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Crystal structure of *N*-(diphenylphosphoryl)-2-methoxybenzamide

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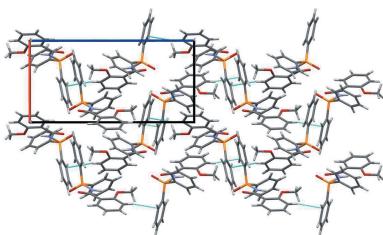
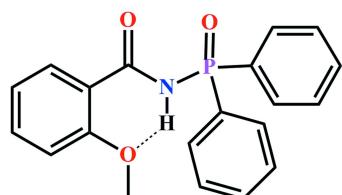
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In the title compound, $C_{20}H_{18}NO_3P$, the $C=O$ and $P=O$ groups of the carbacylamidophosphate (CAPh) fragments are located in a synclinal position relative to each other and are pre-organized for bidentate chelate coordination of metal ions. The $N-H$ group is involved in the formation of an intramolecular hydrogen bond. In the crystal, molecules do not form strong intermolecular interactions but the molecules are linked *via* weak $C-H \cdots \pi$ interactions, forming chains along [001].

1. Chemical context

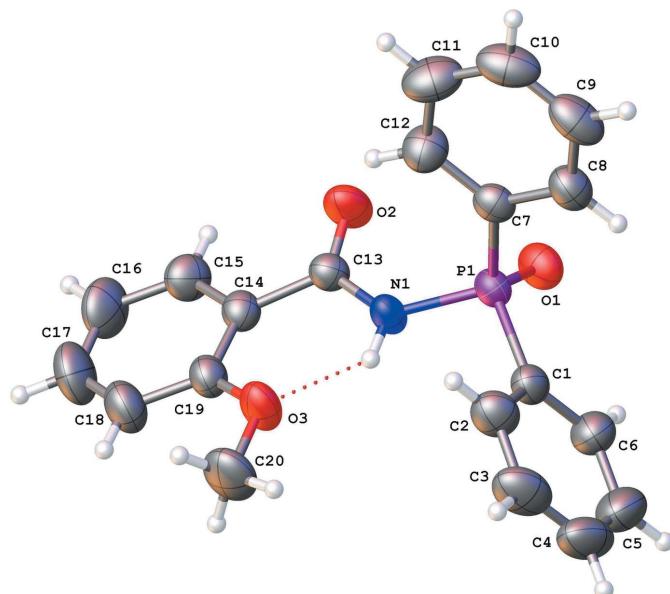
P,N -substituted analogues of β -diketones, carbacylamidophosphates (CAPh) (Amirkhanov *et al.*, 2014) that contain a $C(O)NHP(O)$ structural fragment are known for their wide range of biological activity (Adams *et al.*, 2002; Grimes *et al.*, 2008; Grynyuk *et al.*, 2016). They act as powerful chelating ligands (Skopenko *et al.*, 2004; Amirkhanov *et al.*, 2014) and as lanthanide luminescence sensitizers (Kariaka *et al.*, 2016; Pham *et al.*, 2017; Kariaka *et al.*, 2018). Thus, the syntheses and structure analysis of CAPhs are of increased interest and some structural and conformation studies of related CAPh type molecules were reported recently (Breuer *et al.*, 1990; Amirkhanov *et al.*, 1997; Milton *et al.*, 2004a,b; Kariaka *et al.*, 2014). Herein we report synthesis and crystal structure of a new CAPh, *N*-(diphenylphosphoryl)-2-methoxybenzamide (I).



2. Structural commentary

The molecular structure of the title compound is shown in Fig. 1. The bond lengths in the $C(O)NHP(O)$ fragment are typical for CAPh type ligands. The $C=O$ and $P=O$ groups are located in a synclinal position relatively to each other as evidenced by the $O1-P1-N1-C13$ torsion angle of $-45.5(2)^\circ$, $O2-C13-N1-P1$ torsion angle of $-2.7(3)^\circ$, and the $O1-P1 \cdots C13-O2$ pseudo-torsion angle of $-42.9(2)^\circ$.

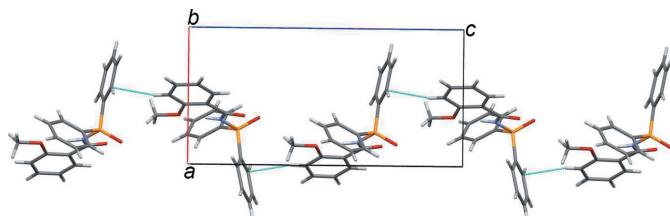
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**Figure 1**

The molecular structure of the title compound with displacement ellipsoids drawn at the 50% probability level.

