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2,3-Dibromo-3-phenylpropionic acid

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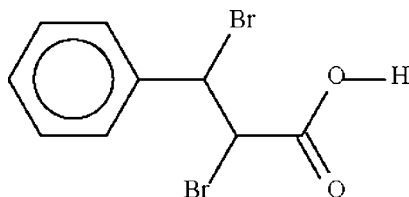
Received 10 September 2008; accepted 11 September 2008

Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.005$ Å; disorder in main residue; R factor = 0.025; wR factor = 0.067; data-to-parameter ratio = 18.1.

In the crystal of the title compound, $\text{C}_9\text{H}_8\text{Br}_2\text{O}_2$, inversion dimers linked by two $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds occur. All of the carbon and oxygen atoms are disordered over two sets of sites in a 2:1 ratio.

Related literature

For *threo*-1,2-diphenyl-2,3-difluoropropionate, see: O'Hagan *et al.* (2006). For *R*-methyl 3-bromo-2-chloro-3-phenylpropionate, see: Shaw *et al.* (1995).



Experimental

Crystal data

 $\text{C}_9\text{H}_8\text{Br}_2\text{O}_2$ $M_r = 307.97$ Orthorhombic, *Pbca* $a = 7.0278$ (1) Å $b = 9.7105$ (1) Å $c = 29.2970$ (4) Å $V = 1999.33$ (4) Å³ $Z = 8$ Mo $K\alpha$ radiation $\mu = 8.07$ mm⁻¹ $T = 100$ (2) K

0.28 × 0.22 × 0.14 mm

Data collection

Bruker SMART APEX
diffractometer
Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.211$, $T_{\max} = 0.398$
(expected range = 0.171–0.323)

17420 measured reflections
2303 independent reflections
2056 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.027$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.025$ $wR(F^2) = 0.066$ $S = 1.06$

2303 reflections

127 parameters

83 restraints

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.61$ e Å⁻³ $\Delta\rho_{\text{min}} = -0.62$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{O1}-\text{H1o}\cdots\text{O2}^i$	0.84	1.86	2.68 (1)	165
$\text{O1}'-\text{H1'o}\cdots\text{O2}^i$	0.84	1.86	2.69 (1)	166

Symmetry code: (i) $-x + 1, -y + 1, -z + 1$.

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *publCIF* (Westrip, 2008).

We thank the University of Malaya for funding this study (FR155/2007 A) and also for the purchase of the diffractometer.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: TK2304).

References

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supporting information

Acta Cryst. (2008). E64, o1946 [doi:10.1107/S160053680802919X]

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S1. Comment

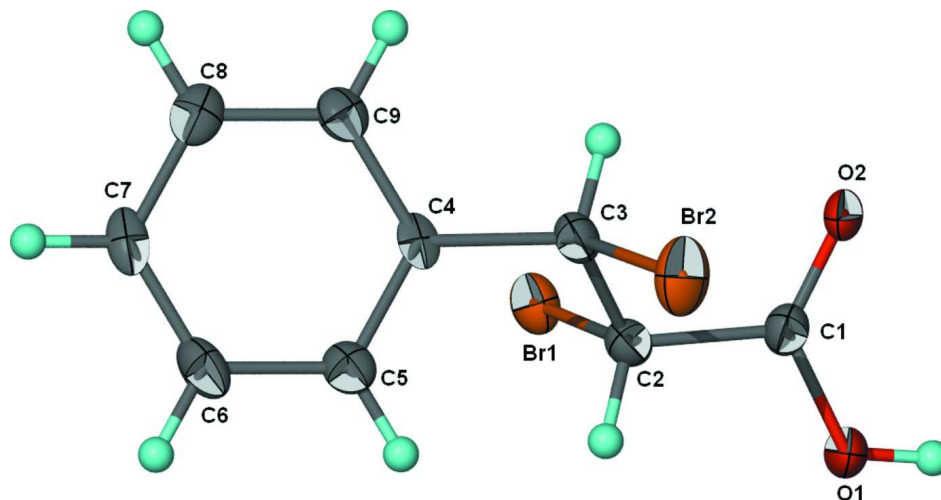
The compound (Scheme I, Fig. 1) was obtained as a side-product from the reaction of cyclopentylidiphenyltin cinnamate and 4-dimethylaminopyridine hydrobromide perbromide. Possibly, bromine added across the double bond of the cinnamate group followed by cleavage of the tin–carbon bond. Only few dihalogenpropionic acid derivatives have been characterized by X-ray crystallography. These are limited to, for example, *threo*-methyl 2,3-difluoropropionate (O'Hagan *et al.*, 2006) and *R*-methyl 3-bromo-2-chloro-3-phenylpropionate (Shaw *et al.*, 1995).

