

X-Ray-Diffraction Study and Determination of Absolute Configuration of the Anticancer Drug S(–)Cyclophosphamide (Endoxan, Cytosan, NSC-26271)

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S(–)Cyclophosphamide, Absolute Configuration, X-Ray Diffraction Methods

(–)Cyclophosphamide (**1**) crystallized from tetrachloromethane in the triclinic space group P1 with cell dimensions $a = 10.500$ (4) Å, $b = 10.490$ (4) Å, $c = 10.761$ (4) Å, $\alpha = 110.0$ (2)°, $\beta = 110.0$ (2)°, $\gamma = 108.9$ (2)°. Three molecules are contained in the unit cell. The X-ray analysis was based on diffractometer measurement of 2635 independent reflections and the structure was solved by Patterson and direct methods. The final R and R_w factors after full-matrix least-squares refinement are 0.0717 and 0.0677, respectively. The absolute configuration is S based on Hamilton's R -factor ratio test. The oxazaphosphorinane ring exists in a chair form with the bis- β -chloroethyl-amino group in equatorial position and about perpendicular to and bisecting the O–P–N plane.

The structure is similar to that found in racemic **1** except a different conformation about one ethyl-amino N–C bond.

Introduction

Cyclophosphamide (**1**, Cytosan, Endoxan, NSC-26271, 2-(bis- β -chloroethyl) amino-2-oxo-1,3,2-oxazaphosphorinane is a powerful, clinically tested and applied cancerostatic agent¹. The stereoselective consumption of **1** by humans² and differentiated cytotoxic activity of its enantiomeric forms³ against ADJ/PC6 plasma cell tumour in mice⁴ suggest that the configuration around phosphorous atom is an important factor in the metabolic process of **1**. These results prompted us to carry out the below described single crystal X-ray analysis of (–)**1** in order to supply information about structure – function relationships. An X-ray study of racemic **1** has been described previously⁵.

Experimental

(–)**1** crystallized from tetrachloromethane in the form of stout, irregular prisms. Space group symmetry and cell constants were derived by photographic and diffractometer techniques and are gathered in Table I. Owing to rather high thermal

Table I. Crystallographic data.

Chemical formula:	$C_7H_{15}N_2O_2Cl_2P$
Space group:	P 1
Cell constants:	$a = 10.500$ (4) Å $b = 10.490$ (4) Å $c = 10.761$ (4) Å $\alpha = 110.0$ (2) Å $\beta = 110.0$ (2) Å $\gamma = 108.9$ (2) Å
Z	3
D_{cal}	1.27 g/cm ³

motion of the molecules in the crystal lattice, the X-ray intensities fell off rapidly with glancing angle Θ and therefore only 2635 data could be collected ($\Theta_{max} = 58^\circ$) using an automated STOE four circle diffractometer equipped with a fine focus tube (Cu K_α , Ni-filter). Data were corrected for geometrical factors but not for absorption (size of crystal only $0.3 \times 0.3 \times 0.5$ mm³), standard deviations were assigned according to counting statistics⁶ and the data were finally converted to normalized structure amplitudes. The structure was solved by direct methods (MULTAN⁷), but since only the positions of P and Cl atoms were apparent from an E-map, they were confirmed against a sharpened Patterson map and then successive Fourier maps were calculated which finally allowed us to deduce the whole structure. In order to establish the absolute con-

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figuration of (–)1, coordinates and anisotropic temperature factors were refined by least squares techniques⁸ for both models (x, y, z) and the enantiomorph ($\bar{x}, \bar{y}, \bar{z}$) with proper inclusion of anomalous scattering contributions from Cl and P atoms⁹. Final weighted discrepancy indices

$$R_w = [\sum w(|F_{\text{obs}}| - |F_{\text{calc}}|)^2 / \sum w |F_{\text{obs}}|^2]^{1/2}$$

are 7.28% for model (x, y, z) and 6.77% for model ($\bar{x}, \bar{y}, \bar{z}$)*. On the basis of Hamilton's R -factor ratio test¹⁰, model (x, y, z) can be excluded at a significance level < 0.005 , see Table II.

Table II. Determination of the absolute configuration of (–)1 using Hamilton's significance test.

