(2)$ – $56.3(2)^\circ$  for (1) and  $54.2(2)$ – $56.7(2)^\circ$  for (2).

In both structures, all available hydrogen-bond donors of the hydantoin, indole and water are involved in intermolecular hydrogen bonds; these are detailed in Tables 2 and 4. In both, the carbonyl O1 atom accepts a hydrogen bond from water. In (1), N1'—H donates to a carbonyl O atom and N3'—H donates to water, while in (2), these roles are reversed. In (1), the carbonyl O2 atom accepts from N—H, while in (2), it accepts from water. In (1), all hydrogen bonds are linear within experimental error, while in (2), they are less so, with angles about hydrogen varying from  $115(2)$  to  $174(2)^\circ$ .

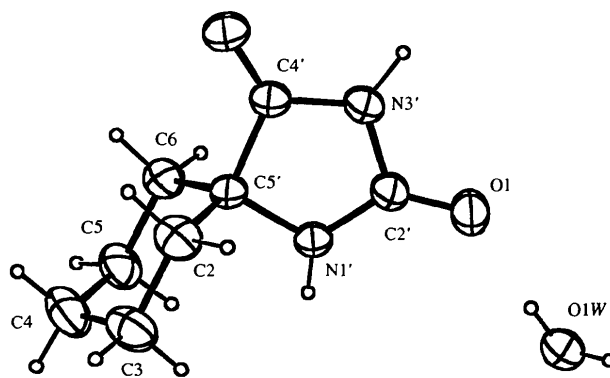


Fig. 1. The molecular structure of (1)· $H_2O$ , with 40% probability ellipsoids. H atoms are shown as circles of arbitrary radii.

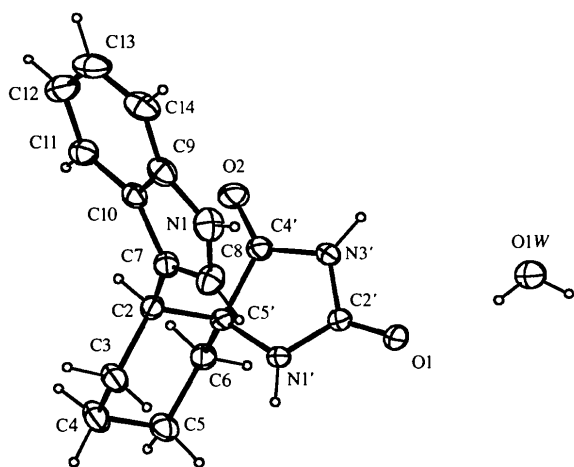


Fig. 2. The molecular structure of (2)·H<sub>2</sub>O, with 30% probability ellipsoids. H atoms are shown as circles of arbitrary radii.

### Refinement

Refinement on  $F^2$

$R(F) = 0.038$

$wR(F^2) = 0.039$

$S = 1.466$

1722 reflections

174 parameters

All H atoms refined

$w = 4F_o^2 / [\sigma^2(F_o^2) + 0.0004F_o^4]$

$(\Delta/\sigma)_{\max} = 0.005$

$\Delta\rho_{\max} = 0.23 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -0.30 \text{ e } \text{\AA}^{-3}$

Extinction correction:

isotropic (Zachariasen, 1963)

Extinction coefficient:

$8.5(2) \times 10^{-6}$

Scattering factors from *International Tables for X-ray Crystallography* (Vol. IV)

Absolute configuration:

refinement of the antipodal structure was inconclusive; for the reported absolute structure,  $R = 0.03803$ ,  $wR = 0.03908$ ,  $S = 1.466$ , while the alternate structure yielded  $R = 0.03801$ ,  $wR = 0.03910$ ,  $S = 1.467$ .

## Experimental

Compound (1) was prepared by reacting cyclohexanone with KCN and ammonium carbonate, according to a procedure reported by Tsang, Schmied, Nyfeler & Goodman (1984). Crystals of the monohydrate were grown by refrigeration of a solution in methanol–water. Compound (2) was prepared via a Bücherer–Bergs synthesis on 2-(3-indolyl)cyclohexanone under an inert atmosphere with heating. The crude product was dissolved in basic water and the pH was adjusted to 7. Crystals of the monohydrate formed within 48 h.

