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Diethylammonium anilino(methoxy)-phosphinate

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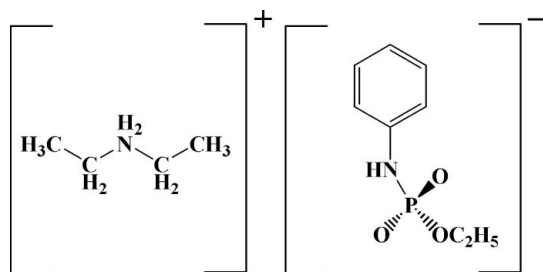
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 Key indicators: single-crystal X-ray study; $T = 173$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.039; wR factor = 0.081; data-to-parameter ratio = 14.9.

The title compound, $[\text{Et}_2\text{NH}_2][(\text{EtO})\text{PO}_2(\text{C}_6\text{H}_5\text{NH})]$ or $\text{C}_4\text{H}_{12}\text{N}^+\cdot\text{C}_8\text{H}_{11}\text{NO}_3\text{P}^-$, is a molecular salt with two anions containing PO_3N groupings and two cations in the asymmetric unit. A network of $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds link the cations and anions into a two-dimensional network.

Related literature

For the use of N -substituted phosphoramidic acids in the synthesis of pyrophosphate groups, see: Quin & Jankowski (1994). For a corresponding dimer complex with similar $\text{P}-\text{O}$ and $\text{P}-\text{N}$ connections, see: Andrianov et al. (1977).



Experimental

Crystal data

 $\text{C}_4\text{H}_{12}\text{N}^+\cdot\text{C}_8\text{H}_{11}\text{NO}_3\text{P}^-$
 $M_r = 274.29$
 Orthorhombic, $Pna2_1$
 $a = 14.341$ (3) Å
 $b = 12.785$ (2) Å
 $c = 15.997$ (3) Å

 $V = 2933.0$ (9) Å³
 $Z = 8$
 Mo $K\alpha$ radiation
 $\mu = 0.19$ mm⁻¹
 $T = 173$ (2) K
 $0.40 \times 0.20 \times 0.08$ mm

Data collection

 Bruker SMART CCD diffractometer
 Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.957$, $T_{\max} = 0.989$

 5062 measured reflections
 4847 independent reflections
 4043 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.017$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.039$
 $wR(F^2) = 0.081$
 $S = 1.03$
 4847 reflections
 325 parameters
 1 restraint

 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.16$ e Å⁻³
 $\Delta\rho_{\min} = -0.23$ e Å⁻³
 Absolute structure: Flack (1983),
 2649 Friedel pairs
 Flack parameter: 0.06 (4)

Table 1

Selected bond lengths (Å).

N1—P1	1.670 (2)	N2—P2	1.651 (2)
P1—O1	1.4798 (19)	P2—O5	1.4897 (18)
P1—O3	1.4994 (17)	P2—O4	1.4983 (19)
P1—O2	1.5979 (18)	P2—O6	1.5969 (17)

Table 2

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1B \cdots O4	0.86	2.13	2.954 (3)	161
N2—H2B \cdots O1	0.86	2.04	2.895 (3)	173
N3—H3C \cdots O4	0.90	1.88	2.742 (3)	160
N4—H4A \cdots O3	0.90	1.94	2.792 (3)	158
N3—H3B \cdots O3 ⁱ	0.90	1.89	2.788 (3)	174
N4—H4B \cdots O5 ⁱⁱ	0.90	1.74	2.637 (3)	172

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

Data collection: *SMART* (Bruker, 1996); cell refinement: *SAINTE* (Bruker, 1996); data reduction: *SAINTE*; 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*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: EZZ138).

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supporting information

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Diethylammonium anilino(methoxy)phosphinate

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S1. Comment

N-substituted phosphoramidic acids are useful agents in the synthesis of pyrophosphate groups (Quin & Jankowski, 1994). However the structures of these kinds of materials have not been well characterized. By the use of the O,*N*-substituted phosphoramidic acids as ligands in the preparation of new metal phosphate frameworks, we obtained the title compound as a salt of the O,*N*-substituted phosphoramidic acid.

