

## 4,4'-Dichloro-2,2'-[imidazolidine-1,3-diylbis(methylene)]diphenol

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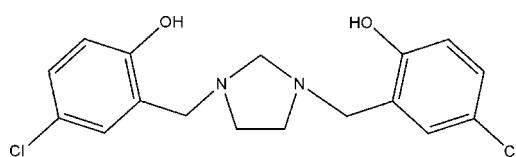
Received 25 August 2011; accepted 1 September 2011

Key indicators: single-crystal X-ray study;  $T = 120\text{ K}$ ; mean  $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$ ;  $R$  factor = 0.035;  $wR$  factor = 0.101; data-to-parameter ratio = 14.4.

The imidazolidine ring in the title compound,  $\text{C}_{17}\text{H}_{18}\text{Cl}_2\text{N}_2\text{O}_2$ , adopts a twist conformation. The observed conformation is stabilized by two intramolecular  $\text{O}-\text{H}\cdots\text{N}$  hydrogen bonds, with both N atoms acting as hydrogen-bond acceptors. The phenyl substituents are aligned at  $70.0(1)$  and  $76.6(1)^\circ$  with respect to the best plane through the five atoms of the imidazolidine ring. Weak intermolecular  $\text{C}-\text{H}\cdots\text{O}$  interactions stabilize the crystal packing.

### Related literature

For the preparation of the title compound, see: Rivera *et al.* (1993). For synthetic applications of these di-Mannich bases, see: Rivera & Quevedo (2004); Rivera *et al.* (2004). For a closely related structure, see: Rivera *et al.* (2010). For puckering parameters, see: Cremer & Pople (1975). For applications of tetrahydrosalens and heterocalixarenes in medicine and metal-complex catalysis, see: Balsells & Walsh (2000); Weber *et al.* (1996).



### Experimental

#### Crystal data

$\text{C}_{17}\text{H}_{18}\text{Cl}_2\text{N}_2\text{O}_2$   
 $M_r = 353.23$   
Monoclinic,  $P2_1/n$   
 $a = 10.8640(2)\text{ \AA}$   
 $b = 9.6125(2)\text{ \AA}$   
 $c = 16.7242(4)\text{ \AA}$   
 $\beta = 106.608(2)^\circ$

$$V = 1673.65(6)\text{ \AA}^3$$

$$Z = 4$$

Cu  $K\alpha$  radiation

$$\mu = 3.58\text{ mm}^{-1}$$

$$T = 120\text{ K}$$

$$0.42 \times 0.37 \times 0.25\text{ mm}$$

### Data collection

Oxford Diffraction Xcalibur Atlas Gemini ultra diffractometer  
Absorption correction: analytical (*CrysAlis PRO*; Oxford Diffraction, 2010)  
 $T_{\min} = 0.669$ ,  $T_{\max} = 0.777$

19547 measured reflections  
2994 independent reflections  
2772 reflections with  $I > 2\sigma(I)$ ,  
 $R_{\text{int}} = 0.040$

### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.035$   
 $wR(F^2) = 0.101$   
 $S = 1.04$   
2994 reflections

208 parameters  
H-atom parameters constrained  
 $\Delta\rho_{\max} = 0.29\text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.30\text{ e \AA}^{-3}$

**Table 1**  
Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O1—H1O1 $\cdots$ N1	0.97	1.77	2.6524 (17)	149
O2—H1O2 $\cdots$ N2	0.96	1.77	2.6515 (17)	150
C4—H4B $\cdots$ O2 <sup>i</sup>	0.97	2.52	3.466 (2)	163
C9—H9 $\cdots$ O2 <sup>ii</sup>	0.93	2.47	3.395 (2)	172
C11—H11B $\cdots$ O1 <sup>iii</sup>	0.97	2.58	3.482 (2)	154

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

Data collection: *CrysAlis PRO* (Oxford Diffraction, 2010); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *DIAMOND* (Brandenburg & Putz, 2005); software used to prepare material for publication: *publCIF* (Westrip, 2010).

We acknowledge the Dirección de Investigaciones, Sede Bogotá (DIB) de la Universidad Nacional de Colombia for financial support of this work, as well as the Institutional research plan No. AVOZ10100521 of the Institute of Physics and the project Praemium Academiae of the Academy of Science of the Czech Republic. JSB acknowledges the Vice-rectoría Académica de la Universidad Nacional de Colombia for a fellowship.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT5631).

