

catena-Poly[[[dichloridomercury(II)]- μ -1,4-bis(3-pyridylaminomethyl)benzene- κ^2 N:N'] N,N-dimethylformamide monosolvate]

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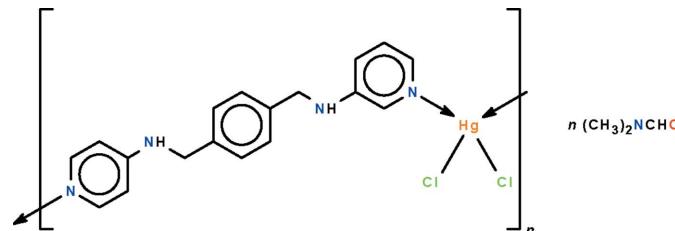
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Key indicators: single-crystal X-ray study; $T = 293\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.013\text{ \AA}$; disorder in solvent or counterion; R factor = 0.051; wR factor = 0.167; data-to-parameter ratio = 18.9.

The crystal structure of the polymeric title compound, $\{[\text{HgCl}_2(\text{C}_{18}\text{H}_{18}\text{N}_4)]\cdot\text{C}_3\text{H}_7\text{NO}\}_n$, features an *N*-heterocyclic ligand which links adjacent HgCl_2 units into a helical chain running along the *b* axis. The coordination geometry of the Hg^{II} atom is a distorted tetrahedron. The dimethylformamide molecule is disordered over two positions in a 1:1 ratio, and is linked to the complex molecules *via* $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

For the structure of the *N*-heterocyclic ligand, see: Zhu *et al.* (2007).



Experimental

Crystal data

$[\text{HgCl}_2(\text{C}_{18}\text{H}_{18}\text{N}_4)]\cdot\text{C}_3\text{H}_7\text{NO}$
 $M_r = 634.95$
Monoclinic, $P2_1/n$
 $a = 8.4851(9)\text{ \AA}$
 $b = 15.1215(14)\text{ \AA}$
 $c = 19.490(2)\text{ \AA}$
 $\beta = 103.826(2)^\circ$

$V = 2428.2(4)\text{ \AA}^3$
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 6.58\text{ mm}^{-1}$
 $T = 293\text{ K}$
 $0.15 \times 0.11 \times 0.11\text{ mm}$

Data collection

Rigaku R-AXIS RAPID IP diffractometer
Absorption correction: multi-scan (*ABSCOR*; Higashi, 1995)
 $T_{\min} = 0.439$, $T_{\max} = 0.531$

22980 measured reflections
5479 independent reflections
2593 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.089$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.051$
 $wR(F^2) = 0.167$
 $S = 1.05$
5479 reflections
290 parameters

42 restraints
H-atom parameters constrained
 $\Delta\rho_{\max} = 1.26\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -1.31\text{ e \AA}^{-3}$

Table 1
Selected bond lengths (\AA).

Hg1—N1	2.395 (7)	Hg1—Cl1	2.355 (3)
Hg1—N4 ⁱ	2.308 (6)	Hg1—Cl2	2.391 (3)

Symmetry code: (i) $-x + \frac{5}{2}, y + \frac{1}{2}, -z + \frac{3}{2}$.

Table 2
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N2—H2N \cdots O1	0.88	2.15	3.03 (3)	174
N2—H2N \cdots O1'	0.88	2.11	2.99 (3)	180
N3—H3N \cdots O1 ⁱⁱ	0.88	2.14	3.01 (3)	166
N3—H3N \cdots O1' ⁱⁱ	0.88	2.13	2.98 (3)	162

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

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalClear* (Rigaku/MSC, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *publCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: XU5257).

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supporting information

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catena-Poly[[[dichloridomercury(II)]- μ -1,4-bis(3-pyridylaminomethyl)benzene- $\kappa^2N:N'$] N,N-dimethylformamide monosolvate]

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S1. Comment

1,4-Bis(2-pyridylaminomethyl)benzene is a flexible *N*-heterocycle whose pyridyl and amino *N*-atoms are capable for forming coordination polymers (Zhu *et al.*, 2007). The crystal structure of $HgCl_2(C_{18}H_{18}N_4)$ DMF features the *N*-heterocycle linking adjacent $HgCl_2$ units into a helical chain (Scheme I, Fig. 1). The geometry of Hg^{II} is a tetrahedron. The lattice DMF molecule is disordered in two positions in a 1:1 ratio. The *N*-heterocycle forms an $N\cdots H\cdots O$ hydrogen bond to the solvent molecule at an $N\cdots O$ distance of 2.99 (3) and 3.03 (3) Å; the hydrogen bond probably stabilizes the solvent molecule so that it is not lost during crystallization.

