

Table 1. Fractional atomic coordinates and equivalent isotropic thermal parameters (\AA^2)
$$U_{eq} = \frac{1}{3} \sum_i \sum_j U_{ij} a_i^* a_j^* \mathbf{a}_i \cdot \mathbf{a}_j$$

	x	y	z	U_{eq}
Hg1	0.0000	0.0000	0.0000	0.044 (1)
Mn1	0.8145 (1)	0.1510 (1)	0.8800 (1)	0.040 (1)
C1	0.6810 (8)	0.2585 (10)	0.7969 (3)	0.058 (3)
O1	0.5967 (8)	0.3256 (9)	0.7458 (3)	0.086 (3)
C2	0.7599 (9)	0.3781 (10)	0.9364 (3)	0.056 (3)
O2	0.7271 (9)	0.5218 (7)	0.9720 (3)	0.085 (4)
C3	1.0673 (8)	0.2717 (9)	0.8676 (3)	0.054 (3)
O3	1.2240 (6)	0.3474 (8)	0.8584 (3)	0.076 (3)
C4	0.8944 (8)	-0.1007 (11)	0.8390 (3)	0.058 (4)
O4	0.9455 (8)	-0.2528 (8)	0.8135 (3)	0.083 (3)
C5	0.5827 (8)	0.0083 (9)	0.9062 (3)	0.049 (3)
O5	0.4400 (7)	-0.0809 (8)	0.9212 (3)	0.066 (3)

Table 2. Geometric parameters (\AA , $^\circ$)

Hg1—Mn1	2.614 (1)	Mn1—C3	1.837 (5)
Mn1—C1	1.829 (6)	Mn1—C4	1.853 (6)
Mn1—C2	1.830 (6)	Mn1—C5	1.855 (5)
Mn1—Hg1—Mn1a	180.0	Hg1—Mn1—C3	85.6 (2)
Hg1—Mn1—C1	179.0 (2)	Hg1—Mn1—C4	85.6 (2)
Hg1—Mn1—C2	83.4 (2)	Hg1—Mn1—C5	84.8 (2)

$V = 755.4(2) \text{\AA}^3$
 $Z = 2$
 $D_x = 2.596 \text{ Mg m}^{-3}$

Prism
 $0.21 \times 0.18 \times 0.15 \text{ mm}$
 Yellow

Data collection

Siemens R3m/V diffractometer
 ω -2 θ scans
 Absorption correction:
 empirical via ψ scans
 $T_{\min} = 0.14$, $T_{\max} = 0.41$
 3934 measured reflections
 1749 independent reflections
 1491 observed reflections
 $[F > 4\sigma(F)]$

$R_{\text{int}} = 0.019$
 $\theta_{\text{max}} = 27.5^\circ$
 $h = -8 \rightarrow 8$
 $k = 0 \rightarrow 8$
 $l = 0 \rightarrow 25$
 3 standard reflections
 monitored every 400
 reflections
 intensity variation: 0%

Refinement

Refinement on F
 Final $R = 0.027$
 $wR = 0.029$
 $S = 1.543$
 1491 reflections
 107 parameters
 Calculated weights
 $w = 1/[\Sigma^2(F) + 0.0001F^2]$
 $(\Delta/\sigma)_{\text{max}} = 0.001$

$\Delta\rho_{\text{max}} = 0.85 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.88 \text{ e \AA}^{-3}$
 Atomic scattering factors
 from *SHELXTL-Plus*
 (Sheldrick, 1990)
 Data collection, all calculations and graphics:
SHELXTL-Plus.

The structure was refined by full-matrix least squares with *SHELXTL-Plus*. All atoms were assigned anisotropic displacement parameters in the refinement.

