

N-Benzyl-P-(2-ethylphenyl)-P-phenyl-phosphinic amide

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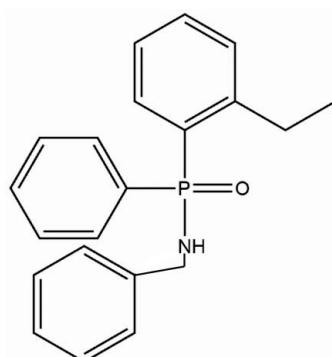
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Key indicators: single-crystal X-ray study; $T = 100\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.044; wR factor = 0.118; data-to-parameter ratio = 13.5.

In the crystal structure of the title compound, $\text{C}_{21}\text{H}_{22}\text{NOP}$, the amine H atom is involved in $\text{N}-\text{H}\cdots\text{O}$ hydrogen-bonding interactions, resulting in chains along the c axis. The crystal lattice is consolidated by weak intermolecular $\text{C}-\text{H}\cdots\pi$ interactions.

Related literature

For the uses of phosphinamides, see: Wuts & Greene (2006); Burgos *et al.* (2008); Popovici *et al.* (2010). For related compounds, see: Priya *et al.* (2005); Fei *et al.* (2004); Gaw *et al.* (1999).



Experimental

Crystal data

$\text{C}_{21}\text{H}_{22}\text{NOP}$

$M_r = 335.37$

Monoclinic, $P2_1/c$

$a = 12.9259(3)\text{ \AA}$

$b = 15.7098(3)\text{ \AA}$

$c = 9.1007(2)\text{ \AA}$

$\beta = 107.578(1)^\circ$
 $V = 1761.73(7)\text{ \AA}^3$
 $Z = 4$
 $\text{Cu } K\alpha$ radiation

$\mu = 1.42\text{ mm}^{-1}$
 $T = 100\text{ K}$
 $0.48 \times 0.08 \times 0.02\text{ mm}$

Data collection

Bruker X8 APEXII 4K KappaCCD diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2008)
 $T_{\min} = 0.549$, $T_{\max} = 0.972$

13806 measured reflections
2946 independent reflections
2772 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.034$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.118$
 $S = 1.05$
2946 reflections

218 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.57\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.57\text{ e \AA}^{-3}$

Table 1

Hydrogen-bond geometry (\AA , $^\circ$).

$Cg2$ and $Cg3$ are the centroids of the C8–C13 and C14–C19 rings, respectively.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1–H1 \cdots O1 ⁱ	0.88	2.09	2.742 (2)	131
C18–H18 \cdots Cg2 ⁱⁱ	0.95	2.98	3.756 (2)	139
C21–H21C \cdots Cg3 ⁱⁱ	0.98	2.83	3.636 (3)	140

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

Data collection: *APEX2* (Bruker, 2008); cell refinement: *SAINT-Plus* (Bruker, 2008); data reduction: *SAINT-Plus* and *XPREP* (Bruker, 2008); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

We thank NRF, THRIP and University of Johannesburg for financial support.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HG5139).

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supporting information

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S1. Comment

Phosphinamides are important functional groups in organic synthesis. They have been employed as amine protecting groups and as substrates for imine activation (Wuts *et al.*, 2006). Besides functioning as protective groups, phosphinamides are also used as catalysts for enantioselective reduction of ketones as building blocks for the synthesis of peptidomimetics *via* phosphinamide-directed benzylic lithiation (Burgos *et al.*, 2008) and also as chiral ligands (Popovici *et al.*, 2010). Herein, we have synthesized a racemic mixture of a chiral phosphinamide and report its crystal structure.

