

5-Methoxytropolone

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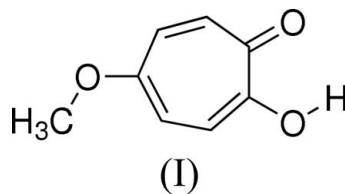
Key indicators

Single-crystal X-ray study
 $T = 153\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$
 R factor = 0.032
 wR factor = 0.081
Data-to-parameter ratio = 12.0For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title compound [systematic name: 2-hydroxy-5-methoxy-2,4,6-cycloheptatrien-1-one], $\text{C}_8\text{H}_8\text{O}_3$, contains intra- and intermolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds, resulting in inversion-symmetry generated dimers. The crystal packing is consolidated by $\pi-\pi$, $\text{C}-\text{H}\cdots\pi$ and $\text{C}-\text{H}\cdots\text{O}$ interactions.

Comment

Troponoids, being a remarkable class of non-benzenoid π -conjugated systems, have been used as the building blocks of various molecular assemblies such as liquid crystals and organogelators (Kubo, 2004). Recently, we have prepared liquid crystals with a troponoid core, *e.g.* 5-alkoxytropolone (Mori *et al.*, 1990), 5-alkanoyloxytropolone (Mori *et al.*, 2001), 5-cyanotropolone (Hashimoto *et al.*, 2001) and 5-nitrotropolone (Kubo *et al.*, 2002). The crystal structures of tropolone (Shimanouchi & Sasada, 1973), 5-cyano- and 5-nitrotropolones (Kubo *et al.*, 2001), and 5-acetyltropolone (Kubo, Yamamoto *et al.*, 2006) have been reported. We now report the structure of the title compound, (I), with the aim of contributing to a deeper understanding of troponoids and their molecular assemblies.



Compound (I) forms $\text{O}-\text{H}\cdots\text{O}$ hydrogen-bonded dimers about inversion centres, involving the OH group and an intermolecular carbonyl $\text{O}1^i$ acceptor (see Table 1 for symmetry code). An intramolecular $\text{O}-\text{H}\cdots\text{O}$ bond to $\text{O}1$ occurs simultaneously (Table 1, Fig. 1). The intermolecular $\text{O}2\cdots\text{O}1^i$ distance of 2.710 (2) \AA is close to that in tropolone (2.746 \AA ; Shimanouchi & Sasada, 1973), where a similar dimerization occurs.

Intermolecular $\pi-\pi$ interactions occur in (I) between the tropolone ring planes, as shown in Fig. 2. The $\text{C}3\cdots\text{C}6^v$ [symmetry code: $(v) \frac{3}{2} - x, \frac{1}{2} + y, \frac{3}{2} - z$] separation is 3.2559 (16) \AA , which is within the range associated with $\pi-\pi$ interactions (3.3–3.8 \AA ; Kubo *et al.*, 2001; Kubo *et al.*, 2002). This value is shorter than that of tropolone (3.418 \AA ; Shimanouchi & Sasada, 1973) and 5-(acetoxy)tropolone [3.455 (2) \AA ; Kubo, Yamamoto *et al.*, 2006].

Intermolecular $\text{C}-\text{H}\cdots\pi$ and $\text{C}-\text{H}\cdots\text{O}$ interactions are observed in the crystal structure of (I) (Table 1, Fig. 2), with distances typical for these types of interactions: $\text{C}-\text{H}\cdots\pi =$

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2.8–3.1 Å (Kubo, Fukeda *et al.*, 2006; Kubo, Yamamoto *et al.*, 2006) and C–H...O = 2.5–2.7 Å (Kubo, Fukeda *et al.*, 2006; Kubo, Matsumoto *et al.*, 2006; Kubo, Yamamoto *et al.*, 2006).

The combination of all of the above interactions builds up a three-dimensional network.

Experimental

Compound (I) was prepared by the reaction of 5-hydroxytropolone with dimethylsulfate (Nozoe *et al.*, 1953). Crystals of (I) were grown from a methanol solution by slow evaporation.

