

2-Methyl-3-(3-methylphenyl)acrylic acid

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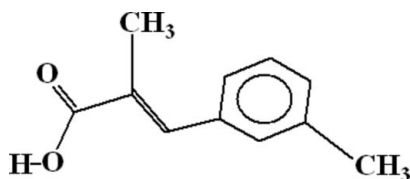
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.056; wR factor = 0.185; data-to-parameter ratio = 23.5.

The crystal structure of the title compound, $\text{C}_{11}\text{H}_{12}\text{O}_2$, consists of dimers which are formed due to intermolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonding. The dimers are linked to each other by $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds, where $\text{C}-\text{H}$ belongs to the benzene ring and the O atom is of a carbonyl group of an adjoining molecule. There exist two intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds which form five-membered rings. There exist two $\pi-\pi$ interactions between the benzene rings. The perpendicular distance in these interactions are 3.006 and 3.396 Å. There also exist $\text{C}-\text{H}\cdots\pi$ and $\text{C}-\text{O}\cdots\pi$ interactions.

Related literature

For related literature, see: Bernstein *et al.* (1995); Liu *et al.* (1999); Muhammad *et al.* (2007); Natella *et al.* (1999); Niaz *et al.* (2008); Parez-Alvarez *et al.* (2001); Wiesner *et al.* (2001).



Experimental

Crystal data

$\text{C}_{11}\text{H}_{12}\text{O}_2$	$V = 959.0$ (2) Å ³
$M_r = 176.21$	$Z = 4$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 7.4430$ (9) Å	$\mu = 0.08$ mm ⁻¹
$b = 13.4094$ (16) Å	$T = 296$ (2) K
$c = 10.2746$ (12) Å	$0.26 \times 0.18 \times 0.15$ mm
$\beta = 110.745$ (4)°	

Data collection

Bruker Kappa APEXII CCD diffractometer	Absorption correction: multi-scan (SADABS; Bruker, 2005)
	$T_{\min} = 0.980$, $T_{\max} = 0.986$
	11342 measured reflections

2820 independent reflections
1075 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.047$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.056$	120 parameters
$wR(F^2) = 0.185$	H-atom parameters constrained
$S = 1.02$	$\Delta\rho_{\text{max}} = 0.23$ e Å ⁻³
2820 reflections	$\Delta\rho_{\text{min}} = -0.20$ e Å ⁻³

Table 1

Hydrogen-bond geometry (Å, °).

C_g is the centroid of the C4–C9 benzene ring.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{O1}-\text{H1}\cdots\text{O2}^{\text{i}}$	0.82	1.83	2.611 (3)	160
$\text{C3}-\text{H3}\cdots\text{O1}$	0.93	2.27	2.703 (3)	108
$\text{C8}-\text{H8}\cdots\text{O1}^{\text{ii}}$	0.93	2.59	3.394 (3)	145
$\text{C10}-\text{H10A}\cdots\text{O2}$	0.96	2.31	2.783 (3)	109
$\text{C10}-\text{H10C}\cdots\text{C}_g^{\text{iii}}$	0.96	2.75	3.610 (3)	149
$\text{C1}-\text{O2}\cdots\text{C}_g^{\text{iv}}$	1.25 (1)	3.57 (1)	3.895 (3)	95 (1)

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

Data collection: APEX2 (Bruker, 2007); cell refinement: APEX2; data reduction: SAINT (Bruker, 2007); program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997) and PLATON (Spek, 2003); software used to prepare material for publication: WinGX publication routines (Farrugia, 1999) and PLATON.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: AT2580).

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supporting information

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2-Methyl-3-(3-methylphenyl)acrylic acid

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S1. Comment

Cinnamic acids and their derivatives have been studied for their pharmacological properties, including hepatoprotective (Parez-Alvarez *et al.*, 2001), antimalarial (Wiesner *et al.*, 2001), antioxidant (Natella *et al.*, 1999) and antihyperglycemic activities (Liu *et al.*, 1999). In continuation of our efforts to synthesize various derivatives of cinnamic acids (Niaz *et al.*, 2008) and their complexes, we herein report the structure of the title compound (I).