For both models 1 ($\bar{x}, \bar{y}, \bar{z}$) and 2 (x, y, z) the R -factors are:

$$\text{Model 1. } R_1 = 0.0717 \quad R_{w1} = 0.0677$$

$$\text{Model 2. } R_2 = 0.0766 \quad R_{w2} = 0.0728$$

$$\text{where } R = \sum ||F_{\text{obs}}| - |F_{\text{calc}}|| / \sum |F_{\text{obs}}|$$

$$R_w = [\sum w(|F_{\text{obs}}| - |F_{\text{calc}}|)^2 / \sum w |F_{\text{obs}}|^2]^{1/2}$$

R and R_w are a measure of the agreement between a set of observed structure amplitudes F_{obs} and a set of calculated structure amplitudes F_{calc} ; w is a set of weights; $w = 1/\sigma^2$, where σ^2 is the variance of F_{obs} .

$$\text{The } R\text{-factor ratios are: } R = R_2/R_1 = 1.06834$$

$$R_w = R_{w2}/R_{w1} = 1.07533$$

In our case for the "dimension of hypothesis", $b = 1$; the "number of reflections", $n = 2635$; the "number of refined parameters", $n = 376$; and "significance level", $\alpha = 0.005$.

$$R_{b, n-m, \alpha} = R_{1, 2259, 0.005} = 1.0018.$$

From the "table of significance points of the R -factor ratio"¹⁰ for $b = 1$, $\alpha = 0.005$, $N = 2000$ and $N_2 = 3000$ we get:

$$R_{1, 2000, 0.005} = 1.0020$$

$$R_{1, 3000, 0.005} = 1.0013$$

Since $N_1 = m - n = 2259$ is between 2000 and 3000 the value 1.0018 was obtained using the following formula¹⁰: for $N_0 < N_1 < N_2$

$$R_{b, N_1, \alpha} \simeq \frac{1}{N_1(N_2 - N_0)} \times [(N_1 - N_0)N_2 R_{b, N_2, \alpha} + (N_2 - N_1)N_0 R_{b, N_0, \alpha}].$$

Both R -factor ratios R and R_w are greater than the R -factor ratio of 1.0018. Therefore for $N_1 = 2259$ and $\alpha = 0.005$, model 2 can be rejected at a level < 0.005 (99.5% probability).

Results and Discussion

Final atomic coordinates for (–)1 are listed in Table III. Tables IV to VI display geometrical data calculated from the entries in Table III. The structure

of (–)1 is presented in Fig. 1. The configuration around the phosphorous atom is S , the complete description of the molecule is $S(-)$ cyclophosphamide. The packing of the molecules within the unit cell is described in the stereo view Fig. 2.

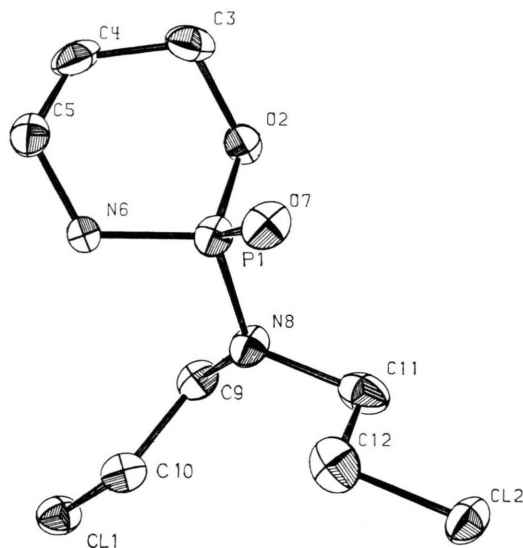


Fig. 1. A plot of $S(-)$ 1 showing thermal ellipsoids of the atoms the 20% probability level.

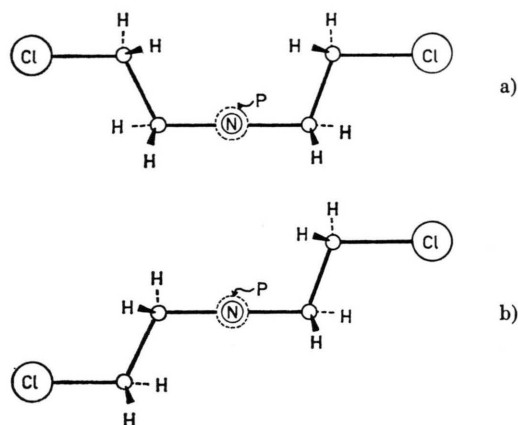


Fig. 2. Schematic drawing of a view along the N–P bond in a) $S(-)$ 1 of this crystal structure and in b) the structure of racemic ($\pm$)1. Note that in a) the two β -chloroethyl groups are related by m symmetry while diad symmetry is observed in b).

