

4-*tert*-Butylpyridinium chloride–4,4'-(propane-2,2-diyl)bis(2,6-dimethylphenol)–toluene (2/2/1)

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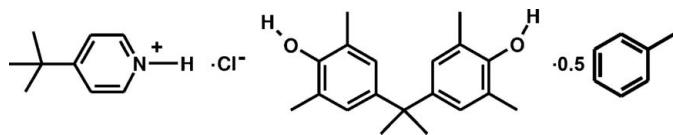
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Key indicators: single-crystal X-ray study; $T = 150\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$; disorder in solvent or counterion; R factor = 0.052; wR factor = 0.114; data-to-parameter ratio = 13.2.

In the title solvated salt, $\text{C}_9\text{H}_{14}\text{N}^+\cdot\text{Cl}^-\cdot\text{C}_{19}\text{H}_{24}\text{O}_2\cdot0.5\text{C}_7\text{H}_7$, two molecules of 4,4'-(propane-2,2-diyl)bis(2,6-dimethylphenol) are linked via $\text{O}-\text{H}\cdots\text{Cl}$ hydrogen bonds to two chloride ions, each of which is also engaged in $\text{N}-\text{H}\cdots\text{Cl}$ hydrogen bonding to a 4-*tert*-butylpyridinium cation, giving a cyclic hydrogen-bonded entity centred at $\frac{1}{2}, \frac{1}{2}, \frac{1}{2}$. The toluene solvent molecule resides in the lattice and resides on an inversion centre; the disorder of the methyl group requires it to have a site-occupancy factor of 0.5. No crystal packing channels are observed.

Related literature

For general background to hydrogen-bond structural information, see: Hamilton & Ibers (1968). For hydrogen bonding between phenols and nitrogen bases, see: Coupar *et al.* (1997); Steiner *et al.* (2000). For hydrogen bonding in phenol molecules, see: Prout *et al.* (1988); Zierner & Surygina, (2000). For the structure of a related bis-phenol molecule, see: Okada (1996). For hydrogen bonds between pyridinium hydrochloride and OH-containing molecules, see: Sykora & Cioffi (2007); Hossain *et al.* (1988). For structural data pertaining to pyridinium hydrohalides, see: Faber *et al.* (1999); Hensen *et al.* (1988); Mootz & Hocken (1989); van de Streek *et al.* (2010).



Experimental

Crystal data

$\text{C}_9\text{H}_{14}\text{N}^+\cdot\text{Cl}^-\cdot\text{C}_{19}\text{H}_{24}\text{O}_2\cdot0.5\text{C}_7\text{H}_7$	$V = 2840.0 (3)\text{ \AA}^3$
$M_r = 501.60$	$Z = 4$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
$a = 13.5656 (9)\text{ \AA}$	$\mu = 0.16\text{ mm}^{-1}$
$b = 14.3215 (9)\text{ \AA}$	$T = 150\text{ K}$
$c = 15.787 (1)\text{ \AA}$	$0.39 \times 0.28 \times 0.02\text{ mm}$
$\beta = 112.186 (1)^{\circ}$	

Data collection

Siemens SMART diffractometer	14535 measured reflections
Absorption correction: multi-scan (Blessing, 1995)	4989 independent reflections
$T_{\min} = 0.700$, $T_{\max} = 0.821$	3253 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.053$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.052$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.114$	$\Delta\rho_{\text{max}} = 0.22\text{ e \AA}^{-3}$
$S = 1.08$	$\Delta\rho_{\text{min}} = -0.20\text{ e \AA}^{-3}$
4989 reflections	
379 parameters	

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N20—H20 \cdots Cl	1.06 (4)	1.98 (4)	3.035 (3)	173 (3)
O2—H2 \cdots Cl ⁱ	0.85 (3)	2.30 (3)	3.079 (2)	153 (3)
O1—H1 \cdots Cl	0.90 (4)	2.27 (4)	3.118 (2)	157 (3)

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

Data collection: SMART (Siemens, 1995); cell refinement: SAINT (Siemens, 1995); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 2012); software used to prepare material for publication: SHELXL97.

