

Tetrakis(benzylamino)phosphonium chloride

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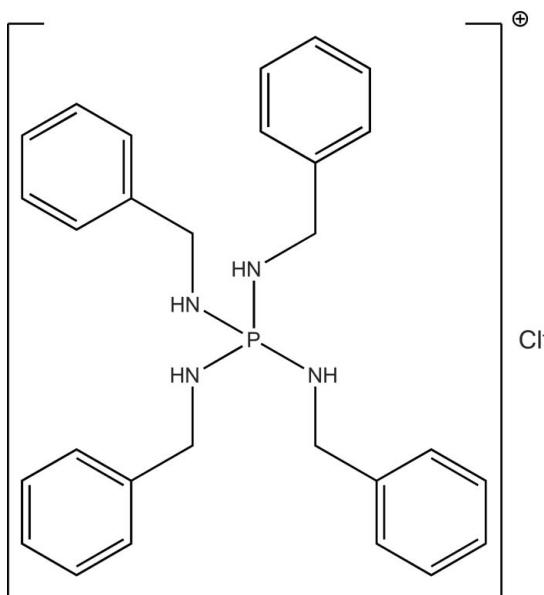
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Key indicators: single-crystal X-ray study; $T = 193\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.036; wR factor = 0.094; data-to-parameter ratio = 20.1.

The title salt, $[\text{P}(\text{NHCH}_2\text{C}_6\text{H}_5)_4]^+\cdot\text{Cl}^-$, crystallizes with the P atom and Cl^- anion lying on a twofold rotation axis. The P atom has a slightly distorted tetrahedral geometry with two classes of N–P–N angles [101.15 (10) and 100.55 (11) $^\circ$ and 113.07 (9) and 114.83 (8) $^\circ$] and the environments of sp^2 -hybridized N atoms are essentially planar (sum of angles = 359.9 and 360.1 $^\circ$). In the crystal, the phosphonium ion interacts with each neighboring chloride ion via two approximately equal N–H \cdots Cl interactions, forming parallel chains along the c axis.

Related literature

For background information on phosphonium salts, see: Hart & Sisler (1964); Levason *et al.* (2006); Schiemenz *et al.* (2003). For related structures, see: Bickley *et al.* (2004); Horstmann & Schnick (1996).



Experimental

Crystal data

$\text{C}_{28}\text{H}_{32}\text{N}_4\text{P}^+\cdot\text{Cl}^-$	$V = 1283.1 (6)\text{ \AA}^3$
$M_r = 491.00$	$Z = 2$
Orthorhombic, $P2_12_12$	Mo $K\alpha$ radiation
$a = 11.359 (3)\text{ \AA}$	$\mu = 0.24\text{ mm}^{-1}$
$b = 14.258 (4)\text{ \AA}$	$T = 193\text{ K}$
$c = 7.923 (3)\text{ \AA}$	$0.50 \times 0.40 \times 0.25\text{ mm}$

Data collection

Rebuilt Syntex P2 ₁ /Siemens P3 four-circle diffractometer	$R_{\text{int}} = 0.024$
6760 measured reflections	2 standard reflections every 98
3122 independent reflections	reflections
2677 reflections with $I > 2\sigma(I)$	intensity decay: 2%

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.036$	$\Delta\rho_{\text{max}} = 0.35\text{ e \AA}^{-3}$
$wR(F^2) = 0.094$	$\Delta\rho_{\text{min}} = -0.33\text{ e \AA}^{-3}$
$S = 1.01$	Absolute structure: Flack (1983),
3122 reflections	1321 Friedel pairs
155 parameters	Flack parameter: 0.08 (8)
	H-atom parameters constrained

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1–H1C \cdots Cl1 ⁱ	0.86	2.35	3.1848 (17)	163
N2–H2A \cdots Cl1	0.86	2.35	3.1855 (18)	165

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

Data collection: *P3/PC* (Siemens, 1989); cell refinement: *P3/PC*; data reduction: *P3/PC*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL* and *Mercury* (Macrae *et al.*, 2008); software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: NK2077).

