

C1	0.2729 (3)	0.4366 (6)	0.5650 (2)	0.0506 (8)
C2	0.3708 (3)	0.6132 (7)	0.5642 (2)	0.0577 (9)
C3	0.4664 (3)	0.6578 (7)	0.6318 (2)	0.0574 (9)
C4	0.4651 (3)	0.5107 (7)	0.7043 (2)	0.0574 (9)
C5	0.3671 (4)	0.3358 (7)	0.7083 (2)	0.0644 (10)
C6	0.2726 (4)	0.2998 (7)	0.6398 (2)	0.0611 (10)
C7	0.1738 (3)	0.3958 (7)	0.4898 (2)	0.0539 (9)
C8	0.1638 (4)	0.5693 (8)	0.4155 (2)	0.0709 (12)
C9	0.5682 (4)	0.8561 (9)	0.6283 (3)	0.0799 (13)

Table 4. Selected geometric parameters ( $\text{\AA}$ ,  $^\circ$ ) for (II)

O1—N1	1.431 (3)	C2—C3	1.382 (5)
N1—C7	1.279 (5)	C3—C4	1.397 (5)
O2—C4	1.362 (4)	C3—C9	1.501 (5)
C1—C2	1.386 (5)	C4—C5	1.386 (5)
C1—C6	1.396 (5)	C5—C6	1.379 (5)
C1—C7	1.486 (4)	C7—C8	1.497 (5)
C7—N1—O1	113.8 (3)	O2—C4—C5	122.5 (3)
C2—C1—C6	116.6 (3)	O2—C4—C3	117.3 (3)
C2—C1—C7	121.4 (3)	C5—C4—C3	120.2 (3)
C6—C1—C7	122.0 (3)	C6—C5—C4	120.4 (3)
C3—C2—C1	124.0 (3)	C5—C6—C1	121.3 (3)
C2—C3—C4	117.5 (3)	N1—C7—C1	116.1 (3)
C2—C3—C9	121.8 (3)	N1—C7—C8	123.0 (3)
C4—C3—C9	120.7 (3)	C1—C7—C8	120.8 (3)

Table 5. Comparison of hydrogen-bonding geometries ( $\text{\AA}$ ,  $^\circ$ ) of the two title structures

$D-H \cdots A$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
Compound (I)			
O1—HO1 $\cdots$ O2 <sup>i</sup>	1.82 (3)	2.734 (2)	174 (2)
O2—HO2 $\cdots$ N1 <sup>ii</sup>	1.74 (3)	2.742 (2)	176 (2)
Compound (II)			
O1—HO1 $\cdots$ N1 <sup>iii</sup>	1.75 (4)	2.774 (4)	165 (3)
O2—HO2 $\cdots$ O1 <sup>iv</sup>	1.99 (5)	2.825 (4)	165 (4)

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

All the H atoms in structure (I), including those of the disordered methyl group, were located from the  $\Delta\rho$  map. In structure (II), oxime and hydroxyl H atoms were found in a  $\Delta\rho$  map, while all other H atoms were included in their calculated positions. In the final cycle of refinement of (I), refinement was carried out on the positions of all atoms, with anisotropic displacement parameters for non-H atoms and isotropic displacement coefficients for ordered H atoms. For the disordered H atoms the relative site occupancy [0.60 (2) for atoms *a*, *b* and *c* attached to C8] and a common isotropic displacement parameter were refined. In structure (II), the non-H atoms were refined with anisotropic displacement parameters. Oxime and hydroxyl H-atom positions, the individual isotropic displacement parameters for the benzene-type H atoms, common isotropic displacement parameters and relative site occupancies for disordered methyl H atoms [0.54 (4) for atoms *a*, *b* and *c* attached to C8 and C9] were refined.

There was some trouble with the data collection for (II), caused by the quality of crystals obtained. This, in turn, resulted in a paucity of data and the high values of  $R_{\text{int}}$  and  $R$  but there were no problems in solving and refining the structure.

*SHELXS86* (Sheldrick, 1990) was used for structure solution; refinement was performed using *SHELXL* (Sheldrick, 1994). Molecular graphics were prepared using *ORTEP* (Johnson, 1976). The material for publication was produced using *SHELXL93*.

Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCr (Reference: AB1131). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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## 2'-Hydroxypropiophenone Oxime

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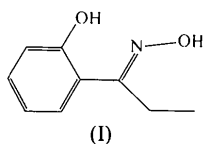
## Abstract

The title structure,  $C_9H_{11}NO_2$ , consists of antiparallel polar chains of molecules, bonded together by a series of intra- and intermolecular hydrogen bonds between oxime and hydroxyl groups. None of the usual oxime–oxime hydrogen bonds has been observed.