We are grateful to Ms Tanya Groutso of the University of Auckland for the data collection.

Supporting information for this paper is available from the IUCr electronic archives (Reference: GG2132).

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supplementary materials

Acta Cryst. (2014). E70, o413–o414 [doi:10.1107/S1600536814004942]

4-*tert*-Butylpyridinium chloride–4,4'-(propane-2,2-diyl)bis(2,6-dimethylphenol)–toluene (2/2/1)

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1. Introduction

2. Experimental

2.1. Synthesis and crystallization

The title compound crystallized from toluene as very thin colourless plates from the reaction between TiCl_4 and 4,4'-(propane-2,2-diyl)bis(2,6-dimethylphenol) in the presence of 4-*tert*-butylpyridine.

2.2. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 1. All H atoms (except H1 and H2) were placed in calculated positions and in the refinement allowed to ride on the C atoms to which they were attached. H1 and H2 were located from a difference map and their positions were then allowed to ride on their oxygen atoms. All H atoms were assigned fixed isotropic thermal parameters. Disorder was associated with *tert*-butyl substituent on the pyridinium ion and two alternative positions were found for this group with site occupancy factors of 0.62 and 0.38. Both were refined as a rigid group. A half-weighted molecule of toluene disordered across a centre of symmetry was identified in the asymmetric unit. The numbering system used for the diphenol and pyridinium moieties is indicated in figure 1, where the linking of the moieties *via* the chloride ion is shown.