S2. Experimental

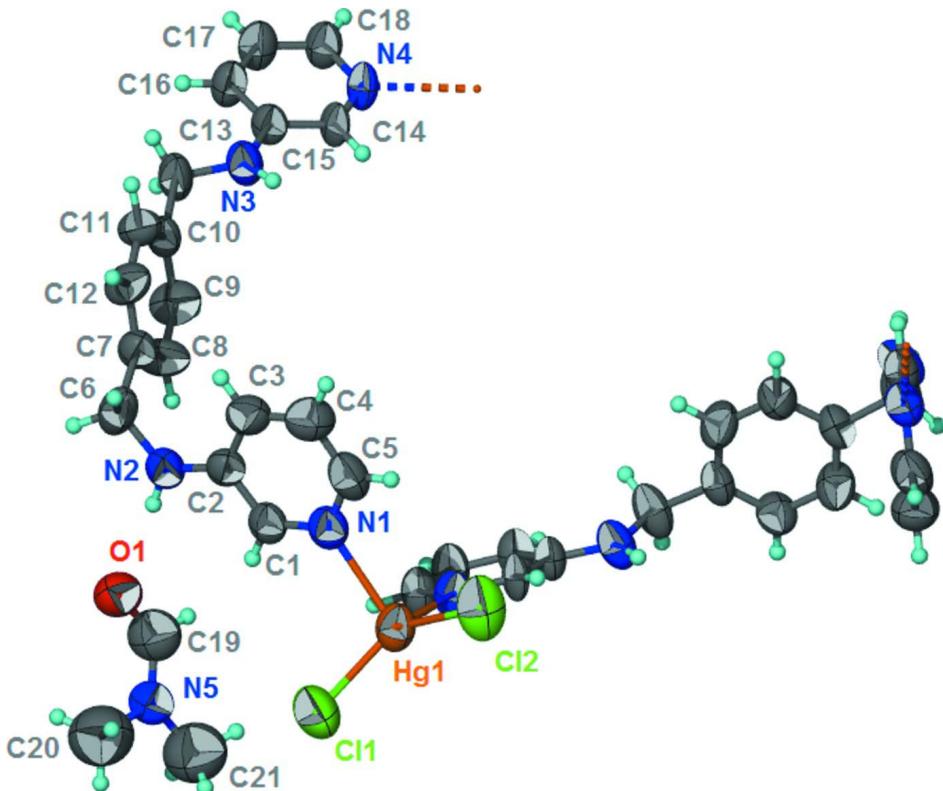
A THF solution (10 ml) of mercuric chloride (2 mmol) was mixed with a DMF solution (5 ml) of 1,4-bis(3-pyridylaminomethyl)benzene (2 mmol). The solution was filtered and sent aside for the growth of colorless crystals.

S3. Refinement

Carbon-bound H-atoms were placed in calculated positions ($C-H$ 0.93 Å) and were included in the refinement in the riding model approximation, with $U(H)$ set to $1.2U(C)$. The amino H-atoms similar treated ($N-H$ 0.86 Å).

The lattice DMF molecule is disordered over two sites; the disorder could not be refined, and was assumed to be a 1:1 type of disorder. The $C-O$ distances were restrained to 1.25 ± 0.01 Å, the $C_{\text{carbonyl}}-N$ distances to 1.35 ± 0.01 Å and the $N-C_{\text{methyl}}$ distances to 1.45 ± 0.01 Å. Each component was restrained to planar, with a maximum deviation of 0.01 Å. The temperature factors of the primed atoms were set to those of the unprimed ones, and the anisotropic temperature factors were restrained to be nearly isotropic.

The final difference Fourier map had peaks/holes in the vicinity of $Hg1$.

**Figure 1**

Thermal ellipsoid plot (Barbour, 2001) of a portion of the polymeric chain structure of $\text{HgCl}_2(\text{C}_{18}\text{H}_{18}\text{N}_4)\text{-DMF}$ at the 50% probability level; hydrogen atoms are drawn as spheres of arbitrary radius.