As shown in Fig. 1, the asymmetric unit of the title compound is composed of two diethyl-amine cations, and two *N*-ethoxyphosphoryl-phenylamide anions. The geometrical parameters of the independent anions are similar. The phosphorus atoms have tetrahedral coordination geometries. The shortest P—O bond lengths correspond to the P=O double bonds. The longest P—O distances are due to the influence of the —OEt group (Andrianov *et al.*, 1977). The O—P—O and O—P—N bond angles range from 102.961 (98)–119.158 (108)° and 105.126 (113)–112.197 (117)°, indicating that the geometries of the tetrahedra are slightly distorted.

Hydrogen bonds exist between the diethyl-amine cations and the *N*-ethoxyphosphoryl-phenylamide anions. The N—H···O connections result a two dimensional packing motif (Fig. 2).

S2. Experimental

A solution of aniline (12.5 mmol) and 12.5 mmol of Et₂NH in 15 ml of ether was added to a solution of 12.5 mmol of ethyl phosphorodichloridate in 15 ml of ether. After 20 h, the solution was filtered and the filtrate was evaporated to give a powder. The powder was dissolved in 30 ml of an acetone-water mixture (1:1) containing 1 g of NaOH. After 10 min, the solvent was evaporated and the residue dried *in vacuo*. Recrystallization of the precipitate from a chloroform solution yielded crystals of the title compound.

S3. Refinement

H atoms were positioned geometrically and refined using a riding model, with N—H = 0.86–0.90 Å, C—H = 0.93–0.97 Å and with $U_{\text{iso}}(\text{H}) = 1.2$ (1.5 for methyl groups) times $U_{\text{eq}}(\text{C})$. >.