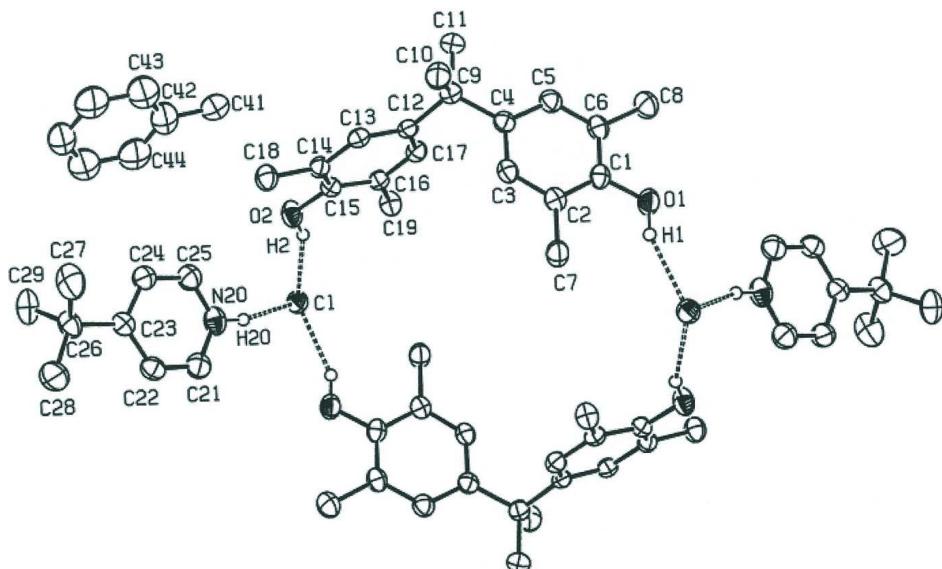
3. Results and discussion

As part of a study into the reactions of TiCl_4 with various phenols in the presence of pyridines, crystals of the title compound were isolated from a toluene solution. X-ray structural analysis revealed two molecules of $(\text{CH}_3)_2\text{C}(\text{C}_6\text{H}_2\text{Me}_2\text{OH})_2$ linked together by hydrogen bonds to 2 chloride ions which in turn are further linked to two 4-*tert*-butyl pyridinium-H ions about a centre of symmetry to form a ring of atoms (Figure 1). No Ti atom was present. Distances of 3.118 (2) and 3.079 (2) Å were observed for $\text{Cl}\cdots\text{O}1$ and $\text{Cl}\cdots\text{O}2$ respectively. The H contact distances and $\text{O}-\text{H}\cdots\text{Cl}$ angles observed were 2.27 (4) Å, 157 (3) ° and 2.30 (3) Å, 153 (3) ° for H1 and H2, respectively. The Cl is also H-bound to H20 of the pyridinium-H ion with $\text{Cl}\cdots\text{N}20$ and $\text{Cl}\cdots\text{H}20$ approaches of 3.035 (3) and 1.98 (4) Å respectively. The angle $\text{N}20-\text{H}20\cdots\text{Cl}$ is 173 (3) °. These distances are typical of such H-bonded distances as described elsewhere (Hamilton & Ibers, 1968). H-bonds between N bases and phenolic moieties are not uncommon (Coupar *et al.* 1997) and the interaction is often strong (Steiner *et al.* 2000). However a Cl ion linking a phenol and an ammine hydrochloride as occurs in the title compound has apparently not been observed before. A single link between pyridinium hydrochloride and the OH hydrogen of triphenylmethanol has been observed (Sykora & Cioffi, 2007) as has a similar link with the OH group of 1,1,3,3-tetraphenyl-1,3-disiloxanediol (Hossain *et al.* 1988).

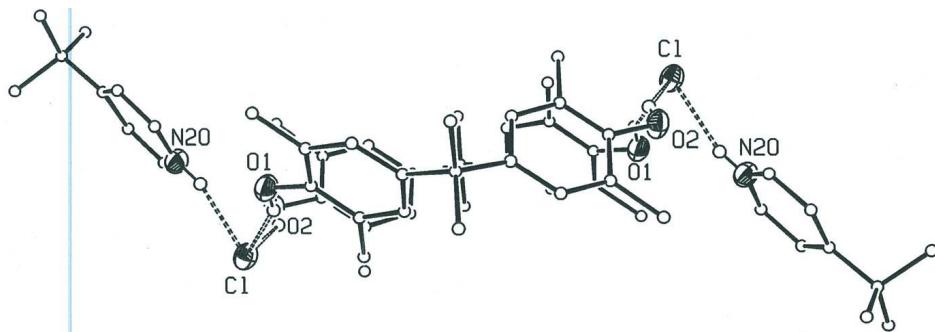
The N20 to Cl separation for the 4-*tert*-butylpyridinium hydrochloride in the present compound [3.035 (3) Å] compares with that found for 4-methylpyridinium hydrochloride itself at 2.999 (2) Å, 2.981 (4) Å in 3-methylpyridinium hydrochloride and 3.162 (2) Å in 4-methylpyridinium hydrobromide (Faber *et al.* 1999). Taking into account the imprecise positioning of hydrogen atoms in X-ray crystal structures, in the latter three compounds the N—H···Cl bond angles are 180 °, 177 (5) ° [and 165 (5) °] and 173 (3) ° whereas in the present compound this angle is 173 (3) °. For $(C_6H_5)_3COH \cdot C_5H_6N^+ \cdot Cl^-$ the N to Cl separation is 3.008 (2) Å and the N—H···Cl bond angle is 169 ° (Sykora & Cioffi, 2007). It is noted here that for pyridinium hydrochloride itself two polymorphs found by crystallography (Hensen *et al.* 1988; Mootz & Hocken, 1989) have been substantiated by dispersion-corrected density functional theory calculations (van de Streek *et al.* 2010).

The O to Cl separations in the title compound are 3.079 (2) and 3.118 (2) Å and these compare with 3.134 (1) Å in $(C_6H_5)_3COH \cdot C_5H_6N^+ \cdot Cl^-$ (Sykora & Cioffi, 2007). The C1—O1 bond length in the present compound [1.382 (3) Å] is not significantly different from the C15—O2 bond length [1.378 (3) Å] at the opposite end of the bis-phenol. In comparison, 2,4,6-trimethylphenol which is structurally similar to the phenolic portion of the present molecule but contains a hydrogen bonded O—H···OH system, has a C—O bond length of 1.386 (2) Å (Ziemer & Surygina, 2000). 2,6-Diisopropylphenol which also has a hydrogen bonded O—H···OH system has C—O bond lengths of 1.480 (7) Å (C—O—H···O section) and 1.382 (8) Å (C—O···H section) (Prout *et al.* 1988).

