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3-(4-Methoxybenzoyl)propionic acid

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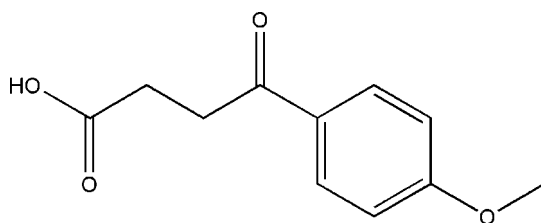
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Key indicators: single-crystal X-ray study; $T = 150$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.048; wR factor = 0.112; data-to-parameter ratio = 16.4.

In the crystal of the title compound, $\text{C}_{11}\text{H}_{12}\text{O}_4$, inversion dimers arise from pairs of intermolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds and $\text{C}-\text{H}\cdots\text{O}$ bonds further consolidate the packing. There is also a $\text{C}-\text{H}\cdots\pi$ contact between the benzene ring and the methylene group.

Related literature

For general background, see: Hashem *et al.* (2007); Husain *et al.* (2005). For bond-length data, see: Allen *et al.* (1987).



Experimental

Crystal data

$\text{C}_{11}\text{H}_{12}\text{O}_4$
 $M_r = 208.21$
 Monoclinic, $P2_1/c$
 $a = 5.0511$ (3) Å
 $b = 10.0219$ (7) Å
 $c = 20.0840$ (12) Å
 $\beta = 90.107$ (6)°

$V = 1016.67$ (11) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.10$ mm⁻¹
 $T = 150$ (1) K
 $0.20 \times 0.18 \times 0.13$ mm

Data collection

Bruker–Nonius KappaCCD area-detector diffractometer
 Absorption correction: integration (Coppens, 1970)
 $T_{\min} = 0.979$, $T_{\max} = 0.987$

8320 measured reflections
 2236 independent reflections
 1662 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.048$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.048$
 $wR(F^2) = 0.112$
 $S = 1.13$
 2236 reflections

136 parameters
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.18$ e Å⁻³
 $\Delta\rho_{\min} = -0.20$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O2}-\text{H2}\cdots\text{O1}^{\text{i}}$	0.82	1.81	2.628 (3)	173
$\text{C6}-\text{H6}\cdots\text{O3}^{\text{ii}}$	0.93	2.34	3.247 (3)	164
$\text{C11}-\text{H11B}\cdots\text{O4}^{\text{iii}}$	0.96	2.60	3.328 (3)	133
$\text{C3}-\text{H3B}\cdots\text{Cg1}^{\text{iv}}$	0.97	2.74	3.591 (3)	146

Symmetry codes: (i) $-x, -y, -z$; (ii) $-x, y - \frac{1}{2}, -z + \frac{1}{2}$; (iii) $-x + 2, -y, -z + 1$; (iv) $x + 1, y, z$. Cg1 is the centroid of the phenyl ring.

Data collection: *COLLECT* (Hooft, 1998); cell refinement: *COLLECT* and *DENZO* (Otwinowski & Minor, 1997); data reduction: *COLLECT* and *DENZO*; program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HK2556).

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supplementary materials

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3-(4-Methoxybenzoyl)propionic acid

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Comment

Benzoyl propionic acids are important intermediates in heterocyclic chemistry and have been used for the synthesis of various biologically active five -membered heterocycles such as butenolides, pyrrolones (Husain *et al.*, 2005), oxadiazoles and triazoles (Hashem *et al.*, 2007). In view of the versatility of these compounds, we synthesized the title compound and reported herein its crystal structure.

In the title compound (Fig. 1), the bond lengths (Allen *et al.*, 1987) and angles are within normal ranges. O3, O4, C2, C3 and C4 atoms are 0.067 (3), -0.003 (3), -0.163 (4), -0.013 (3) and 0.016 (3) Å away from the phenyl plane, respectively.

In the crystal structure, intermolecular O-H...O and C-H...O hydrogen bonds (Table 1) link the molecules (Fig. 2), in which they may be effective in the stabilization of the structure. There also exist a C—H... π contact (Table 1) between the phenyl ring and the methylene group.

