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Key indicators

Single-crystal X-ray study
 $T = 173$ K
Mean $\sigma(\text{C}-\text{C}) = 0.005$ Å
 R factor = 0.064
 wR factor = 0.216
Data-to-parameter ratio = 15.6

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

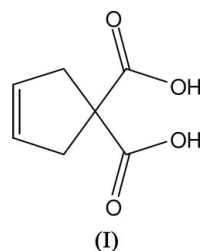
Cyclopent-3-ene-1,1-dicarboxylic acid

The cyclopentene ring of the title compound, $\text{C}_7\text{H}_8\text{O}_4$, is essentially planar. The molecules form zigzag chains *via* $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds.

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Comment

The title compound, (I), is the product of a ring-closing metathesis (Rölle & Grubbs, 2002). A perspective view of (I) is shown in Fig. 1. Bond lengths and angles can be regarded as normal (Cambridge Structural Database, Version 5.27 plus one update; *MOGUL* Version 1.1; Allen, 2002). The cyclopentene ring is essentially planar (r.m.s. deviation of the ring atoms = 0.031 Å). The two carboxyl groups form dihedral angles of 67.2 (3) and 63.4 (4)° [for C11/O11/O12 and C21/O21/O22, respectively], with the cyclopentene ring. The molecules form zigzag chains *via* $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds (Fig. 2).

**Experimental**

The title compound was synthesized according to the procedure described by Rölle & Grubbs (2002). Colourless crystals suitable for X-ray diffraction were grown from a methanol solution of (I).

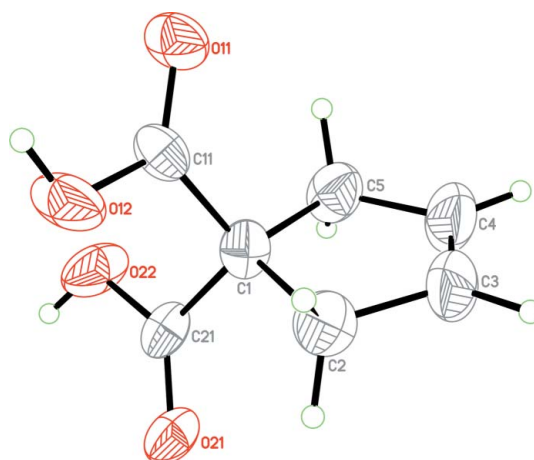


Figure 1
Perspective view of the title compound with the atom-numbering scheme; displacement ellipsoids are drawn at the 30% probability level.

Crystal data

C₇H₈O₄
M_r = 156.13
 Monoclinic, *C*2/*c*
a = 20.543 (8) Å
b = 6.179 (2) Å
c = 12.394 (5) Å
 β = 98.35 (3)°
V = 1556.6 (10) Å³
Z = 8

D_x = 1.333 Mg m⁻³
 Mo *K*α radiation
 Cell parameters from 4322 reflections
 θ = 3.2–24.3°
 μ = 0.11 mm⁻¹
T = 173 (2) K
 Block, colourless
 0.30 × 0.20 × 0.20 mm

Data collection

Bruker SMART CCD three-circle diffractometer
 ω scans
 Absorption correction: none
 9705 measured reflections
 1589 independent reflections

930 reflections with *I* > 2σ(*I*)
*R*_{int} = 0.057
 θ_{max} = 26.4°
h = -25 → 25
k = -7 → 7
l = -15 → 15

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.064
wR(*F*²) = 0.217
S = 0.95
 1589 reflections
 102 parameters
 H-atom parameters constrained

w = 1/[σ²(*F_o*²) + (0.1161*P*)² + 1.782*P*]
 where *P* = (*F_o*² + 2*F_c*²)/3
 (Δ/*σ*)_{max} < 0.001
 Δρ_{max} = 0.28 e Å⁻³
 Δρ_{min} = -0.21 e Å⁻³

Table 1

Selected geometric parameters (Å, °).

C11—O11	1.208 (4)	C21—O21	1.210 (4)
C11—O12	1.294 (4)	C21—O22	1.315 (4)
O11—C11—O12	123.6 (3)	O21—C21—O22	124.2 (3)
O11—C11—C1	123.6 (3)	O21—C21—C1	124.5 (3)
O12—C11—C1	112.7 (3)	O22—C21—C1	111.3 (3)

Table 2

Hydrogen-bond geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
O12—H12...O11 ⁱ	0.84	1.79	2.619 (3)	168
O22—H22...O21 ⁱⁱ	0.84	1.81	2.650 (4)	173

Symmetry codes: (i) -*x*, -*y* + 2, -*z* + 1; (ii) -*x* + ½, -*y* + ¾, -*z* + 1.

H atoms were located in a difference map, but refined with fixed individual displacement parameters [*U*_{iso}(H) = 1.2*U*_{eq}(C)] using a

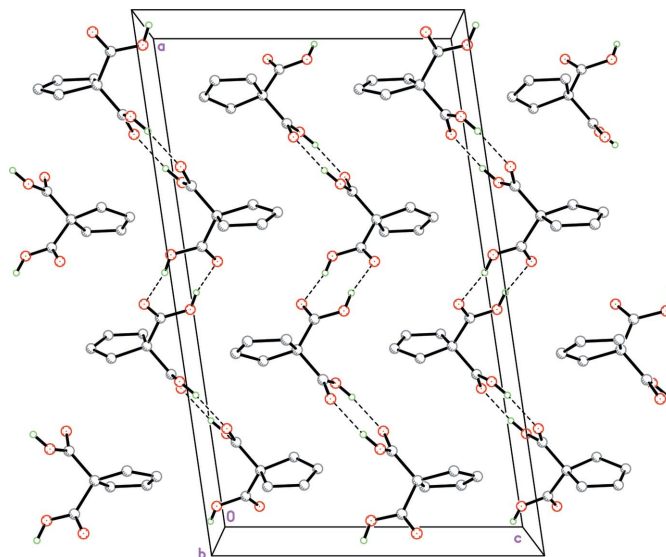


Figure 2

Packing diagram of the title compound, showing the hydrogen bonds as dashed lines. The view is projected on to the *ac* plane and H atoms not involved in hydrogen bonding have been omitted for clarity.

riding model, with O—H = 0.84 Å, and C—H = 0.95 and 0.99 Å for *Csp*² and methylene C, respectively. In addition, the torsion angles about the C—O bonds of the hydroxyl groups were refined.

Data collection: *SMART* (Siemens, 1995); cell refinement: *SAINT* (Siemens, 1995); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* in *SHELXTL-Plus* (Sheldrick, 1991); software used to prepare material for publication: *SHELXL97*.

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