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N,N'-Dibenzyl-*N''*-(2,6-difluorobenzoyl)-*N,N'*-dimethylphosphoric triamide

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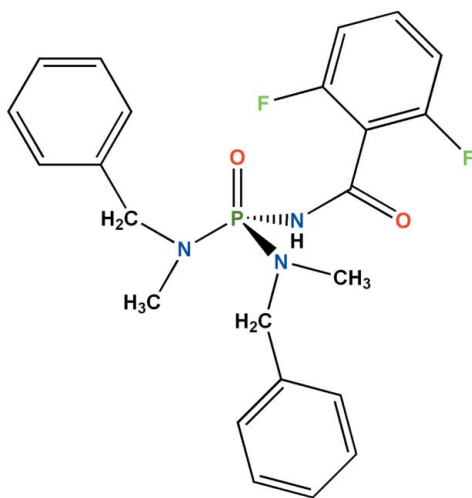
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Key indicators: single-crystal X-ray study; $T = 200$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.048; wR factor = 0.126; data-to-parameter ratio = 18.1.

The phosphoryl and carbonyl groups in the title compound, $\text{C}_{23}\text{H}_{24}\text{F}_2\text{N}_3\text{O}_2\text{P}$, are *anti* to each other. The P atom is in a tetrahedral coordination environment and the environment of each N atom is essentially planar, the average bond angles at the two N atoms being 119.9 and 119.1°. The H atom of the $\text{C}(=\text{O})\text{NHP}(=\text{O})$ group is involved in an intermolecular $\text{P}=\text{O} \cdots \text{H}-\text{N}-$ hydrogen bond, forming centrosymmetric dimers.

Related literature

For related structures, see: Pourayoubi & Sabbaghi (2009); Sabbaghi *et al.* (2010).



Experimental

Crystal data

$\text{C}_{23}\text{H}_{24}\text{F}_2\text{N}_3\text{O}_2\text{P}$
 $M_r = 443.42$
 Triclinic, $P\bar{1}$
 $a = 9.9370$ (11) Å
 $b = 11.1093$ (15) Å
 $c = 11.5902$ (14) Å
 $\alpha = 89.101$ (4)°
 $\beta = 67.826$ (4)°

$\gamma = 71.664$ (4)°
 $V = 1116.9$ (2) Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 0.16$ mm⁻¹
 $T = 200$ K
 $0.30 \times 0.25 \times 0.20$ mm

Data collection

Bruker SMART X2S benchtop
 CCD area-detector
 diffractometer
 Absorption correction: multi-scan
SADABS (Bruker, 2005)
 $T_{\min} = 0.952$, $T_{\max} = 0.968$

13602 measured reflections
 5173 independent reflections
 3851 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.036$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.048$
 $wR(F^2) = 0.126$
 $S = 1.03$
 5173 reflections
 286 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.37$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.32$ e Å⁻³

Table 1

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—H1A···O2 ⁱ	0.83 (2)	1.92 (2)	2.752 (2)	173 (2)

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

Data collection: *SMART* (Bruker, 2005); cell refinement: *SAINT* (Bruker, 2005); data reduction: *SAINT*; 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. The authors wish to thank Bruker AXS, Inc. for the use of one of their SMART X2S benchtop instruments.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: NG5027).

References

- Bruker (2005). *SADABS*, *SMART* and *SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
 Pourayoubi, M. & Sabbaghi, F. (2009). *J. Chem. Crystallogr.* **39**, 874–880.
 Sabbaghi, F., Pourayoubi, M., Toghraee, M. & Divjakovic, V. (2010). *Acta Cryst. E* **66**, o344.
 Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.

supporting information

Acta Cryst. (2010). E66, o2524 [doi:10.1107/S1600536810035725]

