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 Diphenyl (*p*-tolylamido)phosphate

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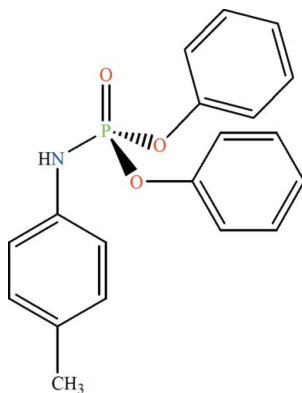
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 Key indicators: single-crystal X-ray study; $T = 120$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.045; wR factor = 0.099; data-to-parameter ratio = 17.2.

 The P atom in the title compound, $\text{C}_{19}\text{H}_{18}\text{NO}_3\text{P}$, exhibits a distorted tetrahedral configuration while the N atom shows a planar coordination. In the crystal, intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds form centrosymmetric dimers.

Related literature

 The reaction of compounds having phosphorus-halide bonds with primary and secondary amines results in formation of phosphorus-nitrogen compounds, see: Chivers *et al.* (2003). For amidophosphoric acid esters (APEs), see: Gholivand *et al.* (2007); Ghadimi *et al.* (2007). For applications of APEs, see: Bao *et al.* (1993); Ghadimi *et al.* (2008); Nguyen & Kim (2008).


Experimental

Crystal data

 $\text{C}_{19}\text{H}_{18}\text{NO}_3\text{P}$
 $M_r = 339.31$

 Triclinic, $P\bar{1}$
 $a = 9.7406$ (10) Å
 $b = 9.9653$ (10) Å
 $c = 11.1788$ (12) Å
 $\alpha = 96.337$ (2)°
 $\beta = 109.303$ (2)°
 $\gamma = 117.827$ (2)°

 $V = 858.93$ (15) Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 0.18$ mm⁻¹
 $T = 120$ K
 $0.24 \times 0.21 \times 0.11$ mm

Data collection

 Bruker SMART 1000 CCD area-detector diffractometer
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.980$, $T_{\max} = 0.989$

 8165 measured reflections
 3744 independent reflections
 3232 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.020$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.045$
 $wR(F^2) = 0.099$
 $S = 1.01$
 3744 reflections

 218 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.31$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.57$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N1}-\text{H1}\cdots\text{O1}^i$	0.79	2.07	2.840 (3)	167

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

Data collection: SMART (Bruker, 1998); cell refinement: SAINT-Plus (Bruker, 1998); data reduction: SAINT-Plus; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

Support of this investigation by Ferdowsi University of Mashhad is gratefully acknowledged.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: NG5073).

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supporting information

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Diphenyl (*p*-tolylamido)phosphate

Mehrdad Pourayoubi, Poorya Zargaran, Shahriar Ghamamy and Hossein Eshtiagh-Hosseini

S1. Comment

The reaction of compounds having phosphorus-halide bonds with primary and secondary amines results in formation of phosphorus-nitrogen compounds (Chivers *et al.*, 2003). Amidophosphoric acid esters (APEs), a family of these compounds, contain an (O)(O)P(=O)(N) or (O)P(=O)(N)(N) skeleton which may be obtained from some initial phosphorus substances such as (RO)P(O)Cl₂ and (NR¹R²)(RO)P(O)Cl (Gholivand *et al.*, 2007; Ghadimi *et al.*, 2007). APEs have attracted attention because of the flame retardancy for their bisphosphorus derivatives (Nguyen & Kim, 2008), chiral catalyst preparation (Bao *et al.*, 1993) and biological properties (Ghadimi *et al.*, 2008).

Here, we report on the synthesis and single-crystal X-ray determination of title APE compound (Fig. 1); single crystals were obtained from CHCl₃/n-C₇H₁₆ at r. t.

The phosphorus atom has a distorted tetrahedral configuration with the bond angles in the range of 94.65 (7)° [O(2)–P(1)–O(3)] to 117.14 (8)° [O(1)–P(1)–O(2)]. In crystal lattice, the H-bonded centrosymmetric dimer is formed *via* an intermolecular PO⋯HN hydrogen bond (N⋯O = 2.840 (3) Å). A fragment of unit cell packing showing the hydrogen bond is presented in Fig. 2.

