

## 3,3-Bis(2-furylmethyl)-2-methyl-4-oxocyclohex-1-ene-1-carboxylic acid

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## Key indicators

Single-crystal X-ray study

T = 298 K

Mean  $\sigma(\text{C}-\text{C}) = 0.002 \text{ \AA}$ 

R factor = 0.048

wR factor = 0.139

Data-to-parameter ratio = 18.1

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

In the title molecule,  $\text{C}_{18}\text{H}_{18}\text{O}_5$ , the cyclohexene ring adopts an envelope conformation. The dihedral angle between the two furan rings is  $65.9(1)^\circ$ . In the crystal structure, the carboxyl groups of inversion-related molecules are involved in the formation of dimers through  $\text{O}-\text{H}\cdots\text{O}$  hydrogen bonds.

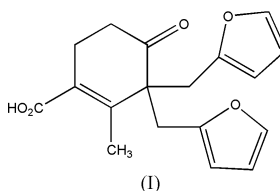
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## Comment

The title compound, (I), is a novel bisalkylation product of Hagemann's ester. Many natural and non-natural products involve spiro junctions. Recently, we obtained (I) as a by-product in the alkylation of Hagemann's ester with <sup>t</sup>BuOK/<sup>t</sup>BuOH/2-furylmethyl chloride (Chakraborty *et al.*, 1997). We report here the crystal and molecular structure of (I).



The title molecule contains one cyclohexene ring (C1–C6) and two furan rings (C8–C11/O1 and C13–C16/O2; Fig. 1). The dihedral angle between the two furan rings is  $65.9(1)^\circ$ . The cyclohexene ring adopts an envelope conformation, with atom C5 deviating from the mean plane passing through the remaining atoms in that ring by  $0.670(3) \text{ \AA}$ . The plane of the carboxyl group is twisted from the C1–C4/C6 plane by  $14.1(3)^\circ$ . The C2–C3 distance of  $1.346(2) \text{ \AA}$  confirms the double-bond character of this linkage. The molecular structure is stabilized by  $\text{C}-\text{H}\cdots\text{O}$  interactions (Table 2) involving the carbonyl O atoms. As usual, the carboxyl group serves, simultaneously, as hydrogen-bond donor and acceptor, to form dimers through a centrosymmetric pair of  $\text{O}-\text{H}\cdots\text{O}$  hydrogen bonds (Table 2 and Fig. 2). The geometry of this hydrogen bond agrees well with those reported by Jeffrey & Saenger (1994).

## Experimental

Hagemann's ester, on reaction with <sup>t</sup>BuOK in <sup>t</sup>BuOH, followed by refluxing with 2-furylmethyl chloride, afforded the monoalkylated Hagemann's ester along with the bisalkylated product (Chakraborty *et al.*, 1997). The crude reaction mixture, on subsequent hydrolysis with KOH/EtOH–H<sub>2</sub>O, furnished 2-(2-furylmethyl)-3-methylcyclohex-2-en-1-one and the title compound. Single crystals of the latter were grown by slow evaporation of an ethyl acetate solution of the title compound [m.p.: 415–416 K ( $\text{CH}_3\text{COOCH}_2\text{CH}_3$ )]. IR spectroscopy (KBr,  $\text{cm}^{-1}$ ): 3140, 1718, 1684; <sup>1</sup>H NMR (200 MHz,  $\text{CDCl}_3$ ):  $\delta$  (p.p.m.) 2.00–2.16 (*m*, 4 H), 2.26 (*s*, 3 H), 2.98 (*d*, 2 H,  $J = 14.5 \text{ Hz}$ ), 3.33(*d*, 2 H,  $J = 14.5 \text{ Hz}$ ), 5.93(*d*, 2 H,  $J = 3.25 \text{ Hz}$ ), 6.20 (*dd*, 2 H,  $J =$

3.25 and 1.9 Hz), 7.23 (*d*, 2 H, *J* = 1.9 Hz); <sup>13</sup>C NMR (50 MHz, CDCl<sub>3</sub>): δ (p.p.m.) 16.72, 23.53, 35.60, 38.06, 58.29, 107.79, 110.27, 127.61, 141.66, 148.80, 150.79, 172.99, 211.88; MS (*m/z*): 314 (M<sup>+</sup>), 233, 148, 81 (base peak), 53.

Crystal data

C<sub>18</sub>H<sub>18</sub>O<sub>5</sub> *D<sub>x</sub>* = 1.300 Mg m<sup>-3</sup>  
*M<sub>r</sub>* = 314.32 Mo *Kα* radiation  
 Monoclinic, *P*<sub>2</sub><sub>1</sub>/*n* Cell parameters from 5554 reflections  
*a* = 7.125 (1) Å reflections  
*b* = 8.798 (1) Å *θ* = 2.0–28.0°  
*c* = 25.625 (2) Å *μ* = 0.10 mm<sup>-1</sup>  
*β* = 91.214 (1)° *T* = 298 (2) K  
*V* = 1606.0 (3) Å<sup>3</sup> Block, colourless  
*Z* = 4 0.55 × 0.55 × 0.55 mm