The asymmetric unit of (I), (Fig. 1) contains one molecule. The phosphorus is in a tetrahedral environment as in other phosphine oxides. The P=O, P–N and P–C bond distances are comparable to similar compounds in literature (Priya *et al.*, 2005; Fei *et al.*, 2004; Gaw *et al.*, 1999). In the crystal of (I), O atom is involved in N–H···O=P hydrogen bonding interaction (Table 1), thus resulting in chains that run in the crystallographic *c* direction. The crystal lattice is consolidated by a pair of weak C–H···π intermolecular interactions [$C18\cdots Cg = 3.756(2)$ Å, $\angle C18\text{--}H18\cdots Cg = 139$ ° $C21\cdots Cg = 3.636(3)$ Å, $\angle C21\text{--}H21c\cdots Cg = 140$ °] (Fig. 2).

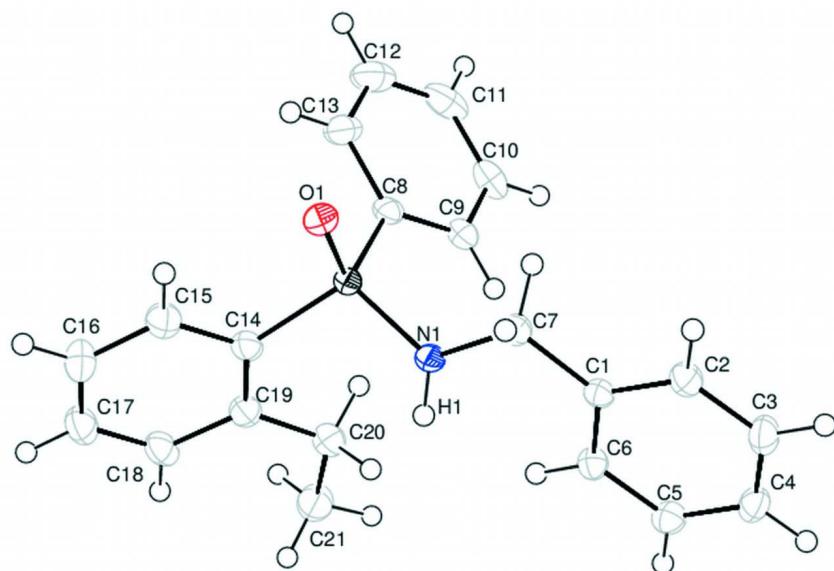
S2. Experimental

To a solution of 1-bromo-2-ethylbenzene (1 ml, 7.3 mmol) in dry THF (10 ml) were added Mg (263 mg, 11 mmol) and a catalytic amount of iodine crystals. The resulting mixture was refluxed overnight under nitrogen atmosphere. The resulting Grignard reagent was added drop wise to a solution of PhPCl₂ (1.2 ml, 8.31 mmol) in THF (10 ml) at -70 °C and stirred for 3 h followed by addition of benzylamine (1.8 ml, 16.62 mmol). After stirring the reaction mixture for 4 h at -70 °C under nitrogen atmosphere, 30% aq hydrogen peroxide (5 ml) was added at 0 °C and stirred at this temperature for an additional 1 hr. The reaction mixture was then allowed to warm up to room temperature, ethylacetate (20 ml) was added and the resulting solution was washed with water (3 x 20 ml). The ethylacetate layer was dried over Na₂SO₄, filtered, and evaporated. The residue product was purified by crystallization from a 1:2 mixture of DCM and hexane to afford the title compound in 55% yield as white crystals; mp 104–107 °C;