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supporting information

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S1. Comment

Phosphonium salts, $[PR_4]^+$, are widely used as large cations to stabilize a variety of anionic species and to phase-transfer anions into low polarity organic media (Levason *et al.*, 2006). A few examples of tetraamino phosphonium salts exist in the literature (Hart & Sisler, 1964) and some of them are used as catalysis for preparing fluorine-containing compounds by a halogen/fluorine exchange reaction (Schiemenz *et al.*, 2003). Also, the crystal structure of $P(NH_2)_4Cl$ (Horstmann & Schnick, 1996) and $[P(NHPh)_4]Cl$ (Bickley *et al.*, 2004) have been reported. In an effort to further investigation into these types of compounds, the crystal structure of $[P(NHCH_2C_6H_5)_4]^+.Cl^-$ is presented.

The title salt, $[P(NHCH_2C_6H_5)_4]^+.Cl^-$, crystallizes with the P atom and Cl^- anion lying on a two-fold rotation axis in the space group $P2_{1}2_{1}2$. The supramolecular structure of compound exhibits a polymeric chain of alternating phosphonium and chloride ions in the solid state (Fig. 2) with $P\cdots P$ distances of 7.923 Å. The phosphonium ion interacts with each neighboring chloride ion *via* two equal N—H \cdots Cl interactions and six-membered rings around each phosphorus atom are formed. The centroid–centroid distance between adjacent phenyl rings is 4.009 Å, indicating no strong π – π stacking interactions exist in compound.

The four P—N bonds are of almost equal lengths 1.6166 (14) and 1.6184 (14) Å and are similar to those found in $[P(NHPh)_4]Cl$ (Bickley *et al.*, 2004) and $P(NH_2)_4Cl$ (Horstmann & Schnick, 1996). These P—N bonds are shorter than the typical P—N single bond length (1.77 Å). As observed in $[P(NHPh)_4]Cl$, there are two classes of N—P—N angles resulting in a distorted tetrahedral environment for P1. More acute angles of 101.15 (10) and 100.55° are observed within the hydrogen bonding chelates of the N1—P1—N1ⁱ and N2—P1—N2ⁱ (*i*: -*x*, -*y*, *z*) units respectively, whereas the remaining four N—P—N angles are 113.07 (9) and 114.83 (8)°. Although the main cause for the distortion from the ideal tetrahedral geometry is unclear, it seems to be partly controlled by hydrogen bonding. The sum of angles around nitrogen atoms are exactly 360° and these atoms are sp^2 hybridized.

S2. Experimental

All the reagents and solvents were used as obtained without further purification. Phosphorus pentachloride (2 mmol) was added drop wise with constant stirring to a toluene solution (30 ml) of benzylamine (8 mmol) at 0°C. After an hour stirring, the reaction mixture was refluxed for 3 h. The solid formed during the reaction was filtered and then washed with distilled water, toluene, chloroform and dried. Single crystals suitable for single-crystal X-ray diffraction analysis were obtained from the mixture of MeOH and CH₃CN solution.

S3. Refinement

The hydrogen atoms of NH groups were found in difference Fourier synthesis, the H(C) atom positions were calculated. All hydrogen atoms were refined in isotropic approximation in riding model with the $U_{iso}(H)$ parameters equal to 1.2 $U_{eq}(X)$, where $U(X)$ are the equivalent thermal parameters of the atoms to which corresponding H atoms are bonded.

The absolute structure parameter (Flack, 1983) is refined based on 1321 Friedel pairs.