Lists of structure factors, anisotropic thermal parameters and complete geometry have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 55174 (11 pp.). Copies may be obtained through The Technical Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England. [CIF reference: NA1007]

References

- Clegg, W. & Wheatley, P. J. (1971). *J. Chem. Soc. A*, pp. 3572–3574.
 Gäde, W. & Weiss, E. (1981). *Angew. Chem.* **93**, 796–797.
 Jaitner, P. E., Peringer, P., Huttner, G. & Zolnai, L. (1981). *Transition Met. Chem.* **6**, 86–90.
 Katcher, M. L. & Simon, G. L. (1972). *Inorg. Chem.* **11**, 1651–1657.
 Kuzmina, L. G., Ginzburg, A. G. G., Struchkov, Yu. T. & Kursanov, D. N. (1983). *J. Organomet. Chem.* **253**, 329–337.
 Mirsky, K. (1978). In *Computing in Crystallography*, edited by H. Schenk, R. Olthof-Hazekamp, H. van Koningsveld & G. C. Bassi, pp. 169–182. Delft Univ. Press.
 Onaka, S., Kondo, Y., Yamashita, M., Tatematsu, Y., Kato, Y., Goto, M. & Ito, T. (1985). *Inorg. Chem.* **24**, 1070–1076.
 Sheldrick, G. M. (1990). *SHELXTL-Plus. Structure Determination Software Programs*. Siemens Analytical X-ray Instruments, Inc., Madison, Wisconsin, USA.
 Williams, D. E. (1983). *PCK83*. Crystal molecular packing analysis program. QCPE Program No. 481, Univ. of Indiana, Indiana, USA.

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trans-1,3-Bis(diphenylphosphinoyl)cyclohexane

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Abstract

The relative configuration of the two diphenylphosphinoyl groups was determined as *trans* by X-ray structure analysis. The cyclohexane ring has a chair conformation with one diphenylphosphinoyl group in an equatorial position and the other in an axial position.

Comment

As part of synthetic studies of the Lewis acid-promoted double addition reactions of secondary phosphines with α,β -unsaturated aldehydes and ketones, the title compound (1) was obtained from diphenylphosphine and 2-cyclohexenone in the presence of catalytic amounts of NbCl_5 and stoichiometric $\text{BF}_3 \cdot \text{OEt}_2$ (Hashimoto, Maeta, Matsumoto, Morooka, Ohba & Suzuki, 1992). Its melting point is 555–556 K; NMR and IR data are given in Hashimoto *et al.* (1992). The O1—P1—C1—H(C1) and O2—P2—C3—H(C3) torsion angles are nearly 180° . There is an imbalance in the P—C—C bond angles for the Ph groups of $4.7(2)$ – $7.3(2)^\circ$, which may be the result of non-bonded H···H repulsions. The H atoms of C1, C8 and C18 and those of C3, C24 and C26 are arranged in

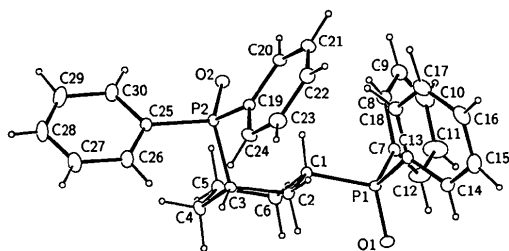
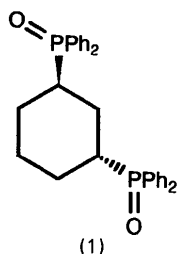


Fig. 1. The molecular structure with thermal ellipsoids at 10% probability level. H atoms are represented by circles of radius 0.1 Å.

approximately equilateral triangles with H...H distances ranging from 2.35(5) to 2.60(3) Å.