S2. Experimental

To a solution of (C₆H₅O)₂P(O)Cl in chloroform, a solution of *p*-toluidine and triethylamine (1:1:1 mole ratio) in chloroform was added at 273 K. After 4 h stirring, the solvent was removed and product was washed with distilled water and recrystallized from chloroform/n-heptane at room temperature. IR (KBr, cm⁻¹): 3170.6 (NH), 3057.7, 2939.8, 2866.6, 2713.5, 2618.1, 2385.3, 1952.3, 1884.4, 1786.4, 1729.0, 1597.2, 1490.3, 1393.0, 1279.7, 1229.7, 1175.6, 1069.5, 974.8, 891.6, 819.2, 762.4, 681.6. Raman (cm⁻¹): 3066.5, 3018.3, 2933.4, 1616.1, 1589.1, 1382.7, 1288.2, 1253.5, 1216.9, 1191.8, 1170.6, 1068.4, 1027.9, 1006.6, 941.1, 904.5, 837.0, 781.0, 759.8, 730.9, 636.4, 615.2, 590.1, 368.3, 333.6. ³¹P{¹H} NMR (202.45 MHz, DMSO-d₆, 300.0 K, H₃PO₄ external): -6.36 p.p.m. (*s*). ¹H NMR (500.13 MHz, DMSO-d₆, 300.0 K, TMS): 2.21 (*s*, 3H, CH₃), 7.09 (*s*, 4H, Ar–H), 7.19–7.23 (*m*, 6H, Ar–H), 7.37–7.40 (*m*, 4H, Ar–H), 8.70 p.p.m. (*d*, ²J(P,H) = 10.6 Hz, 1H, NH). ¹³C NMR (125.75 MHz, DMSO-d₆, 300.0 K, TMS): 20.16 (*s*, 1 C, CH₃), 117.86 (*d*, ³J(P,C) = 7.7 Hz, 2 C, C_{ortho}), 120.05 (*d*, ³J(P,C) = 4.8 Hz, 4 C, C_{ortho}), 125.14 (*s*), 129.60 (*s*), 129.87 (*s*), 130.44 (*s*), 137.09 (*s*), 150.11 p.p.m. (*d*, ²J(P,C) = 6.3 Hz, 2 C, C_{ipso}).

S3. Refinement

The hydrogen atom of NH group was found in difference Fourier synthesis. The H(C) atom positions were calculated. The H(N) atom was refined in isotropic approximation in riding model, the H(C) atoms were refined in isotropic approximation in riding model with the *U*_{iso}(H) parameters equal to 1.2 *U*_{eq}(Xi) or 1.5 *U*_{eq}(Cii), where *U*(Xi) are the equivalent thermal parameters of the NH and CH atoms and *U*(Cii) are the ones of the CH₃ carbon atoms to which the corresponding H atoms are bonded.

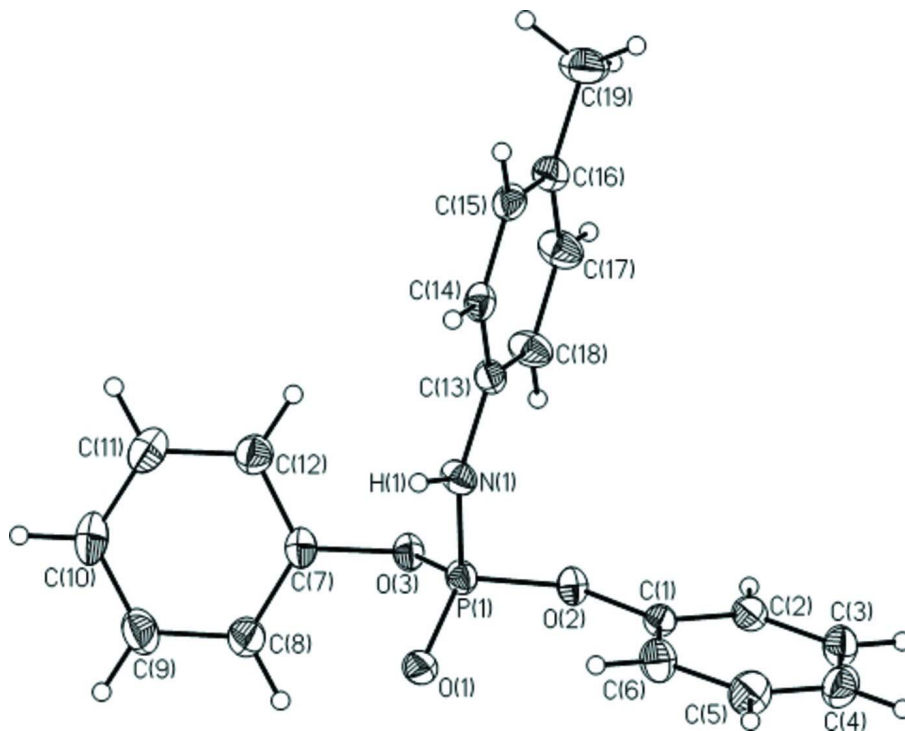


Figure 1

Molecular view with the atom labeling scheme, displacement ellipsoids are drawn at the 50% probability level. H atoms are represented as small spheres of arbitrary radii.