## Comment

As a continuation of studies of oxime molecules with a second functional group, possessing donor–acceptor capabilities (Maurin, Paul & Curtin, 1992, 1994; Maurin, Winnicka-Maurin & Leś, 1994), several hydroxy–oxime structures have been determined (Maurin, 1994; Maurin, Wieteska & Winnicka-Maurin, 1994). 2'-Hydroxypropiophenone oxime, (I), is the first hydroxy–oxime structure in which intra- and intermolecular hydrogen bonds are combined in a chain formation. The conformation of the molecules and the numbering of the atoms are given in Fig. 1. The hydrogen-

bonding scheme between two independent molecules is also shown.



The crystal structure is made up of antiparallel molecular chains. The chains are held together by a series of intra- and intermolecular hydrogen bonds, each chain having alternate *A* and *B* molecules. The hydrogen-bonding pattern in each chain is OH<sub>hydroxyl</sub>⋯N<sub>oxime</sub> (intramolecular), OH<sub>oxime</sub>⋯O<sub>hydroxyl</sub> (intermolecular), OH<sub>hydroxyl</sub>⋯N<sub>oxime</sub> (intramolecular), etc. The dimensions of the hydrogen-bond geometries are given in Table 3. Both oxime–hydroxyl hydrogen bonds are slightly longer than those observed in 4'-hydroxyacetophenone oxime [2.734 (2) Å; Maurin, 1994]. The intramolecular hydroxyl–oxime hydrogen bonds, however, are considerably shorter than the corresponding intermolecular hydrogen bonds [2.741 (2) Å]. The structure of 2'-hydroxypropiophenone oxime has more in common with oxime–carboxyl structures (Maurin, Paul & Curtin, 1992, 1994), where separated antiparallel polar chains have been observed, than with the previously reported oxime–hydroxyl structures (Maurin, Wieteska & Winnicka-Maurin, 1994; Maurin, 1994). There is a series of intermolecular oxime–hydroxyl hydrogen bonds constructing the infinite polar chains of molecules. The same role in oxime–carboxyl structures is played by

oxime–carboxyl and carboxyl–oxime pairs. On the other hand, there is a lack of those hydroxyl–oxime intermolecular hydrogen bonds (they are intramolecular in the present structure) that cause cross-chain interactions in some oxime–hydroxyl structures (Maurin, Winnicka-Maurin & Leś, 1994; Maurin, 1994).

In both molecules of the title compound, the C7 atom lies almost in the plane of the bonded phenyl ring, but the oxime group is twisted slightly around the C1–C7 bond. The dihedral angles between the best plane of the benzene ring and the plane of the oxime group (C1–C7–N1–O1) are 4.69 (18) and 11.55 (20)° for molecules *A* and *B*, respectively. The hydroxyl group of molecule *A* is almost coplanar with the phenyl ring, while that in molecule *B* is slightly out of the plane. In both molecules, the C1–C2 distance is longer than the other phenyl C–C distances, suggesting some repulsion of the *ortho*-substituents. However, the shortening of the C2–C3, C3–C4, C4–C5 and C5–C6 bonds is probably caused by thermal motion. No close intermolecular contacts other than the hydrogen bonds were observed.

## Experimental

The title compound was synthesized by the reaction of 2'-hydroxypropiophenone and hydroxylamine hydrochloride in alkaline solution. The recrystallization of the solid product from ethanol–water solution gave colourless prismatic crystals. A fragment cut from a larger crystal was used for data collection.

### Crystal data

C<sub>9</sub>H<sub>11</sub>NO<sub>2</sub>  
*M<sub>r</sub>* = 165.19  
 Monoclinic  
*P*2<sub>1</sub>/*n*  
*a* = 9.274 (2) Å  
*b* = 14.753 (2) Å  
*c* = 13.290 (2) Å  
 $\beta$  = 108.40 (2)°  
*V* = 1725.4 (5) Å<sup>3</sup>  
*Z* = 8  
*D<sub>x</sub>* = 1.272 Mg m<sup>-3</sup>

Mo *K*α radiation  
 $\lambda$  = 0.71073 Å  
 Cell parameters from 25 reflections  
 $\theta$  = 9.5–10.9°  
 $\mu$  = 0.090 mm<sup>-1</sup>  
*T* = 253 (2) K  
 Prismatic, transparent  
 0.60 × 0.45 × 0.40 mm  
 Colourless