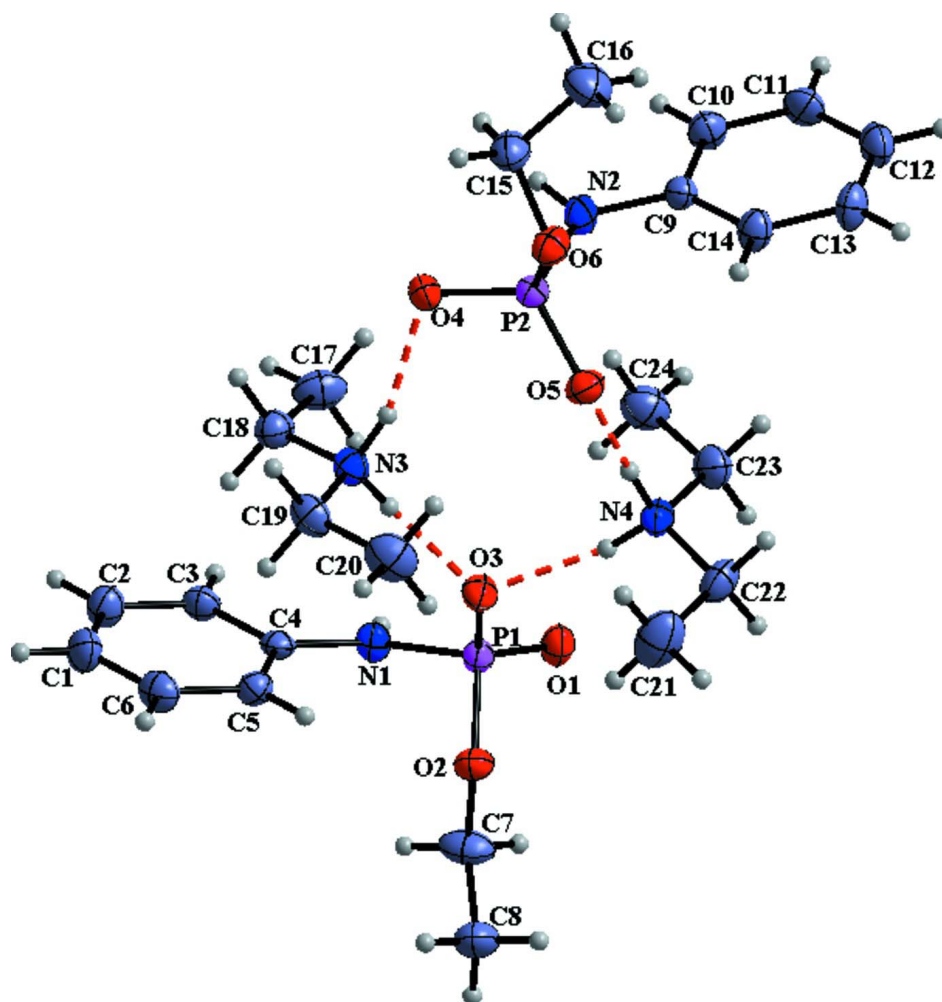
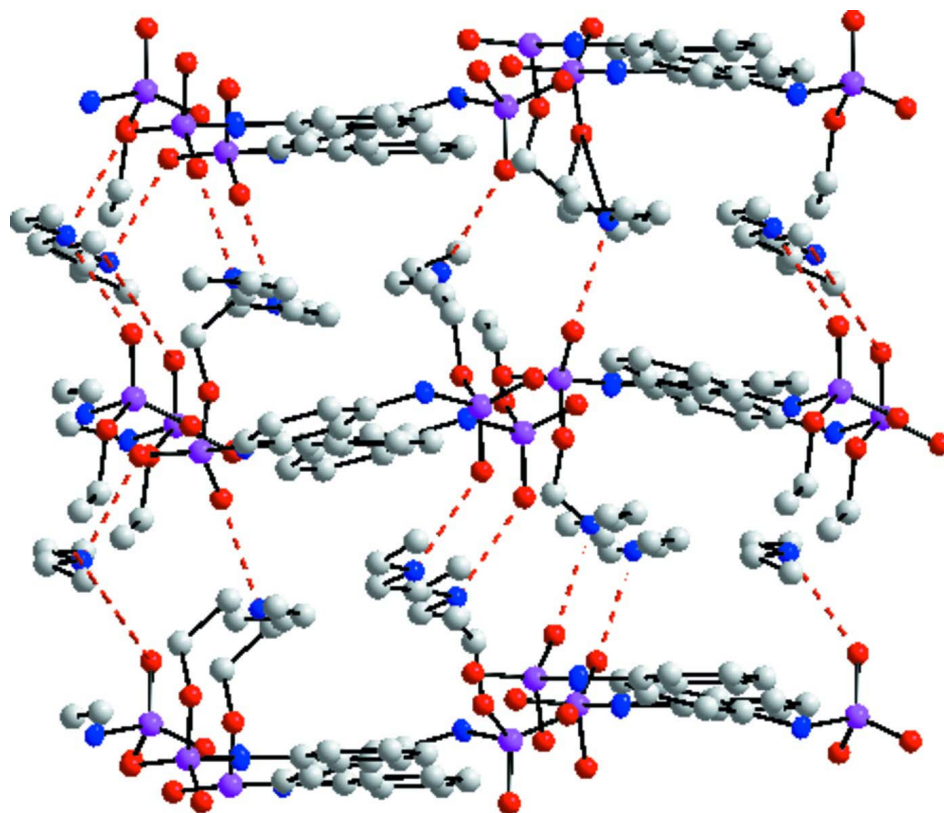


Figure 1

The molecular structure of (I), with atom labels and 50% probability displacement ellipsoids for non-H atoms. The red dashed lines indicate the hydrogen bonds.

**Figure 2**

The packing of (I), viewed down the b axis, showing the N—H \cdots O hydrogen bonds between the diethylamine cations and the N -ethoxy phosphoryl-phenylamide anions. H-atoms have been omitted for clarity.

Diethylammonium anilino(methoxy)phosphinate

Crystal data

$C_4H_{12}N^+ \cdot C_8H_{11}NO_3P^-$

$M_r = 274.29$

Orthorhombic, $Pna2_1$

Hall symbol: P2c-2n

$a = 14.341(3) \text{ \AA}$

$b = 12.785(2) \text{ \AA}$

$c = 15.997(3) \text{ \AA}$

$V = 2933.0(9) \text{ \AA}^3$

$Z = 8$

$F(000) = 1184$

$D_x = 1.242 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 4847 reflections

$\theta = 3.3\text{--}25.0^\circ$

$\mu = 0.19 \text{ mm}^{-1}$

$T = 173 \text{ K}$

Needle-like, colorless

$0.4 \times 0.2 \times 0.08 \text{ mm}$

Data collection

Bruker SMART CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 1996)