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$M_r = 634.95$

Monoclinic, $P2_1/n$

Hall symbol: -P 2yn

$a = 8.4851 (9)$ Å

$b = 15.1215 (14)$ Å

$c = 19.490 (2)$ Å

$\beta = 103.826 (2)^\circ$

$V = 2428.2 (4)$ Å³

$Z = 4$

Data collection

Rigaku R-AXIS RAPID IP
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scan

Absorption correction: multi-scan
(*ABSCOR*; Higashi, 1995)

$T_{\min} = 0.439$, $T_{\max} = 0.531$

$F(000) = 1232$

$D_x = 1.737$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 10259 reflections

$\theta = 3.2\text{--}27.5^\circ$

$\mu = 6.58$ mm⁻¹

$T = 293$ K

Prism, colorless

$0.15 \times 0.11 \times 0.11$ mm

22980 measured reflections

5479 independent reflections

2593 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.089$

$\theta_{\max} = 27.5^\circ$, $\theta_{\min} = 3.2^\circ$

$h = -11 \rightarrow 11$

$k = -19 \rightarrow 19$

$l = -25 \rightarrow 22$

*Refinement*Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.051$ $wR(F^2) = 0.167$ $S = 1.05$

5479 reflections

290 parameters

42 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

 $w = 1/[\sigma^2(F_o^2) + (0.0656P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} = 0.001$ $\Delta\rho_{\max} = 1.26 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{\min} = -1.31 \text{ e } \text{\AA}^{-3}$ *Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
Hg1	0.65054 (5)	0.63785 (2)	0.73300 (2)	0.0944 (2)	
Cl1	0.3880 (4)	0.5916 (3)	0.7371 (2)	0.1488 (12)	
Cl2	0.7868 (4)	0.77413 (15)	0.72364 (19)	0.1366 (11)	
N1	0.7052 (9)	0.5581 (4)	0.6351 (4)	0.0845 (19)	
N2	0.5616 (9)	0.3645 (4)	0.5347 (5)	0.097 (2)	
H2N	0.4912	0.3482	0.5589	0.116*	
N3	1.2859 (9)	0.1242 (4)	0.5624 (4)	0.089 (2)	
H3N	1.2888	0.1819	0.5685	0.106*	
N4	1.6587 (8)	0.0700 (4)	0.6783 (3)	0.0778 (18)	
C1	0.6262 (10)	0.4841 (5)	0.6124 (5)	0.077 (2)	
H1	0.5528	0.4619	0.6366	0.092*	
C2	0.6491 (10)	0.4383 (5)	0.5535 (5)	0.078 (2)	
C3	0.7569 (11)	0.4741 (7)	0.5156 (5)	0.097 (3)	
H3	0.7720	0.4457	0.4753	0.117*	
C4	0.8377 (13)	0.5496 (7)	0.5381 (6)	0.105 (3)	
H4	0.9108	0.5734	0.5144	0.126*	
C5	0.8089 (11)	0.5909 (6)	0.5976 (6)	0.098 (3)	
H5	0.8631	0.6435	0.6127	0.118*	
C6	0.5810 (12)	0.3099 (7)	0.4743 (5)	0.108 (3)	
H6A	0.5769	0.3485	0.4342	0.130*	
H6B	0.4894	0.2698	0.4617	0.130*	
C7	0.7345 (10)	0.2565 (5)	0.4868 (5)	0.077 (2)	
C8	0.7897 (12)	0.2164 (6)	0.5502 (6)	0.100 (3)	
H8	0.7379	0.2257	0.5866	0.120*	
C9	0.9204 (13)	0.1631 (7)	0.5601 (5)	0.101 (3)	
H9	0.9559	0.1349	0.6035	0.121*	
C10	1.0050 (10)	0.1482 (5)	0.5074 (5)	0.073 (2)	
C11	0.9536 (11)	0.1896 (6)	0.4447 (5)	0.085 (2)	
H11	1.0070	0.1811	0.4087	0.102*	
C12	0.8207 (12)	0.2447 (6)	0.4349 (5)	0.093 (3)	
H12	0.7875	0.2749	0.3923	0.111*	
C13	1.1469 (10)	0.0856 (6)	0.5190 (6)	0.098 (3)	
H13A	1.1204	0.0317	0.5408	0.118*	
H13B	1.1685	0.0701	0.4738	0.118*	