S2. Experimental

The compound was obtained as a side-product from the reaction of cyclopentylidiphenyltin cinnamate (0.3 g, 0.6 mmol) and 4-dimethylaminopyridine hydrobromide perbromide (0.25 g) in a mixture of chloroform and ethanol.

S3. Refinement

The structure is disordered over two positions with respect of the non-bromide atoms. The Br1 atom is connected to the carbon atom in the 2-position in the major component but is connected to the carbon atom in the 3-position in the minor component. Conversely, the Br2 atom is connected to the carbon atom in the 3-position in the major component but is connected to the carbon atom in the 2-position in the minor component. All distances in the major component were restrained to within 0.01 Å of their equivalents in the minor component. The phenyl rings were restrained into rigid hexagons of 1.39 Å sides. Additionally, the four-atom carboxyl and seven-atom benzyl units were each restrained to be nearly flat. The anisotropic displacement parameters of the primed atoms were restrained to be equal to those of the unprimed atoms; these were also restrained to be nearly isotropic. In the initial stages of the refinement, the occupancy refined to an approximate 2:1 ratio. However, with the inclusion of hydrogen atoms, the refinement was unstable. Ratios that were either slightly smaller or slightly larger than 2:1 did not yield any significant differences in the final residual index. The ratio was then fixed to 2:1. Oxygen and carbon-bound H-atoms were placed in calculated positions (C—H 0.95 to 1.00 Å, O—H 0.84 Å) and were included in the riding model approximation, with $U(\text{H})$ set to 1.2 to $1.5U_{\text{eq}}(\text{C}, \text{O})$.

**Figure 1**

70% Probability thermal ellipsoid plot (Barbour, 2001) of $C_9H_8Br_2O_2$. Hydrogen atoms are drawn as spheres of arbitrary radii. For clarity, the minor component is not shown.

2,3-Dibromo-3-phenylpropionic acid

Crystal data

$C_9H_8Br_2O_2$

$M_r = 307.97$

Orthorhombic, *Pbca*

Hall symbol: -P 2ac 2ab

$a = 7.0278$ (1) Å

$b = 9.7105$ (1) Å

$c = 29.2970$ (4) Å

$V = 1999.33$ (4) Å³

$Z = 8$

$F(000) = 1184$

$D_x = 2.046$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 8127 reflections

$\theta = 2.2$ – 28.2°

$\mu = 8.07$ mm⁻¹

$T = 100$ K

Block, colorless

$0.28 \times 0.22 \times 0.14$ mm

Data collection

Bruker SMART APEX
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 1996)

$T_{\min} = 0.211$, $T_{\max} = 0.398$

17420 measured reflections

2303 independent reflections

2056 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.027$

$\theta_{\max} = 27.5^\circ$, $\theta_{\min} = 2.8^\circ$

$h = -9 \rightarrow 9$

$k = -12 \rightarrow 12$

$l = -37 \rightarrow 38$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.025$

$wR(F^2) = 0.066$

$S = 1.06$

2303 reflections

127 parameters

83 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

$w = 1/[\sigma^2(F_o^2) + (0.0337P)^2 + 3.2334P]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.61$ e Å⁻³