$T_{\min} = 0.957$, $T_{\max} = 0.989$

5062 measured reflections

4847 independent reflections

4043 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.002$

$\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 3.3^\circ$

$h = -17 \rightarrow 17$

$k = -15 \rightarrow 15$

$l = -19 \rightarrow 18$

*Refinement*Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.039$ $wR(F^2) = 0.081$ $S = 1.03$

4847 reflections

325 parameters

1 restraint

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.0363P)^2 + 0.5497P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} = 0.001$ $\Delta\rho_{\max} = 0.16 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{\min} = -0.23 \text{ e } \text{\AA}^{-3}$ Absolute structure: Flack (1983), 2649 Friedel
pairs

Absolute structure parameter: 0.06 (4)

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)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	0.50725 (13)	0.16964 (16)	0.52519 (13)	0.0230 (5)
H1B	0.5374	0.2242	0.5414	0.028*
N4	0.30305 (14)	-0.01440 (16)	0.74774 (12)	0.0265 (5)
H4A	0.3436	-0.0013	0.7061	0.032*
H4B	0.2451	-0.0069	0.7270	0.032*
C1	0.4316 (2)	0.1694 (2)	0.27143 (19)	0.0359 (7)
H1A	0.4151	0.1696	0.2152	0.043*
C2	0.45479 (18)	0.2624 (2)	0.31172 (18)	0.0336 (7)
H2A	0.4541	0.3251	0.2823	0.040*
C3	0.47881 (17)	0.2616 (2)	0.39546 (16)	0.0269 (6)
H3A	0.4937	0.3242	0.4219	0.032*
C4	0.48107 (16)	0.1682 (2)	0.44095 (16)	0.0220 (6)
C5	0.45765 (17)	0.0754 (2)	0.39961 (16)	0.0242 (6)
H5A	0.4583	0.0122	0.4285	0.029*
C6	0.43349 (18)	0.0772 (2)	0.31564 (18)	0.0315 (7)
H6A	0.4183	0.0150	0.2888	0.038*
C7	0.63999 (18)	-0.0209 (2)	0.5543 (2)	0.0407 (7)
H7A	0.6703	0.0043	0.6046	0.049*
H7B	0.6556	0.0261	0.5087	0.049*
C8	0.6729 (2)	-0.1300 (2)	0.53435 (19)	0.0415 (7)
H8A	0.7393	-0.1297	0.5266	0.062*
H8B	0.6432	-0.1540	0.4841	0.062*
H8C	0.6572	-0.1760	0.5797	0.062*