As a result the CAPh fragment conformation is pre-organized for bidentate chelate coordination of metal ions.

The conjugation between the carbamide group and the anisole substituent is broken, as evidenced by the value of the C13–C14 bond length of 1.496 (3) Å, which is comparable to the mean value for non-conjugated C_{ar}–Csp² bonds (1.488 Å; Burgi & Dunitz, 1994). At the same time, the anisole and carbamide fragments are slightly non-coplanar. The C19–C14–C13–N1 torsion angle is 13.6 (3)° despite the formation of the N1–H1···O3 strong enough hydrogen bond (the H1···O3 distance is 1.93 Å and the N1–H1···O3 angle is 137°; Table 1). The methyl group of the methoxy substituent lies in the plane of the attached benzene ring despite the significant steric repulsion [the shortened intramolecular contacts are: H20A···H18 = 2.26 Å (the sum of the vdW radii is 2.32 Å; Zefirov, 1997), H20C···H18 = 2.28 Å and C20···H18 = 2.48 Å (the sum of the vdW radii is 2.87 Å)]. The phosphorus atom environment has a distorted tetrahedral configuration. The C1–C6 phenyl ring is almost coplanar with the P=O bond [the C6–C1–P1–O1 torsion angle is –5.7 (2)°] while the C7–C12 phenyl ring is rotated significantly relatively to the P=O bond as defined by the C8–C7–P1–O1 torsion angle of –72.7 (2)°.

**Figure 2**

A section of a chain along the *c* axis formed by weak C–H···π interactions (shown as blue dotted lines).

Table 1
Hydrogen-bond geometry (Å, °).

D–H···A	D–H	H···A	D···A	D–H···A
N1–H1···O3	0.86	1.93	2.628 (2)	137
C18–H18···Cg ⁱ	0.93	2.99	3.864 (3)	158

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

3. Supramolecular features

It has been shown that CAPhs display different solid-state motifs (Breuer *et al.*, 1990; Amirkhanov *et al.*, 1997; Milton *et al.*, 2004a,b; Kariaka *et al.*, 2014), *i.e.* chains, dimers and more seldom monomers. These motifs are realized through existence of hydrogen bonds with participation of the –N–H group. In crystal of (I), the –N–H group participates in an intramolecular hydrogen bond. There are no strong proton donors capable of forming intermolecular hydrogen bonds in this molecule. Thus the title molecules form only weak C–H···π interactions leading to chains of molecules along the *c*-axis direction (Figs. 2 and 3).

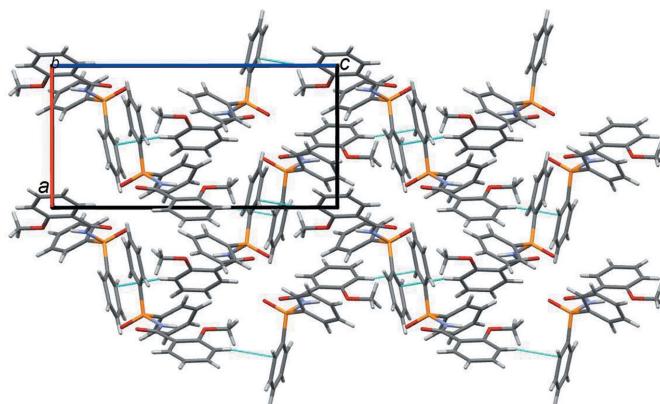
4. Database survey

A search of the Cambridge Structural Database (CSD, Version 5.40, update of November 2018; Groom *et al.*, 2016) for complexes containing CAPh ligands yielded 189 hits. In the CAPh fragments, the mean C=O bond length is 1.222 Å, the mean C–N bond length is 1.364 Å, the mean N–P bond length is 1.686 Å and the mean P=O bond length is 1.504 Å.

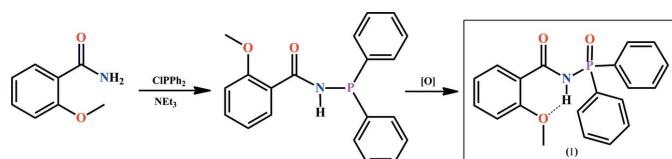
5. Synthesis and crystallization

N-(diphenylphosphoryl)-2-methoxybenzamide (I) was prepared according to a two-step reaction (Fig. 4).