Crystal data

$C_8H_6O_3$	$Z = 4$
$M_r = 152.15$	$D_x = 1.401 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/n$	Cu $K\alpha$ radiation
$a = 8.434 (2) \text{ \AA}$	$\mu = 0.91 \text{ mm}^{-1}$
$b = 7.265 (2) \text{ \AA}$	$T = 153.1 \text{ K}$
$c = 12.308 (3) \text{ \AA}$	Prism, colourless
$\beta = 106.976 (15)^\circ$	$0.38 \times 0.19 \times 0.13 \text{ mm}$
$V = 721.3 (3) \text{ \AA}^3$	

Data collection

Rigaku R-Axis RAPID diffractometer	6663 measured reflections
ω scans	1272 independent reflections
Absorption correction: multi-scan (ABSCOR; Higashi, 1995)	1169 reflections with $F^2 > 2\sigma(F^2)$
$T_{\min} = 0.573$, $T_{\max} = 0.888$	$R_{\text{int}} = 0.036$
	$\theta_{\text{max}} = 68.2^\circ$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.03P)^2 + 0.1908P]$
$R[F^2 > 2\sigma(F^2)] = 0.032$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.081$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.13$	$\Delta\rho_{\text{max}} = 0.20 \text{ e \AA}^{-3}$
1272 reflections	$\Delta\rho_{\text{min}} = -0.14 \text{ e \AA}^{-3}$
106 parameters	Extinction correction: SHELXL97 (Sheldrick 1997)
H atoms treated by a mixture of independent and constrained refinement	Extinction coefficient: 0.0228 (14)

Table 1

Hydrogen-bond geometry (Å, °).

D–H...A	D–H	H...A	D...A	D–H...A
O2–H1...O1	0.92 (2)	2.07 (2)	2.605 (1)	115 (1)
O2–H1...O1 ⁱ	0.92 (2)	1.96 (2)	2.710 (2)	138 (2)
C6–H4...O2 ⁱⁱ	0.95	2.44	3.267 (2)	146
C7–H5...O1 ⁱⁱⁱ	0.95	2.61	3.540 (2)	165
C8–H6...C7 ^{iv}	0.98	2.89	3.530 (2)	124

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

The H atom of the OH group was located in a difference map and its position and U_{iso} value were freely refined. All C-bound H atoms were placed in idealized locations (C–H = 0.95–0.98 Å) and refined as riding, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $1.5U_{\text{eq}}(\text{methyl C})$.

Data collection: *CrystalClear* (Rigaku, 1999); cell refinement: *CrystalClear*; data reduction: *CrystalStructure* (Rigaku/MS, 2004); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick 1997); molecular graphics: *ORTEP III* (Burnett & Johnson, 1996); software used to prepare material for publication: *CrystalStructure*.

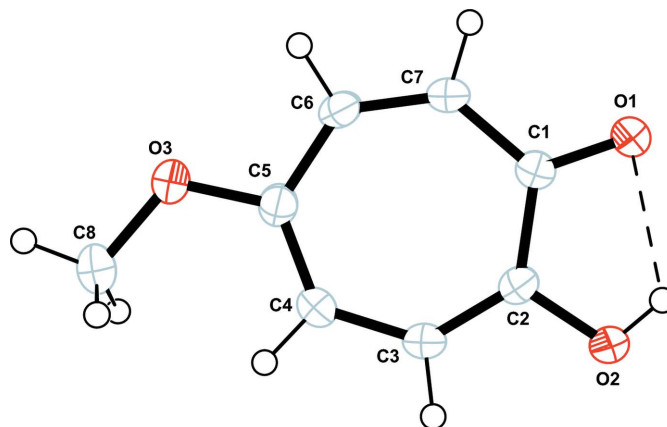


Figure 1

The molecular structure of (I), showing 50% probability displacement ellipsoids. The hydrogen bond is indicated by a dashed line.

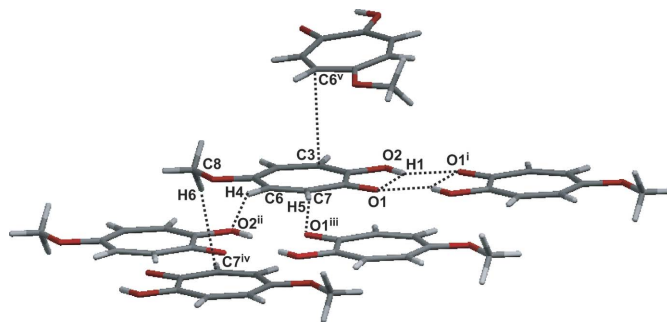


Figure 2

Intermolecular O–H...O, C–H...O, π – π and C–H... π interactions in (I). Dotted lines indicate these interactions. [Symmetry codes: (i) $2 - x, 1 - y, 1 - z$; (ii) $x, y - 1, z$; (iii) $2 - x, -y, 1 - z$; (iv) $1 - x, -y, 1 - z$; (v) $\frac{3}{2} - x, \frac{1}{2} + y, \frac{3}{2} - z$.]

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