C21	0.3111 (3)	-0.1992 (2)	0.7038 (2)	0.0540 (9)
H21A	0.3195	-0.2693	0.7240	0.081*
H21B	0.3596	-0.1828	0.6645	0.081*
H21C	0.2516	-0.1935	0.6769	0.081*
C22	0.3154 (2)	-0.1243 (2)	0.77572 (17)	0.0351 (7)
H22A	0.2671	-0.1417	0.8157	0.042*
H22B	0.3752	-0.1312	0.8036	0.042*
C23	0.3172 (2)	0.0639 (2)	0.81480 (18)	0.0359 (7)
H23A	0.3785	0.0545	0.8391	0.043*
H23B	0.2713	0.0529	0.8585	0.043*
C24	0.3086 (2)	0.1737 (2)	0.7821 (2)	0.0434 (8)
H24A	0.3185	0.2224	0.8269	0.065*
H24B	0.2474	0.1838	0.7591	0.065*
H24C	0.3544	0.1851	0.7393	0.065*
P1	0.48758 (4)	0.07915 (5)	0.59834 (4)	0.02230 (16)
O1	0.52914 (12)	0.12003 (13)	0.67646 (11)	0.0297 (4)
O2	0.54000 (11)	-0.02352 (13)	0.56616 (11)	0.0259 (4)
O3	0.38708 (11)	0.04682 (12)	0.59813 (12)	0.0272 (4)
N2	0.55256 (14)	0.33041 (17)	0.73990 (14)	0.0283 (5)
H2B	0.5502	0.2687	0.7187	0.034*
N3	0.72854 (14)	0.40960 (16)	0.50343 (13)	0.0267 (5)
H3B	0.7820	0.4225	0.5310	0.032*
H3C	0.6826	0.4046	0.5416	0.032*
C9	0.55185 (17)	0.3353 (2)	0.82782 (16)	0.0233 (6)
C10	0.55401 (18)	0.2428 (2)	0.87430 (17)	0.0302 (7)
H10A	0.5560	0.1789	0.8466	0.036*
C11	0.55331 (19)	0.2445 (2)	0.96026 (18)	0.0343 (7)
H11A	0.5552	0.1820	0.9899	0.041*
C12	0.54987 (18)	0.3381 (2)	1.00280 (18)	0.0346 (7)
H12A	0.5492	0.3391	1.0609	0.042*
C13	0.5474 (2)	0.4306 (2)	0.95808 (18)	0.0336 (7)
H13A	0.5451	0.4941	0.9863	0.040*
C14	0.54840 (19)	0.4294 (2)	0.87124 (17)	0.0298 (7)
H14A	0.5467	0.4922	0.8419	0.036*
C15	0.37515 (18)	0.4351 (2)	0.66248 (18)	0.0357 (7)
H15A	0.3879	0.3613	0.6542	0.043*
H15B	0.3479	0.4624	0.6115	0.043*
C16	0.3085 (2)	0.4486 (3)	0.7328 (2)	0.0479 (8)
H16A	0.2516	0.4121	0.7205	0.072*
H16B	0.2955	0.5216	0.7404	0.072*
H16C	0.3353	0.4207	0.7831	0.072*
C17	0.7673 (2)	0.2234 (2)	0.5187 (2)	0.0412 (7)
H17A	0.7725	0.1585	0.4889	0.062*
H17B	0.7220	0.2164	0.5625	0.062*
H17C	0.8267	0.2413	0.5424	0.062*
C18	0.73711 (19)	0.3085 (2)	0.45931 (17)	0.0315 (6)
H18A	0.7824	0.3149	0.4146	0.038*
H18B	0.6775	0.2900	0.4347	0.038*