To a solution of *o*-methoxybenzamide (1.51 g, 0.01 mol) and triethylamine (2.03 g, 2.8 ml, 0.02 mol) in 20 ml of dioxane was added chlorodiphenylphosphine (2.2 g, 1.79 ml, 0.01 mol) under an inert atmosphere. The mixture was stirred under reflux for 60 min and evaporated to dryness to give a pasty residue, which was mixed with 20 ml of acetone and then a

**Figure 3**

The crystal packing of the title compound viewed along the *b* axis.

**Figure 4**

The two-step reaction for the preparation of the title compound (I).

solution of 0.01 mol of H_2O_2 in 5 ml of acetone was added dropwise under vigorous stirring at 273 K. The brownish solution was evaporated and the residue was mixed with 50 ml of 10% aqueous HOAc. The solid precipitate was filtered, washed with cold water (2×20 ml) and recrystallized from *i*-PrOH [2.8 g (80%)]. Single crystals suitable for X-ray diffraction were grown from dilute *i*-PrOH solution by slow evaporation after one week.

M.p. 431–434 K. IR (KBr pellet, cm^{-1}): 3271 ν (NH), 3059 w , 3011 w , 2985 w , 2949 w , 2924 w , 2843 w , 1671 vs [$\nu(\text{CO})$], 1601 m , 1486 w , 1461 vs (Amide-II), 1436 s , 1294 m , 1242 m , 1225 vs [$\nu(\text{PO})$], 1181 m , 1160 w , 1125 s , 1109 m , 1045 w , 1012 m , 868 w [(PN)], 840 m , 786 m , 767 m , 754 s , 726 m , 702 m , 668 w , 634 w , 592 w , 543 m , 524 s , 512 m , 487 m , 442 w . ^1H NMR (DMSO-*d*6): C—H 3.95 (*s*, 3H), 7.07 (*t*, 1H), 7.22 (*d*, 1H), 7.58 (*t*, 7H), 7.67 (*d*, 1H), 7.88 (*m*, 4H), N—H 9.90 (*d*, 1H) ppm. UV–Vis abs. (CH_2Cl_2 , λ_{max} , nm): 240, 295.

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. H atoms were placed in calculated positions ($\text{N}—\text{H} = 0.86$, $\text{C}—\text{H} = 0.93$ – 0.96 Å) and refined as riding with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$ or $1.5U_{\text{eq}}(\text{C}-\text{methyl})$.

References

Table 2 Experimental details.	
Crystal data	$\text{C}_{20}\text{H}_{18}\text{NO}_3\text{P}$
Chemical formula	M_r 351.32
	Orthorhombic, $P2_12_12_1$
Crystal system, space group	Temperature (K) 294
a, b, c (Å)	a, b, c (Å) 8.317 (2), 12.657 (2), 16.763 (3)
V (Å 3)	V (Å 3) 1764.6 (6)
Z	Z 4
Radiation type	Mo $K\alpha$
μ (mm $^{-1}$)	μ (mm $^{-1}$) 0.17
Crystal size (mm)	Crystal size (mm) $0.5 \times 0.4 \times 0.3$
Data collection	
Diffractometer	Rigaku Oxford Diffraction Xcalibur, Sapphire3
Absorption correction	Multi-scan (<i>CrysAlis PRO</i> ; Rigaku OD, 2018)
T_{\min}, T_{\max}	T_{\min}, T_{\max} 0.986, 1.000
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	No. of measured, independent and observed [$I > 2\sigma(I)$] reflections 18281, 5691, 4247
R_{int}	R_{int} 0.032
(sin θ/λ) $_{\text{max}}$ (Å $^{-1}$)	(sin θ/λ) $_{\text{max}}$ (Å $^{-1}$) 0.749
Refinement	
$R[F^2 > 2\sigma(F^2)]$, $wR(F^2)$, S	$R[F^2 > 2\sigma(F^2)]$, $wR(F^2)$, S 0.043, 0.113, 1.11
No. of reflections	No. of reflections 5691
No. of parameters	No. of parameters 227
H-atom treatment	H-atom treatment H-atom parameters constrained
$\Delta\rho_{\text{max}}, \Delta\rho_{\text{min}}$ (e Å $^{-3}$)	$\Delta\rho_{\text{max}}, \Delta\rho_{\text{min}}$ (e Å $^{-3}$) 0.23, –0.32
Absolute structure	Absolute structure Flack x determined using 1430 quotients $[(I^+)-(I^-)]/[(I^+)+(I^-)]$ (Parsons <i>et al.</i> , 2013)
Absolute structure parameter	Absolute structure parameter 0.01 (4)
Computer programs: <i>CrysAlis PRO</i> (Rigaku OD, 2018), <i>SHELXT</i> (Sheldrick, 2015a), <i>SHELXL</i> (Sheldrick, 2015b) and <i>OLEX2</i> (Dolomanov <i>et al.</i> , 2009).	
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supporting information