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# supporting information

*Acta Cryst.* (2011). E67, o2581 [https://doi.org/10.1107/S1600536811035677]

## 4,4'-Dichloro-2,2'-[imidazolidine-1,3-diylbis(methylene)]diphenol

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### S1. Comment

In connection with our synthetic studies on heterocyclic compounds we earlier synthesized a series of *di*-Mannich bases named 2,2'-(imidazolidine-1,3-diyldimethanediyl)bis(4-substitutedphenol) by reaction of appropriate phenols with 1,3,6,8-tetraazatricyclo[4.4.1.1<sup>3,8</sup>]dodecane (Rivera *et al.*, 1993). They are promising synthetic intermediates for the synthesis of tetrahydrosalens (Rivera, Quevedo, Navarro & Maldonado, 2004) and heterocalixarenes (Rivera & Quevedo, 2004), which find wide use in both medicine and metal-complex catalysis (Balsells & Walsh, 2000; Weber *et al.* 1996). These Mannich bases are convenient models for studying the nature of hydrogen bonding and weak noncovalent interactions, which play a key role in biological processes and design of complex structures.

We report here the structure of the title compound (**I**) (Fig. 1), which was prepared according to the previously reported procedure (Rivera *et al.*, 1993) but using the intriguing aminal 1,3,6,8-tetraazatricyclo[4.3.1.1<sup>3,8</sup>]undecane.

Recrystallization from methanol by slow evaporation over a period of one week affording crystals suitable for X-ray analysis.

The asymmetric unit of (**I**), Fig 1, contains one independent 2,2'-(imidazolidine-1,3-diylbis(methylene))bis(4-chlorophenol) molecule. Distances and angles are similar to those observed before in the closely related structure 4,4'-dichloro-2,2'-(3aR,7aR/3aS,7aS)-2,3,3a,4,5,6,7,7a-octahydro-1*H*-1,3-benzimidazole-1,3-diylbis(methylene)]diphenol (Rivera *et al.*, 2010). The imidazolidine ring is in a twist conformation on C1–N2 with Q(2) 0.402 (2) Å and  $\varphi$  52.7 (2) $^\circ$  (Cremer & Pople, 1975). Its central ring makes an angle of 70.0 (1) $^\circ$  and 76.6 (1) $^\circ$  with the planar phenyl rings (C5—C10) and (C12—C17) respectively. The crystal structure has two intramolecular hydrogen bonds and three C—H···O intermolecular hydrogen bonds (Table 1). The unit cell contains four molecules of the title compound (**I**), which form pairs of hydrogen bonded dimers (Table 1, Figs. 2). Neighboring pairs of these dimers are orthogonally arranged with respect to each other. Lattice binding is provided principally by C—H···O interactions, shown in Figure 2. The chains, aligned along the *c* axis, are further linked together *via* cross-linking weaker C—H···O interactions (Table 1).

### S2. Experimental

For the originally reported synthesis, see: Rivera *et al.* (1993)

### S3. Refinement

All H atoms could be located in a difference Fourier synthesis. Nevertheless, H atoms were refined as riding with H bonded to O at the positions where they were found and C—H distances of 0.93 Å for aromatic H and C—H = 0.97 Å for methylene groups. All H atoms were refined with displacement coefficients  $U_{\text{iso}}(\text{H})$  set to  $1.5U_{\text{eq}}(\text{O})$  for hydroxyl groups and to  $1.2U_{\text{eq}}(\text{C})$  for the CH- and CH<sub>2</sub>- groups.