It is remarkable that within the unit cell (= the asymmetric unit in this space group) three independent molecules are assembled which show identical conformations with torsion angles not deviating more

* A list of observed and calculated structure factors may be obtained from the authors.

than 3σ . The molecules are linked via hydrogen bonds

$N(6) - H \cdots O(7'')$, 2.891 Å;

$N(6') - H \cdots O(7)$, 2.893 Å and

$N(6'') - H \cdots O(7')$, 2.878 Å,

i. e. each molecule acts simultaneously as a hydrogen bond donor and acceptor.

The cyclophosphamide molecules in the present investigation have similar geometry as discussed for the crystal structure of the racemate⁵ with only one torsion angle different. The 1,3,2-oxazaphosphorinane rings display chair conformation with the bis- β -chloroethylamino substituent in equatorial

orientation. The latter is approximately perpendicular to the $O(2) - P(1) - N(6)$ plane, the torsion angles with the $C(9) - N(8) - C(11)$ planes being 83.4° , 85.6° , and 86.1° for molecules, A, B and C. This conformation results from partial double bond character of the $P - N(8)$ bond which is shortened by 0.119 Å with respect to a normal single bond of 1.769 Å. Similarly, the configuration about $N(8)$ is nearly planar, with the sum of angles around $N(8)$ at 357.5° . Further it is noticeable that the $P(1) - N(6)$ distance, 1.636 Å, indicates partial double bond character as well, *i. e.* the hydrogen atom at $N(6)$ is rather acidic.

Table III. Fractional atomic coordinates and anisotropic temperature factors of the non-hydrogen atoms with standard deviations in parentheses. Temperature factors are in the form $T = \exp [-(\beta_{11}h^2 + \beta_{22}k^2 + \beta_{33}l^2 + 2\beta_{12}hk + 2\beta_{13}hl + 2\beta_{23}kl)]$.