C19	0.70839 (19)	0.4987 (2)	0.44723 (17)	0.0326 (6)
H19A	0.6487	0.4876	0.4201	0.039*
H19B	0.7559	0.5023	0.4041	0.039*
C20	0.7060 (2)	0.6010 (2)	0.4943 (2)	0.0475 (8)
H20A	0.6928	0.6570	0.4562	0.071*
H20B	0.7653	0.6128	0.5204	0.071*
H20C	0.6583	0.5981	0.5364	0.071*
P2	0.55732 (4)	0.42782 (5)	0.67233 (4)	0.02279 (16)
O4	0.56495 (11)	0.37700 (13)	0.58821 (11)	0.0282 (4)
O5	0.62869 (12)	0.50728 (13)	0.69568 (10)	0.0306 (4)
O6	0.46099 (11)	0.48965 (13)	0.68041 (12)	0.0294 (4)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0269 (11)	0.0204 (12)	0.0219 (11)	−0.0046 (9)	−0.0028 (9)	−0.0019 (9)
N4	0.0218 (12)	0.0335 (13)	0.0241 (11)	0.0008 (9)	−0.0012 (9)	0.0045 (10)
C1	0.0436 (17)	0.0395 (18)	0.0247 (15)	0.0050 (14)	−0.0059 (13)	−0.0015 (14)
C2	0.0404 (16)	0.0300 (16)	0.0304 (16)	0.0006 (13)	−0.0034 (12)	0.0025 (14)
C3	0.0305 (14)	0.0226 (15)	0.0276 (16)	0.0018 (12)	0.0013 (11)	−0.0045 (12)
C4	0.0181 (13)	0.0249 (15)	0.0231 (15)	0.0032 (11)	0.0024 (10)	−0.0017 (11)
C5	0.0255 (14)	0.0204 (14)	0.0268 (16)	−0.0006 (11)	−0.0014 (11)	−0.0014 (12)
C6	0.0319 (16)	0.0342 (16)	0.0284 (16)	−0.0013 (12)	−0.0040 (12)	−0.0077 (14)
C7	0.0223 (15)	0.0437 (18)	0.0561 (19)	0.0006 (13)	0.0027 (13)	−0.0137 (15)
C8	0.0409 (17)	0.0400 (18)	0.0435 (18)	0.0134 (14)	0.0066 (14)	−0.0004 (14)
C21	0.073 (2)	0.0354 (18)	0.054 (2)	−0.0014 (17)	−0.0147 (17)	0.0019 (16)
C22	0.0366 (16)	0.0322 (16)	0.0365 (17)	0.0011 (13)	−0.0049 (13)	0.0116 (14)
C23	0.0386 (17)	0.0401 (17)	0.0291 (15)	0.0011 (13)	−0.0048 (13)	−0.0042 (14)
C24	0.0464 (18)	0.0368 (17)	0.0470 (19)	−0.0006 (14)	0.0049 (15)	−0.0052 (14)
P1	0.0261 (4)	0.0211 (3)	0.0197 (3)	0.0005 (3)	−0.0011 (3)	0.0006 (3)
O1	0.0411 (11)	0.0250 (9)	0.0230 (10)	0.0006 (8)	−0.0046 (9)	0.0006 (9)
O2	0.0239 (10)	0.0239 (10)	0.0298 (11)	0.0019 (7)	0.0000 (7)	−0.0027 (8)
O3	0.0225 (9)	0.0320 (10)	0.0272 (10)	−0.0021 (7)	0.0011 (9)	0.0028 (9)
N2	0.0400 (13)	0.0201 (12)	0.0247 (13)	−0.0001 (10)	0.0015 (10)	−0.0041 (9)
N3	0.0250 (12)	0.0307 (13)	0.0245 (12)	0.0010 (9)	−0.0009 (9)	0.0021 (10)
C9	0.0230 (14)	0.0255 (15)	0.0213 (15)	−0.0023 (11)	−0.0003 (11)	−0.0002 (12)
C10	0.0362 (16)	0.0238 (15)	0.0306 (17)	−0.0010 (12)	0.0000 (12)	−0.0010 (13)
C11	0.0428 (18)	0.0298 (16)	0.0304 (17)	−0.0056 (14)	−0.0008 (12)	0.0090 (13)
C12	0.0383 (17)	0.0454 (19)	0.0202 (16)	−0.0005 (13)	−0.0033 (12)	0.0007 (13)
C13	0.0422 (18)	0.0308 (17)	0.0279 (16)	0.0054 (13)	−0.0027 (12)	−0.0089 (13)
C14	0.0374 (16)	0.0256 (15)	0.0264 (16)	0.0020 (12)	−0.0050 (12)	0.0008 (12)
C15	0.0281 (14)	0.0456 (17)	0.0334 (16)	−0.0015 (13)	−0.0007 (14)	−0.0054 (14)
C16	0.0383 (18)	0.060 (2)	0.045 (2)	−0.0006 (16)	0.0093 (15)	−0.0028 (16)
C17	0.0384 (16)	0.0291 (16)	0.056 (2)	−0.0014 (13)	0.0010 (14)	0.0036 (14)
C18	0.0276 (15)	0.0327 (16)	0.0341 (16)	−0.0017 (12)	0.0008 (11)	−0.0017 (13)
C19	0.0342 (16)	0.0357 (17)	0.0277 (14)	0.0065 (13)	0.0007 (11)	0.0078 (13)
C20	0.054 (2)	0.0384 (18)	0.050 (2)	0.0042 (15)	0.0052 (16)	0.0090 (15)
P2	0.0246 (3)	0.0225 (3)	0.0213 (3)	−0.0017 (3)	0.0002 (3)	−0.0004 (3)

O4	0.0334 (10)	0.0304 (10)	0.0208 (10)	-0.0043 (8)	0.0031 (8)	-0.0011 (9)
O5	0.0292 (10)	0.0307 (10)	0.0320 (11)	-0.0062 (8)	-0.0044 (8)	0.0011 (8)
O6	0.0264 (9)	0.0299 (10)	0.0319 (10)	0.0012 (8)	-0.0018 (8)	-0.0045 (9)

Geometric parameters (Å, °)