The crystal structure of 3-(4-bromophenyl)-2-methylacrylic acid (Muhammad *et al.*, 2007) has been reported. The title compound (I) have a similar environment about the carboxylate group but the attachment of methyl instead of Br-atom is at *meta*-position instead of *para*-position.

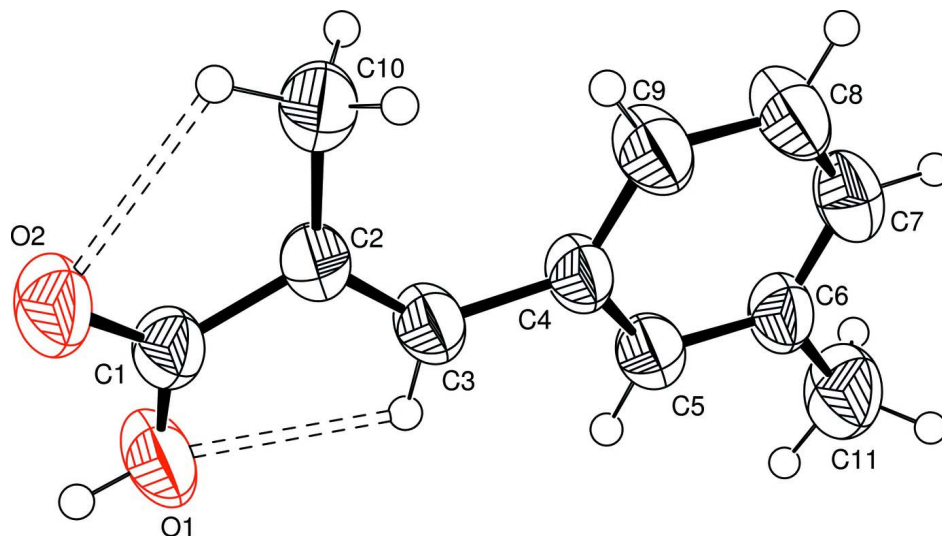
In the crystal structure of the title compound, the C—C bonds are in the range 1.462 (3)–1.500 (3) Å, and C=C have a value of 1.339 (3) Å. The resonant C—O bonds have values of 1.250 (3) and 1.271 (3) Å. In the asymmetric unit, there are two intermolecular H-bonds of C—H \cdots O type (Table 2, Fig 1). Due to these H-bonds, two five membered rings (O1/C1/C2/C3/H3 \cdots O1) and (O2/C1/C2/C10/H10A \cdots O2) are formed. The intermolecular (O1—H1 \cdots O2ⁱ [symmetry code: i = -x, -y, -z + 1]) hydrogen bond forces the molecules into centrosymmetric dimers, forming a $R_2^2(8)$ motif (Bernstein *et al.* 1995). These dimers are linked to each other by the second intermolecular H-bonding, C8—H8 \cdots O2ⁱⁱ [symmetry code: ii = -x + 1, y + 1/2, -z + 1/2] as shown in Fig 2. There exist π - π interactions between the centroids (Cg) of benzene (C4—C9) rings of adjacent molecules. The $Cg\cdots Cg^{\text{iii}}$ [symmetry code: iii = x, -y + 1/2, z - 1/2] and $Cg\cdots Cg^{\text{iv}}$ [symmetry code: iv = x, -y + 1/2, z + 1/2] have a perpendicular distance of 3.006 and 3.396 Å, respectively. There exist also C10—H10C $\cdots Cg^{\text{iv}}$ interaction, with a distance of 3.610 (3) Å between C10 and Cg^{iv} . Similarly another π -interaction is present between C1—O2 and Cg^{v} [symmetry code: v = x - 1, y, z] with a distance of 3.895 (3) Å between C1 and Cg^{v} . The detail of these interactions is also included in Table 1.