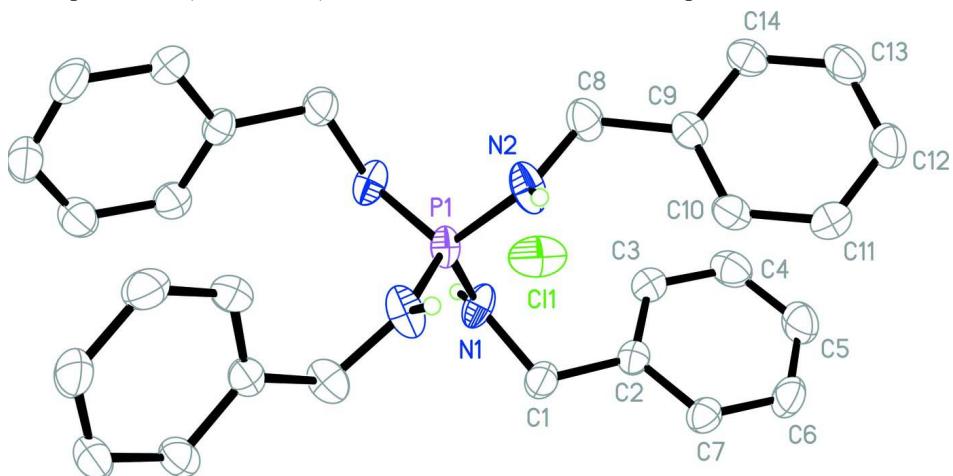


Figure 1

The structure of the title compound with 50% displacement ellipsoids and C-bound H atoms omitted. Unlabeled atoms are related to labeled atoms by a two-fold rotation.

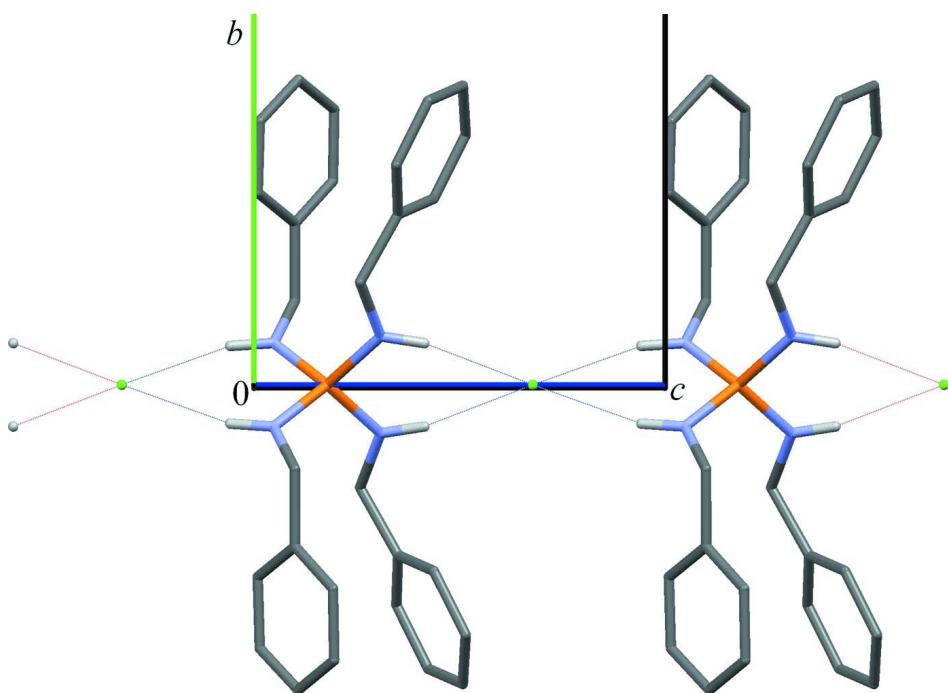


Figure 2

An *a*-axis projection showing hydrogen bonding dotted lines. The long *b* axis is truncated.