Data collection

Bruker SMART CCD area-detector 3824 independent reflections  
 diffractometer 3057 reflections with *I* > 2σ(*I*)  
*ω* scans *R*<sub>int</sub> = 0.020  
 Absorption correction: multi-scan *θ*<sub>max</sub> = 28.3°  
 (SADABS; Sheldrick, 1996) *h* = -9 → 9  
*T*<sub>min</sub> = 0.866, *T*<sub>max</sub> = 0.950 *k* = -7 → 11  
 9924 measured reflections *l* = -34 → 34

Refinement

Refinement on *F*<sup>2</sup> *w* = 1/[σ<sup>2</sup>(*F<sub>o</sub>*<sup>2</sup>) + (0.0606*P*)<sup>2</sup> + 0.5648*P*]  
*R*[*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.048 where *P* = (*F<sub>o</sub>*<sup>2</sup> + 2*F<sub>c</sub>*<sup>2</sup>)/3  
*wR*(*F*<sup>2</sup>) = 0.139 (Δ/σ)<sub>max</sub> < 0.001  
*S* = 1.06 Δ*ρ*<sub>max</sub> = 0.44 e Å<sup>-3</sup>  
 3824 reflections Δ*ρ*<sub>min</sub> = -0.29 e Å<sup>-3</sup>  
 211 parameters Extinction correction: *SHELXL97*  
 H-atom parameters constrained Extinction coefficient: 0.036 (3)

Table 1 Selected geometric parameters (Å, °).

O3–C18	1.216 (2)	C8–C9	1.343 (2)
O4–C18	1.299 (2)	C10–C11	1.320 (3)
O5–C6	1.212 (2)	C13–C14	1.345 (3)
C2–C3	1.346 (2)	C15–C16	1.316 (3)
C2–C3–C18–O3	11.8 (3)	C2–C3–C18–O4	-167.8 (2)
C4–C3–C18–O3	-172.2 (2)	C4–C3–C18–O4	8.2 (2)

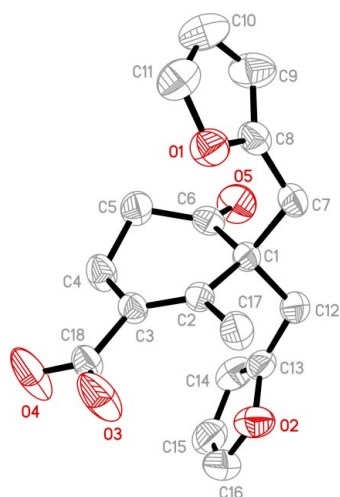


Figure 1 A view of the title molecule showing 50% probability displacement ellipsoids and the atomic numbering scheme. H atoms have been omitted for clarity.

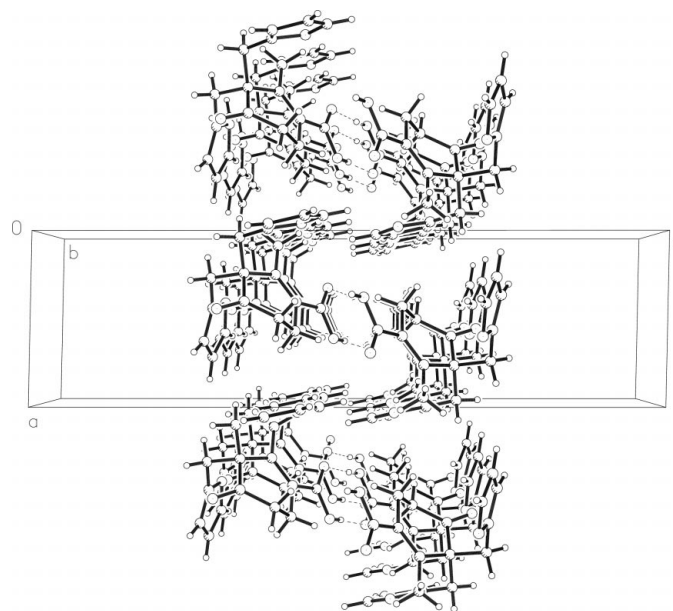


Figure 2 Part of the molecular packing, showing the dimer formation.

Table 2 Hydrogen-bonding geometry (Å, °).

<i>D</i> – <i>H</i> ··· <i>A</i>	<i>D</i> – <i>H</i>	<i>H</i> ··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> – <i>H</i> ··· <i>A</i>
O4–H4···O3 <sup>i</sup>	0.82	1.83	2.649 (2)	176
C12–H12A···O5	0.97	2.47	2.828 (2)	101
C17–H17C···O3	0.96	2.40	2.777 (2)	103

Symmetry codes: (i) -*x*, 1 - *y*, -*z*.

All the H atoms were positioned geometrically and allowed to ride on the parent atoms, with aromatic C–H = 0.93 Å, methylene C–H = 0.97 Å, methyl C–H = 0.96 Å and O–H = 0.82 Å. The *U*<sub>iso</sub>(H) values were set at 1.5*U*<sub>eq</sub>(C) for the methyl H atoms and at 1.2*U*<sub>eq</sub>(C) for other C-bound H atoms; *U*<sub>iso</sub>(H) for the carboxyl H atom was set equal to 1.5*U*<sub>eq</sub>(O). A rotating group model was used for the methyl group. Owing to the large fraction of weak data at higher angles, the completeness is slightly low (96%).

Data collection: *SMART* (Siemens, 1995); cell refinement: *SAINT* (Siemens, 1995); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXTL*.

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