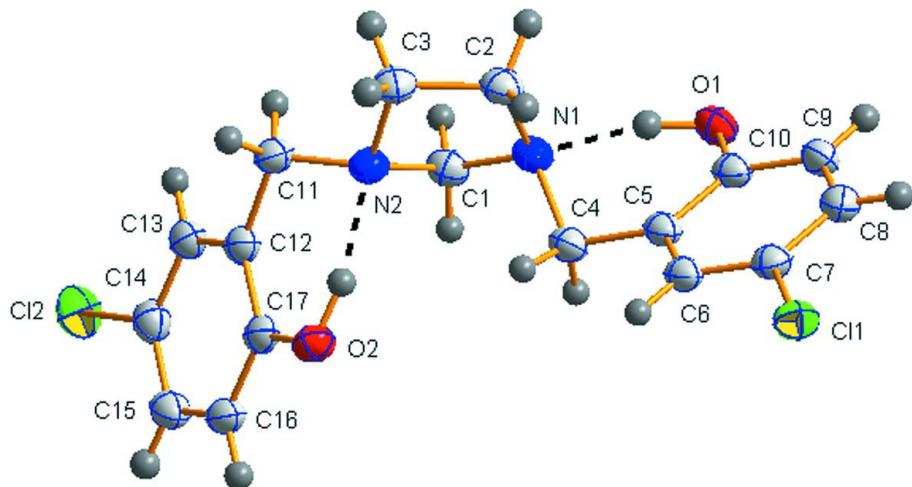
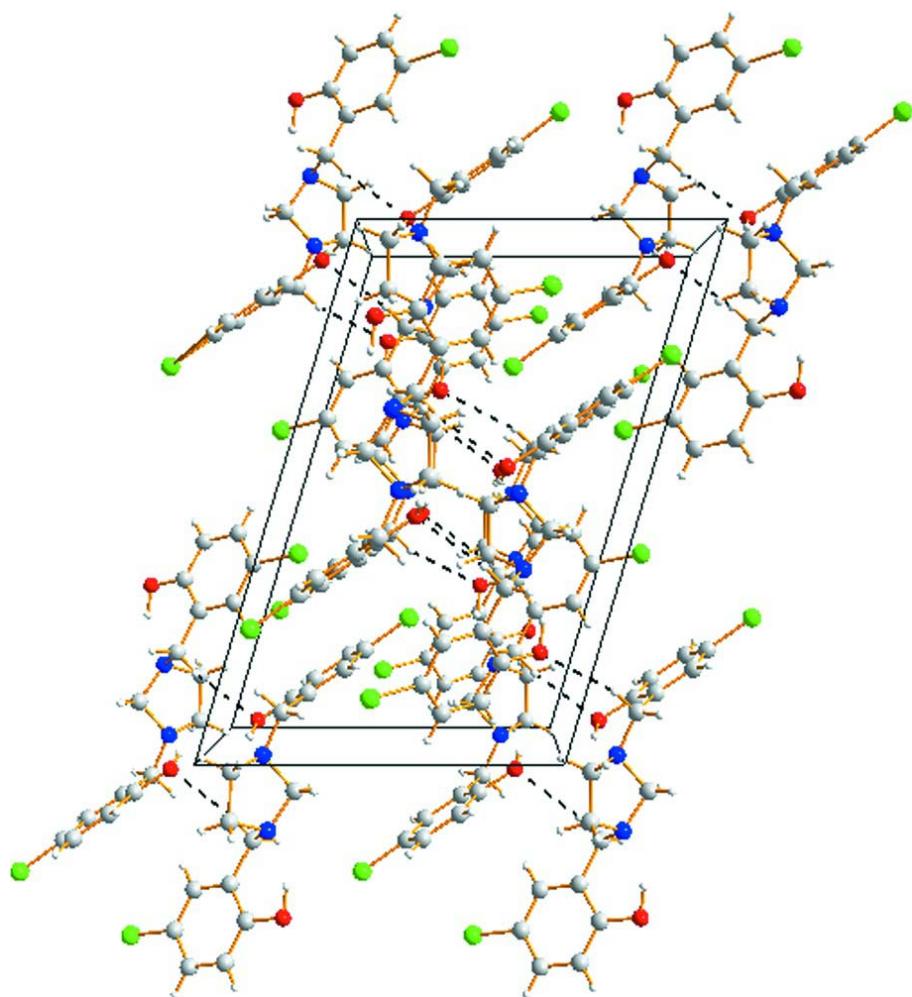


Figure 1

A view of (I) with the numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.



**Figure 2**

Packing of the molecules of the title compound view along  $b$  axis.

**4,4'-Dichloro-2,2'-[imidazolidine-1,3-diylbis(methylene)]diphenol***Crystal data*

$C_{17}H_{18}Cl_2N_2O_2$   
 $M_r = 353.23$   
Monoclinic,  $P2_1/n$   
Hall symbol: -P 2yn  
 $a = 10.8640 (2)$  Å  
 $b = 9.6125 (2)$  Å  
 $c = 16.7242 (4)$  Å  
 $\beta = 106.608 (2)^\circ$   
 $V = 1673.65 (6)$  Å<sup>3</sup>  
 $Z = 4$