C14	1.5277 (9)	0.1115 (5)	0.6468 (4)	0.0675 (19)	
H14	1.5119	0.1688	0.6610	0.081*	
C15	1.4147 (9)	0.0755 (5)	0.5950 (4)	0.0700 (19)	
C16	1.4362 (11)	-0.0123 (5)	0.5778 (5)	0.083 (2)	
H16	1.3588	-0.0406	0.5429	0.100*	
C17	1.5701 (12)	-0.0560 (6)	0.6123 (5)	0.092 (3)	
H17	1.5835	-0.1152	0.6020	0.111*	
C18	1.6827 (11)	-0.0152 (5)	0.6607 (5)	0.083 (2)	
H18	1.7778	-0.0446	0.6825	0.100*	
O1	0.300 (4)	0.314 (2)	0.6094 (14)	0.110 (3)	0.50
N5	0.181 (3)	0.3560 (13)	0.7007 (11)	0.085 (3)	0.50
C19	0.303 (3)	0.3275 (14)	0.6732 (14)	0.128 (6)	0.50
H19	0.4011	0.3163	0.7053	0.154*	0.50
C20	0.025 (3)	0.375 (2)	0.6545 (18)	0.197 (11)	0.50
H20A	0.0333	0.4262	0.6267	0.295*	0.50
H20B	-0.0525	0.3856	0.6823	0.295*	0.50
H20C	-0.0096	0.3252	0.6239	0.295*	0.50
C21	0.180 (5)	0.380 (2)	0.7725 (13)	0.185 (9)	0.50
H21A	0.1230	0.4349	0.7723	0.278*	0.50
H21B	0.2899	0.3868	0.7998	0.278*	0.50
H21C	0.1273	0.3346	0.7931	0.278*	0.50
O1'	0.323 (4)	0.309 (2)	0.6172 (14)	0.110 (3)	0.50
N5'	0.181 (3)	0.3701 (11)	0.6889 (12)	0.085 (3)	0.50
C19'	0.200 (3)	0.3494 (13)	0.6242 (13)	0.128 (6)	0.50
H19'	0.1203	0.3654	0.5844	0.154*	0.50
C20'	0.052 (4)	0.416 (2)	0.712 (2)	0.197 (11)	0.50
H20D	-0.0339	0.3751	0.7128	0.295*	0.50
H20E	0.0113	0.4629	0.6793	0.295*	0.50
H20F	0.0940	0.4398	0.7580	0.295*	0.50
C21'	0.308 (3)	0.344 (2)	0.7492 (15)	0.185 (9)	0.50
H21D	0.3351	0.2828	0.7443	0.278*	0.50
H21E	0.2704	0.3509	0.7917	0.278*	0.50
H21F	0.4020	0.3798	0.7517	0.278*	0.50

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Hg1	0.0921 (3)	0.0868 (3)	0.0909 (3)	0.00588 (18)	-0.0045 (2)	-0.00621 (18)
C11	0.096 (2)	0.202 (3)	0.142 (3)	-0.024 (2)	0.017 (2)	-0.030 (3)
Cl2	0.139 (3)	0.0719 (13)	0.172 (3)	0.0028 (14)	-0.015 (2)	0.0193 (16)
N1	0.071 (5)	0.083 (4)	0.092 (5)	0.005 (4)	0.007 (4)	-0.001 (4)
N2	0.065 (5)	0.099 (5)	0.123 (7)	0.002 (4)	0.014 (5)	-0.031 (5)
N3	0.066 (5)	0.072 (4)	0.116 (6)	0.005 (3)	-0.003 (4)	-0.016 (4)
N4	0.079 (5)	0.073 (4)	0.065 (4)	-0.009 (3)	-0.014 (3)	-0.005 (3)
C1	0.062 (5)	0.080 (5)	0.081 (5)	0.007 (4)	0.002 (4)	0.002 (4)
C2	0.062 (5)	0.078 (5)	0.089 (6)	0.013 (4)	0.010 (5)	-0.010 (5)
C3	0.078 (7)	0.114 (8)	0.097 (7)	0.020 (5)	0.014 (6)	-0.012 (6)
C4	0.107 (8)	0.103 (7)	0.113 (8)	0.004 (6)	0.041 (7)	0.005 (6)