Experimental

Crystal data

$C_{30}H_{30}O_2P_2$

$M_r = 484.5$

Triclinic

$P\bar{1}$

$a = 11.476$ (1) Å

$b = 13.177$ (1) Å

$c = 9.267$ (1) Å

$\alpha = 106.19$ (1)°

$\beta = 96.38$ (1)°

$\gamma = 96.00$ (1)°

$V = 1323.9$ (2) Å³

$Z = 2$

Data collection

Rigaku AFC-5 four-circle diffractometer

θ - 2θ scans

Absorption correction: none

6361 measured reflections

6059 independent reflections

3866 observed reflections

$[|F_o| > 3\sigma(|F_o|)]$

Refinement

Refinement on F

Final $R = 0.046$

$wR = 0.037$

$S = 2.787$

3866 reflections

427 parameters

All H-atom parameters refined

Calculated weights, $w=1/\sigma$

$D_x = 1.215$ Mg m⁻³

Mo $K\alpha$ radiation

$\lambda = 0.71073$ Å

Cell parameters from 38 reflections

$\theta = 12$ – 15°

$\mu = 0.183$ mm⁻¹

$T = 299$ K

Prism

$0.45 \times 0.40 \times 0.26$ mm

Colourless

$R_{int} = 0.012$

$\theta_{max} = 27.5^\circ$

$h = 0 \rightarrow 14$

$k = -17 \rightarrow 16$

$l = -12 \rightarrow 11$

5 standard reflections

monitored every 100

reflections

intensity variation: 1%

$(\Delta/\sigma)_{max} = 0.115$

$\Delta\rho_{max} = 0.241$ e Å⁻³

$\Delta\rho_{min} = -0.296$ e Å⁻³

Atomic scattering factors from *International Tables for X-ray Crystallography* (1974, Vol. IV, Table 2.2B)

Table 1. Fractional atomic coordinates and equivalent isotropic thermal parameters (Å²)

$$U_{eq} = \frac{1}{3} \sum_i \sum_j U_{ij} a_i^* a_j^* a_i \cdot a_j$$

	x	y	z	U_{eq}
P1	0.12729 (6)	0.25541 (5)	0.31258 (8)	0.0376 (3)
P2	0.30505 (6)	0.64368 (5)	0.46583 (8)	0.0386 (3)
O1	-0.0029 (2)	0.2198 (1)	0.2794 (2)	0.0495 (9)
O2	0.3919 (2)	0.5884 (1)	0.3759 (2)	0.050 (1)
C1	0.1697 (2)	0.3870 (2)	0.2926 (3)	0.036 (1)
C2	0.1244 (3)	0.4719 (2)	0.4146 (3)	0.038 (1)
C3	0.1501 (2)	0.5849 (2)	0.4006 (3)	0.039 (1)
C4	0.1077 (3)	0.5877 (3)	0.2377 (4)	0.053 (2)
C5	0.1519 (3)	0.5016 (3)	0.1176 (4)	0.056 (2)
C6	0.1195 (3)	0.3910 (2)	0.1343 (3)	0.050 (2)
C7	0.2069 (2)	0.1659 (2)	0.1878 (3)	0.041 (1)
C8	0.3276 (3)	0.1850 (3)	0.1870 (4)	0.059 (2)
C9	0.3825 (4)	0.1142 (3)	0.0874 (5)	0.077 (2)
C10	0.3192 (4)	0.0245 (3)	-0.0123 (5)	0.086 (3)
C11	0.2013 (4)	0.0041 (4)	-0.0136 (5)	0.110 (3)
C12	0.1447 (3)	0.0755 (3)	0.0858 (4)	0.083 (2)
C13	0.1861 (2)	0.2610 (2)	0.5045 (3)	0.040 (1)
C14	0.1157 (3)	0.2116 (2)	0.5844 (4)	0.057 (2)
C15	0.1587 (5)	0.2129 (3)	0.7314 (4)	0.075 (2)
C16	0.2704 (4)	0.2618 (3)	0.7962 (5)	0.077 (2)
C17	0.3408 (4)	0.3128 (3)	0.7210 (4)	0.073 (2)
C18	0.2986 (3)	0.3114 (3)	0.5740 (4)	0.058 (2)
C19	0.3326 (2)	0.6451 (2)	0.6623 (3)	0.039 (1)
C20	0.4285 (3)	0.5969 (2)	0.7041 (4)	0.049 (2)
C21	0.4550 (3)	0.5954 (3)	0.8532 (4)	0.063 (2)
C22	0.3876 (3)	0.6392 (3)	0.9585 (4)	0.070 (2)
C23	0.2907 (3)	0.6863 (3)	0.9195 (4)	0.070 (2)
C24	0.2639 (3)	0.6880 (3)	0.7706 (4)	0.057 (2)
C25	0.3076 (3)	0.7804 (2)	0.4618 (3)	0.043 (1)
C26	0.2322 (3)	0.8467 (3)	0.5301 (4)	0.070 (2)
C27	0.2341 (4)	0.9490 (3)	0.5156 (6)	0.094 (3)
C28	0.3107 (5)	0.9829 (4)	0.4305 (6)	0.104 (3)
C29	0.3858 (5)	0.9189 (4)	0.3620 (5)	0.100 (3)
C30	0.3861 (4)	0.8174 (3)	0.3788 (4)	0.072 (2)