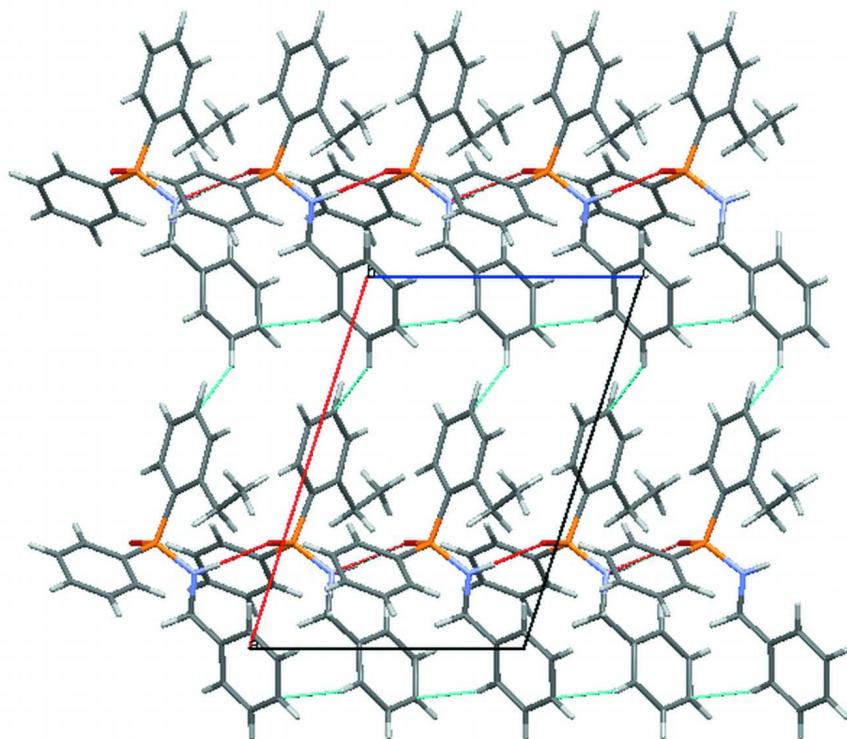
¹H NMR (CDCl₃, 400 MHz): δ 7.90–7.71 (m, 3H), 7.49–7.13 (m, 11H), 4.92–4.70 (m, NH peak), 4.22 (d, $J = 8.0$ Hz, 2H), 3.06 (q, $J = 7.4$ and 15.0 Hz, 2H), 1.10 (t, $J = 7.6$ Hz, 3H); ¹³C NMR (CDCl₃, 75 MHz): δ 149.2, 149.0, 139.7, 139.6, 133.1, 132.9, 132.2, 133.0, 131.8, 131.7, 130.2, 130.0, 128.7, 128.6, 128.4, 127.8, 127.3, 125.4, 125.3, 44.8, 27.1, 15.5; ³¹P NMR (CDCl₃, 400 MHz): δ 27.03.

S3. Refinement

Carbon-bound H-atoms were placed in calculated positions [C—H = 0.98 Å for Me H atoms, 0.99 Å for Methylene H atoms and 0.95 Å for aromatic H atoms; $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ (1.5 for Me groups)] and were included in the refinement in the riding model approximation. The nitrogen proton was located in a difference map and constrained with N—H = 0.88 Å ($U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$).

**Figure 1**

The asymmetric unit of the title compound with displacement ellipsoids drawn at the 50% probability level.

**Figure 2**

Packing diagram of the crystal of (I) showing chains of N—H···O intermolecular interactions as viewed down the crystallographic *c* direction.

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 $\beta = 107.578 (1)^\circ$
 $V = 1761.73 (7) \text{ \AA}^3$
 $Z = 4$

$F(000) = 712$
 $D_x = 1.264 \text{ Mg m}^{-3}$
Cu $K\alpha$ radiation, $\lambda = 1.54178 \text{ \AA}$
Cell parameters from 14165 reflections
 $\theta = 3.6\text{--}66.2^\circ$
 $\mu = 1.42 \text{ mm}^{-1}$
 $T = 100 \text{ K}$
Needle, colourless
 $0.48 \times 0.08 \times 0.02 \text{ mm}$

Data collection

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 φ and ω scans
Absorption correction: multi-scan
(SADABS; Bruker, 2008)
 $T_{\min} = 0.549$, $T_{\max} = 0.972$
13806 measured reflections

2946 independent reflections
2772 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.034$
 $\theta_{\max} = 66.2^\circ$, $\theta_{\min} = 3.6^\circ$
 $h = -15 \rightarrow 14$
 $k = -18 \rightarrow 18$
 $l = -9 \rightarrow 10$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.118$
 $S = 1.05$
2946 reflections
218 parameters
0 restraints
Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map
Hydrogen site location: inferred from
neighbouring sites
H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0573P)^2 + 1.6926P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.57 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.57 \text{ e \AA}^{-3}$

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.