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Crystal structure of *N*-(diphenylphosphoryl)-2-methoxybenzamide

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Computing details

Data collection: *CrysAlis PRO* (Rigaku OD, 2018); cell refinement: *CrysAlis PRO* (Rigaku OD, 2018); data reduction: *CrysAlis PRO* (Rigaku OD, 2018); program(s) used to solve structure: *SHELXT* (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL* (Sheldrick, 2015b); molecular graphics: *OLEX2* (Dolomanov *et al.*, 2009); software used to prepare material for publication: *OLEX2* (Dolomanov *et al.*, 2009).

N-(Diphenylphosphoryl)-2-methoxybenzamide

Crystal data

$C_{20}H_{18}NO_3P$
 $M_r = 351.32$
Orthorhombic, $P2_12_12_1$
 $a = 8.317$ (2) Å
 $b = 12.657$ (2) Å
 $c = 16.763$ (3) Å
 $V = 1764.6$ (6) Å³
 $Z = 4$
 $F(000) = 736$

$D_x = 1.322$ Mg m⁻³
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 5505 reflections
 $\theta = 3.2\text{--}30.6^\circ$
 $\mu = 0.17$ mm⁻¹
 $T = 294$ K
Block, colourless
0.5 × 0.4 × 0.3 mm

Data collection

Rigaku Oxford Diffraction Xcalibur, Sapphire3 diffractometer
Radiation source: fine-focus sealed X-ray tube,
Enhance (Mo) X-ray Source
Graphite monochromator
Detector resolution: 16.1827 pixels mm⁻¹
 ω scans
Absorption correction: multi-scan
(CrysAlis PRO; Rigaku OD, 2018)

$T_{\min} = 0.986$, $T_{\max} = 1.000$
18281 measured reflections
5691 independent reflections
4247 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.032$
 $\theta_{\max} = 32.2^\circ$, $\theta_{\min} = 3.2^\circ$
 $h = -12 \rightarrow 12$
 $k = -17 \rightarrow 16$
 $l = -23 \rightarrow 24$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.113$
 $S = 1.11$
5691 reflections
227 parameters
0 restraints
Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0551P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.23$ e Å⁻³
 $\Delta\rho_{\min} = -0.32$ e Å⁻³
Absolute structure: Flack x determined using
1430 quotients $[(I^+)-(I^-)]/[(I^+)+(I^-)]$ (Parsons *et al.*, 2013)
Absolute structure parameter: 0.01 (4)