Atom	X	Y	Z	B (11) * 10000.	B (22) * 10000.	B (33) * 10000.	B (12) * 10000.	B (13) * 10000.	B (23) * 10000.
P1	-.7657 (4)	-.5713 (3)	-.9948 (4)	166 (5)	117 (4)	148 (4)	45 (4)	63 (4)	69 (4)
O2	-.6915 (8)	-.6604 (9)	-1.0670 (9)	153 (14)	213 (15)	189 (15)	96 (12)	83 (12)	111 (13)
C3	-.6630 (15)	-.6303 (16)	-1.1798 (17)	182 (23)	234 (25)	245 (27)	79 (20)	144 (22)	116 (22)
C4	-.8147 (17)	-.6862 (17)	-1.3133 (15)	226 (27)	265 (27)	154 (21)	36 (22)	115 (21)	106 (21)
C5	-.8973 (14)	-.6040 (14)	-1.2721 (15)	174 (19)	172 (19)	187 (22)	56 (17)	74 (18)	109 (18)
N6	-.9152 (9)	-.6063 (10)	-1.1468 (9)	130 (14)	213 (16)	143 (13)	89 (12)	69 (11)	100 (12)
O7	-.6499 (9)	-.4074 (8)	-.8771 (9)	259 (16)	127 (12)	202 (15)	37 (11)	100 (14)	74 (12)
N8	-.8408 (10)	-.6672 (9)	-.9211 (10)	173 (15)	137 (13)	154 (14)	22 (12)	60 (13)	84 (12)
C9	-.9719 (13)	-.8284 (11)	-1.0202 (12)	211 (19)	141 (15)	161 (17)	90 (14)	103 (16)	89 (14)
C10	-1.1221 (14)	-.8246 (13)	-1.0330 (14)	224 (22)	125 (17)	201 (21)	83 (16)	126 (19)	82 (16)
C11	-.7349 (13)	-.6050 (13)	-.7409 (14)	258 (19)	288 (19)	419 (21)	185 (16)	228 (17)	255 (17)
C12	-.7952 (19)	-.5209 (17)	-.6862 (20)	311 (32)	227 (27)	355 (39)	184 (26)	165 (31)	171 (29)
CL1	-1.2856 (4)	-1.0212 (4)	-1.1546 (4)	183 (5)	164 (5)	182 (5)	56 (4)	100 (5)	79 (5)
CL2	-.6636 (5)	-.4271 (4)	-.4655 (4)	260 (8)	234 (7)	157 (6)	82 (6)	86 (6)	92 (5)
P1'	-.6221 (4)	-.0035 (4)	-.7584 (4)	122 (4)	184 (5)	147 (5)	75 (4)	65 (4)	80 (4)
O2'	-.5346 (9)	.1617 (9)	-.7392 (9)	175 (14)	170 (14)	152 (14)	108 (13)	66 (13)	86 (13)
C3'	-.5674 (16)	.1563 (17)	-.8871 (18)	197 (24)	260 (28)	291 (32)	145 (23)	119 (24)	208 (26)
C4'	-.5063 (16)	.0658 (18)	-.9593 (15)	196 (23)	297 (31)	164 (22)	127 (23)	90 (20)	165 (23)
C5'	-.5898 (14)	-.1014 (16)	-1.0035 (13)	159 (20)	233 (25)	102 (17)	97 (19)	32 (16)	78 (18)
N6'	-.5872 (10)	-.1151 (11)	-.8708 (11)	138 (15)	172 (16)	144 (16)	76 (13)	58 (13)	74 (14)
O7'	-.7904 (9)	-.0523 (12)	-.8048 (11)	107 (13)	380 (23)	230 (19)	115 (15)	83 (14)	157 (18)
N8'	-.5279 (11)	.0191 (13)	-.5880 (11)	123 (15)	288 (22)	118 (15)	115 (15)	66 (13)	108 (16)
C9'	-.3660 (12)	.0456 (14)	-.5249 (13)	112 (16)	181 (20)	139 (19)	47 (15)	58 (15)	66 (17)
C10'	-.3711 (12)	-.1018 (14)	-.5427 (14)	96 (16)	195 (21)	179 (22)	51 (16)	47 (16)	92 (19)
C11'	-.5897 (17)	.0595 (17)	-.4749 (19)	203 (27)	219 (28)	228 (29)	71 (23)	7 (24)	106 (26)
C12'	-.6776 (18)	-.0911 (15)	-.5012 (21)	273 (31)	143 (22)	304 (36)	70 (22)	105 (29)	100 (24)
CL1'	-.1734 (4)	-.0739 (4)	-.4683 (4)	152 (5)	238 (6)	212 (6)	101 (5)	97 (5)	141 (5)
CL2'	-.7670 (4)	-.0450 (5)	-.3733 (4)	214 (7)	361 (9)	212 (7)	144 (6)	147 (6)	160 (7)
P1''	-.1898 (0)	-.4279 (0)	-.0884 (0)	202 (5)	151 (4)	196 (5)	110 (4)	130 (5)	109 (4)
O2''	-.3558 (9)	-.5052 (9)	-.2388 (9)	159 (13)	180 (14)	154 (13)	75 (11)	83 (11)	73 (12)
C3''	-.3500 (16)	-.5338 (14)	-.3797 (14)	255 (25)	150 (20)	149 (21)	97 (19)	56 (19)	44 (17)
C4''	-.2604 (19)	-.3813 (17)	-.3631 (15)	343 (35)	261 (27)	173 (23)	199 (27)	177 (25)	138 (22)
C5''	-.0921 (16)	-.2964 (14)	-.2361 (15)	238 (25)	172 (21)	219 (24)	119 (19)	151 (22)	111 (19)
N6''	-.0780 (10)	-.2770 (10)	-.0888 (11)	152 (15)	146 (15)	184 (17)	69 (12)	89 (14)	89 (13)
O7''	-.1452 (11)	-.5477 (11)	-.0871 (11)	307 (18)	219 (16)	284 (19)	187 (15)	211 (17)	191 (15)
N8''	-.2111 (12)	-.3538 (11)	.0605 (11)	257 (20)	182 (16)	202 (17)	149 (16)	169 (16)	135 (15)
C9''	-.2376 (13)	-.2223 (14)	.0943 (14)	173 (18)	189 (19)	192 (20)	109 (17)	126 (17)	126 (17)
C10''	-.0904 (14)	-.0715 (14)	.2284 (14)	197 (21)	184 (20)	148 (20)	107 (17)	89 (18)	62 (17)
C11''	-.2508 (17)	-.4564 (19)	.1332 (18)	198 (27)	363 (38)	210 (28)	115 (27)	96 (24)	4 (27)
C12''	-.1051 (16)	-.3988 (17)	.2539 (20)	186 (24)	263 (28)	271 (31)	103 (22)	104 (24)	88 (26)
CL1''	-.1203 (4)	.0912 (4)	.2703 (4)	230 (6)	186 (5)	194 (5)	124 (5)	110 (5)	110 (4)
CL2''	-.1485 (5)	-.5305 (5)	.3384 (5)	313 (8)	293 (8)	284 (8)	192 (7)	190 (7)	214 (7)