C5	0.072 (6)	0.086 (6)	0.126 (8)	-0.002 (5)	0.004 (6)	0.003 (6)
C6	0.096 (7)	0.108 (7)	0.101 (7)	0.026 (6)	-0.016 (6)	-0.034 (6)
C7	0.064 (5)	0.081 (5)	0.082 (6)	0.013 (4)	0.005 (5)	-0.010 (4)
C8	0.092 (7)	0.113 (7)	0.101 (7)	0.036 (6)	0.037 (6)	0.020 (6)
C9	0.113 (8)	0.122 (7)	0.066 (6)	0.029 (6)	0.017 (6)	0.021 (5)
C10	0.057 (5)	0.070 (4)	0.080 (6)	0.007 (4)	-0.006 (4)	-0.007 (4)
C11	0.083 (6)	0.100 (6)	0.075 (6)	0.016 (5)	0.023 (5)	-0.007 (5)
C12	0.104 (8)	0.107 (6)	0.063 (5)	0.025 (5)	0.012 (5)	-0.002 (5)
C13	0.066 (6)	0.090 (6)	0.119 (8)	0.007 (5)	-0.017 (5)	-0.029 (5)
C14	0.058 (5)	0.068 (4)	0.056 (4)	0.010 (3)	-0.027 (3)	-0.004 (3)
C15	0.063 (5)	0.069 (4)	0.068 (5)	0.012 (4)	-0.004 (4)	0.003 (4)
C16	0.086 (6)	0.067 (4)	0.083 (6)	0.007 (4)	-0.006 (5)	-0.011 (4)
C17	0.094 (7)	0.070 (5)	0.099 (7)	0.009 (5)	-0.006 (6)	-0.006 (5)
C18	0.082 (6)	0.070 (5)	0.083 (6)	0.007 (4)	-0.009 (5)	-0.003 (4)
O1	0.108 (7)	0.101 (4)	0.125 (6)	0.006 (5)	0.035 (5)	-0.011 (4)
N5	0.083 (5)	0.087 (6)	0.082 (6)	0.010 (4)	0.015 (5)	0.016 (5)
C19	0.125 (10)	0.134 (9)	0.122 (10)	0.015 (7)	0.023 (8)	0.009 (8)
C20	0.188 (13)	0.210 (14)	0.193 (14)	0.023 (9)	0.047 (10)	-0.009 (9)
C21	0.189 (12)	0.195 (12)	0.177 (12)	0.005 (9)	0.054 (9)	0.000 (9)
O1'	0.108 (7)	0.101 (4)	0.125 (6)	0.006 (5)	0.035 (5)	-0.011 (4)
N5'	0.083 (5)	0.087 (6)	0.082 (6)	0.010 (4)	0.015 (5)	0.016 (5)
C19'	0.125 (10)	0.134 (9)	0.122 (10)	0.015 (7)	0.023 (8)	0.009 (8)
C20'	0.188 (13)	0.210 (14)	0.193 (14)	0.023 (9)	0.047 (10)	-0.009 (9)
C21'	0.189 (12)	0.195 (12)	0.177 (12)	0.005 (9)	0.054 (9)	0.000 (9)

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

Hg1—N1	2.395 (7)	C11—H11	0.9300
Hg1—N4 ⁱ	2.308 (6)	C12—H12	0.9300
Hg1—Cl1	2.355 (3)	C13—H13A	0.9700
Hg1—Cl2	2.391 (3)	C13—H13B	0.9700
N1—C1	1.325 (10)	C14—C15	1.331 (10)
N1—C5	1.364 (11)	C14—H14	0.9300
N2—C2	1.342 (10)	C15—C16	1.392 (10)
N2—C6	1.479 (11)	C16—C17	1.346 (12)
N2—H2N	0.8800	C16—H16	0.9300
N3—C15	1.345 (10)	C17—C18	1.324 (11)
N3—C13	1.404 (10)	C17—H17	0.9300
N3—H3N	0.8800	C18—H18	0.9300
N4—C14	1.296 (9)	O1—C19	1.253 (10)
N4—C18	1.361 (9)	N5—C19	1.344 (10)
N4—Hg1 ⁱⁱ	2.308 (6)	N5—C20	1.440 (10)
C1—C2	1.392 (11)	N5—C21	1.448 (10)
C1—H1	0.9300	C19—H19	0.9300
C2—C3	1.414 (12)	C20—H20A	0.9600
C3—C4	1.351 (13)	C20—H20B	0.9600
C3—H3	0.9300	C20—H20C	0.9600
C4—C5	1.390 (13)	C21—H21A	0.9600