The chlorines in the present lattice form a pyramidal structure in the interactions with their 3 H-bonded neighbours (Figure 2). Angles are 118, 79 and 69 ° for O1···Cl···O2, O1···Cl···N20 and O2···Cl···N20 respectively and 117, 82 and 76 ° for H1···Cl···H2, H1···Cl···H20 and H2···Cl···H20 respectively. Within the phenolic moiety and the pyridinium-H ion bond distances and angles are unremarkable. The two phenyl rings of the former (C1—C6 and C12—C17) are inclined to one another at an angle of 84.78 (8) °, a value which is similar to the values reported [86.9 (2), 83.6 (2) and 79.2 (2) °] for the three independent molecules in the asymmetric unit of $(CH_3)_2C(C_6H_4OH)_2$ (*bisphenol A*) (Okada, 1996). Two disordered sites were found for the terminal tertiary butyl substituent on the pyridinium-H group. Toluene solvent molecules occupy space between the H-bound units in the crystal lying across centres of symmetry. No close approaches of this solvent to the other molecules are observed. The crystal packing shows that no channels are developed in the overall structure.

**Figure 1**

ORTEP diagram of the title compound showing the numbering system and the H-bonding network. H not involved in this bonding have been omitted for reasons of clarity. Note also that only one of the disordered toluene methyl substituents has been included. Thermal ellipsoids have been drawn at the 50% probability level.

**Figure 2**

Edge-on ORTEP diagram showing the pyramidal structure about the Cl ions

4-*tert*-Butylpyridinium chloride–4,4'-(propane-2,2-diyl)bis(2,6-dimethylphenol)–toluene (2/2/1)

Crystal data



$M_r = 501.60$

Monoclinic, $P2_1/n$

Hall symbol: -P 2yn

$a = 13.5656 (9) \text{ \AA}$

$b = 14.3215 (9) \text{ \AA}$

$c = 15.787 (1) \text{ \AA}$

$\beta = 112.186 (1)^\circ$

$V = 2840.0 (3) \text{ \AA}^3$

$Z = 4$

$F(000) = 1082$

$D_x = 1.173 \text{ Mg m}^{-3}$

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

Cell parameters from 7233 reflections

$\theta = 2.0\text{--}25.1^\circ$

$\mu = 0.16 \text{ mm}^{-1}$

$T = 150 \text{ K}$

Thin plate, colourless

$0.39 \times 0.28 \times 0.02 \text{ mm}$

Data collection

Siemens SMART
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
area-detector ω scan
Absorption correction: multi-scan
(Blessing, 1995)
 $T_{\min} = 0.700$, $T_{\max} = 0.821$
14535 measured reflections
4989 independent reflections
3253 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.053$
 $\theta_{\max} = 25.1^\circ$, $\theta_{\min} = 2.0^\circ$
 $h = -16 \rightarrow 10$
 $k = -17 \rightarrow 15$
 $l = -18 \rightarrow 18$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.052$
 $wR(F^2) = 0.114$
 $S = 1.08$
4989 reflections
379 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
 $w = 1/[\sigma^2(F_o^2) + (0.0296P)^2 + 1.4177P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.009$
 $\Delta\rho_{\max} = 0.22 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.20 \text{ e } \text{\AA}^{-3}$
Extinction correction: *SHELXL97* (Sheldrick,
2008), $F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$
Extinction coefficient: 0.0025 (5)

