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endo-1,4,4-Tribromo-3-methyltricyclo-[5.2.1.0^{2,6}]dec-8-ene-5,10-dione 10-Ethylene Acetal

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Abstract

The molecular geometry of the title compound, $C_{13}H_{13}Br_3O_3$, is similar to that observed for the 3-chloro- analogue, *endo-*1,4,4-tribromo-3-methyltricyclo[5.2.1.0^{2,6}]dec-8-ene-5,10-dione 10-ethylene acetal.

Comment

We have recently described the preparation of dimethyl 2-methylcubane-1,4-dicarboxylate, (I) (Lowe et al., 1994), through an indirect approach, which featured the gem-dibromoketone (II). The structure of (II) was originally assigned on the basis of NOE difference

measurements and the X-ray structural analysis reported here was performed to confirm the assignment.

$$MeO_2C$$
 CO_2Me
 Br
 Br
 Br
 Br
 Br

The molecular geometry is similar to that observed for the 3-chloro- analogue, *endo*-1,4,4-tribromo-3-chlorotricyclo[5.2.1.0^{2,6}]dec-8-ene-5,10-dione 10-ethylene ketal (Gable, Parker & Tsanaktsidis, 1994).

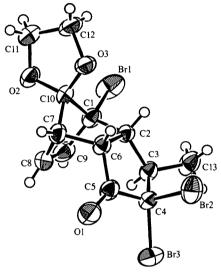


Fig. 1. ORTEPII (Johnson, 1976) drawing of (II). Displacement ellipsoids are drawn at the 50% probability level. H atoms are given arbitrary radii of 0.1 Å.

Experimental

Compound (II) was prepared as described by Lowe *et al.* (1994) and crystals were obtained by slow evaporation from diethyl ether.

Crystal data

 D_m not measured

 $C_{13}H_{13}Br_3O_3$ Mo $K\alpha$ radiation $\lambda = 0.71073 \text{ Å}$ $M_r = 456.96$ Triclinic Cell parameters from 25 $P\bar{1}$ reflections $\theta = 11.3 - 21.2^{\circ}$ a = 6.6161(13) Å $\mu = 8.314 \text{ mm}^{-1}$ b = 9.633(2) ÅT = 293(1) Kc = 12.692(3) Å $\alpha = 98.27 (2)^{\circ}$ Block $0.40 \times 0.37 \times 0.37 \text{ mm}$ $\beta = 104.84 (2)^{\circ}$ $\gamma = 106.74(2)^{\circ}$ Colourless $V = 727.8(3) \text{ Å}^3$ Z = 2 $D_x = 2.085 \text{ Mg m}^{-3}$

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Data collection

Enraf-Nonius CAD-4-MachS diffractometer $\omega/2\theta$ scans
Absorption correction:
Gaussian by integration (Sheldrick, 1976) $T_{\min} = 0.041$, $T_{\max} = 0.166$ 4197 measured reflections
3332 independent reflections

2462 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.028$ $\theta_{\text{max}} = 27.47^{\circ}$ $h = -1 \rightarrow 8$ $k = -12 \rightarrow 12$ $l = -16 \rightarrow 16$ 3 standard reflections frequency: 160 min intensity decay: 18%

Refinement

Refinement on F^2 $(\Delta/\sigma)_{\rm max} = 0.001$ $\Delta \rho_{\text{max}} = 0.790 \text{ e Å}^{-3}$ $R[F^2 > 2\sigma(F^2)] = 0.045$ $wR(F^2) = 0.119$ $\Delta \rho_{\min} = -0.667 \text{ e Å}^{-3}$ S = 1.055Extinction correction: 3332 reflections SHELXL93 208 parameters Extinction coefficient: Methine H atoms refined, 0.0116(15)others riding Scattering factors from $w = 1/[\sigma^2(F_o^2) + (0.0563P)^2]$ International Tables for Crystallography (Vol. C) + 1.0706P1 where $P = (F_o^2 + 2F_c^2)/3$

Table 1. Selected geometric parameters (Å, °)

		-	
Br1C1	1.927 (5)	C2—C3	1.541 (6)
Br2—C4	1.952 (5)	C3—C4	1.522 (6)
Br3—C4	1.921 (5)	C4—C5	1.529 (6)
C1C2	1.545 (6)	C7—C10	1.551 (7)
C1C10	1.557 (6)	C8—C9	1.308 (9)
C9C1C2	109.2 (4)	C3C4C5	105.4 (3)
C9C1C10	99.8 (4)	C3C4Br3	112.7 (3)
C2C1C10	100.7 (3)	C5—C4—Br3	112.9 (3)
C9C1Br1	115.8 (4)	C3—C4—Br2	113.1 (3)
C2—C1—Br1	113.5 (3)	C5—C4—Br2	104.0 (3)
C10—C1—BrI	116.0 (3)	Br3C4Br2	108.5 (2)
C13C3C4	116.8 (4)	C7C10C1	92.1 (3)
C13—C3—C2	115.0 (4)		

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989). Cell refinement: *CAD-4 Software*. Data reduction: *CAD-4 Software*. Program(s) used to solve structure: *SHELXS*86 (Sheldrick, 1990). Program(s) used to refine structure: *SHELXL*93 (Sheldrick, 1993). Molecular graphics: *ORTEPII* (Johnson, 1976). Software used to prepare material for publication: *SHELXL*93.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: TA1187). Services for accessing these data are described at the back of the journal.

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Spirapril Hydrochloride Hydrate

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Abstract

The crystal structure of the monohydrate form of spirapril hydrochloride, $(8S-\{7[R^*(R^*)],8R^*\})-7-(2-\{[1-(ethoxycarbonyl)-3-phenylpropyl]amino\}-1-oxopropyl)-1,4-dithia-7-azaspiro[4.4]nonane-8-carboxylic acid hydrochloride, <math>C_{22}H_{31}N_2O_5S_2^{\dagger}.Cl^-.H_2O$, was determined. The spirapril molecules adopt a *trans* conformation along the amide bond. The incorporated water molecules form hydrogen bonds with the host molecules.

Comment

Spirapril hydrochloride, (I), belongs to a class of *N*-carboxyalkyl dipeptide ACE inhibitors. Like many other of these dipeptide drugs, it is not very stable in the solid state. The major degradation pathway involves an intramolecular cyclization reaction (Strickley, Visor, Lin & Gu, 1989; Gu, Strickley, Chi & Chowhan, 1990). In our study of the chemical stability of pharmaceutical solids, we chose spirapril hydrochloride as a model compound to investigate the effect of molecular mobility on solid-state reactivity. The structure of spirapril hydrochloride monohydrate was determined and the information on the molecular geometry will aid in the understanding of the molecular details of the cyclization reaction.

An *ORTEPII* (Johnson, 1976) drawing of spirapril hydrochloride is presented in Fig. 1. For the sake of simplicity, the protonated spirapril moiety is referred to as the cation. It adopts a *trans* conformation at the amide bond since C5 and C8 (higher priority than C11) are on the opposite side of the amide bond and the torsion angle between C8—N7 and C6—C5 is -175.7 (3)°. The same kind of *trans* conformation is also observed in other *N*-carboxyalkyl dipeptide ACE inhibitors such as quinapril hydrochloride (Hausin & Codding, 1991) and enalapril maleate (Precigoux, Geoffre & Leroy,