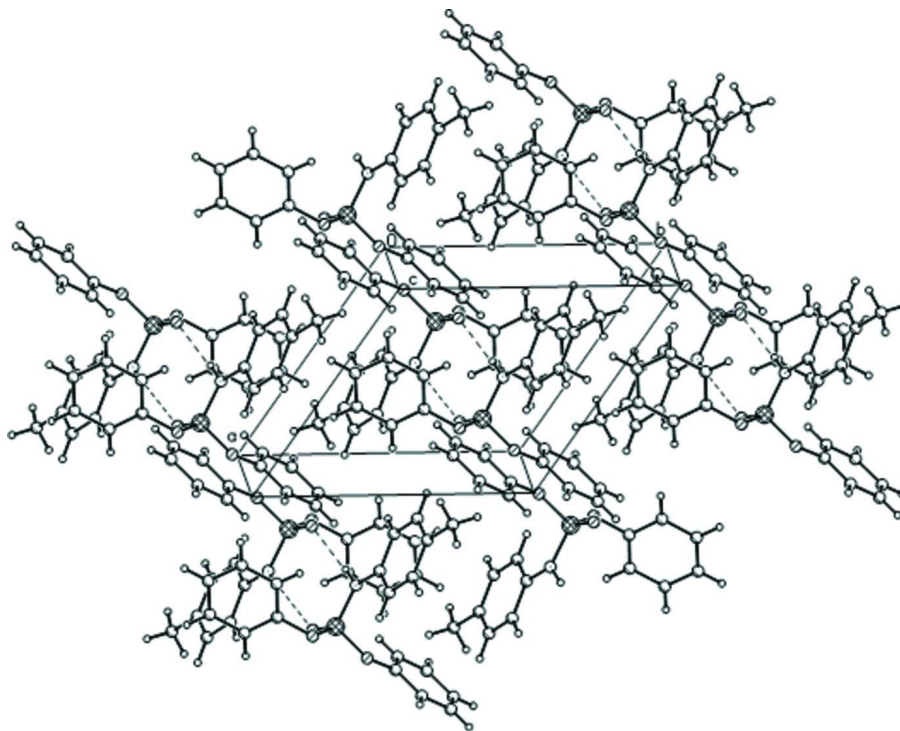


Figure 2

Partial packing view showing the formation of dimer through N–H···O interaction. H bonds are shown as dashed lines.

Diphenyl (*p*-tolylamido)phosphate

Crystal data

$C_{19}H_{18}NO_3P$

$M_r = 339.31$

Triclinic, $P\bar{1}$

$a = 9.7406$ (10) Å

$b = 9.9653$ (10) Å

$c = 11.1788$ (12) Å

$\alpha = 96.337$ (2)°

$\beta = 109.303$ (2)°

$\gamma = 117.827$ (2)°

$V = 858.93$ (15) Å³

$Z = 2$

$F(000) = 356$

$D_x = 1.312$ Mg m⁻³

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

Cell parameters from 948 reflections

$\theta = 3$ –29°

$\mu = 0.18$ mm⁻¹

$T = 120$ K

Prism, colorless

0.24 × 0.21 × 0.11 mm

Data collection

Bruker SMART 1000 CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

phi and ω scans

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 1996)

$T_{\min} = 0.980$, $T_{\max} = 0.989$

8165 measured reflections

3744 independent reflections

3232 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.020$

$\theta_{\max} = 27.0^\circ$, $\theta_{\min} = 2.0^\circ$

$h = -12 \rightarrow 12$

$k = -12 \rightarrow 12$

$l = -14 \rightarrow 14$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.045$

$wR(F^2) = 0.099$

$S = 1.01$

3744 reflections

218 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: mixed

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.010P)^2 + 1.4P]$

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

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

$\Delta\rho_{\max} = 0.31$ e Å⁻³

$\Delta\rho_{\min} = -0.57$ e Å⁻³

Special details

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

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
P1	0.30226 (6)	0.32453 (5)	0.30029 (5)	0.01976 (12)