N1—C4	1.399 (3)	N2—C9	1.408 (3)
N1—P1	1.670 (2)	N2—P2	1.651 (2)
N1—H1B	0.8600	N2—H2B	0.8600
N4—C23	1.481 (3)	N3—C18	1.478 (3)
N4—C22	1.485 (3)	N3—C19	1.480 (3)
N4—H4A	0.9000	N3—H3B	0.9000
N4—H4B	0.9000	N3—H3C	0.9000
C1—C6	1.374 (4)	C9—C14	1.391 (4)
C1—C2	1.393 (4)	C9—C10	1.397 (4)
C1—H1A	0.9300	C10—C11	1.375 (4)
C2—C3	1.383 (4)	C10—H10A	0.9300
C2—H2A	0.9300	C11—C12	1.378 (4)
C3—C4	1.399 (4)	C11—H11A	0.9300
C3—H3A	0.9300	C12—C13	1.382 (4)
C4—C5	1.399 (3)	C12—H12A	0.9300
C5—C6	1.387 (4)	C13—C14	1.389 (4)
C5—H5A	0.9300	C13—H13A	0.9300
C6—H6A	0.9300	C14—H14A	0.9300
C7—O2	1.447 (3)	C15—O6	1.444 (3)
C7—C8	1.507 (4)	C15—C16	1.487 (4)
C7—H7A	0.9700	C15—H15A	0.9700
C7—H7B	0.9700	C15—H15B	0.9700
C8—H8A	0.9600	C16—H16A	0.9600
C8—H8B	0.9600	C16—H16B	0.9600
C8—H8C	0.9600	C16—H16C	0.9600
C21—C22	1.498 (4)	C17—C18	1.507 (4)
C21—H21A	0.9600	C17—H17A	0.9600
C21—H21B	0.9600	C17—H17B	0.9600
C21—H21C	0.9600	C17—H17C	0.9600
C22—H22A	0.9700	C18—H18A	0.9700
C22—H22B	0.9700	C18—H18B	0.9700
C23—C24	1.504 (4)	C19—C20	1.509 (4)
C23—H23A	0.9700	C19—H19A	0.9700
C23—H23B	0.9700	C19—H19B	0.9700
C24—H24A	0.9600	C20—H20A	0.9600
C24—H24B	0.9600	C20—H20B	0.9600
C24—H24C	0.9600	C20—H20C	0.9600
P1—O1	1.4798 (19)	P2—O5	1.4897 (18)
P1—O3	1.4994 (17)	P2—O4	1.4983 (19)
P1—O2	1.5979 (18)	P2—O6	1.5969 (17)
C4—N1—P1	128.36 (18)	C9—N2—P2	128.40 (19)