***N,N'*-Dibenzyl-*N''*-(2,6-difluorobenzoyl)-*N,N'*-dimethylphosphoric triamide**

Mehrdad Pourayoubi, Atekeh Tarahhomi, Arnold L. Rheingold and James A. Golen

S1. Comment

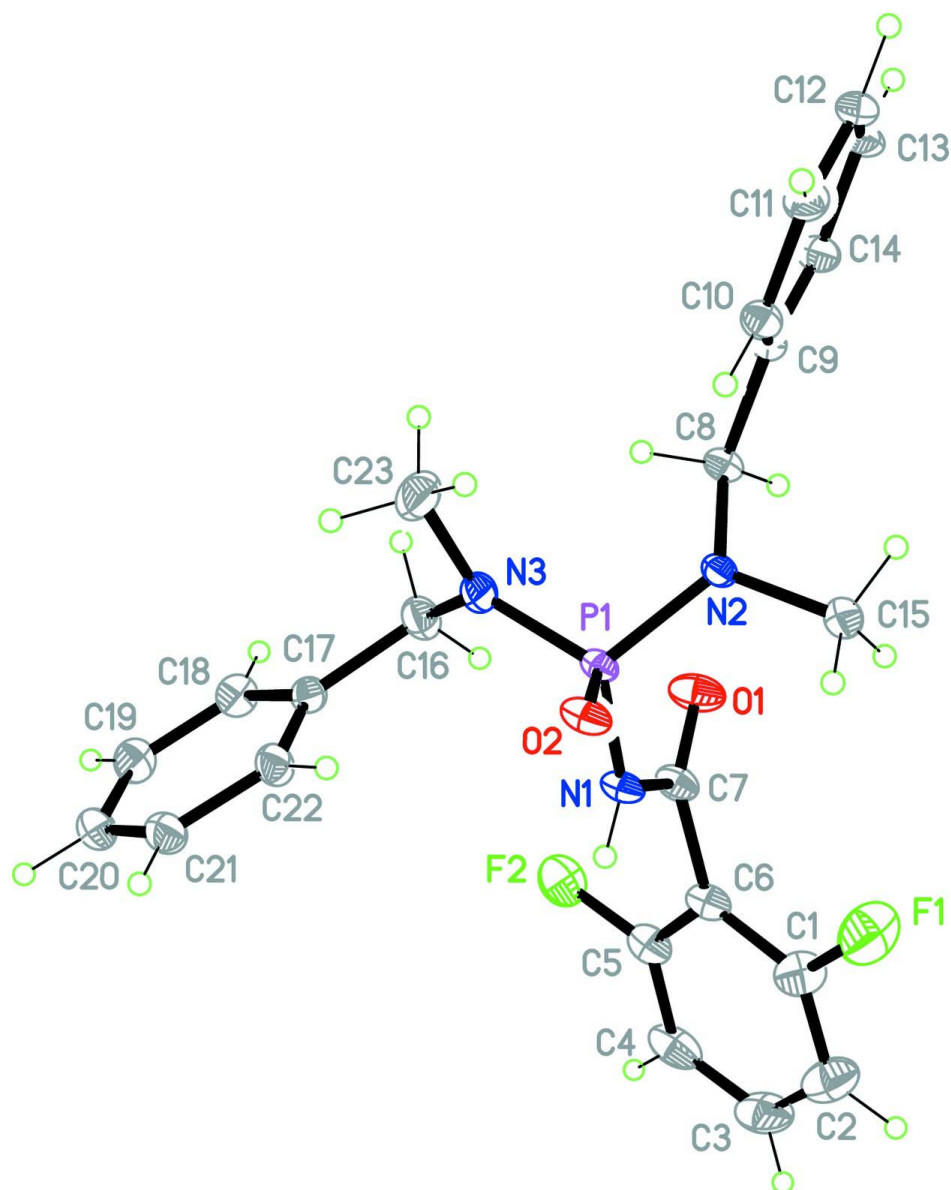
Following the previous works about carbacylamidophosphates with a C(=O)NHP(=O) skeleton such as P(O)[NHC(O)C₆H₄(4-NO₂)]₂[N(CH(CH₃)₂)(CH₂C₆H₅)]₂ (Pourayoubi & Sabbaghi, 2009) and P(O)[NHC(O)C₆H₄(4-NO₂)]₂[NHC₆H₁₁]₂ (Sabbaghi *et al.*, 2010), here, we report the synthesis and crystal structure of title compound, P(O)[NHC(O)C₆H₃(2,6-F₂)]₂[N(CH₃)(CH₂C₆H₅)]₂. The phosphoryl and carbonyl groups are *anti* to each other and the phosphorus atom has a slightly distorted tetrahedral configuration (Fig. 1). The bond angles around the P atom are in the range of 107.11 (8)°–114.81 (8)°. The P1–N2 and P1–N3 bond lengths (1.6405 (14) Å and 1.6266 (16) Å) are shorter than the P1–N1 bond (1.6886 (15) Å). The environment of the nitrogen atoms is essentially planar; the angles C23–N3–P1, C16–N3–C23 and C16–N3–P1 are 117.53 (14)°, 116.19 (17)° and 126.13 (14)°, respectively (with average = 119.9°). A similar result was obtained for the bond angles around N2 atom (average = 119.1°). Furthermore, the angle C7–N1–P1 is 126.79 (13)°. The P=O bond length of 1.4796 (13) Å is standard for phosphoramidate compounds. The hydrogen atom of the C(=O)NHP(=O) group is involved in an intermolecular –P=O⋯H–N– hydrogen bond (see Table 1) to form a centrosymmetric dimeric aggregate.