O1	0.26650 (17)	0.38136 (15)	0.40642 (13)	0.0232 (3)
O2	0.16664 (16)	0.14657 (15)	0.20459 (12)	0.0212 (3)
O3	0.28250 (16)	0.40203 (15)	0.18341 (13)	0.0216 (3)
N1	0.4912 (2)	0.34721 (19)	0.36132 (15)	0.0214 (3)
H1	0.5535	0.4125	0.4324	0.026*
C1	0.1005 (2)	0.0096 (2)	0.24317 (19)	0.0203 (4)
C2	-0.0075 (2)	-0.1338 (2)	0.1388 (2)	0.0235 (4)
H2A	-0.0313	-0.1345	0.0492	0.028*
C3	-0.0801 (3)	-0.2761 (2)	0.1673 (2)	0.0296 (4)
H3A	-0.1534	-0.3750	0.0968	0.036*
C4	-0.0461 (3)	-0.2743 (3)	0.2985 (2)	0.0319 (5)
H4A	-0.0961	-0.3719	0.3179	0.038*
C5	0.0611 (3)	-0.1297 (3)	0.4011 (2)	0.0325 (5)
H5A	0.0837	-0.1289	0.4907	0.039*
C6	0.1362 (3)	0.0147 (2)	0.3750 (2)	0.0276 (4)
H6A	0.2097	0.1137	0.4454	0.033*
C7	0.3697 (2)	0.5708 (2)	0.22208 (19)	0.0227 (4)
C8	0.2885 (3)	0.6432 (2)	0.2520 (2)	0.0300 (4)
H8A	0.1772	0.5812	0.2487	0.036*
C9	0.3736 (3)	0.8088 (3)	0.2869 (2)	0.0347 (5)
H9A	0.3204	0.8612	0.3081	0.042*
C10	0.5357 (3)	0.8980 (2)	0.2909 (2)	0.0335 (5)
H10A	0.5929	1.0112	0.3143	0.040*
C11	0.6148 (3)	0.8228 (3)	0.2610 (2)	0.0327 (5)
H11A	0.7263	0.8845	0.2646	0.039*
C12	0.5309 (3)	0.6567 (2)	0.2257 (2)	0.0272 (4)
H12A	0.5838	0.6039	0.2046	0.033*
C13	0.5718 (2)	0.3089 (2)	0.29081 (18)	0.0204 (4)
C14	0.7373 (2)	0.3403 (2)	0.36440 (19)	0.0216 (4)
H14A	0.7917	0.3854	0.4590	0.026*
C15	0.8219 (2)	0.3056 (2)	0.2994 (2)	0.0238 (4)
H15A	0.9342	0.3281	0.3504	0.029*
C16	0.7450 (2)	0.2383 (2)	0.1608 (2)	0.0255 (4)
C17	0.5803 (3)	0.2073 (3)	0.0896 (2)	0.0294 (4)
H17A	0.5251	0.1608	-0.0049	0.035*
C18	0.4949 (3)	0.2425 (3)	0.1528 (2)	0.0275 (4)
H18A	0.3832	0.2212	0.1015	0.033*
C19	0.8337 (3)	0.1978 (3)	0.0886 (2)	0.0363 (5)
H19A	0.7571	0.1485	-0.0074	0.054*
H19B	0.8614	0.1228	0.1227	0.054*
H19C	0.9401	0.2957	0.1035	0.054*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
P1	0.0190 (2)	0.0172 (2)	0.0197 (2)	0.00924 (19)	0.00640 (18)	0.00282 (18)
O1	0.0218 (6)	0.0218 (7)	0.0239 (7)	0.0107 (6)	0.0102 (5)	0.0030 (5)
O2	0.0227 (6)	0.0166 (6)	0.0198 (6)	0.0095 (5)	0.0069 (5)	0.0035 (5)