Table 2. Geometric parameters (Å, °)

P1—O1	1.489 (2)	P2—O2	1.489 (2)
P1—C1	1.817 (3)	P2—C3	1.827 (3)
P1—C7	1.807 (3)	P2—C19	1.807 (3)
P1—C13	1.809 (3)	P2—C25	1.810 (3)
O1—P1—C1	113.5 (1)	P1—C1—C2	110.1 (2)
O1—P1—C7	112.0 (1)	P1—C1—C6	110.3 (2)
O1—P1—C13	111.5 (1)	P2—C3—C2	112.7 (2)
C1—P1—C7	106.2 (1)	P2—C3—C4	112.2 (2)
C1—P1—C13	106.6 (1)	P1—C7—C8	123.5 (2)
C7—P1—C13	106.6 (1)	P1—C7—C12	118.6 (2)
O2—P2—C3	115.2 (1)	P1—C13—C14	118.3 (2)
O2—P2—C19	111.6 (1)	P1—C13—C18	123.0 (2)
O2—P2—C25	112.2 (1)	P2—C19—C20	117.0 (2)
C3—P2—C19	105.3 (1)	P2—C19—C24	124.3 (2)
C3—P2—C25	104.0 (1)	P2—C25—C26	123.4 (3)
C19—P2—C25	107.9 (1)	P2—C25—C30	117.6 (2)

Program(s) used to solve structure: *Xtal3.0 SIMPEL*, *GENTAN* (Hall & Stewart, 1990). Program(s) used to refine structure: *Xtal3.0 CRYLSQ* (Hall & Stewart, 1990). Molecular graphics: *Xtal3.0 ORTEP* (Hall & Stewart, 1990). Software used to prepare material for publication: *Xtal3.0 BONDLA*, *CIFIO* (Hall & Stewart, 1990).

Lists of structure factors, anisotropic thermal parameters, H-atom coordinates and complete geometry have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 55206 (18 pp.). Copies may be obtained through The Technical Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England. [CIF reference: AS1010]

References

- Hall, S. R. & Stewart, J. M. (1990). Editors. *Xtal3.0 Reference Manual*. Univs. of Western Australia, Australia, and Maryland, USA.
 Hashimoto, T., Maeta, H., Matsumoto, T., Morooka, M., Ohba, S. & Suzuki, K. (1992). *Synlett*, pp. 340–342.

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Structure of Oxatamide Monohydrate: an Anti-Allergic Drug

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Abstract

The crystal structure of the monohydrate of oxatamide, 1-{3-[4-(diphenylmethyl)-1-piperazinyl]propyl}-1,3-dihydro-2*H*-benzimidazol-2-one, has been determined at 100 K. The oxatamide molecule adopts an extended conformation with a planar benzimidazolone fragment. The water molecule has a cohesive function, connecting three oxatamide molecules by intermolecular hydrogen bonds.