S2. Experimental

The reaction of phosphorus pentachloride (3.478 g, 16.7 mmol) and 2,6-difluorobenzamide (2.624 g, 16.7 mmol) in dry CCl₄ at 358 K (3 h) and then the treatment of formic acid (0.769 g, 16.7 mmol) at ice bath temperature leads to 2,6-F₂–C₆H₃C(O)NHP(O)Cl₂. To a solution of 2,6-F₂–C₆H₃C(O)NHP(O)Cl₂ (0.411 g, 1.5 mmol) in dry CHCl₃, a solution of *N*-methylbenzylamine (0.727 g, 6 mmol) in dry CHCl₃ was added dropwise at 273 K. After 4 h stirring, the solvent was evaporated in vacuum. The solid was washed with distilled water. Single crystals were obtained from a solution of the title compound in CH₃OH and *n*-C₇H₁₄ (5:1) after slow evaporation at room temperature. IR (KBr, cm⁻¹): 3446, 3042, 2867, 1690, 1607, 1465, 1369, 1210, 1150, 1019, 865, 823, 763, 725.

S3. Refinement

APEX2 software was used for preliminary determination of the unit cell. Determination of integral intensities and unit cell refinement were performed using *SAINTE* and data were corrected for absorption using *SADABS*. Structure was solved by direct methods and all non-hydrogen atoms were refined as anisotropic by Fourier full matrix least squares. Hydrogen H1A was found from a Fourier difference map and was allowed to refine and all other hydrogen atoms were placed in calculated positions with appropriate riding factors.

**Figure 1**

An ORTEP-style plot of title compound. Ellipsoids are given at the 50% probability level.

***N,N'*-Dibenzyl-*N''*-(2,6-difluorobenzoyl)-*N,N'*-dimethylphosphoric triamide**

Crystal data

$C_{23}H_{24}F_2N_3O_2P$
 $M_r = 443.42$
 Triclinic, $P\bar{1}$
 $a = 9.9370$ (11) Å
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 $c = 11.5902$ (14) Å
 $\alpha = 89.101$ (4)°
 $\beta = 67.826$ (4)°
 $\gamma = 71.664$ (4)°
 $V = 1116.9$ (2) Å³

$Z = 2$
 $F(000) = 464$
 $D_x = 1.318$ Mg m⁻³
 Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
 Cell parameters from 5531 reflections
 $\theta = 2.4$ – 27.8 °
 $\mu = 0.16$ mm⁻¹
 $T = 200$ K
 BLOCK, colorless
 $0.30 \times 0.25 \times 0.20$ mm