O3	0.0215 (6)	0.0166 (6)	0.0225 (6)	0.0093 (5)	0.0070 (5)	0.0038 (5)
N1	0.0191 (7)	0.0228 (8)	0.0174 (7)	0.0118 (7)	0.0039 (6)	0.0000 (6)
C1	0.0181 (8)	0.0185 (8)	0.0253 (9)	0.0105 (7)	0.0094 (7)	0.0068 (7)
C2	0.0224 (9)	0.0226 (9)	0.0250 (9)	0.0123 (8)	0.0101 (8)	0.0051 (8)
C3	0.0259 (10)	0.0201 (9)	0.0355 (11)	0.0099 (8)	0.0108 (9)	0.0035 (8)
C4	0.0300 (11)	0.0246 (10)	0.0435 (12)	0.0140 (9)	0.0178 (10)	0.0162 (9)
C5	0.0370 (12)	0.0332 (11)	0.0314 (11)	0.0197 (10)	0.0166 (9)	0.0159 (9)
C6	0.0310 (10)	0.0241 (10)	0.0236 (10)	0.0140 (9)	0.0090 (8)	0.0057 (8)
C7	0.0240 (9)	0.0174 (9)	0.0229 (9)	0.0105 (8)	0.0071 (7)	0.0060 (7)
C8	0.0246 (10)	0.0244 (10)	0.0399 (12)	0.0135 (8)	0.0129 (9)	0.0090 (9)
C9	0.0365 (12)	0.0258 (10)	0.0448 (13)	0.0208 (10)	0.0152 (10)	0.0100 (9)
C10	0.0350 (11)	0.0187 (9)	0.0351 (11)	0.0108 (9)	0.0079 (9)	0.0092 (8)
C11	0.0270 (10)	0.0268 (10)	0.0369 (12)	0.0093 (9)	0.0128 (9)	0.0132 (9)
C12	0.0264 (10)	0.0252 (10)	0.0300 (10)	0.0136 (8)	0.0122 (8)	0.0098 (8)
C13	0.0198 (9)	0.0176 (8)	0.0229 (9)	0.0105 (7)	0.0082 (7)	0.0048 (7)
C14	0.0193 (9)	0.0165 (8)	0.0215 (9)	0.0077 (7)	0.0044 (7)	0.0038 (7)
C15	0.0166 (8)	0.0198 (9)	0.0300 (10)	0.0086 (7)	0.0068 (8)	0.0067 (8)
C16	0.0221 (9)	0.0235 (9)	0.0303 (10)	0.0126 (8)	0.0112 (8)	0.0054 (8)
C17	0.0286 (10)	0.0363 (11)	0.0223 (10)	0.0200 (9)	0.0082 (8)	0.0033 (8)
C18	0.0228 (9)	0.0351 (11)	0.0240 (10)	0.0188 (9)	0.0061 (8)	0.0037 (8)
C19	0.0263 (10)	0.0434 (13)	0.0379 (12)	0.0191 (10)	0.0148 (9)	0.0040 (10)

Geometric parameters (Å, °)

P1—O1	1.4691 (14)	C8—H8A	0.9500
P1—O2	1.5793 (13)	C9—C10	1.385 (3)
P1—O3	1.5926 (14)	C9—H9A	0.9500
P1—N1	1.6279 (16)	C10—C11	1.386 (3)
O2—C1	1.399 (2)	C10—H10A	0.9500
O3—C7	1.414 (2)	C11—C12	1.394 (3)
N1—C13	1.417 (2)	C11—H11A	0.9500
N1—H1	0.7881	C12—H12A	0.9500
C1—C6	1.387 (3)	C13—C18	1.388 (3)
C1—C2	1.388 (3)	C13—C14	1.403 (3)
C2—C3	1.387 (3)	C14—C15	1.391 (3)
C2—H2A	0.9500	C14—H14A	0.9500
C3—C4	1.390 (3)	C15—C16	1.395 (3)
C3—H3A	0.9500	C15—H15A	0.9500
C4—C5	1.386 (3)	C16—C17	1.393 (3)
C4—H4A	0.9500	C16—C19	1.509 (3)
C5—C6	1.393 (3)	C17—C18	1.387 (3)
C5—H5A	0.9500	C17—H17A	0.9500
C6—H6A	0.9500	C18—H18A	0.9500
C7—C8	1.383 (3)	C19—H19A	0.9800
C7—C12	1.377 (3)	C19—H19B	0.9800
C8—C9	1.389 (3)	C19—H19C	0.9800
O1—P1—O2	117.14 (8)	C10—C9—H9A	119.8

