

## 2-Methoxy-2-phenylacetic acid

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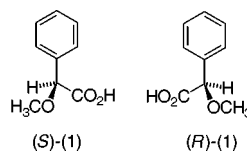
## Key indicators

Single-crystal X-ray study  
 $T = 299$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å  
 $R$  factor = 0.055  
 $wR$  factor = 0.174  
Data-to-parameter ratio = 13.0For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.

2-Methoxy-2-phenylacetic acid,  $\text{C}_9\text{H}_{10}\text{O}_3$ , forms helical columns of single enantiomers linked by hydrogen bonding between the acidic proton of one molecule and the methoxy O atom of a neighbouring molecule, to give an overall racemic structure.

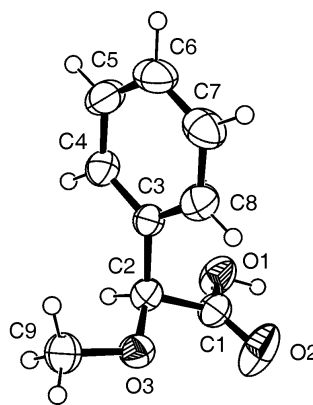
## Comment

2-Methoxy-2-phenylacetic acid, (1), is a chiral compound with one stereogenic center. The racemate of (1) has been crystallized. It forms helical columns of single enantiomers linked by hydrogen bonding between the acidic proton of one molecule and the methoxy O atom of a neighbouring molecule in the solid state [ $\text{O1}\cdots\text{O3}^i = 2.725(2)$  Å,  $\text{O1}-\text{H1O}\cdots\text{O3}^i = 162(3)$  Å; symmetry code as in Table 1] (Figs. 1 and 2). Atom H1O is located in the plane defined by atoms O1, O2<sup>i</sup>, and O3<sup>i</sup> [the deviation is only  $0.05(3)$  Å]. This arrangement leads to a packing of molecules of (1) (Fig. 3) similar to the structure that has been reported for its acidic sodium acid salt (Moore *et al.*, 1980).

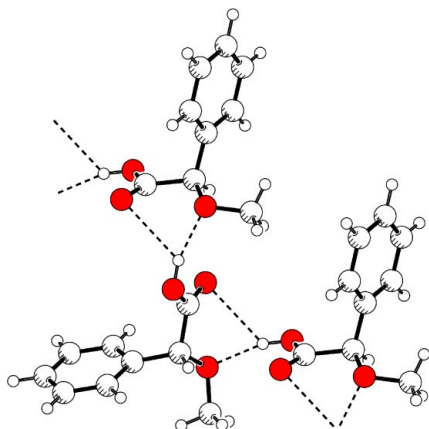


## Experimental

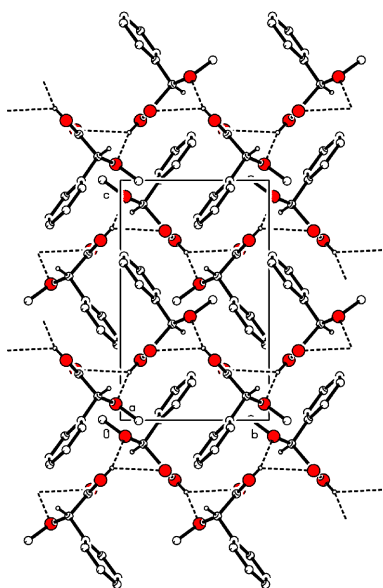
The racemate of 2-methoxy-2-phenylacetic acid, (1), is commercially available. The crystal analyzed in this study was grown from a solution of *rac*-2-(2-methoxy-2-phenylacetylsulfanyl)pyridine (Gottwald *et al.*, 2004) in  $\text{CH}_2\text{Cl}_2/n$ -hexane (3:1 *v/v*), which was kept in the dark for 7 d at 293 K.

**Figure 1**

The molecular structure of (*S*)-(1) which was arbitrarily selected from the racemate that is present in the unit cell of (1). Displacement ellipsoids are drawn at the 50% probability level.



**Figure 2**  
Hydrogen bonding of (1) in the solid state.



**Figure 3**  
(*P*)-helical packing of (*S*)-(1) and (*M*)-helical arrangement of (*R*)-(1) in the unit cell. View along [100].

#### Crystal data

$C_9H_{10}O_3$   
 $M_r = 166.17$   
 Monoclinic,  $P2_1/n$   
 $a = 10.093$  (3) Å  
 $b = 7.123$  (2) Å  
 $c = 12.270$  (3) Å  
 $\beta = 110.40$  (1)°  
 $V = 826.8$  (4) Å<sup>3</sup>  
 $Z = 4$

$D_x = 1.335$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 Cell parameters from 877  
 reflections  
 $\theta = 3.3$ – $20.9^\circ$   
 $\mu = 0.10$  mm<sup>-1</sup>  
 $T = 299$  (2) K  
 Prism, colorless  
 $0.60 \times 0.48 \times 0.16$  mm

#### Data collection

Oxford Diffraction Xcalibur CCD  
 diffractometer  
 $\omega$  rotation scans  
 Absorption correction: none  
 4076 measured reflections  
 1665 independent reflections

994 reflections with  $I > 2\sigma(I)$   
 $R_{int} = 0.035$   
 $\theta_{max} = 26.4^\circ$   
 $h = -12 \rightarrow 12$   
 $k = -8 \rightarrow 8$   
 $l = -15 \rightarrow 8$

#### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.055$   
 $wR(F^2) = 0.174$   
 $S = 0.88$   
 1665 reflections  
 128 parameters  
 H atoms treated by a mixture of  
 independent and constrained  
 refinement

$w = 1/[\sigma^2(F_o^2) + (0.1149P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{max} = 0.045$   
 $\Delta\rho_{max} = 0.25$  e Å<sup>-3</sup>  
 $\Delta\rho_{min} = -0.23$  e Å<sup>-3</sup>  
 Extinction correction: *SHELXL97*  
 and Larson (1970)  
 Extinction coefficient: 0.051 (11)

**Table 1**

Selected geometric parameters (Å, °).

C1—O2	1.187 (3)	C2—C3	1.502 (3)
C1—O1	1.306 (3)	C9—O3	1.422 (3)
C1—C2	1.512 (3)	O1—H1O	0.90 (3)
C2—O3	1.408 (3)		
O2—C1—O1	124.6 (2)	O3—C2—C1	106.16 (18)
O2—C1—C2	123.8 (2)	C1—O1—H1O	104.9 (18)
O1—C1—C2	111.6 (2)	C2—O3—C9	112.89 (17)
O2—C1—C2—O3	-16.5 (3)	O1—C1—C2—C3	-74.3 (3)
O1—C1—C2—O3	163.59 (19)	C3—C2—O3—C9	70.6 (2)
O2—C1—C2—C3	105.6 (3)	C1—C2—O3—C9	-168.23 (18)

**Table 2**

Hydrogen-bonding 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—H1O...O3 <sup>i</sup>	0.90 (3)	1.85 (3)	2.725 (2)	162 (3)
O1—H1O...O2 <sup>i</sup>	0.90 (3)	2.57 (3)	3.190 (3)	127 (2)

Symmetry code: (i)  $\frac{3}{2} - x, \frac{1}{2} + y, \frac{3}{2} - z$ .

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2001); cell refinement: *CrysAlis RED* (Oxford Diffraction, 2001); data reduction: *CrysAlis RED*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *PLATON2002* (Spek, 2002); software used to prepare material for publication: *SHELXL97*.

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