>>> The Following Model ALERTS were generated - (Acta-Mode) <<< Format: alert-number_ALERT_alert-type_alert-level text 414_ALERT_2_C Short Intra D—H..H—X H1.. H6.. 1.98 Å ng. 414_ALERT_2_C Short Intra D—H..H—X H1.. H20A.. 1.96 Å ng. 911_ALERT_3_C Missing # FCF Refl Between THmin & STh/L= 0.594 147 793_ALERT_4_G The Model has Chirality at P1 (Verify) ... R 802_ALERT_4_G CIF Input Record(s) with more than 80 Characters ! 909_ALERT_3_G Percentage of Observed Data at Theta(Max) still 89 Perc. Noted:

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

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.98660 (15)	0.18270 (12)	0.8933 (2)	0.0197 (4)

C2	1.09281 (16)	0.21294 (13)	0.9436 (2)	0.0250 (4)
H2	1.1152	0.2556	0.8857	0.03*
C3	1.16615 (17)	0.18153 (14)	1.0770 (2)	0.0286 (5)
H3	1.2384	0.2026	1.1098	0.034*
C4	1.13436 (16)	0.11951 (13)	1.1627 (2)	0.0258 (5)
H4	1.1847	0.0976	1.2538	0.031*
C5	1.02898 (16)	0.08980 (13)	1.1148 (2)	0.0245 (4)
H5	1.0065	0.0477	1.1736	0.029*
C6	0.95550 (15)	0.12126 (12)	0.9808 (2)	0.0222 (4)
H6	0.8831	0.1004	0.9489	0.027*
C7	0.91011 (15)	0.21860 (13)	0.7457 (2)	0.0236 (4)
H7A	0.9099	0.2815	0.7531	0.028*
H7B	0.9373	0.2033	0.6585	0.028*
C8	0.75770 (15)	0.10376 (13)	0.4367 (2)	0.0215 (4)
C9	0.82914 (15)	0.04046 (13)	0.5135 (2)	0.0248 (4)
H9	0.8565	0.0413	0.6228	0.03*
C10	0.86062 (17)	-0.02365 (14)	0.4320 (3)	0.0325 (5)
H10	0.9095	-0.0665	0.4852	0.039*
C11	0.82036 (19)	-0.02514 (16)	0.2719 (3)	0.0382 (6)
H11	0.8413	-0.0692	0.2153	0.046*
C12	0.7494 (2)	0.03806 (16)	0.1952 (3)	0.0383 (6)
H12	0.7225	0.0372	0.0858	0.046*
C13	0.71770 (17)	0.10198 (14)	0.2759 (2)	0.0295 (5)
H13	0.6688	0.1447	0.2223	0.035*
C14	0.58144 (16)	0.17310 (13)	0.5493 (2)	0.0237 (4)
C15	0.50497 (17)	0.23235 (14)	0.4639 (2)	0.0291 (5)
H15	0.528	0.2767	0.41	0.035*
C16	0.39696 (17)	0.22687 (15)	0.4572 (3)	0.0338 (5)
H16	0.346	0.2675	0.4004	0.041*
C17	0.36414 (17)	0.16148 (15)	0.5343 (3)	0.0313 (5)
H17	0.2899	0.1569	0.5296	0.038*
C18	0.43787 (16)	0.10283 (14)	0.6179 (2)	0.0278 (5)
H18	0.4133	0.0584	0.6699	0.033*
C19	0.54819 (16)	0.10689 (13)	0.6284 (2)	0.0247 (4)
C20	0.62697 (16)	0.04282 (14)	0.7234 (2)	0.0288 (5)
H20A	0.6801	0.0735	0.8083	0.035*