### Compound (1)·H<sub>2</sub>O

#### Crystal data

C<sub>8</sub>H<sub>12</sub>N<sub>2</sub>O<sub>2</sub>·H<sub>2</sub>O

$M_r = 186.21$

Orthorhombic

$Pna2_1$

$a = 16.889(1) \text{ \AA}$

$b = 9.2407(5) \text{ \AA}$

$c = 6.2644(6) \text{ \AA}$

$V = 977.6(2) \text{ \AA}^3$

$Z = 4$

$D_x = 1.265 \text{ Mg m}^{-3}$

$D_m$  not measured

Cu  $K\alpha$  radiation

$\lambda = 1.54184 \text{ \AA}$

Cell parameters from 25 reflections

$\theta = 14\text{--}30^\circ$

$\mu = 0.78 \text{ mm}^{-1}$

$T = 296 \text{ K}$

Fragment

$0.33 \times 0.10 \times 0.10 \text{ mm}$

Colorless

1722 reflections with  $I > \sigma(I)$

$R_{\text{int}} = 0.034$

$\theta_{\max} = 75.0^\circ$

$h = 0 \rightarrow 21$

$k = -11 \rightarrow 11$

$l = -7 \rightarrow 7$

3 standard reflections

frequency: 120 min

intensity decay: 0.4%

Table 1. Selected geometric parameters ( $\text{\AA}$ ,  $^\circ$ ) for (1)·H<sub>2</sub>O

|             |           |             |           |
|-------------|-----------|-------------|-----------|
| O1—C2'      | 1.218 (2) | N3'—C2'     | 1.397 (2) |
| O2—C4'      | 1.220 (2) | N3'—C4'     | 1.345 (2) |
| N1'—C2'     | 1.326 (2) | C4'—C5'     | 1.518 (2) |
| N1'—C5'     | 1.461 (2) |             |           |
| C2'—N1'—C5' | 113.4 (1) | O2—C4'—N3'  | 126.6 (2) |
| C2'—N3'—C4' | 111.4 (1) | O2—C4'—C5'  | 125.3 (1) |
| O1—C2'—N1'  | 129.4 (2) | N3'—C4'—C5' | 108.1 (1) |
| O1—C2'—N3'  | 123.3 (1) | N1'—C5'—C4' | 99.9 (1)  |
| N1'—C2'—N3' | 107.3 (1) |             |           |

Table 2. Hydrogen-bonding geometry ( $\text{\AA}$ ,  $^\circ$ ) for (1)·H<sub>2</sub>O

| D—H...A                     | D—H      | H...A    | D...A     | D—H...A |
|-----------------------------|----------|----------|-----------|---------|
| N1'—H1'...O2'               | 0.82 (2) | 2.09 (2) | 2.908 (2) | 176 (2) |
| N3'—H3'...O1W <sup>i</sup>  | 0.95 (2) | 1.82 (2) | 2.760 (2) | 175 (2) |
| O1W—H2W...O1                | 0.96 (2) | 1.83 (2) | 2.788 (2) | 177 (2) |
| O1W—H1W...O1 <sup>iii</sup> | 0.88 (2) | 1.98 (2) | 2.852 (2) | 178 (3) |

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

### Compound (2)·H<sub>2</sub>O

#### Crystal data

C<sub>16</sub>H<sub>17</sub>N<sub>3</sub>O<sub>2</sub>·H<sub>2</sub>O

$M_r = 301.35$

Monoclinic

$P2_1/c$

$a = 11.991(1) \text{ \AA}$

$b = 8.3535(7) \text{ \AA}$

$c = 15.159(1) \text{ \AA}$

$\beta = 94.731(8)^\circ$

$V = 1513.3(4) \text{ \AA}^3$

$Z = 4$

$D_x = 1.323 \text{ Mg m}^{-3}$

$D_m$  not measured

Cu  $K\alpha$  radiation

$\lambda = 1.54184 \text{ \AA}$

Cell parameters from 25 reflections

$\theta = 6\text{--}17^\circ$

$\mu = 0.72 \text{ mm}^{-1}$

$T = 296 \text{ K}$

Prism

$0.20 \times 0.14 \times 0.13 \text{ mm}$

Colorless

#### Data collection

Enraf–Nonius CAD-4 diffractometer

$\omega/2\theta$  scans

Absorption correction: none

3469 measured reflections

3113 independent reflections

2560 reflections with

$I > \sigma(I)$

$R_{\text{int}} = 0.032$

$\theta_{\max} = 75.0^\circ$

$h = 0 \rightarrow 15$

$k = 0 \rightarrow 10$

$l = -18 \rightarrow 17$

3 standard reflections

frequency: 120 min

intensity decay: 0.3%

## Refinement

Refinement on  $F^2$  $R(F) = 0.054$  $wR(F^2) = 0.088$  $S = 1.533$ 

2560 reflections

276 parameters

All H atoms refined

 $w = 4F_o^2/[\sigma^2(F_o^2)$   
 $+ 0.0004F_o^4]$  $(\Delta/\sigma)_{\max} = 0.049$  $\Delta\rho_{\max} = 0.20 \text{ e } \text{\AA}^{-3}$  $\Delta\rho_{\min} = -0.11 \text{ e } \text{\AA}^{-3}$ 