Data collection

Bruker SMART X2S benchtop CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Curved silicon crystal monochromator

phi and ω scans

Absorption correction: multi-scan

SADABS (Bruker, 2005)

$T_{\min} = 0.952$, $T_{\max} = 0.968$

13602 measured reflections

5173 independent reflections

3851 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.036$

$\theta_{\max} = 27.8^\circ$, $\theta_{\min} = 2.4^\circ$

$h = -13 \rightarrow 13$

$k = -14 \rightarrow 14$

$l = -15 \rightarrow 15$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.048$

$wR(F^2) = 0.126$

$S = 1.03$

5173 reflections

286 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 atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0603P)^2 + 0.2218P]$

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

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

$\Delta\rho_{\max} = 0.37 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -0.32 \text{ e } \text{\AA}^{-3}$

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.

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	0.66485 (5)	0.42270 (4)	0.29906 (4)	0.02461 (13)
F1	0.41237 (17)	0.18210 (16)	0.50507 (15)	0.0723 (5)
F2	0.16095 (14)	0.43844 (12)	0.27352 (13)	0.0532 (4)
O1	0.50833 (16)	0.27969 (15)	0.19272 (13)	0.0445 (4)
O2	0.68882 (13)	0.46617 (12)	0.40757 (12)	0.0323 (3)
N1	0.48905 (16)	0.40601 (15)	0.35526 (15)	0.0277 (3)
N2	0.79630 (15)	0.28474 (13)	0.22927 (13)	0.0246 (3)
N3	0.66719 (17)	0.52094 (15)	0.19333 (15)	0.0358 (4)
C1	0.2810 (2)	0.2390 (2)	0.4858 (2)	0.0449 (5)
C2	0.1464 (3)	0.2189 (3)	0.5632 (2)	0.0583 (7)
H2A	0.1443	0.1678	0.6299	0.070*
C3	0.0152 (3)	0.2755 (3)	0.5402 (2)	0.0573 (7)
H3A	-0.0785	0.2634	0.5924	0.069*
C4	0.0174 (2)	0.3490 (2)	0.4434 (2)	0.0485 (6)
H4A	-0.0733	0.3880	0.4282	0.058*