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Crystal data

$C_{28}H_{32}N_4P^+\cdot Cl^-$
 $M_r = 491.00$
Orthorhombic, $P2_12_12$
Hall symbol: P 2 2ab

$a = 11.359 (3) \text{ \AA}$
 $b = 14.258 (4) \text{ \AA}$
 $c = 7.923 (3) \text{ \AA}$
 $V = 1283.1 (6) \text{ \AA}^3$

$Z = 2$
 $F(000) = 520$
 $D_x = 1.271 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Cell parameters from 24 reflections

$\theta = 10\text{--}11^\circ$
 $\mu = 0.24 \text{ mm}^{-1}$
 $T = 193 \text{ K}$
Plate, colourless
 $0.50 \times 0.40 \times 0.25 \text{ mm}$

Data collection

Rebuilt Syntex P2₁/Siemens P3 four-circle diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 $\theta/2\theta$ scans
6760 measured reflections
3122 independent reflections
2677 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.024$
 $\theta_{\text{max}} = 28.1^\circ, \theta_{\text{min}} = 2.3^\circ$
 $h = -15 \rightarrow 15$
 $k = -18 \rightarrow 18$
 $l = -10 \rightarrow 10$
2 standard reflections every 98 reflections
intensity decay: 2%

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.094$
 $S = 1.01$
3122 reflections
155 parameters
0 restraints
Primary atom site location: structure-invariant direct methods

Secondary atom site location: difference Fourier map
Hydrogen site location: difference Fourier map
H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0604P)^2 + 0.1221P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.35 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.33 \text{ e \AA}^{-3}$
Absolute structure: Flack (1983), 1321 Friedel pairs
Absolute structure parameter: 0.08 (8)

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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C11	0.0000	0.0000	0.67845 (6)	0.03585 (15)
P1	0.0000	0.0000	0.17782 (6)	0.02407 (13)
N1	0.08364 (13)	0.05683 (9)	0.04823 (17)	0.0305 (3)
H1C	0.0684	0.0528	-0.0578	0.037*
N2	-0.07306 (14)	0.06507 (10)	0.30836 (17)	0.0347 (3)
H2A	-0.0623	0.0560	0.4146	0.042*
C1	0.18316 (15)	0.11449 (11)	0.0998 (2)	0.0327 (4)
H1A	0.2487	0.1016	0.0248	0.039*
H1B	0.2067	0.0958	0.2125	0.039*
C2	0.16072 (13)	0.21915 (10)	0.1000 (2)	0.0255 (3)