### Data collection

CAD-4  $\kappa$ -axis four-circle diffractometer  
 Profile data from  $\omega/\theta$  scans  
 Absorption correction: none  
 4398 measured reflections  
 4152 independent reflections  
 2222 observed reflections [*I* > 2σ(*I*)]

*R*<sub>int</sub> = 0.013  
 $\theta_{\max}$  = 27.97°  
*h* = -12 → 11  
*k* = 0 → 19  
*l* = 0 → 17  
 3 standard reflections monitored every 200 reflections  
 intensity variation: none

### Refinement

Refinement on *F*<sup>2</sup>  
*R*[*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.0441  
*wR*(*F*<sup>2</sup>) = 0.1095

( $\Delta/\sigma$ )<sub>max</sub> < 0.001  
 $\Delta\rho_{\max}$  = 0.130 e Å<sup>-3</sup>  
 $\Delta\rho_{\min}$  = -0.188 e Å<sup>-3</sup>

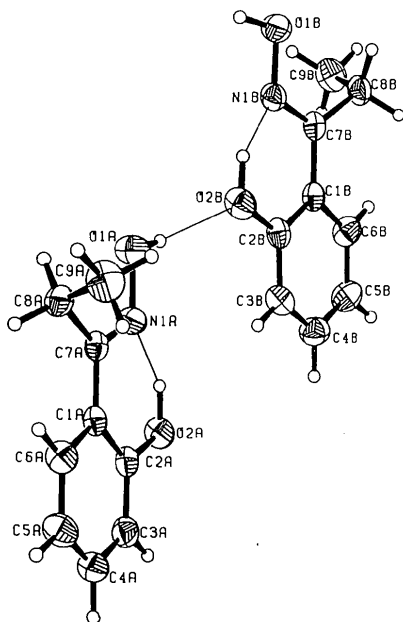


Fig. 1. Conformation of 2'-hydroxypropiophenone oxime molecules and the atomic numbering scheme. Hydrogen bonds are marked as thin lines.

S = 0.910  
4152 reflections  
305 parameters  
All H-atom parameters  
refined  
Calculated weights  
 $w = 1/[\sigma^2(F_o^2) + (0.0572P)^2]$   
where  $P = (F_o^2 + 2F_c^2)/3$

Extinction correction: none  
Atomic scattering factors  
from *International Tables  
for Crystallography* (1992),  
Vol. C, Tables 4.2.6.8 and  
6.1.1.4)

Table 3. Selected intra- and intermolecular hydrogen-bond distances (Å) and angles (°)

D	H	A	D...A	D—H...A
O2A	H2AO	N1A	2.534 (2)	146 (2)
O2B	H2BO	N1B	2.542 (2)	151 (2)
O1B	H1BO	O2A'	2.797 (2)	166 (2)
O1A	H1AO	O2B''	2.788 (2)	154 (2)

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

Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (Å<sup>2</sup>)

$$U_{eq} = (1/3)\sum_i \sum_j U_{ij} a_i^* a_j^* \mathbf{a}_i \cdot \mathbf{a}_j$$

	x	y	z	$U_{eq}$
C1A	-0.0716 (2)	0.55378 (9)	0.74258 (10)	0.0395 (3)
C2A	-0.1922 (2)	0.51035 (9)	0.66521 (11)	0.0424 (4)
C3A	-0.3378 (2)	0.51229 (11)	0.67073 (14)	0.0524 (4)
C4A	-0.3687 (2)	0.55541 (12)	0.75322 (15)	0.0579 (5)
C5A	-0.2542 (2)	0.59798 (13)	0.83043 (15)	0.0605 (5)
C6A	-0.1087 (2)	0.59697 (11)	0.82466 (13)	0.0522 (4)
C7A	0.0854 (2)	0.55290 (9)	0.73940 (10)	0.0407 (4)
C8A	0.2110 (2)	0.60172 (12)	0.82110 (13)	0.0507 (4)
C9A	0.2879 (3)	0.5422 (2)	0.9151 (2)	0.0776 (7)
N1A	0.10925 (15)	0.50514 (9)	0.66502 (9)	0.0465 (3)
O1A	0.26130 (13)	0.50491 (9)	0.66789 (10)	0.0610 (3)
O2A	-0.16895 (14)	0.46549 (8)	0.58157 (8)	0.0542 (3)
C1B	0.2082 (2)	0.74457 (9)	1.07936 (11)	0.0423 (4)
C2B	0.2675 (2)	0.79876 (10)	1.01492 (11)	0.0443 (4)
C3B	0.4197 (2)	0.79548 (12)	1.02358 (14)	0.0547 (4)
C4B	0.5176 (2)	0.74053 (13)	1.09804 (15)	0.0606 (5)
C5B	0.4639 (2)	0.68842 (13)	1.16442 (15)	0.0629 (5)
C6B	0.3118 (2)	0.69006 (12)	1.15399 (13)	0.0551 (4)
C7B	0.0454 (2)	0.74464 (9)	1.06989 (11)	0.0431 (4)
C8B	-0.0204 (2)	0.67514 (11)	1.12598 (14)	0.0519 (4)
C9B	-0.0252 (3)	0.7087 (2)	1.2325 (2)	0.0677 (6)
N1B	-0.03563 (15)	0.80753 (8)	1.01210 (10)	0.0473 (3)
O1B	-0.18876 (14)	0.80397 (8)	1.00698 (10)	0.0622 (4)
O2B	0.1770 (2)	0.85614 (8)	0.94125 (9)	0.0578 (3)