S2. Experimental

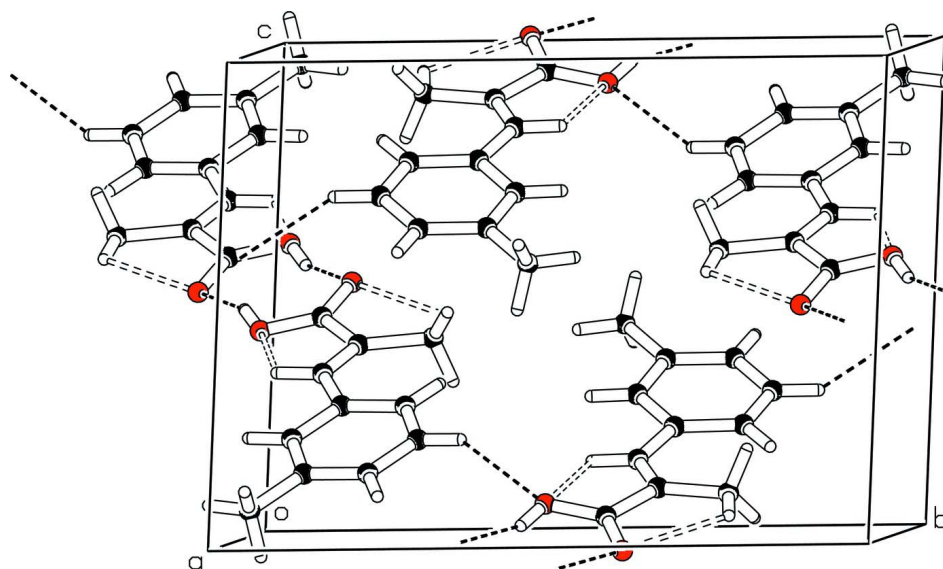
Compound (I) was prepared according to our previously reported method (Muhammad *et al.*, 2007). A mixture of 3-methylbenzaldehyde (10 mmol, 1.18 ml), methylmalonic acid (2.36 g, 20 mmol) and piperidine (20 mmol, 1.98 ml) in pyridine (12.5 ml) solution was heated on a steam-bath for 24 h. The reaction mixture was cooled and added to a mixture of 25 ml of concentrated HCl and 50 g of ice. The precipitate formed in the acidified mixture was filtered off and washed with ice-cold water. The product was recrystallized from ethanol [yield; 90%, m.p. 321 K].

S3. Refinement

All H-atoms were positioned geometrically, with C—H = 0.93 and 0.96 Å for aromatic and methyl H-atoms, respectively, and O—H = 0.82 Å for hydroxyl O-atom, and constrained to ride on their parent atoms. The thermal parameters of methyl and hydroxyl H-atoms was taken 1.5 times while for all other H-atoms it was taken 1.2 times of the parent atoms.

**Figure 1**

ORTEP drawing of the title compound, with the atom numbering scheme. The thermal ellipsoids are drawn at the 50% probability level. H-atoms are shown by small circles of arbitrary radii. The intramolecular H-bonds are shown by dotted lines.

**Figure 2**

The packing figure (*PLATON*: Spek, 2003) which shows the dimeric nature of the compound and the interlinkages of the dimers.

2-Methyl-3-(3-methylphenyl)acrylic acid

Crystal data

$C_{11}H_{12}O_2$

$M_r = 176.21$

Monoclinic, $P2_1/c$

Hall symbol: $-P 2_1/c$

$a = 7.4430 (9) \text{ \AA}$

$b = 13.4094 (16) \text{ \AA}$

$c = 10.2746 (12) \text{ \AA}$

$\beta = 110.745 (4)^\circ$

$V = 959.0 (2) \text{ \AA}^3$

$Z = 4$

$F(000) = 376$
 $D_x = 1.220 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
 Cell parameters from 2820 reflections
 $\theta = 2.6\text{--}30.2^\circ$

$\mu = 0.08 \text{ mm}^{-1}$
 $T = 296 \text{ K}$
 Prismatic, colourless
 $0.26 \times 0.18 \times 0.15 \text{ mm}$

Data collection

Bruker KAPPA APEXII CCD
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 Detector resolution: $7.2 \text{ pixels mm}^{-1}$
 ω scans
 Absorption correction: multi-scan
 (SADABS; Bruker, 2005)
 $T_{\min} = 0.980$, $T_{\max} = 0.986$