C4—N1—H1B	115.8	C9—N2—H2B	115.8
P1—N1—H1B	115.8	P2—N2—H2B	115.8
C23—N4—C22	113.87 (19)	C18—N3—C19	113.6 (2)
C23—N4—H4A	108.8	C18—N3—H3B	108.9
C22—N4—H4A	108.8	C19—N3—H3B	108.9
C23—N4—H4B	108.8	C18—N3—H3C	108.9
C22—N4—H4B	108.8	C19—N3—H3C	108.9
H4A—N4—H4B	107.7	H3B—N3—H3C	107.7
C6—C1—C2	119.3 (3)	C14—C9—C10	117.9 (2)
C6—C1—H1A	120.4	C14—C9—N2	122.5 (2)
C2—C1—H1A	120.4	C10—C9—N2	119.6 (2)
C3—C2—C1	120.1 (3)	C11—C10—C9	121.2 (3)
C3—C2—H2A	120.0	C11—C10—H10A	119.4
C1—C2—H2A	120.0	C9—C10—H10A	119.4
C2—C3—C4	121.1 (3)	C10—C11—C12	120.6 (3)
C2—C3—H3A	119.5	C10—C11—H11A	119.7
C4—C3—H3A	119.5	C12—C11—H11A	119.7
C3—C4—N1	119.7 (2)	C11—C12—C13	119.2 (3)
C3—C4—C5	118.2 (2)	C11—C12—H12A	120.4
N1—C4—C5	122.1 (2)	C13—C12—H12A	120.4
C6—C5—C4	120.2 (2)	C12—C13—C14	120.6 (3)
C6—C5—H5A	119.9	C12—C13—H13A	119.7
C4—C5—H5A	119.9	C14—C13—H13A	119.7
C1—C6—C5	121.2 (3)	C13—C14—C9	120.6 (3)
C1—C6—H6A	119.4	C13—C14—H14A	119.7
C5—C6—H6A	119.4	C9—C14—H14A	119.7
O2—C7—C8	108.4 (2)	O6—C15—C16	110.0 (2)
O2—C7—H7A	110.0	O6—C15—H15A	109.7
C8—C7—H7A	110.0	C16—C15—H15A	109.7
O2—C7—H7B	110.0	O6—C15—H15B	109.7
C8—C7—H7B	110.0	C16—C15—H15B	109.7
H7A—C7—H7B	108.4	H15A—C15—H15B	108.2
C7—C8—H8A	109.5	C15—C16—H16A	109.5
C7—C8—H8B	109.5	C15—C16—H16B	109.5
H8A—C8—H8B	109.5	H16A—C16—H16B	109.5
C7—C8—H8C	109.5	C15—C16—H16C	109.5
H8A—C8—H8C	109.5	H16A—C16—H16C	109.5
H8B—C8—H8C	109.5	H16B—C16—H16C	109.5
C22—C21—H21A	109.5	C18—C17—H17A	109.5
C22—C21—H21B	109.5	C18—C17—H17B	109.5
H21A—C21—H21B	109.5	H17A—C17—H17B	109.5
C22—C21—H21C	109.5	C18—C17—H17C	109.5
H21A—C21—H21C	109.5	H17A—C17—H17C	109.5
H21B—C21—H21C	109.5	H17B—C17—H17C	109.5
N4—C22—C21	111.6 (2)	N3—C18—C17	110.7 (2)
N4—C22—H22A	109.3	N3—C18—H18A	109.5
C21—C22—H22A	109.3	C17—C18—H18A	109.5
N4—C22—H22B	109.3	N3—C18—H18B	109.5

C21—C22—H22B	109.3	C17—C18—H18B	109.5
H22A—C22—H22B	108.0	H18A—C18—H18B	108.1
N4—C23—C24	111.6 (2)	N3—C19—C20	111.6 (2)
N4—C23—H23A	109.3	N3—C19—H19A	109.3
C24—C23—H23A	109.3	C20—C19—H19A	109.3
N4—C23—H23B	109.3	N3—C19—H19B	109.3
C24—C23—H23B	109.3	C20—C19—H19B	109.3
H23A—C23—H23B	108.0	H19A—C19—H19B	108.0
C23—C24—H24A	109.5	C19—C20—H20A	109.5
C23—C24—H24B	109.5	C19—C20—H20B	109.5
H24A—C24—H24B	109.5	H20A—C20—H20B	109.5
C23—C24—H24C	109.5	C19—C20—H20C	109.5
H24A—C24—H24C	109.5	H20A—C20—H20C	109.5
H24B—C24—H24C	109.5	H20B—C20—H20C	109.5
O1—P1—O3	119.11 (11)	O5—P2—O4	118.09 (10)
O1—P1—O2	111.87 (10)	O5—P2—O6	103.68 (10)
O3—P1—O2	103.01 (9)	O4—P2—O6	110.52 (10)
O1—P1—N1	106.21 (10)	O5—P2—N2	112.25 (11)
O3—P1—N1	110.60 (10)	O4—P2—N2	105.31 (11)
O2—P1—N1	105.30 (10)	O6—P2—N2	106.54 (10)
C7—O2—P1	119.30 (16)	C15—O6—P2	118.84 (16)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1 <i>B</i> ...O4	0.86	2.13	2.954 (3)	161
N2—H2 <i>B</i> ...O1	0.86	2.04	2.895 (3)	173
N3—H3 <i>C</i> ...O4	0.90	1.88	2.742 (3)	160
N4—H4 <i>A</i> ...O3	0.90	1.94	2.792 (3)	158
N3—H3 <i>B</i> ...O3 ⁱ	0.90	1.89	2.788 (3)	174
N4—H4 <i>B</i> ...O5 ⁱⁱ	0.90	1.74	2.637 (3)	172

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