C3	0.06933 (14)	0.25994 (12)	0.0093 (2)	0.0309 (3)
H3A	0.0193	0.2226	-0.0548	0.037*
C4	0.05265 (16)	0.35639 (13)	0.0142 (3)	0.0396 (4)
H4A	-0.0090	0.3832	-0.0461	0.047*
C5	0.12664 (17)	0.41274 (12)	0.1078 (3)	0.0435 (5)
H5A	0.1142	0.4772	0.1120	0.052*
C6	0.21920 (16)	0.37315 (12)	0.1953 (3)	0.0394 (4)
H6A	0.2703	0.4111	0.2564	0.047*
C7	0.23623 (14)	0.27655 (11)	0.1922 (2)	0.0301 (3)
H7A	0.2984	0.2502	0.2520	0.036*
C8	-0.15598 (16)	0.13807 (12)	0.2581 (2)	0.0342 (4)
H8A	-0.2355	0.1160	0.2787	0.041*
H8B	-0.1480	0.1491	0.1379	0.041*
C9	-0.13829 (14)	0.22969 (11)	0.3506 (2)	0.0265 (3)
C10	-0.03555 (14)	0.25145 (12)	0.4376 (2)	0.0311 (3)
H10A	0.0248	0.2075	0.4437	0.037*
C11	-0.02213 (15)	0.33760 (13)	0.5152 (3)	0.0392 (4)
H11A	0.0472	0.3514	0.5725	0.047*
C12	-0.11172 (17)	0.40365 (12)	0.5081 (3)	0.0419 (4)
H12A	-0.1024	0.4617	0.5597	0.050*
C13	-0.21480 (16)	0.38267 (12)	0.4238 (3)	0.0374 (4)
H13A	-0.2752	0.4266	0.4191	0.045*
C14	-0.22821 (14)	0.29623 (12)	0.3465 (2)	0.0317 (3)
H14A	-0.2982	0.2824	0.2909	0.038*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.0282 (3)	0.0634 (4)	0.0159 (2)	-0.0003 (3)	0.000	0.000
P1	0.0382 (3)	0.0190 (2)	0.0151 (2)	0.0001 (2)	0.000	0.000
N1	0.0447 (8)	0.0280 (6)	0.0188 (6)	-0.0127 (6)	-0.0030 (6)	-0.0009 (5)
N2	0.0570 (9)	0.0301 (6)	0.0169 (6)	0.0133 (7)	0.0002 (6)	-0.0021 (5)
C1	0.0334 (8)	0.0262 (7)	0.0387 (9)	-0.0025 (6)	-0.0071 (7)	-0.0007 (7)
C2	0.0281 (7)	0.0251 (7)	0.0234 (7)	-0.0037 (6)	0.0038 (6)	-0.0003 (6)
C3	0.0296 (7)	0.0322 (8)	0.0309 (8)	-0.0019 (6)	-0.0011 (6)	-0.0003 (7)
C4	0.0343 (8)	0.0371 (9)	0.0473 (11)	0.0062 (7)	0.0032 (8)	0.0078 (8)
C5	0.0431 (10)	0.0251 (8)	0.0624 (13)	-0.0012 (7)	0.0105 (9)	0.0014 (8)
C6	0.0394 (9)	0.0304 (8)	0.0483 (11)	-0.0120 (7)	0.0038 (9)	-0.0070 (8)
C7	0.0288 (7)	0.0311 (8)	0.0305 (8)	-0.0046 (6)	0.0020 (7)	-0.0007 (7)
C8	0.0399 (9)	0.0340 (8)	0.0288 (8)	0.0069 (7)	-0.0046 (7)	-0.0056 (7)
C9	0.0297 (7)	0.0273 (7)	0.0225 (8)	0.0029 (6)	0.0037 (6)	0.0023 (6)
C10	0.0258 (7)	0.0320 (7)	0.0354 (8)	0.0049 (6)	0.0031 (7)	-0.0015 (7)
C11	0.0277 (8)	0.0379 (9)	0.0520 (11)	-0.0061 (7)	0.0008 (8)	-0.0068 (8)
C12	0.0424 (10)	0.0238 (8)	0.0595 (12)	-0.0031 (7)	0.0106 (9)	-0.0052 (8)
C13	0.0357 (9)	0.0284 (8)	0.0481 (11)	0.0093 (7)	0.0061 (8)	0.0055 (7)
C14	0.0299 (7)	0.0330 (8)	0.0323 (9)	0.0064 (6)	-0.0019 (7)	0.0041 (7)

Geometric parameters (\AA , \circ)