Table IV. Fractional atomic coordinates for the hydrogen atoms.

Atom	X	Y	Z
H3 1	-.6097	-.6991	-1.2222
H3 2	-.5829	-.5072	-1.1303
H4 1	-.8899	-.8106	-1.3616
H4 2	-.7947	-.6682	-1.4000
H5 1	-.8292	-.4813	-1.2374
H5 2	-1.0111	-.6567	-1.3711
H6 1	-1.0299	-.6323	-1.1534
H9 1	-.9870	-.8741	-1.1371
H9 2	-.9507	-.8987	-.9749
H11 1	-.6109	-.5324	-.6988
H11 2	-.7543	-.7020	-.7194
H12 1	.0739	.3800	.2696
H12 2	.2232	-.4200	.2854
H10 1	-.1378	.2600	-.0972
H10 2	-.1311	.2200	.0622
H3' 1	-.5080	.2762	-.8645
H3' 2	-.6927	.1033	-.9619
H4' 1	-.3830	.1149	-.8808
H4' 2	-.5213	.0686	-1.0636
H5' 1	-.5345	-.1613	-1.0476
H5' 2	-.7112	-.1561	-1.0924
H6' 1	-.5605	-.2016	-.8497
H9' 1	-.2926	.1375	-.4073
H9' 2	-.3164	.0791	-.5933
H11' 1	-.4944	.1469	-.3546
H11' 2	-.6688	.1084	-.5083
H10' 1	-.4191	-.1357	-.4770
H10' 2	-.4423	-.1938	-.6627
H12' 1	.3452	-.1820	.4251
H3'' 1	-.2927	-.6017	-.3986
H3'' 2	-.4681	-.5950	-.4778
H4'' 1	-.3125	-.3091	-.3346
H4'' 2	-.2643	-.4001	-.4700
H5'' 1	-.0301	-.1838	-.2224
H5'' 2	-.0381	-.3664	-.2681
H6'' 1	.0066	-.1612	.0191
H9'' 1	-.3301	-.2439	.1209
H9'' 2	-.2737	-.2108	-.0079
H10'' 1	-.0550	-.0849	.3278
H10'' 2	.0011	-.0515	.1991
H11'' 1	-.3289	-.4416	.1736
H11'' 2	-.2996	-.5812	.0511
H12'' 1	-.0582	-.2730	.3720

The bis- β -chloroethylamino group displays nearly mirror-symmetry, Fig. 2. The torsion angles about the N–C bonds, P(1)–N(8)–C(9)–C(10) and P(1)–N(8)–C(11)–C(12) are opposite, at -104° and 97° but the orientations about the ethyl bonds, N–CH₂–CH₂–Cl, are both *trans*.

From the close structural relationship between the three S(–) **1** molecules one could conclude that they exist in a rather constrained conformation. A comparison with the crystal structure of the racemic ($\pm$) **1** (Tables V–VII) shows that the six-membered oxazaphosphorinane ring and its substituents N(8), C(9) and C(11) have nearly identical geometry but the conformation about the N(8)–C(9) bond is different, with the torsion angle P(1)–N(8)–C(9)–C(10) at 103° . This brings the two β -chloroethyl groups into a propeller-like orientation with nearly twofold rotational symmetry, Fig. 2. Thus the oxazaphosphorinane moiety appears to be rather rigid but rotation does occur about the N–C bonds.