O1—P1—O3	114.32 (8)	C8—C9—H9A	119.8
O2—P1—O3	94.65 (7)	C11—C10—C9	120.22 (19)
O1—P1—N1	111.29 (8)	C11—C10—H10A	119.9
O2—P1—N1	107.89 (8)	C9—C10—H10A	119.9
O3—P1—N1	110.35 (8)	C10—C11—C12	120.1 (2)
C1—O2—P1	126.66 (12)	C10—C11—H11A	119.9
C7—O3—P1	116.80 (11)	C12—C11—H11A	119.9
C13—N1—P1	127.92 (13)	C7—C12—C11	118.45 (19)
C13—N1—H1	115.8	C7—C12—H12A	120.8
P1—N1—H1	112.7	C11—C12—H12A	120.8
C6—C1—C2	121.82 (18)	C18—C13—C14	118.81 (17)
C6—C1—O2	123.22 (16)	C18—C13—N1	123.00 (16)
C2—C1—O2	114.95 (16)	C14—C13—N1	118.18 (16)
C3—C2—C1	119.12 (18)	C15—C14—C13	120.21 (17)
C3—C2—H2A	120.4	C15—C14—H14A	119.9
C1—C2—H2A	120.4	C13—C14—H14A	119.9
C4—C3—C2	120.23 (19)	C14—C15—C16	121.26 (17)
C4—C3—H3A	119.9	C14—C15—H15A	119.4
C2—C3—H3A	119.9	C16—C15—H15A	119.4
C3—C4—C5	119.69 (19)	C17—C16—C15	117.64 (18)
C3—C4—H4A	120.2	C17—C16—C19	120.30 (19)
C5—C4—H4A	120.2	C15—C16—C19	122.06 (18)
C6—C5—C4	121.1 (2)	C18—C17—C16	121.78 (19)
C6—C5—H5A	119.5	C18—C17—H17A	119.1
C4—C5—H5A	119.5	C16—C17—H17A	119.1
C1—C6—C5	118.05 (19)	C17—C18—C13	120.29 (18)
C1—C6—H6A	121.0	C17—C18—H18A	119.9
C5—C6—H6A	121.0	C13—C18—H18A	119.9
C8—C7—C12	122.44 (18)	C16—C19—H19A	109.5
C8—C7—O3	118.89 (17)	C16—C19—H19B	109.5
C12—C7—O3	118.65 (17)	H19A—C19—H19B	109.5
C7—C8—C9	118.42 (19)	C16—C19—H19C	109.5
C7—C8—H8A	120.8	H19A—C19—H19C	109.5
C9—C8—H8A	120.8	H19B—C19—H19C	109.5
C10—C9—C8	120.3 (2)		
O1—P1—O2—C1	-50.89 (16)	C12—C7—C8—C9	0.0 (3)
O3—P1—O2—C1	-171.29 (14)	O3—C7—C8—C9	-178.54 (18)
N1—P1—O2—C1	75.58 (15)	C7—C8—C9—C10	0.2 (3)
O1—P1—O3—C7	50.54 (15)	C8—C9—C10—C11	-0.4 (3)
O2—P1—O3—C7	173.16 (13)	C9—C10—C11—C12	0.5 (3)
N1—P1—O3—C7	-75.81 (14)	C8—C7—C12—C11	0.0 (3)
O1—P1—N1—C13	-179.07 (15)	O3—C7—C12—C11	178.59 (17)
O2—P1—N1—C13	51.11 (18)	C10—C11—C12—C7	-0.3 (3)
O3—P1—N1—C13	-51.04 (18)	P1—N1—C13—C18	-0.3 (3)
P1—O2—C1—C6	5.8 (3)	P1—N1—C13—C14	179.39 (14)
P1—O2—C1—C2	-175.33 (13)	C18—C13—C14—C15	0.1 (3)
C6—C1—C2—C3	-0.8 (3)	N1—C13—C14—C15	-179.59 (17)

O2—C1—C2—C3	-179.63 (17)	C13—C14—C15—C16	-0.4 (3)
C1—C2—C3—C4	0.6 (3)	C14—C15—C16—C17	0.1 (3)
C2—C3—C4—C5	-0.1 (3)	C14—C15—C16—C19	-179.19 (19)
C3—C4—C5—C6	-0.2 (3)	C15—C16—C17—C18	0.4 (3)
C2—C1—C6—C5	0.4 (3)	C19—C16—C17—C18	179.8 (2)
O2—C1—C6—C5	179.22 (18)	C16—C17—C18—C13	-0.7 (3)
C4—C5—C6—C1	0.0 (3)	C14—C13—C18—C17	0.4 (3)
P1—O3—C7—C8	-86.63 (19)	N1—C13—C18—C17	-179.86 (19)
P1—O3—C7—C12	94.78 (19)		

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>
N1—H1 \cdots O1 ⁱ	0.79	2.07	2.840 (3)	167

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