$\Delta\rho_{\min} = -0.62$ e Å⁻³

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
Br1	0.48392 (4)	0.31033 (3)	0.364350 (9)	0.02208 (9)	
Br2	-0.06355 (4)	0.45596 (3)	0.436758 (9)	0.03094 (10)	
O1	0.4216 (6)	0.3279 (4)	0.48256 (13)	0.0308 (10)	0.67
H1O	0.4765	0.3569	0.5061	0.046*	0.67
O2	0.4214 (10)	0.5338 (5)	0.4476 (3)	0.0209 (10)	0.67
C1	0.3847 (4)	0.4118 (4)	0.44922 (12)	0.0189 (8)	0.67
C2	0.2859 (5)	0.3388 (4)	0.40959 (12)	0.0176 (7)	0.67
H2	0.2344	0.2480	0.4200	0.021*	0.67
C3	0.1313 (5)	0.4223 (4)	0.38839 (12)	0.0183 (7)	0.67
H3	0.1859	0.5130	0.3790	0.022*	0.67
C4	0.0358 (4)	0.3575 (5)	0.34721 (14)	0.0133 (8)	0.67
C5	-0.0307 (8)	0.2228 (5)	0.3486 (3)	0.0186 (11)	0.67
H5	-0.0167	0.1697	0.3756	0.022*	0.67
C6	-0.1176 (9)	0.1657 (9)	0.3104 (4)	0.0224 (6)	0.67
H6	-0.1631	0.0736	0.3114	0.027*	0.67
C7	-0.1381 (9)	0.2433 (15)	0.2709 (3)	0.0181 (8)	0.67
H7	-0.1976	0.2043	0.2448	0.022*	0.67
C8	-0.0716 (13)	0.3781 (14)	0.26947 (16)	0.0206 (7)	0.67
H8	-0.0856	0.4311	0.2424	0.025*	0.67
C9	0.0154 (11)	0.4352 (8)	0.3076 (2)	0.0183 (9)	0.67
H9	0.0608	0.5273	0.3067	0.022*	0.67
O1'	0.3463 (14)	0.3474 (10)	0.4912 (3)	0.0308 (10)	0.33
H1'O	0.4202	0.3715	0.5122	0.046*	0.33
O2'	0.380 (2)	0.5445 (11)	0.4542 (7)	0.0209 (10)	0.33
C1'	0.3118 (9)	0.4305 (9)	0.4576 (2)	0.0189 (8)	0.33
C2'	0.1737 (9)	0.3684 (7)	0.4228 (2)	0.0176 (7)	0.33
H2'	0.1633	0.2666	0.4274	0.021*	0.33
C3'	0.2239 (9)	0.3993 (8)	0.3748 (2)	0.0183 (7)	0.33
H3'	0.2355	0.5012	0.3709	0.022*	0.33
C4'	0.0862 (9)	0.3432 (12)	0.3394 (3)	0.0133 (8)	0.33
C5'	0.0090 (19)	0.2117 (12)	0.3425 (6)	0.0186 (11)	0.33
H5'	0.0421	0.1536	0.3674	0.022*	0.33
C6'	-0.1164 (19)	0.165 (2)	0.3092 (8)	0.0224 (6)	0.33
H6'	-0.1692	0.0754	0.3113	0.027*	0.33
C7'	-0.1648 (18)	0.250 (3)	0.2729 (6)	0.0181 (8)	0.33
H7'	-0.2505	0.2185	0.2501	0.022*	0.33
C8'	-0.088 (3)	0.382 (3)	0.2698 (4)	0.0206 (7)	0.33
H8'	-0.1207	0.4398	0.2449	0.025*	0.33
C9'	0.038 (2)	0.4282 (17)	0.3031 (5)	0.0183 (9)	0.33
H9'	0.0906	0.5181	0.3010	0.022*	0.33

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.02081 (14)	0.02271 (15)	0.02272 (15)	-0.00251 (10)	-0.00187 (10)	-0.00479 (10)