$F(000) = 736$   
 $D_x = 1.402 \text{ Mg m}^{-3}$   
Cu  $K\alpha$  radiation,  $\lambda = 1.5418$  Å  
Cell parameters from 9456 reflections  
 $\theta = 4.2\text{--}67.1^\circ$   
 $\mu = 3.58 \text{ mm}^{-1}$   
 $T = 120$  K  
Prism, colourless  
 $0.42 \times 0.37 \times 0.25$  mm

*Data collection*

Oxford Diffraction Xcalibur Atlas Gemini ultra diffractometer  
Radiation source: Enhance Ultra (Cu) X-ray Source  
Mirror monochromator  
Detector resolution: 10.3784 pixels mm<sup>-1</sup>  
Rotation method data acquisition using  $\omega$  scans  
Absorption correction: analytical  
(CrysAlis PRO; Oxford Diffraction, 2010)

$T_{\min} = 0.669$ ,  $T_{\max} = 0.777$   
19547 measured reflections  
2994 independent reflections  
2772 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.040$   
 $\theta_{\max} = 67.2^\circ$ ,  $\theta_{\min} = 4.4^\circ$   
 $h = -12 \rightarrow 12$   
 $k = -11 \rightarrow 11$   
 $l = -19 \rightarrow 17$

*Refinement*

Refinement on  $F^2$   
Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.035$   
 $wR(F^2) = 0.101$   
 $S = 1.04$   
2994 reflections  
208 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-atom parameters constrained  
 $w = 1/[\sigma^2(F_o^2) + (0.0604P)^2 + 0.5161P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.29 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.30 \text{ e } \text{\AA}^{-3}$

*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. The H atoms were all located in a difference map, but those attached to carbon atoms were repositioned geometrically. The isotropic temperature parameters of hydrogen atoms were calculated as  $1.2 * U_{\text{eq}}$  of the parent atom. The distance between hydrogen and oxygen atom in hydroxyl group was fixed to the distance 0.87 Å.

*Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.21837 (14)	0.21659 (16)	0.42784 (10)	0.0322 (3)
H1A	0.1749	0.1280	0.4260	0.039*
H1B	0.1608	0.2905	0.4337	0.039*
C2	0.36940 (16)	0.33596 (16)	0.38001 (10)	0.0368 (4)
H2A	0.3437	0.4280	0.3576	0.044*
H2B	0.4409	0.3050	0.3607	0.044*
C3	0.40707 (16)	0.33848 (17)	0.47479 (10)	0.0372 (4)
H3A	0.4991	0.3275	0.4981	0.045*
H3B	0.3809	0.4250	0.4949	0.045*
C4	0.29989 (15)	0.10433 (15)	0.32188 (10)	0.0320 (3)
H4A	0.2325	0.0354	0.3154	0.038*
H4B	0.3766	0.0692	0.3621	0.038*
C5	0.32658 (14)	0.12712 (15)	0.23931 (9)	0.0303 (3)
C6	0.42918 (14)	0.06128 (15)	0.22089 (9)	0.0312 (3)
H6	0.4843	0.0051	0.2607	0.037*
C7	0.44951 (15)	0.07915 (16)	0.14360 (10)	0.0326 (3)
C8	0.37107 (16)	0.16444 (16)	0.08356 (10)	0.0354 (4)
H8	0.3864	0.1763	0.0320	0.043*
C9	0.26963 (16)	0.23171 (16)	0.10149 (10)	0.0351 (4)
H9	0.2168	0.2901	0.0619	0.042*
C10	0.24590 (14)	0.21281 (15)	0.17805 (10)	0.0316 (3)
C11	0.31970 (15)	0.23442 (16)	0.58004 (10)	0.0330 (3)
H11A	0.2605	0.3103	0.5793	0.040*
H11B	0.4013	0.2578	0.6198	0.040*
C12	0.26818 (14)	0.10368 (16)	0.60816 (9)	0.0299 (3)
C13	0.18268 (15)	0.11162 (16)	0.65594 (9)	0.0328 (3)
H13	0.1516	0.1976	0.6667	0.039*
C14	0.14386 (17)	-0.00807 (17)	0.68735 (11)	0.0358 (4)
C15	0.18658 (17)	-0.13748 (17)	0.67129 (10)	0.0387 (4)
H15	0.1597	-0.2173	0.6929	0.046*
C16	0.26995 (16)	-0.14685 (17)	0.62260 (10)	0.0361 (4)
H16	0.2985	-0.2336	0.6109	0.043*
C17	0.31142 (15)	-0.02753 (16)	0.59103 (10)	0.0311 (3)
N1	0.25961 (12)	0.23659 (13)	0.35315 (8)	0.0322 (3)
N2	0.33788 (12)	0.21975 (13)	0.49679 (8)	0.0300 (3)
O1	0.14312 (10)	0.27831 (11)	0.19253 (7)	0.0364 (3)
H1O1	0.1592	0.2802	0.2528	0.044*
O2	0.39647 (10)	-0.04067 (12)	0.54511 (7)	0.0348 (3)
H1O2	0.4004	0.0494	0.5208	0.042*
Cl1	0.57874 (4)	-0.00643 (4)	0.12223 (3)	0.03906 (15)
Cl2	0.04238 (5)	0.00586 (5)	0.75129 (3)	0.04919 (16)

Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C1	0.0287 (7)	0.0310 (8)	0.0365 (8)	0.0010 (6)	0.0086 (6)	0.0027 (6)
C2	0.0402 (9)	0.0281 (8)	0.0418 (9)	-0.0043 (6)	0.0116 (7)	0.0039 (6)
C3	0.0384 (8)	0.0313 (8)	0.0417 (9)	-0.0080 (6)	0.0111 (7)	-0.0005 (6)
C4	0.0335 (8)	0.0265 (7)	0.0346 (8)	0.0011 (6)	0.0073 (6)	0.0047 (6)
C5	0.0323 (7)	0.0237 (7)	0.0317 (8)	-0.0034 (6)	0.0039 (6)	0.0025 (6)
C6	0.0316 (7)	0.0245 (7)	0.0332 (8)	-0.0015 (6)	0.0025 (6)	0.0014 (6)
C7	0.0343 (8)	0.0257 (7)	0.0368 (8)	-0.0026 (6)	0.0083 (6)	-0.0026 (6)
C8	0.0437 (9)	0.0294 (8)	0.0321 (8)	-0.0033 (6)	0.0090 (7)	0.0008 (6)
C9	0.0404 (9)	0.0273 (8)	0.0328 (8)	0.0009 (6)	0.0025 (7)	0.0046 (6)
C10	0.0312 (7)	0.0240 (7)	0.0360 (8)	-0.0020 (6)	0.0043 (6)	0.0008 (6)
C11	0.0367 (8)	0.0263 (7)	0.0355 (8)	-0.0008 (6)	0.0094 (7)	-0.0034 (6)
C12	0.0289 (7)	0.0286 (8)	0.0290 (7)	0.0006 (6)	0.0029 (6)	-0.0009 (6)
C13	0.0337 (8)	0.0322 (8)	0.0301 (8)	0.0025 (6)	0.0052 (6)	0.0002 (6)
C14	0.0339 (8)	0.0402 (9)	0.0307 (8)	-0.0035 (6)	0.0053 (7)	0.0011 (6)
C15	0.0442 (9)	0.0320 (8)	0.0350 (8)	-0.0078 (7)	0.0036 (7)	0.0040 (7)
C16	0.0415 (9)	0.0266 (7)	0.0351 (8)	0.0008 (6)	0.0026 (7)	-0.0015 (6)
C17	0.0299 (7)	0.0296 (7)	0.0291 (8)	0.0007 (6)	0.0008 (6)	-0.0018 (6)
N1	0.0327 (6)	0.0264 (6)	0.0368 (7)	0.0001 (5)	0.0089 (6)	0.0035 (5)
N2	0.0286 (6)	0.0279 (6)	0.0330 (7)	-0.0015 (5)	0.0081 (5)	-0.0007 (5)
O1	0.0341 (6)	0.0341 (6)	0.0385 (6)	0.0057 (4)	0.0063 (5)	0.0060 (5)
O2	0.0353 (6)	0.0299 (5)	0.0395 (6)	0.0045 (4)	0.0109 (5)	-0.0021 (5)
Cl1	0.0404 (3)	0.0345 (2)	0.0437 (3)	0.00186 (14)	0.01431 (19)	-0.00169 (15)
Cl2	0.0477 (3)	0.0585 (3)	0.0470 (3)	-0.00269 (18)	0.0226 (2)	0.00619 (18)