H20B	0.6675	0.0175	0.6578	0.035*
C21	0.57827 (18)	-0.02982 (15)	0.7939 (3)	0.0341 (5)
H21A	0.5395	-0.0061	0.8619	0.051*
H21B	0.6366	-0.0673	0.8533	0.051*
H21C	0.5276	-0.0625	0.7113	0.051*
N1	0.79874 (12)	0.18732 (10)	0.71386 (18)	0.0206 (4)
H1	0.776	0.1667	0.7886	0.025*
O1	0.71769 (11)	0.27168 (9)	0.45065 (15)	0.0254 (3)
P1	0.71741 (4)	0.19123 (3)	0.53744 (5)	0.01919 (17)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0195 (9)	0.0215 (9)	0.0192 (10)	0.0013 (7)	0.0074 (8)	-0.0021 (7)
C2	0.0216 (10)	0.0272 (10)	0.0265 (10)	-0.0034 (8)	0.0078 (8)	0.0019 (8)
C3	0.0194 (10)	0.0337 (11)	0.0295 (11)	-0.0023 (8)	0.0024 (8)	-0.0006 (9)
C4	0.0238 (10)	0.0275 (10)	0.0227 (10)	0.0055 (8)	0.0022 (8)	0.0010 (8)
C5	0.0272 (10)	0.0225 (10)	0.0244 (10)	0.0022 (8)	0.0085 (8)	0.0032 (8)
C6	0.0201 (9)	0.0229 (10)	0.0233 (10)	-0.0007 (8)	0.0060 (8)	0.0000 (8)
C7	0.0185 (9)	0.0301 (10)	0.0221 (10)	-0.0039 (8)	0.0062 (8)	0.0042 (8)
C8	0.0194 (9)	0.0270 (10)	0.0203 (9)	-0.0050 (8)	0.0093 (7)	0.0004 (8)
C9	0.0204 (10)	0.0307 (11)	0.0256 (10)	-0.0017 (8)	0.0102 (8)	0.0004 (8)
C10	0.0240 (11)	0.0325 (12)	0.0458 (13)	-0.0015 (9)	0.0177 (10)	-0.0032 (10)
C11	0.0390 (13)	0.0396 (13)	0.0462 (14)	-0.0108 (10)	0.0282 (11)	-0.0169 (11)
C12	0.0442 (13)	0.0490 (14)	0.0256 (11)	-0.0133 (11)	0.0164 (10)	-0.0079 (10)
C13	0.0327 (11)	0.0356 (12)	0.0211 (10)	-0.0059 (9)	0.0094 (9)	0.0003 (9)
C14	0.0206 (10)	0.0307 (11)	0.0202 (10)	-0.0018 (8)	0.0070 (8)	-0.0044 (8)
C15	0.0276 (11)	0.0306 (11)	0.0296 (11)	0.0007 (9)	0.0097 (9)	0.0033 (9)
C16	0.0239 (11)	0.0370 (12)	0.0372 (12)	0.0061 (9)	0.0043 (9)	0.0032 (10)
C17	0.0210 (10)	0.0375 (12)	0.0354 (12)	-0.0020 (9)	0.0086 (9)	-0.0041 (10)
C18	0.0224 (10)	0.0323 (11)	0.0305 (11)	-0.0058 (9)	0.0106 (8)	-0.0067 (9)
C19	0.0231 (10)	0.0285 (11)	0.0222 (10)	-0.0022 (8)	0.0064 (8)	-0.0047 (8)
C20	0.0239 (10)	0.0356 (12)	0.0278 (11)	-0.0026 (9)	0.0091 (8)	0.0006 (9)
C21	0.0308 (11)	0.0370 (12)	0.0338 (12)	-0.0036 (9)	0.0086 (9)	0.0053 (10)
N1	0.0180 (8)	0.0269 (9)	0.0180 (8)	-0.0029 (6)	0.0069 (7)	0.0036 (6)
O1	0.0255 (7)	0.0291 (8)	0.0218 (7)	0.0018 (6)	0.0075 (6)	0.0047 (6)
P1	0.0162 (3)	0.0251 (3)	0.0166 (3)	-0.00091 (18)	0.00544 (19)	0.00211 (18)