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>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
N20	0.5777 (2)	0.96003 (16)	0.67143 (17)	0.0415 (6)	
Cl	0.37477 (6)	0.87567 (5)	0.53461 (4)	0.0384 (2)	
O1	0.52861 (16)	0.85019 (13)	0.43127 (13)	0.0406 (5)	
C1	0.5296 (2)	0.78792 (17)	0.36475 (17)	0.0305 (6)	
C2	0.4550 (2)	0.71593 (17)	0.33306 (17)	0.0306 (6)	
C3	0.4654 (2)	0.65504 (17)	0.26836 (17)	0.0305 (6)	
H3	0.4158	0.6053	0.2469	0.037*	
C4	0.5451 (2)	0.66375 (17)	0.23373 (17)	0.0299 (6)	
C5	0.6164 (2)	0.73712 (17)	0.26655 (17)	0.0339 (6)	
H5	0.6712	0.7449	0.2435	0.041*	
C6	0.6106 (2)	0.79978 (17)	0.33215 (18)	0.0329 (6)	
C7	0.3669 (2)	0.70383 (19)	0.36795 (19)	0.0400 (7)	
H7A	0.3209	0.6520	0.3355	0.060*	
H7B	0.3249	0.7614	0.3575	0.060*	
H7C	0.3975	0.6902	0.4336	0.060*	

C8	0.6908 (2)	0.8769 (2)	0.3682 (2)	0.0464 (8)
H8A	0.7396	0.8766	0.3355	0.070*
H8B	0.7313	0.8674	0.4337	0.070*
H8C	0.6539	0.9371	0.3588	0.070*
C9	0.5473 (2)	0.59489 (17)	0.15972 (17)	0.0331 (6)
C10	0.4500 (2)	0.61622 (19)	0.07168 (17)	0.0428 (7)
H10A	0.4526	0.6815	0.0542	0.064*
H10B	0.3845	0.6054	0.0827	0.064*
H10C	0.4511	0.5753	0.0223	0.064*
C11	0.6480 (2)	0.60593 (19)	0.13774 (19)	0.0430 (7)
H11A	0.7112	0.5997	0.1941	0.064*
H11B	0.6478	0.6676	0.1109	0.064*
H11C	0.6490	0.5575	0.0942	0.064*
C12	0.5458 (2)	0.49417 (17)	0.19209 (16)	0.0271 (6)
C13	0.6213 (2)	0.46605 (17)	0.27644 (17)	0.0289 (6)
H13	0.6688	0.5114	0.3141	0.035*
C14	0.62964 (19)	0.37446 (17)	0.30749 (16)	0.0275 (6)
C15	0.5596 (2)	0.30877 (17)	0.25142 (17)	0.0289 (6)
C16	0.4806 (2)	0.33380 (17)	0.16800 (17)	0.0284 (6)
C17	0.4755 (2)	0.42643 (17)	0.14003 (16)	0.0276 (6)
H17	0.4220	0.4441	0.0833	0.033*
C18	0.7107 (2)	0.34792 (18)	0.40014 (17)	0.0355 (7)
H18A	0.7588	0.3004	0.3927	0.053*
H18B	0.6742	0.3229	0.4382	0.053*
H18C	0.7519	0.4033	0.4297	0.053*
C19	0.4016 (2)	0.26264 (18)	0.11083 (18)	0.0381 (7)
H19A	0.3731	0.2274	0.1497	0.057*
H19B	0.4370	0.2197	0.0831	0.057*
H19C	0.3432	0.2945	0.0626	0.057*
O2	0.56282 (16)	0.21611 (12)	0.27588 (14)	0.0407 (5)
C21	0.5685 (2)	0.9709 (2)	0.7521 (2)	0.0462 (8)
H21	0.5092	0.9447	0.7618	0.055*
C22	0.6438 (2)	1.01925 (19)	0.8208 (2)	0.0420 (7)
H22	0.6367	1.0258	0.8781	0.050*
C23	0.7310 (2)	1.05919 (17)	0.80820 (18)	0.0324 (6)
C24	0.7377 (2)	1.04480 (18)	0.72336 (19)	0.0374 (7)
H24	0.7967	1.0691	0.7120	0.045*
C25	0.6597 (2)	0.99569 (19)	0.65561 (19)	0.0424 (7)
H25	0.6646	0.9873	0.5976	0.051*
C26	0.8127 (2)	1.11584 (19)	0.88468 (19)	0.0408 (7)
C27	0.8489 (7)	1.0589 (5)	0.9743 (5)	0.067 (3) 0.619 (7)
H27A	0.7867	1.0424	0.9885	0.100* 0.619 (7)
H27B	0.8982	1.0964	1.0243	0.100* 0.619 (7)
H27C	0.8847	1.0018	0.9670	0.100* 0.619 (7)
C28	0.7510 (4)	1.2038 (3)	0.9005 (4)	0.0549 (19) 0.619 (7)
H28A	0.7999	1.2425	0.9495	0.082* 0.619 (7)
H28B	0.6923	1.1828	0.9177	0.082* 0.619 (7)
H28C	0.7224	1.2404	0.8440	0.082* 0.619 (7)
C29	0.9038 (5)	1.1479 (5)	0.8648 (4)	0.070 (2) 0.619 (7)