11342 measured reflections
 2820 independent reflections
 1075 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.047$
 $\theta_{\max} = 30.2^\circ$, $\theta_{\min} = 2.6^\circ$
 $h = -10 \rightarrow 8$
 $k = -18 \rightarrow 13$
 $l = -10 \rightarrow 14$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.056$
 $wR(F^2) = 0.185$
 $S = 1.02$
 2820 reflections
 120 parameters
 0 restraints
 Primary atom site location: structure-invariant
 direct methods
 Secondary atom site location: difference Fourier
 map

Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0657P)^2 + 0.191P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.24 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.20 \text{ e \AA}^{-3}$
 Extinction correction: empirical,
 $F_c^* = kF_c[1 + 0.001x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$
 Extinction coefficient: $0.006(2)$

Special details

Geometry. Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.1630 (3)	-0.00578 (12)	0.4174 (2)	0.0795 (8)
O2	0.0675 (2)	0.12222 (11)	0.51106 (18)	0.0699 (7)
C1	0.1639 (3)	0.08682 (17)	0.4441 (2)	0.0515 (8)
C2	0.2797 (3)	0.15393 (16)	0.3911 (2)	0.0484 (7)
C3	0.4010 (3)	0.11136 (17)	0.3377 (2)	0.0533 (8)
C4	0.5340 (3)	0.15354 (16)	0.2764 (2)	0.0497 (8)
C5	0.6214 (3)	0.08770 (17)	0.2126 (2)	0.0534 (8)
C6	0.7448 (3)	0.11877 (18)	0.1461 (2)	0.0553 (8)
C7	0.7829 (4)	0.21882 (19)	0.1470 (3)	0.0637 (9)
C8	0.7031 (4)	0.2856 (2)	0.2118 (3)	0.0699 (10)

C9	0.5794 (4)	0.25412 (17)	0.2757 (3)	0.0646 (9)
C10	0.2492 (3)	0.26372 (17)	0.4024 (3)	0.0658 (10)
C11	0.8342 (4)	0.0464 (2)	0.0760 (3)	0.0788 (11)
H1	0.10360	-0.03595	0.45872	0.0954*
H3	0.40046	0.04203	0.33992	0.0639*
H5	0.59610	0.01991	0.21458	0.0640*
H7	0.86445	0.24181	0.10285	0.0764*
H8	0.73305	0.35299	0.21244	0.0838*
H9	0.52556	0.30031	0.31872	0.0774*
H10A	0.15463	0.27413	0.44482	0.0988*
H10B	0.20537	0.29296	0.31123	0.0988*
H10C	0.36802	0.29433	0.45849	0.0988*
H11A	0.79146	-0.01992	0.08522	0.1182*
H11B	0.97157	0.04960	0.11877	0.1182*
H11C	0.79701	0.06301	-0.02083	0.1182*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.1052 (15)	0.0487 (10)	0.1240 (17)	-0.0062 (9)	0.0893 (13)	-0.0004 (10)
O2	0.0786 (12)	0.0602 (11)	0.0958 (13)	-0.0042 (8)	0.0617 (11)	-0.0100 (9)
C1	0.0537 (13)	0.0479 (13)	0.0629 (15)	0.0003 (10)	0.0330 (12)	-0.0028 (11)
C2	0.0460 (12)	0.0467 (12)	0.0570 (14)	-0.0032 (10)	0.0239 (11)	0.0020 (11)
C3	0.0583 (14)	0.0481 (12)	0.0634 (15)	-0.0046 (11)	0.0339 (12)	0.0003 (12)
C4	0.0510 (13)	0.0508 (12)	0.0549 (14)	-0.0052 (10)	0.0282 (12)	0.0009 (11)
C5	0.0525 (14)	0.0513 (13)	0.0611 (15)	-0.0035 (10)	0.0261 (12)	0.0021 (11)
C6	0.0490 (13)	0.0649 (16)	0.0589 (15)	-0.0034 (11)	0.0278 (12)	0.0047 (12)
C7	0.0631 (15)	0.0742 (17)	0.0646 (16)	-0.0139 (13)	0.0358 (14)	0.0053 (13)
C8	0.0812 (19)	0.0581 (15)	0.0849 (19)	-0.0194 (13)	0.0475 (17)	-0.0021 (13)
C9	0.0767 (17)	0.0525 (14)	0.0804 (18)	-0.0098 (12)	0.0475 (15)	-0.0045 (13)
C10	0.0574 (15)	0.0538 (15)	0.097 (2)	-0.0010 (11)	0.0408 (15)	-0.0017 (14)
C11	0.0757 (18)	0.088 (2)	0.091 (2)	0.0014 (15)	0.0521 (17)	-0.0051 (16)