Data collection, cell refinement and data reduction: Enraf-Nonius software. Program(s) used to solve structure: *SHELXS86* (Sheldrick, 1990). Program(s) used to refine structure: *SHELXL* (Sheldrick, 1994). Molecular graphics: *ORTEPII* (Johnson, 1976). Software used to prepare material for publication: *SHELXL*.

Lists of structure factors, anisotropic displacement parameters, H-atom coordinates, complete geometry and a stereoview of the crystal packing have been deposited with the IUCr (Reference: AB1142). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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Table 2. Selected geometric parameters (Å, °)

C1A—C6A	1.397 (2)	C1B—C6B	1.397 (2)
C1A—C2A	1.412 (2)	C1B—C2B	1.404 (2)
C1A—C7A	1.470 (2)	C1B—C7B	1.475 (2)
C2A—O2A	1.368 (2)	C2B—O2B	1.364 (2)
C2A—C3A	1.376 (2)	C2B—C3B	1.380 (2)
C3A—C4A	1.374 (2)	C3B—C4B	1.376 (3)
C4A—C5A	1.374 (3)	C4B—C5B	1.376 (3)
C5A—C6A	1.376 (3)	C5B—C6B	1.373 (3)
C7A—N1A	1.289 (2)	C7B—N1B	1.285 (2)
C7A—C8A	1.502 (2)	C7B—C8B	1.506 (2)
C8A—C9A	1.509 (3)	C8B—C9B	1.513 (3)
N1A—O1A	1.399 (2)	N1B—O1B	1.401 (2)
O1A—H1AO	0.86 (2)	O1B—H1BO	0.91 (2)
O2A—H2AO	0.96 (2)	O2B—H2BO	0.98 (2)
C6A—C1A—C2A	116.21 (15)	C6B—C1B—C2B	116.6 (2)
C6A—C1A—C7A	121.14 (14)	C6B—C1B—C7B	121.12 (14)
C2A—C1A—C7A	122.64 (13)	C2B—C1B—C7B	122.32 (13)
O2A—C2A—C3A	117.54 (14)	O2B—C2B—C3B	117.56 (15)
O2A—C2A—C1A	121.32 (14)	O2B—C2B—C1B	121.38 (15)
C3A—C2A—C1A	121.15 (15)	C3B—C2B—C1B	121.06 (15)
C4A—C3A—C2A	120.5 (2)	C4B—C3B—C2B	120.4 (2)
C5A—C4A—C3A	120.1 (2)	C5B—C4B—C3B	120.0 (2)
C4A—C5A—C6A	119.6 (2)	C6B—C5B—C4B	119.6 (2)
C5A—C6A—C1A	122.4 (2)	C5B—C6B—C1B	122.4 (2)
N1A—C7A—C1A	115.98 (13)	N1B—C7B—C1B	116.20 (13)
N1A—C7A—C8A	122.37 (15)	N1B—C7B—C8B	122.4 (2)
C1A—C7A—C8A	121.60 (14)	C1B—C7B—C8B	121.42 (13)
C7A—C8A—C9A	112.0 (2)	C7B—C8B—C9B	112.13 (15)
C7A—N1A—O1A	113.32 (12)	C7B—N1B—O1B	113.25 (12)

*Acta Cryst.* (1994). **C50**, 1359–1362

## (1S,2S,5R,6S)-(+)-6-Carbanilino-1,5-dimethyltricyclo[3.2.0.0<sup>2,6</sup>]heptane

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## Abstract

The asymmetric unit of the title compound, *N*-phenyl-6-(1,5-dimethyltricyclo[3.2.0.0<sup>2,6</sup>]heptane)carboxamide, C<sub>16</sub>H<sub>19</sub>NO, contains three independent molecules. There are intermolecular hydrogen bonds of the N—H...O type.