C5	0.1560 (2)	0.3646 (2)	0.36861 (19)	0.0370 (5)
C6	0.2916 (2)	0.31110 (18)	0.38644 (17)	0.0315 (4)
C7	0.4403 (2)	0.32953 (18)	0.30107 (17)	0.0309 (4)
C8	0.86123 (19)	0.24379 (17)	0.09447 (16)	0.0280 (4)
H8A	0.8268	0.3173	0.0509	0.034*
H8B	0.8212	0.1771	0.0792	0.034*
C9	1.03733 (19)	0.19106 (16)	0.03916 (16)	0.0250 (4)
C10	1.1236 (2)	0.22336 (17)	0.09620 (17)	0.0296 (4)
H10A	1.0727	0.2797	0.1720	0.035*
C11	1.2842 (2)	0.17395 (19)	0.04339 (18)	0.0341 (4)
H11A	1.3419	0.1964	0.0836	0.041*
C12	1.3599 (2)	0.09219 (18)	-0.06744 (19)	0.0358 (5)
H12A	1.4693	0.0584	-0.1033	0.043*
C13	1.2756 (2)	0.06030 (19)	-0.12524 (19)	0.0393 (5)
H13A	1.3270	0.0049	-0.2017	0.047*
C14	1.1147 (2)	0.10908 (18)	-0.07202 (18)	0.0336 (4)
H14A	1.0575	0.0859	-0.1123	0.040*
C15	0.8149 (2)	0.18153 (18)	0.30902 (18)	0.0351 (4)
H15A	0.9229	0.1261	0.2767	0.053*
H15B	0.7497	0.1313	0.3084	0.053*
H15C	0.7842	0.2184	0.3951	0.053*
C16	0.5493 (2)	0.56887 (19)	0.14191 (18)	0.0382 (5)
H16A	0.4917	0.5083	0.1535	0.046*
H16B	0.6010	0.5705	0.0506	0.046*
C17	0.4345 (2)	0.70157 (17)	0.20021 (17)	0.0290 (4)
C18	0.3442 (2)	0.76890 (19)	0.13718 (19)	0.0360 (5)
H18A	0.3538	0.7308	0.0604	0.043*
C19	0.2409 (2)	0.8908 (2)	0.1858 (2)	0.0402 (5)
H19A	0.1796	0.9355	0.1426	0.048*
C20	0.2266 (2)	0.9478 (2)	0.2971 (2)	0.0414 (5)
H20A	0.1568	1.0319	0.3297	0.050*
C21	0.3144 (2)	0.88166 (19)	0.36020 (18)	0.0399 (5)
H21A	0.3047	0.9202	0.4368	0.048*
C22	0.4176 (2)	0.75831 (18)	0.31215 (18)	0.0344 (4)
H22A	0.4767	0.7130	0.3569	0.041*
C23	0.7939 (3)	0.5743 (2)	0.1510 (3)	0.0603 (7)
H23A	0.8498	0.5537	0.0599	0.091*
H23B	0.8646	0.5376	0.1925	0.091*
H23C	0.7516	0.6673	0.1724	0.091*
H1A	0.429 (2)	0.442 (2)	0.427 (2)	0.038 (6)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
P1	0.0173 (2)	0.0247 (2)	0.0261 (2)	-0.00350 (17)	-0.00496 (17)	-0.00334 (17)
F1	0.0611 (9)	0.0976 (12)	0.0788 (11)	-0.0319 (9)	-0.0460 (8)	0.0389 (9)
F2	0.0449 (7)	0.0539 (8)	0.0632 (9)	-0.0087 (6)	-0.0301 (7)	0.0035 (7)
O1	0.0354 (8)	0.0614 (10)	0.0329 (8)	-0.0183 (7)	-0.0068 (6)	-0.0151 (7)