Geometric parameters (\AA , $^\circ$)

C1—C6	1.386 (3)	C12—C13	1.378 (3)
C1—C2	1.393 (3)	C12—H12	0.95
C1—C7	1.516 (3)	C13—H13	0.95
C2—C3	1.386 (3)	C14—C19	1.404 (3)
C2—H2	0.95	C14—C15	1.409 (3)
C3—C4	1.386 (3)	C14—P1	1.816 (2)
C3—H3	0.95	C15—C16	1.382 (3)
C4—C5	1.380 (3)	C15—H15	0.95
C4—H4	0.95	C16—C17	1.381 (3)
C5—C6	1.391 (3)	C16—H16	0.95
C5—H5	0.95	C17—C18	1.378 (3)
C6—H6	0.95	C17—H17	0.95
C7—N1	1.465 (2)	C18—C19	1.402 (3)
C7—H7A	0.99	C18—H18	0.95
C7—H7B	0.99	C19—C20	1.505 (3)
C8—C9	1.393 (3)	C20—C21	1.534 (3)
C8—C13	1.398 (3)	C20—H20A	0.99
C8—P1	1.813 (2)	C20—H20B	0.99

C9—C10	1.383 (3)	C21—H21A	0.98
C9—H9	0.95	C21—H21B	0.98
C10—C11	1.392 (3)	C21—H21C	0.98
C10—H10	0.95	N1—P1	1.6332 (16)
C11—C12	1.388 (4)	N1—H1	0.88
C11—H11	0.95	O1—P1	1.4910 (14)
C6—C1—C2	118.48 (18)	C8—C13—H13	120.1
C6—C1—C7	122.93 (17)	C19—C14—C15	120.01 (18)
C2—C1—C7	118.59 (17)	C19—C14—P1	126.76 (16)
C3—C2—C1	120.79 (19)	C15—C14—P1	113.20 (15)
C3—C2—H2	119.6	C16—C15—C14	121.0 (2)
C1—C2—H2	119.6	C16—C15—H15	119.5
C4—C3—C2	120.18 (19)	C14—C15—H15	119.5
C4—C3—H3	119.9	C17—C16—C15	119.0 (2)
C2—C3—H3	119.9	C17—C16—H16	120.5
C5—C4—C3	119.48 (18)	C15—C16—H16	120.5
C5—C4—H4	120.3	C18—C17—C16	120.71 (19)
C3—C4—H4	120.3	C18—C17—H17	119.6
C4—C5—C6	120.29 (19)	C16—C17—H17	119.6
C4—C5—H5	119.9	C17—C18—C19	121.8 (2)
C6—C5—H5	119.9	C17—C18—H18	119.1
C1—C6—C5	120.77 (18)	C19—C18—H18	119.1
C1—C6—H6	119.6	C18—C19—C14	117.48 (19)
C5—C6—H6	119.6	C18—C19—C20	120.43 (19)
N1—C7—C1	112.92 (16)	C14—C19—C20	122.08 (18)
N1—C7—H7A	109	C19—C20—C21	116.47 (17)
C1—C7—H7A	109	C19—C20—H20A	108.2
N1—C7—H7B	109	C21—C20—H20A	108.2
C1—C7—H7B	109	C19—C20—H20B	108.2
H7A—C7—H7B	107.8	C21—C20—H20B	108.2
C9—C8—C13	119.37 (19)	H20A—C20—H20B	107.3
C9—C8—P1	122.39 (15)	C20—C21—H21A	109.5
C13—C8—P1	118.21 (16)	C20—C21—H21B	109.5
C10—C9—C8	120.60 (19)	H21A—C21—H21B	109.5
C10—C9—H9	119.7	C20—C21—H21C	109.5
C8—C9—H9	119.7	H21A—C21—H21C	109.5
C9—C10—C11	119.8 (2)	H21B—C21—H21C	109.5
C9—C10—H10	120.1	C7—N1—P1	118.96 (13)
C11—C10—H10	120.1	C7—N1—H1	120.5
C12—C11—C10	119.7 (2)	P1—N1—H1	120.5
C12—C11—H11	120.1	O1—P1—N1	116.67 (8)
C10—C11—H11	120.1	O1—P1—C8	109.12 (8)
C13—C12—C11	120.7 (2)	N1—P1—C8	105.70 (9)
C13—C12—H12	119.6	O1—P1—C14	108.77 (9)
C11—C12—H12	119.6	N1—P1—C14	106.44 (9)
C12—C13—C8	119.8 (2)	C8—P1—C14	110.01 (9)
C12—C13—H13	120.1		