Comment

Oxatamide (1) is a potent broad-scale anti-allergic drug by virtue of its inhibition of both the release and the action of allergic mediators (Awouters *et al.*, 1977). The structure analysis of oxatamide was carried out as part of an investigation into the biologically active conformation of certain H_1 -histamine-receptor agonists and antagonists (Richards, Brogden, Heel, Speight & Avery, 1984). A comparison of the crystal structure conformation with conformations obtained by molecular modelling of the active site could provide more insight into the actions of pharmaceuticals at the molecular level. The two central torsion angles in the propyl residue, N2—C8—C9—C10 and C8—C9—C10—N3, are 178.0(3) and 169.2(3)° respectively, resulting in an extended conformation rather than a folded one. The benzimidazolone moiety is planar and rotated through 78.8(3)° with respect to the almost planar C8—C9—C10—N3 chain. The piperazine ring

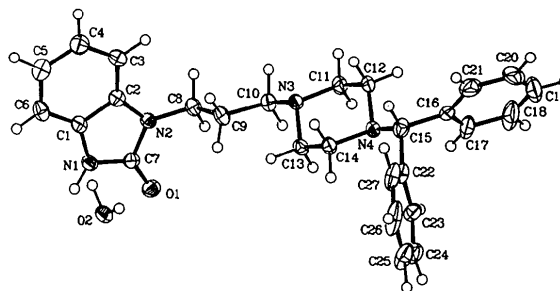
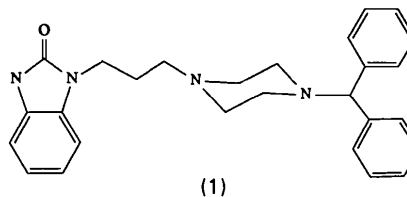


Fig. 1. View (Johnson, 1976) of oxatamide monohydrate showing the labelling of the non-H atoms. Thermal ellipsoids are shown at the 50% probability level and H atoms are drawn as small circles of arbitrary radius.

has a chair conformation, with the substituents on both N atoms in equatorial positions. All hydrogen bonds are intermolecular. The cohesive role of water in the hydrogen-bond framework is embodied by its interaction as a donor to the O1 atom of the carbonyl group of an oxatamide residue at (x, y, z) and to N3 in the piperazine ring of a residue at $(1/2-x, 1/2+y, 1/2-z)$, and as an acceptor of a hydrogen bond donated by N1—H of the residue at $(1/2-x, 1/2-y, 1-z)$. The donor-acceptor distances are 2.801(3), 2.846(4) and 2.756(4) Å respectively, and the donor-hydrogen-acceptor angles are 173(5), 176(4) and 169(4)°.



Experimental

Crystal data

$C_{27}H_{30}N_4O \cdot H_2O$

$M_r = 444.58$

Monoclinic

$C2/c$

$a = 31.1173$ (16) Å

$b = 8.9219$ (6) Å

$c = 19.0721$ (14) Å

$\beta = 112.853$ (5)°

$V = 4879.3$ (6) Å³

$Z = 8$

$D_x = 1.210$ Mg m⁻³

Mo $K\alpha$ radiation

$\lambda = 0.71073$ Å

Cell parameters from 25 reflections

$\theta = 10.84$ – 17.66 °

$\mu = 0.73$ mm⁻¹

$T = 100$ K

Platelet

$0.55 \times 0.20 \times 0.15$ mm

Colourless

Crystal source: Janssen Pharmaceutica, Beerse, Belgium

Data collection

Enraf-Nonius CAD-4

diffractometer

$\omega/2\theta$ scans

4793 measured reflections

4353 independent reflections

3005 observed reflections

$[I > 2.5\sigma(I)]$

$R_{int} = 0.0409$

$\theta_{max} = 25.37$ °

$h = -34 \rightarrow 35$

$k = -10 \rightarrow 0$

$l = -22 \rightarrow 0$

3 standard reflections

frequency: 60 min

intensity variation: 1.2%