Br2	0.03118 (17)	0.04015 (19)	0.02150 (15)	0.00134 (13)	0.00331 (11)	-0.00921 (12)
O1	0.055 (3)	0.0188 (16)	0.0182 (18)	-0.0063 (19)	-0.0143 (18)	0.0006 (13)
O2	0.034 (3)	0.0169 (12)	0.012 (3)	0.0006 (16)	-0.0033 (18)	-0.0032 (12)
C1	0.023 (2)	0.0199 (18)	0.0139 (17)	-0.0030 (19)	-0.0010 (16)	-0.0003 (14)
C2	0.0211 (17)	0.0149 (15)	0.0166 (16)	-0.0017 (14)	0.0007 (13)	-0.0010 (12)
C3	0.021 (2)	0.0159 (16)	0.0175 (17)	-0.0033 (15)	0.0002 (14)	-0.0028 (13)
C4	0.007 (2)	0.0153 (17)	0.017 (2)	0.0062 (19)	0.0015 (16)	-0.0077 (14)
C5	0.016 (3)	0.0174 (16)	0.023 (3)	0.0023 (17)	-0.002 (2)	-0.0009 (15)
C6	0.0180 (12)	0.0183 (13)	0.0310 (15)	-0.0018 (10)	-0.0052 (11)	-0.0067 (11)
C7	0.009 (2)	0.0239 (19)	0.0215 (14)	0.006 (2)	-0.0035 (16)	-0.0106 (11)
C8	0.0169 (19)	0.0259 (15)	0.0190 (12)	0.0082 (15)	0.0007 (11)	-0.0025 (10)
C9	0.017 (2)	0.0167 (14)	0.0211 (18)	0.0027 (13)	0.0015 (14)	-0.0037 (14)
O1'	0.055 (3)	0.0188 (16)	0.0182 (18)	-0.0063 (19)	-0.0143 (18)	0.0006 (13)
O2'	0.034 (3)	0.0169 (12)	0.012 (3)	0.0006 (16)	-0.0033 (18)	-0.0032 (12)
C1'	0.023 (2)	0.0199 (18)	0.0139 (17)	-0.0030 (19)	-0.0010 (16)	-0.0003 (14)
C2'	0.0211 (17)	0.0149 (15)	0.0166 (16)	-0.0017 (14)	0.0007 (13)	-0.0010 (12)
C3'	0.021 (2)	0.0159 (16)	0.0175 (17)	-0.0033 (15)	0.0002 (14)	-0.0028 (13)
C4'	0.007 (2)	0.0153 (17)	0.017 (2)	0.0062 (19)	0.0015 (16)	-0.0077 (14)
C5'	0.016 (3)	0.0174 (16)	0.023 (3)	0.0023 (17)	-0.002 (2)	-0.0009 (15)
C6'	0.0180 (12)	0.0183 (13)	0.0310 (15)	-0.0018 (10)	-0.0052 (11)	-0.0067 (11)
C7'	0.009 (2)	0.0239 (19)	0.0215 (14)	0.006 (2)	-0.0035 (16)	-0.0106 (11)
C8'	0.0169 (19)	0.0259 (15)	0.0190 (12)	0.0082 (15)	0.0007 (11)	-0.0025 (10)
C9'	0.017 (2)	0.0167 (14)	0.0211 (18)	0.0027 (13)	0.0015 (14)	-0.0037 (14)

Geometric parameters (Å, °)

Br1—C2	1.942 (4)	C8—H8	0.9500
Br1—C3'	2.044 (6)	C9—H9	0.9500
Br2—C2'	1.916 (6)	O1'—C1'	1.295 (8)
Br2—C3	1.997 (3)	O1'—H1'O	0.8400
O1—C1	1.298 (4)	O2'—C1'	1.212 (8)
O1—H1O	0.8400	C1'—C2'	1.533 (8)
O2—C1	1.213 (5)	C2'—C3'	1.480 (7)
C1—C2	1.527 (5)	C2'—H2'	1.0000
C2—C3	1.491 (5)	C3'—C4'	1.518 (7)
C2—H2	1.0000	C3'—H3'	1.0000
C3—C4	1.517 (4)	C4'—C5'	1.3900
C3—H3	1.0000	C4'—C9'	1.3900
C4—C5	1.3900	C5'—C6'	1.3900
C4—C9	1.3900	C5'—H5'	0.9500
C5—C6	1.3900	C6'—C7'	1.3900
C5—H5	0.9500	C6'—H6'	0.9500
C6—C7	1.3900	C7'—C8'	1.3900
C6—H6	0.9500	C7'—H7'	0.9500
C7—C8	1.3900	C8'—C9'	1.3900
C7—H7	0.9500	C8'—H8'	0.9500
C8—C9	1.3900	C9'—H9'	0.9500