Experimental

The title compound was synthesized by the condensation of succinic anhydride (2 g, 20 mmol) with anisol (10 ml) in the presence of aluminium chloride (6 g, 42 mmol). The reaction mixture was refluxed for 4 h. After completion of the reaction, excess solvent (anisol) was removed by steam distillation. The resultant solid product was purified by dissolving it in sodium hydroxide solution (5%, w/v), filtering followed by addition of hydrochloric acid. The obtained solid mass was filtered, washed with cold water, dried and crystallized from methanol (yield; 55%, m.p. 419-420 K)

Refinement

H atoms were positioned geometrically, with O-H = 0.82 Å (for OH) and C-H = 0.93, 0.97 and 0.96 Å for aromatic, methylene and methyl H, respectively, and constrained to ride on their parent atoms with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C},\text{O})$.

Figures

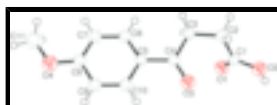


Fig. 1. The molecular structure of the title molecule, with the atom-numbering scheme.

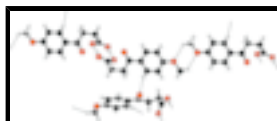


Fig. 2. A partial packing diagram. Hydrogen bonds are shown as dashed lines.



Fig. 3. The formation of the title compound.

3-(4-Methoxybenzoyl)propionic acid

Crystal data

$C_{11}H_{12}O_4$	$F_{000} = 440$
$M_r = 208.21$	$D_x = 1.360 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Melting point: 419(1) K
Hall symbol: -P 2ybc	Mo $K\alpha$ radiation
$a = 5.0511 (3) \text{ \AA}$	$\lambda = 0.71073 \text{ \AA}$
$b = 10.0219 (7) \text{ \AA}$	Cell parameters from 8408 reflections
$c = 20.0840 (12) \text{ \AA}$	$\theta = 1-27.5^\circ$
$\beta = 90.107 (6)^\circ$	$\mu = 0.10 \text{ mm}^{-1}$
$V = 1016.67 (11) \text{ \AA}^3$	$T = 150 (1) \text{ K}$
$Z = 4$	Block, colorless
	$0.20 \times 0.18 \times 0.13 \text{ mm}$

Data collection

Bruker–Nonius KappaCCD area-detector diffractometer	2236 independent reflections
Radiation source: fine-focus sealed tube	1662 reflections with $I > 2\sigma(I)$
Monochromator: graphite	$R_{\text{int}} = 0.048$
Detector resolution: 9.091 pixels mm^{-1}	$\theta_{\text{max}} = 27.5^\circ$
$T = 150(1) \text{ K}$	$\theta_{\text{min}} = 2.0^\circ$
φ and ω scans	$h = -6 \rightarrow 6$
Absorption correction: integration (Coppens, 1970)	$k = -13 \rightarrow 13$
$T_{\text{min}} = 0.979$, $T_{\text{max}} = 0.987$	$l = -26 \rightarrow 22$
8320 measured reflections	

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.048$	H-atom parameters constrained
$wR(F^2) = 0.112$	$w = 1/[\sigma^2(F_o^2) + (0.0363P)^2 + 0.36P]$
$S = 1.13$	where $P = (F_o^2 + 2F_c^2)/3$
2236 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
136 parameters	$\Delta\rho_{\text{max}} = 0.18 \text{ e \AA}^{-3}$
Primary atom site location: structure-invariant direct methods	$\Delta\rho_{\text{min}} = -0.20 \text{ e \AA}^{-3}$
	Extinction correction: none