C4—H4	0.9300	C21—H21B	0.9600
C5—H5	0.9300	C21—H21C	0.9600
C6—C7	1.502 (11)	O1'—C19'	1.250 (10)
C6—H6A	0.9700	N5'—C19'	1.346 (10)
C6—H6B	0.9700	N5'—C20'	1.448 (10)
C7—C8	1.356 (12)	N5'—C21'	1.449 (10)
C7—C12	1.394 (11)	C19'—H19'	0.9300
C8—C9	1.347 (12)	C20'—H20D	0.9600
C8—H8	0.9300	C20'—H20E	0.9600
C9—C10	1.405 (12)	C20'—H20F	0.9600
C9—H9	0.9300	C21'—H21D	0.9600
C10—C11	1.348 (11)	C21'—H21E	0.9600
C10—C13	1.506 (11)	C21'—H21F	0.9600
C11—C12	1.378 (11)		
N4 ⁱ —Hg1—Cl1	110.0 (2)	C10—C11—C12	118.9 (8)
N4 ⁱ —Hg1—Cl2	100.07 (18)	C10—C11—H11	120.6
Cl1—Hg1—Cl2	137.49 (12)	C12—C11—H11	120.6
N4 ⁱ —Hg1—N1	97.9 (2)	C11—C12—C7	122.1 (8)
Cl1—Hg1—N1	104.0 (2)	C11—C12—H12	119.0
Cl2—Hg1—N1	100.6 (2)	C7—C12—H12	119.0
C1—N1—C5	117.9 (8)	N3—C13—C10	110.8 (7)
C1—N1—Hg1	120.9 (6)	N3—C13—H13A	109.5
C5—N1—Hg1	121.1 (6)	C10—C13—H13A	109.5
C2—N2—C6	121.5 (8)	N3—C13—H13B	109.5
C2—N2—H2N	119.2	C10—C13—H13B	109.5
C6—N2—H2N	119.2	H13A—C13—H13B	108.1
C15—N3—C13	121.9 (7)	N4—C14—C15	122.8 (7)
C15—N3—H3N	119.0	N4—C14—H14	118.6
C13—N3—H3N	119.0	C15—C14—H14	118.6
C14—N4—C18	120.1 (7)	C14—C15—N3	119.4 (7)
C14—N4—Hg1 ⁱⁱ	120.6 (5)	C14—C15—C16	117.4 (7)
C18—N4—Hg1 ⁱⁱ	119.3 (5)	N3—C15—C16	123.2 (8)
N1—C1—C2	122.4 (8)	C17—C16—C15	119.4 (8)
N1—C1—H1	118.8	C17—C16—H16	120.3
C2—C1—H1	118.8	C15—C16—H16	120.3
N2—C2—C1	117.5 (8)	C18—C17—C16	120.4 (8)
N2—C2—C3	124.0 (8)	C18—C17—H17	119.8
C1—C2—C3	118.4 (8)	C16—C17—H17	119.8
C4—C3—C2	119.7 (9)	C17—C18—N4	119.7 (8)
C4—C3—H3	120.1	C17—C18—H18	120.2
C2—C3—H3	120.1	N4—C18—H18	120.2
C3—C4—C5	118.3 (9)	C19—N5—C20	120 (3)
C3—C4—H4	120.9	C19—N5—C21	130 (3)
C5—C4—H4	120.9	C20—N5—C21	110 (3)
N1—C5—C4	123.3 (9)	O1—C19—N5	128 (3)
N1—C5—H5	118.4	O1—C19—H19	116.2
C4—C5—H5	118.4	N5—C19—H19	116.2