C1—O1—H1O	120.0	C1'—O1'—H1'O	120.0
O2—C1—O1	126.8 (6)	O2'—C1'—O1'	124.0 (13)
O2—C1—C2	121.4 (6)	O2'—C1'—C2'	123.8 (13)
O1—C1—C2	111.8 (4)	O1'—C1'—C2'	112.3 (8)
C3—C2—C1	113.3 (3)	C3'—C2'—C1'	113.7 (6)
C3—C2—Br1	108.4 (2)	C3'—C2'—Br2	108.7 (4)
C1—C2—Br1	105.0 (2)	C1'—C2'—Br2	103.5 (4)
C3—C2—H2	110.0	C3'—C2'—H2'	110.2
C1—C2—H2	110.0	C1'—C2'—H2'	110.2
Br1—C2—H2	110.0	Br2—C2'—H2'	110.2
C2—C3—C4	115.4 (3)	C2'—C3'—C4'	115.0 (6)
C2—C3—Br2	107.0 (2)	C2'—C3'—Br1	105.6 (4)
C4—C3—Br2	109.20 (19)	C4'—C3'—Br1	108.5 (4)
C2—C3—H3	108.3	C2'—C3'—H3'	109.2
C4—C3—H3	108.3	C4'—C3'—H3'	109.2
Br2—C3—H3	108.3	Br1—C3'—H3'	109.2
C5—C4—C9	120.0	C5'—C4'—C9'	120.0
C5—C4—C3	121.0 (5)	C5'—C4'—C3'	122.3 (11)
C9—C4—C3	119.0 (5)	C9'—C4'—C3'	117.7 (11)
C4—C5—C6	120.0	C4'—C5'—C6'	120.0
C4—C5—H5	120.0	C4'—C5'—H5'	120.0
C6—C5—H5	120.0	C6'—C5'—H5'	120.0
C7—C6—C5	120.0	C7'—C6'—C5'	120.0
C7—C6—H6	120.0	C7'—C6'—H6'	120.0
C5—C6—H6	120.0	C5'—C6'—H6'	120.0
C6—C7—C8	120.0	C6'—C7'—C8'	120.0
C6—C7—H7	120.0	C6'—C7'—H7'	120.0
C8—C7—H7	120.0	C8'—C7'—H7'	120.0
C7—C8—C9	120.0	C7'—C8'—C9'	120.0
C7—C8—H8	120.0	C7'—C8'—H8'	120.0
C9—C8—H8	120.0	C9'—C8'—H8'	120.0
C8—C9—C4	120.0	C8'—C9'—C4'	120.0
C8—C9—H9	120.0	C8'—C9'—H9'	120.0
C4—C9—H9	120.0	C4'—C9'—H9'	120.0
O2—C1—C2—C3	-40.3 (4)	O2'—C1'—C2'—Br2	-77.8 (5)
O1—C1—C2—C3	140.0 (3)	O1'—C1'—C2'—Br2	102.1 (5)
O2—C1—C2—Br1	77.8 (3)	C3—Br2—C2'—C3'	0.9 (4)
O1—C1—C2—Br1	-101.9 (2)	C3—Br2—C2'—C1'	122.1 (7)
C1—C2—C3—C4	176.8 (3)	C1'—C2'—C3'—C4'	-178.0 (6)
Br1—C2—C3—C4	60.7 (4)	Br2—C2'—C3'—C4'	-63.3 (7)
C1—C2—C3—Br2	-61.5 (3)	C1'—C2'—C3'—Br1	62.4 (6)
Br1—C2—C3—Br2	-177.59 (16)	Br2—C2'—C3'—Br1	177.1 (3)
C2—C3—C4—C5	49.8 (4)	C2—Br1—C3'—C2'	0.1 (3)
Br2—C3—C4—C5	-70.8 (3)	C2—Br1—C3'—C4'	-123.7 (8)
C2—C3—C4—C9	-130.2 (4)	C2'—C3'—C4'—C5'	-42.3 (8)
Br2—C3—C4—C9	109.2 (3)	Br1—C3'—C4'—C5'	75.6 (6)
C9—C4—C5—C6	0.0	C2'—C3'—C4'—C9'	137.7 (8)

C3—C4—C5—C6	-179.99 (8)	Br1—C3'—C4'—C9'	-104.3 (6)
C4—C5—C6—C7	0.0	C9'—C4'—C5'—C6'	0.0
C5—C6—C7—C8	0.0	C3'—C4'—C5'—C6'	-179.96 (9)
C6—C7—C8—C9	0.0	C4'—C5'—C6'—C7'	0.0
C7—C8—C9—C4	0.0	C5'—C6'—C7'—C8'	0.0
C5—C4—C9—C8	0.0	C6'—C7'—C8'—C9'	0.0
C3—C4—C9—C8	179.99 (8)	C7'—C8'—C9'—C4'	0.0
O2'—C1'—C2'—C3'	40.0 (6)	C5'—C4'—C9'—C8'	0.0
O1'—C1'—C2'—C3'	-140.1 (6)	C3'—C4'—C9'—C8'	179.96 (8)

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>
O1—H1 _o \cdots O2 ⁱ	0.84	1.86	2.68 (1)	165
O1'—H1' _o \cdots O2 ⁱ	0.84	1.86	2.69 (1)	166

Symmetry code: (i) $-x+1, -y+1, -z+1$.