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

C1—N1	1.455 (2)	C8—H8	0.9300
C1—N2	1.470 (2)	C9—C10	1.388 (2)
C1—H1A	0.9700	C9—H9	0.9300
C1—H1B	0.9700	C10—O1	1.3625 (19)
C2—N1	1.494 (2)	C11—N2	1.468 (2)
C2—C3	1.520 (2)	C11—C12	1.505 (2)
C2—H2A	0.9700	C11—H11A	0.9700
C2—H2B	0.9700	C11—H11B	0.9700
C3—N2	1.470 (2)	C12—C13	1.390 (2)
C3—H3A	0.9700	C12—C17	1.404 (2)
C3—H3B	0.9700	C13—C14	1.380 (2)
C4—N1	1.4873 (19)	C13—H13	0.9300
C4—C5	1.506 (2)	C14—C15	1.381 (2)
C4—H4A	0.9700	C14—Cl2	1.7466 (18)
C4—H4B	0.9700	C15—C16	1.383 (3)
C5—C6	1.391 (2)	C15—H15	0.9300
C5—C10	1.408 (2)	C16—C17	1.390 (2)
C6—C7	1.383 (2)	C16—H16	0.9300
C6—H6	0.9300	C17—O2	1.366 (2)

C7—C8	1.385 (2)	O1—H1O1	0.9725
C7—Cl1	1.7500 (16)	O2—H1O2	0.9623
C8—C9	1.383 (2)		
N1—C1—N2	104.54 (12)	C8—C9—H9	119.7
N1—C1—H1A	110.8	C10—C9—H9	119.7
N2—C1—H1A	110.8	O1—C10—C9	118.76 (14)
N1—C1—H1B	110.8	O1—C10—C5	120.82 (14)
N2—C1—H1B	110.8	C9—C10—C5	120.42 (14)
H1A—C1—H1B	108.9	N2—C11—C12	112.28 (12)
N1—C2—C3	106.13 (12)	N2—C11—H11A	109.1
N1—C2—H2A	110.5	C12—C11—H11A	109.1
C3—C2—H2A	110.5	N2—C11—H11B	109.1
N1—C2—H2B	110.5	C12—C11—H11B	109.1
C3—C2—H2B	110.5	H11A—C11—H11B	107.9
H2A—C2—H2B	108.7	C13—C12—C17	118.97 (14)
N2—C3—C2	104.13 (12)	C13—C12—C11	120.21 (14)
N2—C3—H3A	110.9	C17—C12—C11	120.70 (14)
C2—C3—H3A	110.9	C14—C13—C12	120.02 (15)
N2—C3—H3B	110.9	C14—C13—H13	120.0
C2—C3—H3B	110.9	C12—C13—H13	120.0
H3A—C3—H3B	108.9	C13—C14—C15	121.39 (16)
N1—C4—C5	110.56 (12)	C13—C14—Cl2	118.96 (13)
N1—C4—H4A	109.5	C15—C14—Cl2	119.62 (13)
C5—C4—H4A	109.5	C14—C15—C16	119.09 (15)
N1—C4—H4B	109.5	C14—C15—H15	120.5
C5—C4—H4B	109.5	C16—C15—H15	120.5
H4A—C4—H4B	108.1	C15—C16—C17	120.51 (15)
C6—C5—C10	118.47 (14)	C15—C16—H16	119.7
C6—C5—C4	120.89 (13)	C17—C16—H16	119.7
C10—C5—C4	120.61 (14)	O2—C17—C16	118.82 (14)
C7—C6—C5	120.18 (14)	O2—C17—C12	121.17 (14)
C7—C6—H6	119.9	C16—C17—C12	120.00 (15)
C5—C6—H6	119.9	C1—N1—C4	112.49 (12)
C6—C7—C8	121.45 (14)	C1—N1—C2	103.98 (12)
C6—C7—Cl1	118.97 (12)	C4—N1—C2	111.25 (12)
C8—C7—Cl1	119.57 (12)	C11—N2—C1	114.70 (12)
C9—C8—C7	118.89 (15)	C11—N2—C3	112.33 (12)
C9—C8—H8	120.6	C1—N2—C3	102.70 (12)
C7—C8—H8	120.6	C10—O1—H1O1	106.2
C8—C9—C10	120.56 (14)	C17—O2—H1O2	105.8

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

$D\text{—H}\cdots A$	$D\text{—H}$	$H\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
O1—H1O1…N1	0.97	1.77	2.6524 (17)	149
O2—H1O2…N2	0.96	1.77	2.6515 (17)	150
C4—H4B…O2 <sup>i</sup>	0.97	2.52	3.466 (2)	163

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C9—H9···O2 <sup>ii</sup>	0.93	2.47	3.395 (2)	172
C11—H11B···O1 <sup>iii</sup>	0.97	2.58	3.482 (2)	154

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