Geometric parameters (Å, °)

O1—C1	1.271 (3)	C8—C9	1.373 (4)
O2—C1	1.250 (3)	C3—H3	0.9300
O1—H1	0.8200	C5—H5	0.9300
C1—C2	1.477 (3)	C7—H7	0.9300
C2—C10	1.500 (3)	C8—H8	0.9300
C2—C3	1.339 (3)	C9—H9	0.9300
C3—C4	1.462 (3)	C10—H10A	0.9600
C4—C9	1.391 (3)	C10—H10B	0.9600
C4—C5	1.391 (3)	C10—H10C	0.9600
C5—C6	1.389 (3)	C11—H11A	0.9600
C6—C7	1.371 (4)	C11—H11B	0.9600
C6—C11	1.498 (4)	C11—H11C	0.9600
C7—C8	1.370 (4)		

C1—O1—H1	109.00	C4—C5—H5	119.00
O1—C1—O2	122.0 (2)	C6—C5—H5	119.00
O1—C1—C2	118.3 (2)	C6—C7—H7	119.00
O2—C1—C2	119.7 (2)	C8—C7—H7	119.00
C1—C2—C10	116.4 (2)	C7—C8—H8	120.00
C3—C2—C10	126.3 (2)	C9—C8—H8	120.00
C1—C2—C3	117.2 (2)	C4—C9—H9	120.00
C2—C3—C4	132.0 (2)	C8—C9—H9	120.00
C3—C4—C9	125.5 (2)	C2—C10—H10A	109.00
C5—C4—C9	117.3 (2)	C2—C10—H10B	109.00
C3—C4—C5	117.3 (2)	C2—C10—H10C	109.00
C4—C5—C6	122.9 (2)	H10A—C10—H10B	109.00
C5—C6—C11	121.8 (2)	H10A—C10—H10C	109.00
C7—C6—C11	120.8 (2)	H10B—C10—H10C	109.00
C5—C6—C7	117.5 (2)	C6—C11—H11A	109.00
C6—C7—C8	121.2 (3)	C6—C11—H11B	109.00
C7—C8—C9	120.7 (2)	C6—C11—H11C	110.00
C4—C9—C8	120.4 (2)	H11A—C11—H11B	109.00
C2—C3—H3	114.00	H11A—C11—H11C	109.00
C4—C3—H3	114.00	H11B—C11—H11C	109.00
O1—C1—C2—C3	-10.6 (3)	C9—C4—C5—C6	-2.0 (3)
O1—C1—C2—C10	169.3 (2)	C3—C4—C9—C8	-178.4 (2)
O2—C1—C2—C3	170.7 (2)	C5—C4—C9—C8	1.2 (4)
O2—C1—C2—C10	-9.4 (3)	C4—C5—C6—C7	1.3 (3)
C1—C2—C3—C4	179.6 (2)	C4—C5—C6—C11	-178.9 (2)
C10—C2—C3—C4	-0.3 (4)	C5—C6—C7—C8	0.4 (4)
C2—C3—C4—C5	-171.3 (2)	C11—C6—C7—C8	-179.5 (3)
C2—C3—C4—C9	8.3 (4)	C6—C7—C8—C9	-1.2 (4)
C3—C4—C5—C6	177.54 (19)	C7—C8—C9—C4	0.4 (4)

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>
O1—H1...O2 ⁱ	0.82	1.83	2.611 (3)	160
C3—H3...O1	0.93	2.27	2.703 (3)	108
C8—H8...O1 ⁱⁱ	0.93	2.59	3.394 (3)	145
C10—H10A...O2	0.96	2.31	2.783 (3)	109
C10—H10C...Cg ⁱⁱⁱ	0.9600	2.75	3.610 (3)	149.00
C1—O2...Cg ^{iv}	1.25 (1)	3.57 (1)	3.895 (3)	95 (1)

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