H29A	0.9496	1.1870	0.9153	0.105*	0.619 (7)
H29B	0.8787	1.1843	0.8081	0.105*	0.619 (7)
H29C	0.9444	1.0938	0.8578	0.105*	0.619 (7)
C30	0.8223 (8)	1.2143 (6)	0.8482 (7)	0.069 (4)	0.381 (7)
H30A	0.8707	1.2524	0.8981	0.104*	0.381 (7)
H30B	0.7520	1.2438	0.8237	0.104*	0.381 (7)
H30C	0.8504	1.2089	0.7995	0.104*	0.381 (7)
C31	0.8037 (13)	1.1145 (11)	0.9700 (8)	0.104 (7)	0.381 (7)
H31A	0.8624	1.1501	1.0141	0.156*	0.381 (7)
H31B	0.8065	1.0498	0.9910	0.156*	0.381 (7)
H31C	0.7357	1.1426	0.9647	0.156*	0.381 (7)
C32	0.9270 (7)	1.0667 (8)	0.8952 (7)	0.073 (4)	0.381 (7)
H32A	0.9270	1.0013	0.9134	0.109*	0.381 (7)
H32B	0.9857	1.0999	0.9420	0.109*	0.381 (7)
H32C	0.9361	1.0698	0.8366	0.109*	0.381 (7)
C41	0.3686 (5)	0.8541 (4)	0.9056 (4)	0.0463 (15)	0.50
H41A	0.2966	0.8651	0.9039	0.069*	0.50
H41B	0.3958	0.7950	0.9372	0.069*	0.50
H41C	0.3667	0.8508	0.8430	0.069*	0.50
C42	0.4352 (3)	0.9272 (3)	0.9523 (3)	0.0648 (11)	
C43	0.4328 (3)	1.0136 (3)	0.9105 (3)	0.0702 (11)	
H43	0.3863	1.0227	0.8486	0.084*	
C44	0.5031 (3)	0.9144 (3)	1.0423 (3)	0.0699 (11)	
H44	0.5054	0.8559	1.0714	0.084*	
H20	0.511 (3)	0.927 (2)	0.622 (2)	0.092 (12)*	
H2	0.593 (3)	0.208 (2)	0.333 (2)	0.078 (12)*	
H1	0.477 (3)	0.842 (3)	0.453 (3)	0.099 (15)*	