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
P1	0.75683 (6)	0.34774 (4)	0.68054 (3)	0.03788 (14)
O1	0.8432 (2)	0.33778 (15)	0.75665 (10)	0.0522 (4)
O2	0.8579 (2)	0.57607 (14)	0.70208 (11)	0.0601 (5)
O3	0.8588 (2)	0.48946 (15)	0.46564 (10)	0.0574 (5)
N1	0.8291 (2)	0.43972 (14)	0.61711 (11)	0.0420 (4)
H1	0.839131	0.420776	0.568098	0.050*
C1	0.7720 (3)	0.23277 (18)	0.61801 (14)	0.0454 (5)
C2	0.6930 (4)	0.2254 (2)	0.54479 (17)	0.0585 (7)
H2	0.624578	0.279355	0.528470	0.070*
C3	0.7160 (4)	0.1383 (3)	0.4964 (2)	0.0768 (10)
H3	0.665554	0.134563	0.447009	0.092*
C4	0.8138 (5)	0.0571 (3)	0.5216 (2)	0.0815 (11)
H4	0.827802	-0.001937	0.489292	0.098*
C5	0.8912 (5)	0.0626 (3)	0.5942 (2)	0.0802 (10)
H5	0.956231	0.007064	0.610905	0.096*
C6	0.8722 (3)	0.1509 (2)	0.64224 (18)	0.0591 (6)
H6	0.926469	0.155311	0.690643	0.071*
C7	0.5462 (3)	0.37685 (18)	0.69283 (13)	0.0410 (5)
C8	0.4455 (3)	0.2966 (2)	0.72018 (16)	0.0528 (6)
H8	0.486311	0.229064	0.728194	0.063*
C9	0.2844 (3)	0.3171 (3)	0.73552 (18)	0.0653 (8)
H9	0.217061	0.262777	0.752171	0.078*
C10	0.2247 (3)	0.4165 (3)	0.72626 (17)	0.0669 (8)
H10	0.117415	0.430192	0.738074	0.080*
C11	0.3223 (3)	0.4969 (3)	0.6995 (2)	0.0706 (9)
H11	0.280930	0.564645	0.692976	0.085*
C12	0.4832 (3)	0.4765 (2)	0.68216 (19)	0.0553 (6)
H12	0.548597	0.530641	0.663257	0.066*
C13	0.8734 (3)	0.54108 (18)	0.63499 (14)	0.0405 (5)
C14	0.9427 (3)	0.60615 (19)	0.56888 (14)	0.0417 (5)
C15	1.0164 (3)	0.7003 (2)	0.59123 (18)	0.0557 (6)
H15	1.020894	0.718620	0.644915	0.067*
C16	1.0828 (4)	0.7669 (3)	0.5352 (3)	0.0769 (10)
H16	1.130614	0.830077	0.550753	0.092*
C17	1.0775 (4)	0.7386 (3)	0.4553 (2)	0.0814 (10)
H17	1.124742	0.782560	0.417536	0.098*
C18	1.0042 (4)	0.6472 (3)	0.43069 (19)	0.0662 (7)
H18	0.999839	0.630235	0.376750	0.079*
C19	0.9367 (3)	0.58040 (19)	0.48673 (15)	0.0460 (5)

C20	0.8551 (5)	0.4599 (3)	0.38290 (17)	0.0768 (9)
H20A	0.800672	0.513521	0.352765	0.115*
H20B	0.799139	0.393959	0.377171	0.115*
H20C	0.963101	0.452306	0.363487	0.115*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
P1	0.0406 (3)	0.0420 (3)	0.0310 (2)	-0.0018 (2)	0.0008 (2)	0.0024 (2)
O1	0.0519 (8)	0.0658 (11)	0.0389 (8)	0.0029 (8)	-0.0059 (7)	0.0059 (8)
O2	0.0794 (12)	0.0568 (11)	0.0442 (10)	-0.0133 (9)	0.0112 (9)	-0.0149 (8)
O3	0.0811 (12)	0.0574 (11)	0.0335 (9)	-0.0146 (9)	0.0021 (9)	0.0008 (8)
N1	0.0548 (11)	0.0402 (9)	0.0311 (9)	-0.0092 (8)	0.0059 (8)	-0.0012 (7)
C1	0.0509 (12)	0.0405 (11)	0.0446 (12)	-0.0045 (9)	0.0086 (10)	0.0009 (9)
C2	0.0654 (16)	0.0594 (16)	0.0508 (15)	-0.0080 (12)	0.0012 (12)	-0.0084 (12)
C3	0.089 (2)	0.083 (2)	0.0590 (18)	-0.0218 (19)	0.0111 (16)	-0.0255 (16)
C4	0.097 (3)	0.0608 (19)	0.086 (3)	-0.0094 (17)	0.035 (2)	-0.0242 (18)
C5	0.092 (2)	0.0549 (17)	0.094 (3)	0.0149 (16)	0.027 (2)	-0.0013 (17)
C6	0.0638 (15)	0.0519 (14)	0.0616 (16)	0.0086 (13)	0.0087 (13)	0.0025 (13)
C7	0.0408 (10)	0.0489 (12)	0.0331 (11)	-0.0032 (8)	-0.0014 (8)	-0.0044 (9)
C8	0.0548 (13)	0.0565 (15)	0.0469 (14)	-0.0089 (12)	0.0052 (11)	0.0011 (11)
C9	0.0491 (14)	0.095 (2)	0.0517 (15)	-0.0219 (14)	0.0053 (11)	-0.0048 (14)
C10	0.0392 (13)	0.104 (2)	0.0573 (17)	0.0016 (15)	-0.0015 (11)	-0.0209 (16)
C11	0.0511 (15)	0.076 (2)	0.085 (2)	0.0148 (14)	-0.0076 (15)	-0.0170 (17)
C12	0.0454 (12)	0.0560 (14)	0.0646 (16)	-0.0004 (10)	-0.0037 (12)	0.0033 (13)
C13	0.0398 (10)	0.0424 (12)	0.0391 (11)	-0.0023 (9)	0.0021 (9)	-0.0032 (9)
C14	0.0362 (10)	0.0422 (11)	0.0466 (13)	-0.0018 (9)	0.0011 (9)	0.0036 (10)
C15	0.0512 (13)	0.0511 (14)	0.0648 (17)	-0.0109 (11)	0.0025 (12)	-0.0049 (12)
C16	0.0706 (19)	0.0583 (18)	0.102 (3)	-0.0270 (15)	0.0064 (18)	0.0094 (17)
C17	0.084 (2)	0.078 (2)	0.083 (3)	-0.0257 (18)	0.0138 (19)	0.0254 (19)
C18	0.0763 (18)	0.0690 (18)	0.0532 (15)	-0.0094 (16)	0.0102 (14)	0.0132 (14)
C19	0.0463 (12)	0.0458 (13)	0.0460 (13)	-0.0034 (10)	0.0053 (10)	0.0063 (10)
C20	0.107 (3)	0.086 (2)	0.0374 (14)	-0.012 (2)	0.0033 (16)	-0.0083 (14)