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.0482 (3)	-0.08173 (13)	0.07156 (6)	0.0440 (4)
O2	-0.2693 (3)	0.06464 (14)	0.04647 (6)	0.0443 (4)
H2	-0.1915	0.0654	0.0107	0.053*
O3	0.1138 (2)	0.13027 (12)	0.19230 (6)	0.0398 (3)
O4	0.7706 (3)	-0.03563 (14)	0.44558 (7)	0.0477 (4)
C1	-0.1507 (3)	-0.01793 (17)	0.08665 (8)	0.0323 (4)
C2	-0.2850 (3)	-0.03301 (19)	0.15227 (8)	0.0355 (4)
H2A	-0.4267	-0.0977	0.1478	0.043*
H2B	-0.3645	0.0517	0.1644	0.043*
C3	-0.1025 (3)	-0.07690 (17)	0.20779 (8)	0.0317 (4)
H3A	-0.0091	-0.1569	0.1941	0.038*
H3B	-0.2075	-0.0989	0.2467	0.038*
C4	0.0958 (3)	0.02952 (16)	0.22605 (8)	0.0297 (4)
C5	0.2686 (3)	0.00865 (16)	0.28458 (8)	0.0287 (4)
C6	0.2583 (3)	-0.10711 (17)	0.32209 (8)	0.0333 (4)
H6	0.1367	-0.1730	0.3108	0.040*
C7	0.4243 (3)	-0.12673 (18)	0.37606 (9)	0.0363 (4)
H7	0.4169	-0.2059	0.4002	0.044*
C8	0.6011 (3)	-0.02784 (18)	0.39356 (8)	0.0343 (4)
C9	0.6134 (4)	0.09014 (18)	0.35680 (9)	0.0365 (4)
H9	0.7319	0.1570	0.3688	0.044*
C10	0.4502 (3)	0.10717 (17)	0.30291 (9)	0.0339 (4)
H10	0.4605	0.1855	0.2782	0.041*
C11	0.7813 (5)	-0.1587 (2)	0.48107 (11)	0.0561 (6)
H11A	0.6127	-0.1758	0.5013	0.067*
H11B	0.9151	-0.1535	0.5150	0.067*
H11C	0.8233	-0.2297	0.4508	0.067*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0500 (8)	0.0510 (8)	0.0311 (7)	0.0157 (7)	0.0057 (6)	-0.0004 (6)

supplementary materials

O2	0.0502 (8)	0.0558 (8)	0.0268 (6)	0.0147 (7)	0.0026 (6)	0.0017 (6)
O3	0.0442 (7)	0.0337 (7)	0.0413 (7)	0.0016 (6)	-0.0026 (6)	0.0066 (6)
O4	0.0528 (8)	0.0466 (8)	0.0436 (8)	-0.0063 (7)	-0.0184 (6)	0.0049 (6)
C1	0.0358 (9)	0.0341 (9)	0.0268 (8)	-0.0008 (8)	-0.0027 (7)	-0.0039 (7)
C2	0.0317 (8)	0.0441 (10)	0.0307 (9)	-0.0015 (8)	0.0018 (7)	-0.0002 (8)
C3	0.0323 (8)	0.0360 (9)	0.0267 (8)	0.0005 (7)	0.0022 (7)	-0.0010 (7)
C4	0.0303 (8)	0.0295 (9)	0.0293 (8)	0.0062 (7)	0.0066 (7)	-0.0007 (7)
C5	0.0285 (8)	0.0288 (8)	0.0288 (8)	0.0017 (7)	0.0039 (6)	-0.0024 (7)
C6	0.0355 (9)	0.0306 (9)	0.0338 (9)	-0.0057 (7)	0.0001 (7)	-0.0005 (7)
C7	0.0408 (9)	0.0334 (9)	0.0348 (9)	-0.0025 (8)	-0.0015 (8)	0.0046 (7)
C8	0.0343 (9)	0.0381 (10)	0.0304 (9)	0.0015 (8)	-0.0025 (7)	-0.0023 (7)
C9	0.0372 (9)	0.0320 (9)	0.0402 (10)	-0.0068 (8)	-0.0040 (8)	-0.0032 (8)
C10	0.0364 (9)	0.0288 (9)	0.0364 (9)	0.0001 (7)	0.0031 (7)	0.0006 (7)
C11	0.0659 (14)	0.0541 (13)	0.0481 (12)	-0.0033 (11)	-0.0225 (10)	0.0106 (10)

Geometric parameters (Å, °)