N2—C6—C7	115.3 (8)	C19'—N5'—C20'	132 (3)
N2—C6—H6A	108.4	C19'—N5'—C21'	118 (2)
C7—C6—H6A	108.4	C20'—N5'—C21'	111 (2)
N2—C6—H6B	108.4	O1'—C19'—N5'	121 (3)
C7—C6—H6B	108.4	O1'—C19'—H19'	119.7
H6A—C6—H6B	107.5	N5'—C19'—H19'	119.7
C8—C7—C12	118.5 (8)	N5'—C20'—H20D	109.5
C8—C7—C6	119.1 (8)	N5'—C20'—H20E	109.5
C12—C7—C6	122.4 (9)	H20D—C20'—H20E	109.5
C9—C8—C7	119.4 (8)	N5'—C20'—H20F	109.5
C9—C8—H8	120.3	H20D—C20'—H20F	109.5
C7—C8—H8	120.3	H20E—C20'—H20F	109.5
C8—C9—C10	122.6 (9)	N5'—C21'—H21D	109.5
C8—C9—H9	118.7	N5'—C21'—H21E	109.5
C10—C9—H9	118.7	H21D—C21'—H21E	109.5
C11—C10—C9	118.5 (7)	N5'—C21'—H21F	109.5
C11—C10—C13	120.1 (8)	H21D—C21'—H21F	109.5
C9—C10—C13	121.4 (8)	H21E—C21'—H21F	109.5
N4 ⁱ —Hg1—N1—C1	90.8 (6)	C8—C9—C10—C13	-177.7 (10)
C11—Hg1—N1—C1	-22.2 (7)	C9—C10—C11—C12	0.1 (13)
C12—Hg1—N1—C1	-167.3 (6)	C13—C10—C11—C12	178.2 (8)
N4 ⁱ —Hg1—N1—C5	-94.4 (7)	C10—C11—C12—C7	-2.3 (15)
C11—Hg1—N1—C5	152.6 (6)	C8—C7—C12—C11	4.0 (15)
C12—Hg1—N1—C5	7.5 (7)	C6—C7—C12—C11	-174.4 (9)
C5—N1—C1—C2	1.8 (12)	C15—N3—C13—C10	162.7 (8)
Hg1—N1—C1—C2	176.8 (6)	C11—C10—C13—N3	105.7 (10)
C6—N2—C2—C1	-178.2 (7)	C9—C10—C13—N3	-76.3 (12)
C6—N2—C2—C3	4.8 (14)	C18—N4—C14—C15	-2.5 (12)
N1—C1—C2—N2	-179.4 (8)	Hg1 ⁱⁱ —N4—C14—C15	179.4 (6)
N1—C1—C2—C3	-2.1 (13)	N4—C14—C15—N3	-176.6 (8)
N2—C2—C3—C4	178.9 (9)	N4—C14—C15—C16	4.0 (12)
C1—C2—C3—C4	1.8 (14)	C13—N3—C15—C14	-165.7 (8)
C2—C3—C4—C5	-1.4 (15)	C13—N3—C15—C16	13.6 (14)
C1—N1—C5—C4	-1.3 (14)	C14—C15—C16—C17	-1.8 (13)
Hg1—N1—C5—C4	-176.3 (7)	N3—C15—C16—C17	178.9 (9)
C3—C4—C5—N1	1.1 (16)	C15—C16—C17—C18	-1.9 (15)
C2—N2—C6—C7	73.2 (12)	C16—C17—C18—N4	3.5 (14)
N2—C6—C7—C8	40.2 (13)	C14—N4—C18—C17	-1.4 (12)
N2—C6—C7—C12	-141.5 (9)	Hg1 ⁱⁱ —N4—C18—C17	176.7 (7)
C12—C7—C8—C9	-3.4 (15)	C20—N5—C19—O1	0.0 (2)
C6—C7—C8—C9	175.0 (10)	C21—N5—C19—O1	-174 (3)
C7—C8—C9—C10	1.3 (17)	C20'—N5'—C19'—O1'	180.0 (4)
C8—C9—C10—C11	0.4 (15)	C21'—N5'—C19'—O1'	0.0 (3)

Symmetry codes: (i) $-x+5/2, y+1/2, -z+3/2$; (ii) $-x+5/2, y-1/2, -z+3/2$.

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
N2—H2N···O1	0.88	2.15	3.03 (3)	174
N2—H2N···O1'	0.88	2.11	2.99 (3)	180
N3—H3N···O1 ⁱⁱⁱ	0.88	2.14	3.01 (3)	166
N3—H3N···O1' ⁱⁱⁱ	0.88	2.13	2.98 (3)	162

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