Geometric parameters (\AA , ^\circ)

P1—O1	1.4695 (17)	C8—C9	1.389 (4)
P1—N1	1.6871 (19)	C9—H9	0.9300
P1—C1	1.798 (2)	C9—C10	1.361 (5)
P1—C7	1.802 (2)	C10—H10	0.9300
O2—C13	1.215 (3)	C10—C11	1.377 (5)
O3—C19	1.368 (3)	C11—H11	0.9300
O3—C20	1.437 (3)	C11—C12	1.394 (4)
N1—H1	0.8600	C12—H12	0.9300
N1—C13	1.368 (3)	C13—C14	1.496 (3)
C1—C2	1.396 (4)	C14—C15	1.392 (3)
C1—C6	1.391 (4)	C14—C19	1.416 (3)
C2—H2	0.9300	C15—H15	0.9300

C2—C3	1.382 (4)	C15—C16	1.377 (4)
C3—H3	0.9300	C16—H16	0.9300
C3—C4	1.377 (5)	C16—C17	1.388 (5)
C4—H4	0.9300	C17—H17	0.9300
C4—C5	1.378 (6)	C17—C18	1.371 (5)
C5—H5	0.9300	C18—H18	0.9300
C5—C6	1.387 (4)	C18—C19	1.382 (4)
C6—H6	0.9300	C20—H20A	0.9600
C7—C8	1.394 (3)	C20—H20B	0.9600
C7—C12	1.378 (3)	C20—H20C	0.9600
C8—H8	0.9300		
O1—P1—N1	115.61 (10)	C9—C10—H10	119.8
O1—P1—C1	113.72 (11)	C9—C10—C11	120.3 (2)
O1—P1—C7	113.17 (10)	C11—C10—H10	119.8
N1—P1—C1	99.56 (10)	C10—C11—H11	120.1
N1—P1—C7	106.09 (11)	C10—C11—C12	119.8 (3)
C1—P1—C7	107.49 (11)	C12—C11—H11	120.1
C19—O3—C20	118.6 (2)	C7—C12—C11	120.5 (3)
P1—N1—H1	116.4	C7—C12—H12	119.7
C13—N1—P1	127.23 (16)	C11—C12—H12	119.7
C13—N1—H1	116.4	O2—C13—N1	121.0 (2)
C2—C1—P1	122.3 (2)	O2—C13—C14	121.7 (2)
C6—C1—P1	118.4 (2)	N1—C13—C14	117.24 (19)
C6—C1—C2	119.3 (2)	C15—C14—C13	116.2 (2)
C1—C2—H2	119.9	C15—C14—C19	118.3 (2)
C3—C2—C1	120.3 (3)	C19—C14—C13	125.4 (2)
C3—C2—H2	119.9	C14—C15—H15	119.4
C2—C3—H3	120.1	C16—C15—C14	121.1 (3)
C4—C3—C2	119.9 (3)	C16—C15—H15	119.4
C4—C3—H3	120.1	C15—C16—H16	120.4
C3—C4—H4	119.7	C15—C16—C17	119.2 (3)
C3—C4—C5	120.6 (3)	C17—C16—H16	120.4
C5—C4—H4	119.7	C16—C17—H17	119.2
C4—C5—H5	120.0	C18—C17—C16	121.5 (3)
C4—C5—C6	120.0 (3)	C18—C17—H17	119.2
C6—C5—H5	120.0	C17—C18—H18	120.3
C1—C6—H6	120.0	C17—C18—C19	119.5 (3)
C5—C6—C1	119.9 (3)	C19—C18—H18	120.3
C5—C6—H6	120.0	O3—C19—C14	117.5 (2)
C8—C7—P1	118.24 (19)	O3—C19—C18	122.1 (2)
C12—C7—P1	122.82 (18)	C18—C19—C14	120.4 (2)
C12—C7—C8	118.8 (2)	O3—C20—H20A	109.5
C7—C8—H8	119.8	O3—C20—H20B	109.5
C9—C8—C7	120.3 (3)	O3—C20—H20C	109.5
C9—C8—H8	119.8	H20A—C20—H20B	109.5
C8—C9—H9	119.9	H20A—C20—H20C	109.5
C10—C9—C8	120.2 (3)	H20B—C20—H20C	109.5