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N20	0.0395 (15)	0.0352 (14)	0.0404 (15)	-0.0018 (12)	0.0044 (13)	-0.0008 (11)
C1	0.0462 (4)	0.0377 (4)	0.0290 (4)	-0.0071 (3)	0.0116 (3)	-0.0025 (3)
O1	0.0424 (12)	0.0351 (11)	0.0439 (12)	-0.0056 (9)	0.0160 (10)	-0.0110 (9)
C1	0.0322 (15)	0.0251 (14)	0.0295 (15)	0.0055 (12)	0.0064 (12)	0.0006 (12)
C2	0.0269 (14)	0.0276 (14)	0.0339 (15)	0.0018 (12)	0.0077 (12)	0.0022 (12)
C3	0.0317 (15)	0.0253 (14)	0.0302 (14)	-0.0028 (12)	0.0068 (12)	0.0008 (11)
C4	0.0343 (15)	0.0258 (14)	0.0281 (14)	0.0027 (12)	0.0103 (12)	0.0048 (11)
C5	0.0360 (16)	0.0313 (15)	0.0348 (16)	0.0003 (13)	0.0139 (13)	0.0028 (13)
C6	0.0321 (15)	0.0275 (14)	0.0342 (15)	0.0002 (12)	0.0069 (13)	0.0015 (12)
C7	0.0374 (17)	0.0352 (16)	0.0476 (18)	-0.0033 (13)	0.0163 (14)	-0.0082 (13)
C8	0.0418 (18)	0.0444 (18)	0.0527 (19)	-0.0125 (15)	0.0174 (15)	-0.0088 (15)
C9	0.0410 (16)	0.0316 (15)	0.0288 (14)	-0.0016 (13)	0.0154 (13)	-0.0004 (12)
C10	0.059 (2)	0.0350 (16)	0.0285 (15)	0.0023 (15)	0.0094 (14)	0.0037 (13)
C11	0.059 (2)	0.0370 (17)	0.0422 (17)	-0.0069 (14)	0.0301 (15)	0.0006 (13)
C12	0.0317 (14)	0.0270 (14)	0.0266 (14)	0.0015 (12)	0.0154 (12)	-0.0001 (11)
C13	0.0254 (14)	0.0324 (15)	0.0275 (14)	-0.0024 (12)	0.0085 (12)	-0.0045 (12)
C14	0.0266 (14)	0.0306 (14)	0.0251 (13)	0.0009 (12)	0.0097 (11)	-0.0002 (12)
C15	0.0320 (15)	0.0267 (14)	0.0288 (14)	-0.0008 (12)	0.0122 (12)	0.0009 (11)
C16	0.0283 (14)	0.0285 (14)	0.0285 (14)	-0.0017 (12)	0.0108 (12)	-0.0037 (11)

C17	0.0280 (14)	0.0324 (15)	0.0218 (13)	0.0024 (12)	0.0089 (11)	0.0015 (11)
C18	0.0333 (15)	0.0319 (15)	0.0341 (15)	0.0006 (12)	0.0046 (12)	0.0006 (12)
C19	0.0379 (16)	0.0365 (16)	0.0341 (16)	-0.0034 (13)	0.0069 (13)	-0.0035 (13)
O2	0.0527 (13)	0.0278 (11)	0.0323 (12)	-0.0025 (9)	0.0053 (10)	0.0012 (9)
C21	0.0412 (18)	0.0473 (19)	0.052 (2)	-0.0104 (15)	0.0204 (16)	-0.0086 (16)
C22	0.0415 (17)	0.0466 (18)	0.0400 (17)	-0.0110 (15)	0.0177 (14)	-0.0086 (14)
C23	0.0287 (15)	0.0292 (14)	0.0366 (16)	0.0022 (12)	0.0093 (13)	0.0041 (12)
C24	0.0367 (16)	0.0351 (16)	0.0416 (17)	0.0004 (13)	0.0160 (14)	0.0060 (13)
C25	0.054 (2)	0.0356 (17)	0.0353 (17)	0.0034 (15)	0.0147 (15)	0.0055 (13)
C26	0.0346 (16)	0.0393 (17)	0.0404 (17)	-0.0090 (14)	0.0051 (13)	-0.0003 (14)
C27	0.074 (6)	0.055 (4)	0.040 (4)	-0.011 (3)	-0.014 (3)	0.004 (3)
C28	0.049 (3)	0.041 (3)	0.060 (4)	-0.002 (3)	0.004 (3)	-0.016 (3)
C29	0.052 (4)	0.093 (6)	0.068 (4)	-0.038 (4)	0.025 (3)	-0.024 (4)
C30	0.062 (7)	0.044 (6)	0.073 (7)	-0.004 (5)	-0.007 (6)	-0.003 (5)
C31	0.115 (14)	0.155 (17)	0.045 (7)	-0.097 (13)	0.032 (9)	-0.043 (11)
C32	0.042 (6)	0.060 (7)	0.086 (8)	-0.008 (5)	-0.009 (5)	-0.008 (6)
C41	0.040 (4)	0.054 (4)	0.046 (4)	0.009 (3)	0.017 (3)	0.002 (3)
C42	0.047 (2)	0.058 (2)	0.105 (3)	-0.0047 (18)	0.047 (2)	-0.027 (2)
C43	0.053 (2)	0.080 (3)	0.093 (3)	0.005 (2)	0.044 (2)	-0.017 (2)
C44	0.054 (2)	0.070 (3)	0.099 (3)	0.004 (2)	0.045 (2)	-0.012 (2)