O2	0.0208 (6)	0.0398 (7)	0.0315 (7)	-0.0084 (5)	-0.0061 (5)	-0.0112 (6)
N1	0.0182 (7)	0.0342 (8)	0.0262 (8)	-0.0076 (6)	-0.0044 (6)	-0.0074 (7)
N2	0.0206 (7)	0.0242 (7)	0.0231 (7)	-0.0022 (6)	-0.0064 (6)	0.0009 (6)
N3	0.0259 (8)	0.0294 (8)	0.0446 (10)	-0.0035 (7)	-0.0107 (7)	0.0096 (7)
C1	0.0398 (12)	0.0603 (14)	0.0426 (12)	-0.0222 (11)	-0.0200 (10)	0.0059 (11)
C2	0.0607 (16)	0.0738 (18)	0.0467 (14)	-0.0402 (14)	-0.0133 (12)	0.0104 (13)
C3	0.0415 (13)	0.0695 (17)	0.0565 (15)	-0.0329 (13)	-0.0022 (11)	-0.0100 (13)
C4	0.0249 (10)	0.0540 (14)	0.0626 (15)	-0.0107 (10)	-0.0140 (10)	-0.0199 (12)
C5	0.0305 (10)	0.0391 (11)	0.0407 (11)	-0.0086 (8)	-0.0152 (9)	-0.0073 (9)
C6	0.0242 (9)	0.0382 (10)	0.0320 (10)	-0.0110 (8)	-0.0100 (8)	-0.0080 (8)
C7	0.0229 (9)	0.0361 (10)	0.0314 (10)	-0.0061 (8)	-0.0110 (8)	-0.0053 (8)
C8	0.0242 (9)	0.0297 (9)	0.0241 (9)	-0.0045 (7)	-0.0065 (7)	-0.0026 (7)
C9	0.0220 (8)	0.0247 (9)	0.0249 (9)	-0.0068 (7)	-0.0062 (7)	0.0030 (7)
C10	0.0291 (9)	0.0305 (9)	0.0268 (9)	-0.0079 (8)	-0.0100 (8)	0.0001 (7)
C11	0.0277 (10)	0.0394 (11)	0.0377 (11)	-0.0139 (8)	-0.0135 (8)	0.0078 (9)
C12	0.0202 (9)	0.0349 (10)	0.0431 (11)	-0.0073 (8)	-0.0042 (8)	0.0062 (9)
C13	0.0289 (10)	0.0364 (11)	0.0366 (11)	-0.0061 (8)	0.0008 (8)	-0.0093 (9)
C14	0.0275 (9)	0.0347 (10)	0.0341 (10)	-0.0113 (8)	-0.0063 (8)	-0.0046 (8)
C15	0.0328 (10)	0.0314 (10)	0.0344 (10)	-0.0050 (8)	-0.0106 (8)	0.0058 (8)
C16	0.0437 (11)	0.0322 (10)	0.0317 (10)	-0.0024 (9)	-0.0158 (9)	0.0019 (8)
C17	0.0267 (9)	0.0288 (9)	0.0299 (9)	-0.0078 (7)	-0.0107 (8)	0.0061 (8)
C18	0.0359 (10)	0.0403 (11)	0.0377 (11)	-0.0136 (9)	-0.0202 (9)	0.0070 (9)
C19	0.0299 (10)	0.0413 (12)	0.0511 (13)	-0.0066 (9)	-0.0222 (9)	0.0147 (10)
C20	0.0324 (10)	0.0326 (11)	0.0463 (12)	-0.0009 (8)	-0.0100 (9)	0.0057 (9)
C21	0.0426 (12)	0.0362 (11)	0.0322 (11)	-0.0068 (9)	-0.0103 (9)	0.0014 (9)
C22	0.0364 (10)	0.0304 (10)	0.0323 (10)	-0.0030 (8)	-0.0158 (8)	0.0060 (8)
C23	0.0362 (12)	0.0495 (14)	0.0853 (19)	-0.0157 (11)	-0.0129 (12)	0.0317 (13)

Geometric parameters (Å, °)

P1—O2	1.4796 (13)	C10—H10A	0.9500
P1—N3	1.6266 (16)	C11—C12	1.385 (3)
P1—N2	1.6405 (14)	C11—H11A	0.9500
P1—N1	1.6886 (15)	C12—C13	1.377 (3)
F1—C1	1.360 (2)	C12—H12A	0.9500
F2—C5	1.359 (2)	C13—C14	1.397 (3)
O1—C7	1.219 (2)	C13—H13A	0.9500
N1—C7	1.362 (2)	C14—H14A	0.9500
N1—H1A	0.83 (2)	C15—H15A	0.9800
N2—C8	1.463 (2)	C15—H15B	0.9800
N2—C15	1.471 (2)	C15—H15C	0.9800
N3—C16	1.462 (2)	C16—C17	1.524 (3)
N3—C23	1.474 (3)	C16—H16A	0.9900
C1—C6	1.383 (3)	C16—H16B	0.9900
C1—C2	1.386 (3)	C17—C22	1.380 (3)
C2—C3	1.380 (4)	C17—C18	1.397 (3)
C2—H2A	0.9500	C18—C19	1.385 (3)
C3—C4	1.375 (3)	C18—H18A	0.9500