C10—C9—H9	119.9		
P1—N1—C13—O2	−2.7 (3)	C3—C4—C5—C6	−0.6 (5)
P1—N1—C13—C14	176.74 (16)	C4—C5—C6—C1	1.5 (5)
P1—C1—C2—C3	176.0 (2)	C6—C1—C2—C3	−0.9 (4)
P1—C1—C6—C5	−177.8 (2)	C7—P1—N1—C13	80.8 (2)
P1—C7—C8—C9	175.8 (2)	C7—P1—C1—C2	51.3 (2)
P1—C7—C12—C11	−174.1 (2)	C7—P1—C1—C6	−131.8 (2)
O1—P1—N1—C13	−45.5 (2)	C7—C8—C9—C10	−2.0 (4)
O1—P1—C1—C2	177.4 (2)	C8—C7—C12—C11	0.9 (4)
O1—P1—C1—C6	−5.7 (2)	C8—C9—C10—C11	1.8 (4)
O1—P1—C7—C8	−72.7 (2)	C9—C10—C11—C12	−0.3 (5)
O1—P1—C7—C12	102.2 (2)	C10—C11—C12—C7	−1.0 (5)
O2—C13—C14—C15	12.0 (3)	C12—C7—C8—C9	0.6 (4)
O2—C13—C14—C19	−166.9 (2)	C13—C14—C15—C16	−179.3 (2)
N1—P1—C1—C2	−59.0 (2)	C13—C14—C19—O3	1.2 (4)
N1—P1—C1—C6	117.90 (19)	C13—C14—C19—C18	179.5 (2)
N1—P1—C7—C8	159.47 (19)	C14—C15—C16—C17	−0.8 (5)
N1—P1—C7—C12	−25.6 (2)	C15—C14—C19—O3	−177.7 (2)
N1—C13—C14—C15	−167.5 (2)	C15—C14—C19—C18	0.6 (4)
N1—C13—C14—C19	13.6 (3)	C15—C16—C17—C18	1.7 (6)
C1—P1—N1—C13	−167.8 (2)	C16—C17—C18—C19	−1.4 (5)
C1—P1—C7—C8	53.7 (2)	C17—C18—C19—O3	178.4 (3)
C1—P1—C7—C12	−131.3 (2)	C17—C18—C19—C14	0.2 (4)
C1—C2—C3—C4	1.8 (5)	C19—C14—C15—C16	−0.3 (4)
C2—C1—C6—C5	−0.7 (4)	C20—O3—C19—C14	−178.1 (3)
C2—C3—C4—C5	−1.0 (5)	C20—O3—C19—C18	3.6 (4)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1···O3	0.86	1.93	2.628 (2)	137
C18—H18···Cg ⁱ	0.93	2.99	3.864 (3)	158

Symmetry code: (i) $-x+3/2, -y+1, z-1/2$.