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#### **Key indicators**

Single-crystal X-ray study T = 153 K Mean  $\sigma$ (C–C) = 0.002 Å R factor = 0.039 wR factor = 0.104 Data-to-parameter ratio = 12.2

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

5-(Acetyloxy)tropolone

The title compound (systematic name: 5-acetoxy-2-hydroxy-2,4,6-cycloheptatrien-1-one), C<sub>9</sub>H<sub>8</sub>O<sub>4</sub>, contains intra- and intermolecular O-H···O hydrogen bonds, resulting in centrosymmetric dimers. The acetoxy group makes an angle of 59.86 (5)° with the seven-membered tropolone ring system. The crystal packing is consolidated by  $\pi$ - $\pi$  and C-H···O interactions.

#### Comment

Troponoids, being a remarkable class of non-benzenoid  $\pi$ conjugated systems, have been used as the building blocks of various molecular assemblies, such as liquid crystals (Kubo, 2004; Kubo, Mori *et al.*, 2005) and organogelators (Kubo & Mori, 2005). Recently, the mesomorphic properties of two substituted monocyclic troponoid mesogens, 2,5-dialkanoyloxytropones and 5-alkanoyloxy-2-alkoxytropones, were reported (Mori *et al.*, 2001). The crystal structure of tropolone (Shimanouchi & Sasada, 1973) and its 5-cyano- and 5-nitrotropolone derivatives (Kubo, Yamamoto & Mori, 2001) have been described. We now report the structure of the title compound, (I) (Fig. 1), with the aim of contributing to a deeper understanding of troponoids and their molecular assemblies.



In the molecule of (I), the dihedral angle between the leastsquares planes A (O1/O2/C1–C7) and B (O3/O4/C8/C9) is 59.86 (5)°, which is different from the equivalent value of 71.8° in 2-butanoyloxy-5-nitrotropone (Kubo, Tsuruta & Mori, 2001).

Compound (I) forms  $O-H\cdots O$  hydrogen-bonded dimers about inversion centres, involving the OH group and an intermolecular carbonyl  $O1^i$  acceptor (see Table 1 for symmetry code). An intramolecular  $O-H\cdots O$  bond to atom O1 occurs simultaneously (Table 1, Fig. 2). The intermolecular  $O2\cdots O1^i$  distance of 2.716 (2) Å is close to that in tropolone (2.746 Å), where a similar dimerization occurs.

Intermolecular  $\pi$ - $\pi$  interactions occur in (I) between the tropolone ring planes, leading to infinite stacks propagating in the [100] direction, with all molecules in a stack aligned in the same way, in a 'head-to-head' arrangement (Fig. 2). The C1-C7<sup>v</sup> [symmetry code: (v) x - 1, y, z] separation is 3.455 (2) Å,

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#### Figure 1

The molecular structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii. The intramolecular hydrogen bond is indicated by dashed lines.



#### Figure 2

A packing diagram for (I), viewed down the a axis, with hydrogen bonds shown as dotted lines.

which is within the range associated with  $\pi$ - $\pi$  interactions [3.3-3.8 Å]. This value is similar to the equivalent intermolecular separation in tropolone (3.418 Å; Shimanouchi & Sasada, 1973). However, in this latter compound, the stacks of molecules are generated by a 21 screw axis and adjacent molecules are aligned in a head-to-tail arrangement.

Various intermolecular  $C-H \cdots O$  interactions (Table 1) help to consolidate the crystal packing of (I). Similar interactions have been observed in related materials (Kubo, Matsumoto et al., 2005; Takahashi et al., 2006). The combination of intermolecular  $O-H \cdots O$  and  $C-H \cdots O$  interactions and  $\pi - \pi$  stacking interactions in (I) builds up a three-dimensional network.

# **Experimental**

Compound (I) was prepared by the esterification of 5-hydroxytropolone with acetyl chloride (Nagao et al., 1989). Crystals of (I) were grown from a methanol solution by slow evaporation.

Z = 4

 $D_r = 1.408 \text{ Mg m}^{-3}$ Cu  $K\alpha$  radiation

 $\mu = 0.96 \text{ mm}^{-1}$ 

Prism, colourless

 $0.33 \times 0.26 \times 0.10 \text{ mm}$ 

7929 measured reflections

1512 independent reflections 1407 reflections with  $F^2 > 2\sigma(F^2)$ 

T = 153.1 K

 $R_{\rm int} = 0.033$ 

 $\theta_{\rm max} = 68.1^\circ$ 

### Crystal data

 $C_9H_8O_4$  $M_r = 180.16$ Monoclinic,  $P2_1/c$ a = 3.8076 (12) Å b = 10.987 (3) Å c = 20.356 (6) Å  $\beta = 93.576 (17)^{\circ}$  $\dot{V} = 849.9$  (4) Å<sup>3</sup>

## Data collection

Rigaku R-AXIS RAPID diffractometer

 $\omega$  scans Absorption correction: multi-scan (ABSCOR; Higashi, 1995)  $T_{\min} = 0.656, \ T_{\max} = 0.909$ 

### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0452P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.039$	+ 0.5664P]
$wR(F^2) = 0.104$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.04	$(\Delta/\sigma)_{\rm max} < 0.001$
1512 reflections	$\Delta \rho_{\rm max} = 0.38 \ {\rm e} \ {\rm \AA}^{-3}$
124 parameters	$\Delta \rho_{\rm min} = -0.39 \text{ e } \text{\AA}^{-3}$
H atoms treated by a mixture of	Extinction correction: SHELXL97
independent and constrained	(Sheldrick 1997)
refinement	Extinction coefficient: 0.0126 (11)

### Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O2−H2···O1	0.90 (2)	2.09 (2)	2.589 (2)	114 (2)
$O2-H2\cdots O1^{i}$	0.90 (2)	1.93 (2)	2.716 (2)	145 (2)
C3−H3···O4 <sup>ii</sup>	0.95	2.53	3.309 (2)	139
C6−H5···O4 <sup>iii</sup>	0.95	2.67	3.433 (2)	138
C7−H6···O3 <sup>iv</sup>	0.95	2.50	3.438 (2)	171
C9−H7···O1 <sup>ii</sup>	0.98	2.64	3.586 (2)	163
6	i. 4.	(n) 1	3. (***)	1

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

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  $U_{iso}(H) = 1.2U_{eq}(C)$  or  $1.5U_{eq}(methyl C)$ .

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

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