C3—H3A	0.9500	C19—C20	1.384 (3)
C4—C5	1.386 (3)	C19—H19A	0.9500
C4—H4A	0.9500	C20—C21	1.378 (3)
C5—C6	1.383 (3)	C20—H20A	0.9500
C6—C7	1.512 (3)	C21—C22	1.396 (3)
C8—C9	1.529 (2)	C21—H21A	0.9500
C8—H8A	0.9900	C22—H22A	0.9500
C8—H8B	0.9900	C23—H23A	0.9800
C9—C14	1.389 (2)	C23—H23B	0.9800
C9—C10	1.389 (2)	C23—H23C	0.9800
C10—C11	1.395 (2)		
O2—P1—N3	114.81 (8)	C12—C11—C10	120.19 (18)
O2—P1—N2	110.54 (7)	C12—C11—H11A	119.9
N3—P1—N2	107.66 (8)	C10—C11—H11A	119.9
O2—P1—N1	107.11 (8)	C13—C12—C11	119.55 (17)
N3—P1—N1	107.24 (8)	C13—C12—H12A	120.2
N2—P1—N1	109.34 (8)	C11—C12—H12A	120.2
C7—N1—P1	126.79 (13)	C12—C13—C14	120.31 (18)
C7—N1—H1A	116.4 (14)	C12—C13—H13A	119.8
P1—N1—H1A	116.5 (14)	C14—C13—H13A	119.8
C8—N2—C15	115.44 (14)	C9—C14—C13	120.73 (18)
C8—N2—P1	125.15 (12)	C9—C14—H14A	119.6
C15—N2—P1	116.63 (11)	C13—C14—H14A	119.6
C16—N3—C23	116.19 (17)	N2—C15—H15A	109.5
C16—N3—P1	126.13 (14)	N2—C15—H15B	109.5
C23—N3—P1	117.53 (14)	H15A—C15—H15B	109.5
F1—C1—C6	117.31 (18)	N2—C15—H15C	109.5
F1—C1—C2	119.0 (2)	H15A—C15—H15C	109.5
C6—C1—C2	123.7 (2)	H15B—C15—H15C	109.5
C3—C2—C1	117.9 (2)	N3—C16—C17	114.79 (16)
C3—C2—H2A	121.1	N3—C16—H16A	108.6
C1—C2—H2A	121.1	C17—C16—H16A	108.6
C4—C3—C2	121.4 (2)	N3—C16—H16B	108.6
C4—C3—H3A	119.3	C17—C16—H16B	108.6
C2—C3—H3A	119.3	H16A—C16—H16B	107.5
C3—C4—C5	118.0 (2)	C22—C17—C18	118.77 (17)
C3—C4—H4A	121.0	C22—C17—C16	122.46 (16)
C5—C4—H4A	121.0	C18—C17—C16	118.76 (17)
F2—C5—C6	117.05 (18)	C19—C18—C17	120.47 (19)
F2—C5—C4	119.21 (19)	C19—C18—H18A	119.8
C6—C5—C4	123.7 (2)	C17—C18—H18A	119.8
C5—C6—C1	115.30 (18)	C20—C19—C18	120.37 (18)
C5—C6—C7	121.65 (18)	C20—C19—H19A	119.8
C1—C6—C7	123.04 (17)	C18—C19—H19A	119.8
O1—C7—N1	123.93 (17)	C21—C20—C19	119.48 (19)
O1—C7—C6	121.61 (16)	C21—C20—H20A	120.3
N1—C7—C6	114.46 (15)	C19—C20—H20A	120.3

N2—C8—C9	112.58 (14)	C20—C21—C22	120.33 (19)
N2—C8—H8A	109.1	C20—C21—H21A	119.8
C9—C8—H8A	109.1	C22—C21—H21A	119.8
N2—C8—H8B	109.1	C17—C22—C21	120.56 (18)
C9—C8—H8B	109.1	C17—C22—H22A	119.7
H8A—C8—H8B	107.8	C21—C22—H22A	119.7
C14—C9—C10	118.49 (16)	N3—C23—H23A	109.5
C14—C9—C8	119.50 (15)	N3—C23—H23B	109.5
C10—C9—C8	122.01 (15)	H23A—C23—H23B	109.5
C9—C10—C11	120.73 (17)	N3—C23—H23C	109.5
C9—C10—H10A	119.6	H23A—C23—H23C	109.5
C11—C10—H10A	119.6	H23B—C23—H23C	109.5

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—H1A \cdots O2 ⁱ	0.83 (2)	1.92 (2)	2.752 (2)	173 (2)

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