C6—C1—C2—C3	0.8 (3)	C17—C18—C19—C20	178.74 (19)
C7—C1—C2—C3	-179.37 (19)	C15—C14—C19—C18	0.1 (3)
C1—C2—C3—C4	-0.2 (3)	P1—C14—C19—C18	-177.67 (15)
C2—C3—C4—C5	-0.5 (3)	C15—C14—C19—C20	-178.96 (19)
C3—C4—C5—C6	0.6 (3)	P1—C14—C19—C20	3.2 (3)
C2—C1—C6—C5	-0.7 (3)	C18—C19—C20—C21	5.3 (3)
C7—C1—C6—C5	179.48 (19)	C14—C19—C20—C21	-175.67 (19)
C4—C5—C6—C1	0.0 (3)	C1—C7—N1—P1	-158.63 (13)
C6—C1—C7—N1	5.2 (3)	C7—N1—P1—O1	-44.63 (17)
C2—C1—C7—N1	-174.62 (17)	C7—N1—P1—C8	76.83 (16)
C13—C8—C9—C10	0.1 (3)	C7—N1—P1—C14	-166.21 (15)
P1—C8—C9—C10	-177.86 (15)	C9—C8—P1—O1	137.49 (16)
C8—C9—C10—C11	-0.2 (3)	C13—C8—P1—O1	-40.47 (17)
C9—C10—C11—C12	0.4 (3)	C9—C8—P1—N1	11.27 (18)
C10—C11—C12—C13	-0.5 (3)	C13—C8—P1—N1	-166.69 (15)
C11—C12—C13—C8	0.4 (3)	C9—C8—P1—C14	-103.26 (17)
C9—C8—C13—C12	-0.2 (3)	C13—C8—P1—C14	78.79 (17)
P1—C8—C13—C12	177.84 (16)	C19—C14—P1—O1	-176.35 (17)
C19—C14—C15—C16	0.5 (3)	C15—C14—P1—O1	5.72 (18)
P1—C14—C15—C16	178.53 (17)	C19—C14—P1—N1	-49.9 (2)
C14—C15—C16—C17	-0.8 (3)	C15—C14—P1—N1	132.20 (15)
C15—C16—C17—C18	0.6 (3)	C19—C14—P1—C8	64.2 (2)
C16—C17—C18—C19	0.0 (3)	C15—C14—P1—C8	-113.75 (16)
C17—C18—C19—C14	-0.4 (3)		

Hydrogen-bond geometry (\AA , $^\circ$)

Cg2 and Cg3 are the centroids of the C8—C13 and C14—C19 rings, respectively.

$D\text{—H}\cdots A$	$D\text{—H}$	$H\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
N1—H1 \cdots O1 ⁱ	0.88	2.09	2.742 (2)	131
C18—H18 \cdots Cg2 ⁱⁱ	0.95	2.98	3.756 (2)	139
C21—H21C \cdots Cg3 ⁱⁱ	0.98	2.83	3.636 (3)	140

Symmetry codes: (i) $x, -y+1/2, z+1/2$; (ii) $-x+1, -y, -z+1$.