Table V. Bond distances (Å), with standard deviations in parentheses.

Bond	Mole- cule A	Mole- cule B	Mole- cule C	Ref. 5
P(1)–O(2)	1.570(11)	1.580(10)	1.613(07)	1.580(4)
O(2)–C(3)	1.453(24)	1.486(23)	1.469(20)	1.466(4)
C(3)–C(4)	1.489(20)	1.465(27)	1.485(22)	1.520(10)
C(4)–C(5)	1.474(26)	1.495(23)	1.520(19)	1.488(8)
C(5)–N(6)	1.431(21)	1.477(21)	1.475(22)	1.481(7)
N(6)–P(1)	1.659(10)	1.606(13)	1.644(11)	1.623(5)
P(1)–O(7)	1.474(06)	1.510(10)	1.477(12)	1.466(4)
P(1)–N(8)	1.650(13)	1.648(13)	1.652(13)	1.623(5)
N(8)–C(9)	1.476(11)	1.487(17)	1.444(20)	1.457(7)
C(9)–C(10)	1.551(22)	1.473(22)	1.530(12)	1.512(8)
C(1)–Cl 1	1.812(10)	1.832(14)	1.770(16)	1.769(6)
N(8)–C(11)	1.622(16)	1.564(25)	1.565(27)	1.483(6)
C(11)–C(12)	1.335(27)	1.418(25)	1.384(21)	1.513(7)
C(12)–Cl 2	1.956(18)	1.931(24)	1.918(23)	1.798(5)

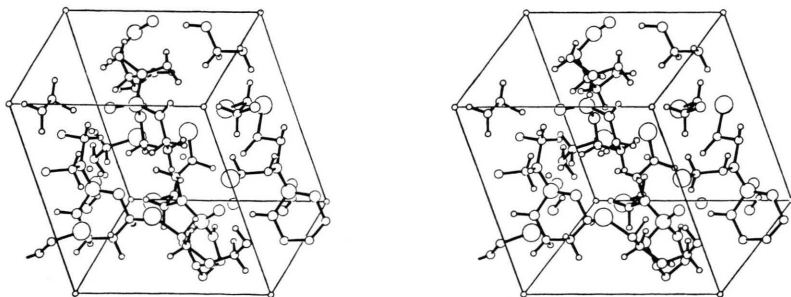


Fig. 3. A stereo view of the structure along a-axis.

Table VI. Bond angles ($^{\circ}$) with standard deviations in parentheses.

Angle	Molecule A	Molecule B	Molecule C	Ref. 5
O(2)–P(1)–N(6)	103.0 (0.5)	105.0 (0.6)	103.6 (0.5)	103.1 (0.2)
O(2)–P(1)–O(7)	112.0 (0.6)	114.4 (0.6)	110.2 (0.5)	113.8 (0.2)
O(2)–P(1)–N(8)	105.8 (0.5)	104.7 (0.6)	106.6 (0.5)	104.4 (0.2)
N(6)–P(1)–O(7)	117.7 (0.6)	117.5 (0.7)	120.1 (0.6)	117.3 (0.3)
N(6)–P(1)–N(8)	106.2 (0.5)	107.2 (0.6)	105.4 (0.6)	106.7 (0.2)
O(7)–P(1)–N(8)	111.1 (0.6)	110.2 (0.7)	110.1 (0.6)	110.4 (0.2)
P(1)–O(2)–C(3)	115.7 (0.9)	113.3 (0.9)	114.7 (0.8)	118.8 (0.4)
O(2)–C(3)–C(4)	108.8 (1.3)	108.9 (1.4)	108.5 (1.2)	108.9 (0.5)
C(3)–C(4)–C(5)	112.6 (1.4)	112.0 (1.4)	111.2 (1.4)	112.1 (0.6)
C(4)–C(5)–N(6)	112.9 (1.2)	110.8 (1.2)	112.5 (1.3)	110.5 (0.5)
P(1)–N(6)–C(5)	119.4 (0.9)	119.8 (1.9)	118.3 (0.9)	121.5 (0.4)
P(1)–N(8)–C(9)	120.8 (0.9)	120.9 (1.0)	120.6 (1.0)	121.7 (0.4)
P(1)–N(8)–C(11)	117.5 (0.8)	118.7 (1.0)	117.7 (1.0)	122.1 (0.4)
C(9)–N(8)–C(11)	118.7 (1.0)	118.2 (1.2)	119.3 (1.2)	116.3 (0.5)
N(8)–C(9)–C(10)	108.2 (1.0)	108.4 (1.1)	111.5 (1.2)	110.6 (0.5)
C(9)–C(10)–Cl 1	109.2 (1.0)	109.6 (1.0)	112.1 (1.0)	110.3 (0.5)
N(8)–C(11)–C(12)	98.4 (1.2)	100.3 (1.4)	101.4 (1.5)	117.7 (0.4)
C(11)–C(12)–Cl 2	101.6 (1.3)	100.7 (1.3)	102.3 (1.3)	107.2 (0.4)