P1—N1 ⁱ	1.6166 (14)	C5—H5A	0.9300
P1—N1	1.6166 (14)	C6—C7	1.391 (2)
P1—N2 ⁱ	1.6184 (14)	C6—H6A	0.9300
P1—N2	1.6184 (14)	C7—H7A	0.9300
N1—C1	1.456 (2)	C8—C9	1.511 (2)
N1—H1C	0.8600	C8—H8A	0.9700
N2—C8	1.459 (2)	C8—H8B	0.9700
N2—H2A	0.8600	C9—C10	1.391 (2)
C1—C2	1.514 (2)	C9—C14	1.395 (2)
C1—H1A	0.9700	C10—C11	1.382 (3)
C1—H1B	0.9700	C10—H10A	0.9300
C2—C3	1.390 (2)	C11—C12	1.388 (3)
C2—C7	1.393 (2)	C11—H11A	0.9300
C3—C4	1.389 (3)	C12—C13	1.380 (3)
C3—H3A	0.9300	C12—H12A	0.9300
C4—C5	1.379 (3)	C13—C14	1.385 (2)
C4—H4A	0.9300	C13—H13A	0.9300
C5—C6	1.380 (3)	C14—H14A	0.9300
N1 ⁱ —P1—N1	101.15 (10)	C5—C6—C7	120.14 (17)
N1 ⁱ —P1—N2 ⁱ	114.83 (8)	C5—C6—H6A	119.9
N1—P1—N2 ⁱ	113.07 (9)	C7—C6—H6A	119.9
N1 ⁱ —P1—N2	113.07 (9)	C6—C7—C2	120.37 (17)
N1—P1—N2	114.83 (8)	C6—C7—H7A	119.8
N2 ⁱ —P1—N2	100.55 (11)	C2—C7—H7A	119.8
C1—N1—P1	124.13 (12)	N2—C8—C9	113.49 (14)
C1—N1—H1C	117.9	N2—C8—H8A	108.9
P1—N1—H1C	117.9	C9—C8—H8A	108.9
C8—N2—P1	124.45 (12)	N2—C8—H8B	108.9
C8—N2—H2A	117.8	C9—C8—H8B	108.9
P1—N2—H2A	117.8	H8A—C8—H8B	107.7
N1—C1—C2	115.22 (14)	C10—C9—C14	118.34 (15)
N1—C1—H1A	108.5	C10—C9—C8	123.02 (14)
C2—C1—H1A	108.5	C14—C9—C8	118.64 (15)
N1—C1—H1B	108.5	C11—C10—C9	120.76 (15)
C2—C1—H1B	108.5	C11—C10—H10A	119.6
H1A—C1—H1B	107.5	C9—C10—H10A	119.6
C3—C2—C7	119.02 (14)	C10—C11—C12	120.27 (17)
C3—C2—C1	122.52 (14)	C10—C11—H11A	119.9
C7—C2—C1	118.45 (15)	C12—C11—H11A	119.9
C4—C3—C2	120.11 (16)	C13—C12—C11	119.65 (16)
C4—C3—H3A	119.9	C13—C12—H12A	120.2
C2—C3—H3A	119.9	C11—C12—H12A	120.2
C5—C4—C3	120.59 (17)	C12—C13—C14	120.01 (16)
C5—C4—H4A	119.7	C12—C13—H13A	120.0
C3—C4—H4A	119.7	C14—C13—H13A	120.0

C4—C5—C6	119.74 (16)	C13—C14—C9	120.96 (16)
C4—C5—H5A	120.1	C13—C14—H14A	119.5
C6—C5—H5A	120.1	C9—C14—H14A	119.5
N1 ⁱ —P1—N1—C1	-174.86 (17)	C5—C6—C7—C2	0.5 (3)
N2 ⁱ —P1—N1—C1	-51.55 (15)	C3—C2—C7—C6	1.0 (2)
N2—P1—N1—C1	63.05 (15)	C1—C2—C7—C6	179.71 (17)
N1 ⁱ —P1—N2—C8	-56.69 (16)	P1—N2—C8—C9	-132.58 (14)
N1—P1—N2—C8	58.69 (17)	N2—C8—C9—C10	17.1 (2)
N2 ⁱ —P1—N2—C8	-179.61 (19)	N2—C8—C9—C14	-163.97 (15)
P1—N1—C1—C2	-101.97 (17)	C14—C9—C10—C11	-1.2 (3)
N1—C1—C2—C3	-19.8 (2)	C8—C9—C10—C11	177.70 (17)
N1—C1—C2—C7	161.45 (15)	C9—C10—C11—C12	0.4 (3)
C7—C2—C3—C4	-1.4 (2)	C10—C11—C12—C13	0.4 (3)
C1—C2—C3—C4	179.90 (17)	C11—C12—C13—C14	-0.3 (3)
C2—C3—C4—C5	0.4 (3)	C12—C13—C14—C9	-0.6 (3)
C3—C4—C5—C6	1.0 (3)	C10—C9—C14—C13	1.4 (2)
C4—C5—C6—C7	-1.5 (3)	C8—C9—C14—C13	-177.61 (16)

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

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

$D\text{—H}\cdots A$	$D\text{—H}$	$H\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
N1—H1C \cdots C11 ⁱⁱ	0.86	2.35	3.1848 (17)	163
N2—H2A \cdots C11	0.86	2.35	3.1855 (18)	165

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