Extinction correction:

isotropic (Zachariassen,  
1963)

Scattering coefficient:

 $2.3 (1) \times 10^{-6}$ Scattering factors from *International Tables for X-ray Crystallography* (Vol. IV)Table 3. Selected geometric parameters ( $\text{\AA}$ ,  $^\circ$ ) for (2).H<sub>2</sub>O

|             |           |             |           |
|-------------|-----------|-------------|-----------|
| O1—C2'      | 1.224 (2) | N1—C9       | 1.372 (2) |
| O2—C4'      | 1.208 (2) | N3'—C2'     | 1.386 (2) |
| N1'—C2'     | 1.336 (2) | N3'—C4'     | 1.368 (2) |
| N1'—C5'     | 1.458 (2) | C4'—C5'     | 1.525 (2) |
| N1—C8       | 1.374 (2) |             |           |
| C2'—N1'—C5' | 112.9 (1) | O2—C4'—C5'  | 126.8 (1) |
| C8—N1—C9    | 109.0 (1) | N3'—C4'—C5' | 106.7 (1) |
| C2'—N3'—C4' | 111.8 (1) | N1'—C5'—C4' | 100.9 (1) |
| O2—C4'—N3'  | 126.4 (2) |             |           |

Table 4. Hydrogen-bonding geometry ( $\text{\AA}$ ,  $^\circ$ ) for (2).H<sub>2</sub>O

| D—H...A                      | D—H      | H...A    | D...A     | D—H...A |
|------------------------------|----------|----------|-----------|---------|
| N1—H1N...O1W <sup>i</sup>    | 0.88 (2) | 2.17 (2) | 3.056 (2) | 174 (2) |
| N1'—H1N'...O1W <sup>ii</sup> | 0.88 (2) | 2.10 (2) | 2.972 (2) | 172 (2) |
| N3'—H3N'...O1 <sup>iii</sup> | 0.91 (2) | 1.97 (2) | 2.828 (2) | 157 (2) |
| O1W—H2W...O1                 | 0.90 (3) | 2.04 (3) | 2.856 (2) | 151 (3) |
| O1W—H1W...O2 <sup>ii</sup>   | 0.80 (3) | 2.76 (3) | 3.187 (2) | 115 (2) |

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

For (1).H<sub>2</sub>O, C—H distances are 0.94 (2)–1.07 (3), N—H 0.82 (2)–0.95 (2) and O—H 0.88 (2)–0.96 (2)  $\text{\AA}$ , while  $U_{\text{iso}}$  values for H atoms are in the range 0.054 (5)–0.119 (9)  $\text{\AA}^2$ . For (2).H<sub>2</sub>O, a series of  $\psi$  scans revealed no measurable change in intensity with rotation angle, thus, no absorption correction was carried out. C—H distances are 0.93 (2)–1.03 (2), N—H 0.88 (2)–0.91 (2) and O—H 0.80 (2)–0.90 (2)  $\text{\AA}$ , while  $U_{\text{iso}}$  values for H atoms are in the range 0.031 (4)–0.19 (1)  $\text{\AA}^2$ .

For both compounds, data collection: *CAD-4 Operations Manual* (Enraf–Nonius, 1977); cell refinement: *CAD-4 Operations Manual*; data reduction: *PROCESS MolEN* (Fair, 1990). Program(s) used to solve structures: direct methods *SHELXS86* (Sheldrick, 1990) for (1).H<sub>2</sub>O; direct methods *MULTAN80* (Main *et al.*, 1980) for (2).H<sub>2</sub>O. For both compounds, program(s) used to refine structures: *LSFM MolEN*; molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *CIF IN MolEN*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: FG1332). Services for accessing these data are described at the back of the journal.

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## Deux Composés Trisubstitués de la 4-Méthylèncyclooctanone

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## Abstract

The structures of two trisubstituted 4-methylenecyclooctanones are reported, namely, di-*tert*-butyl 2-methoxycarbonyl-4-methylene-1-oxocyclooctane-6,6-dicarboxylate (IUPAC numbering: di-*tert*-butyl 5-methoxycarbonyl-3-methylene-6-oxocyclooctane-1,1-dicarboxylate), C<sub>21</sub>H<sub>32</sub>O<sub>7</sub>, (1), and di-*tert*-butyl 2-phenylsulfonyl-4-methylene-1-oxocyclooctane-6,6-dicarboxylate,