Table 7. Torsional angles $^{\circ}$ and standard deviations in parentheses.

Angles	Molecule A	Molecule B	Molecule C	Ref. 5
P(1)–O(2)–C(3)–C(4)	–63.6 (1.5)	–64.5 (1.5)	–65.5 (1.4)	–59.6
P(1)–N(6)–C(5)–C(4)	44.2 (1.6)	44.2 (1.6)	45.6 (1.6)	45.8
P(1)–N(8)–C(9)–C(10)	–104.4 (1.1)	–101.9 (1.2)	–103.7 (1.2)	102.8
P(1)–N(8)–C(11)–C(12)	97.2 (1.3)	96.8 (1.4)	97.4 (1.4)	90.8
O(2)–P(1)–N(6)–C(5)	–40.3 (1.1)	–40.7 (1.2)	–41.3 (1.1)	–38.0
O(2)–P(1)–N(8)–C(9)	–65.2 (1.0)	–68.5 (1.1)	–67.5 (1.1)	–54.4
O(2)–P(1)–N(8)–C(11)	95.0 (0.9)	94.3 (1.2)	94.9 (1.1)	123.6
O(2)–C(3)–C(4)–C(5)	61.9 (1.7)	65.5 (1.7)	64.3 (1.6)	61.9
C(3)–O(2)–P(1)–N(6)	49.6 (1.1)	49.2 (1.1)	50.9 (1.0)	44.9
C(3)–C(2)–P(1)–O(7)	–77.9 (1.1)	–79.0 (1.1)	–78.8 (1.0)	–83.3
C(3)–O(2)–P(1)–N(8)	160.9 (1.0)	161.9 (1.0)	161.8 (1.0)	159.2
C(3)–C(4)–C(5)–N(6)	–53.0 (1.8)	–54.5 (1.8)	–55.0 (1.9)	–55.0
C(5)–N(6)–P(1)–O(7)	83.5 (1.1)	83.7 (1.2)	82.1 (1.1)	88.0
C(5)–N(6)–P(1)–N(8)	–151.4 (1.0)	–151.7 (1.0)	–153.1 (1.0)	–147.6
N(6)–P(1)–N(8)–C(9)	43.9 (1.1)	42.6 (1.2)	42.1 (1.2)	44.9
N(6)–P(1)–N(8)–C(11)	–155.9 (0.9)	–154.6 (1.1)	–155.5 (1.1)	–127.6
O(7)–P(1)–N(8)–C(9)	173.0 (0.9)	171.5 (1.0)	173.1 (1.0)	–177.2
O(7)–P(1)–N(8)–C(11)	–26.8 (1.0)	–25.7 (1.3)	–24.6 (1.3)	0.9
N(8)–C(9)–C(10)–Cl 1	179.9 (0.9)	179.2 (0.9)	179.5 (1.0)	–175.9
N(8)–C(11)–C(12)–Cl 2	–175.9 (0.9)	–176.0 (1.0)	–176.9 (1.0)	176.7
C(9)–N(8)–C(11)–C(12)	–102.1 (1.4)	–99.9 (1.6)	–100.1 (1.6)	–91.1
C(10)–C(9)–N(8)–C(11)	95.6 (1.2)	95.2 (1.5)	94.3 (1.5)	–75.4

* Definition: The angles A–B–C–D are defined zero when the bonds A–B and C–D are cis planar. They are counted positive when, looking along the central B–C bond, the far bond is rotated clockwise with respect to the near bond.

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