O1—C1	1.230 (2)	C5—C6	1.384 (2)
O2—C1	1.301 (2)	C5—C10	1.396 (2)
O2—H2	0.8201	C6—C7	1.383 (2)
O3—C4	1.220 (2)	C6—H6	0.9300
O4—C8	1.352 (2)	C7—H7	0.9301
O4—C11	1.425 (2)	C8—C7	1.379 (2)
C1—C2	1.491 (2)	C8—C9	1.395 (2)
C2—C3	1.511 (2)	C9—H9	0.9300
C2—H2A	0.9700	C10—C9	1.370 (2)
C2—H2B	0.9699	C10—H10	0.9299
C3—H3A	0.9700	C11—H11A	0.9600
C3—H3B	0.9701	C11—H11B	0.9600
C4—C3	1.508 (2)	C11—H11C	0.9600
C5—C4	1.478 (2)		
C1—O2—H2	109.2	C10—C5—C4	119.76 (15)
C8—O4—C11	117.38 (15)	C7—C6—C5	121.48 (16)
O1—C1—O2	123.60 (15)	C7—C6—H6	119.3
O1—C1—C2	122.58 (16)	C5—C6—H6	119.2
O2—C1—C2	113.77 (15)	C8—C7—C6	119.26 (16)
C1—C2—C3	113.83 (14)	C8—C7—H7	120.4
C1—C2—H2A	108.8	C6—C7—H7	120.3
C3—C2—H2A	108.8	O4—C8—C7	124.36 (17)
C1—C2—H2B	108.8	O4—C8—C9	115.40 (16)
C3—C2—H2B	108.7	C7—C8—C9	120.24 (16)
H2A—C2—H2B	107.6	C10—C9—C8	119.74 (16)
C4—C3—C2	112.19 (15)	C10—C9—H9	120.1
C4—C3—H3A	109.3	C8—C9—H9	120.2
C2—C3—H3A	109.2	C9—C10—C5	120.94 (16)
C4—C3—H3B	109.2	C9—C10—H10	119.5
C2—C3—H3B	109.1	C5—C10—H10	119.6
H3A—C3—H3B	107.9	O4—C11—H11A	109.5
O3—C4—C5	120.98 (15)	O4—C11—H11B	109.5

O3—C4—C3	120.04 (15)	H11A—C11—H11B	109.5
C5—C4—C3	118.98 (14)	O4—C11—H11C	109.4
C6—C5—C10	118.33 (15)	H11A—C11—H11C	109.5
C6—C5—C4	121.91 (15)	H11B—C11—H11C	109.5

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
O2—H2 \cdots O1 ⁱ	0.82	1.81	2.628 (3)	173
C6—H6 \cdots O3 ⁱⁱ	0.93	2.34	3.247 (3)	164
C11—H11B \cdots O4 ⁱⁱⁱ	0.96	2.60	3.328 (3)	133
C3—H3B \cdots Cg1 ^{iv}	0.97	2.74	3.591 (3)	146

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

Fig. 1

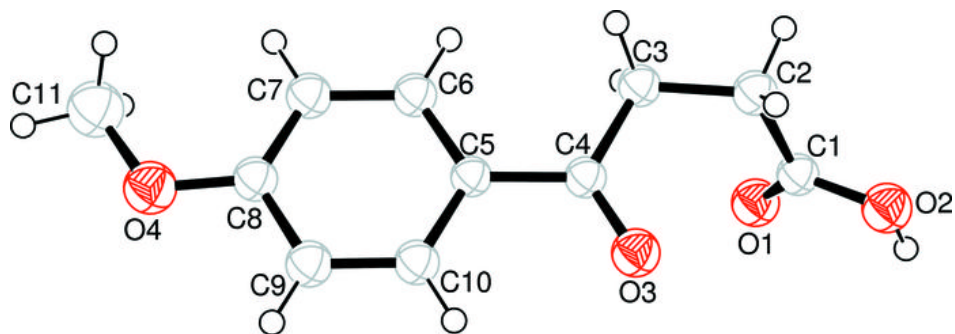


Fig. 2

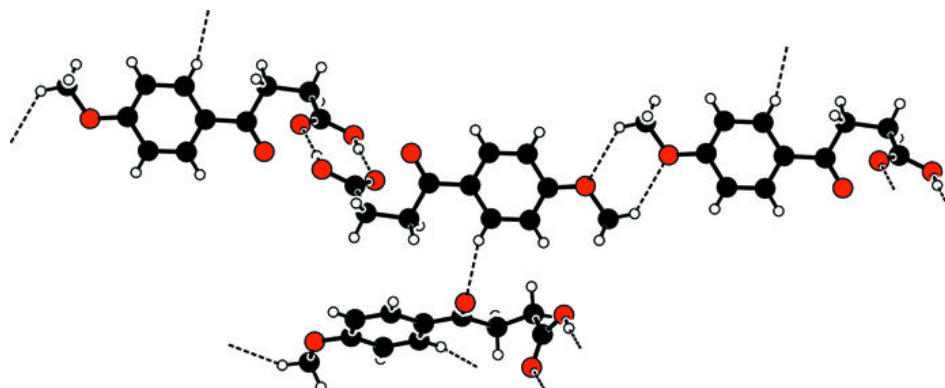


Fig. 3