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

N20—C25	1.331 (4)	C19—H19A	0.9800
N20—C21	1.335 (4)	C19—H19B	0.9800
N20—Cl	3.035 (3)	C19—H19C	0.9800
N20—H20	1.06 (4)	O2—H2	0.85 (3)
Cl—H20	1.98 (4)	C21—C22	1.365 (4)
Cl—H1	2.27 (4)	C21—H21	0.9500
O1—C1	1.382 (3)	C22—C23	1.394 (4)
O1—H1	0.90 (4)	C22—H22	0.9500
C1—C6	1.389 (4)	C23—C24	1.392 (4)
C1—C2	1.398 (3)	C23—C26	1.528 (4)
C2—C3	1.390 (3)	C24—C25	1.379 (4)
C2—C7	1.502 (4)	C24—H24	0.9500
C3—C4	1.389 (4)	C25—H25	0.9500
C3—H3	0.9500	C26—C31	1.398 (12)
C4—C5	1.389 (3)	C26—C29	1.459 (6)
C4—C9	1.538 (3)	C26—C27	1.544 (8)
C5—C6	1.395 (4)	C26—C30	1.547 (9)
C5—H5	0.9500	C26—C28	1.584 (6)
C6—C8	1.504 (3)	C26—C32	1.653 (10)
C7—H7A	0.9800	C27—H27A	0.9800
C7—H7B	0.9800	C27—H27B	0.9800
C7—H7C	0.9800	C27—H27C	0.9800
C8—H8A	0.9800	C28—H28A	0.9800
C8—H8B	0.9800	C28—H28B	0.9800
C8—H8C	0.9800	C28—H28C	0.9800
C9—C12	1.533 (3)	C29—H29A	0.9800
C9—C11	1.539 (4)	C29—H29B	0.9800

C9—C10	1.543 (4)	C29—H29C	0.9800
C10—H10A	0.9800	C30—H30A	0.9800
C10—H10B	0.9800	C30—H30B	0.9800
C10—H10C	0.9800	C30—H30C	0.9800
C11—H11A	0.9800	C31—H31A	0.9800
C11—H11B	0.9800	C31—H31B	0.9800
C11—H11C	0.9800	C31—H31C	0.9800
C12—C17	1.391 (3)	C32—H32A	0.9800
C12—C13	1.398 (3)	C32—H32B	0.9800
C13—C14	1.390 (3)	C32—H32C	0.9800
C13—H13	0.9500	C41—C42	1.398 (6)
C14—C15	1.391 (3)	C41—H41A	0.9800
C14—C18	1.510 (3)	C41—H41B	0.9800
C15—O2	1.378 (3)	C41—H41C	0.9800
C15—C16	1.395 (3)	C42—C44	1.383 (5)
C16—C17	1.392 (3)	C42—C43	1.398 (5)
C16—C19	1.507 (3)	C43—C44 ⁱ	1.373 (5)
C17—H17	0.9500	C43—H43	0.9500
C18—H18A	0.9800	C44—C43 ⁱ	1.373 (5)
C18—H18B	0.9800	C44—H44	0.9500
C18—H18C	0.9800		
C25—N20—C21	121.4 (3)	C16—C19—H19C	109.5
C25—N20—Cl	128.1 (2)	H19A—C19—H19C	109.5
C21—N20—Cl	109.82 (19)	H19B—C19—H19C	109.5
C25—N20—H20	125 (2)	C15—O2—H2	113 (2)
C21—N20—H20	113 (2)	N20—C21—C22	120.4 (3)
H20—Cl—H1	82.2 (14)	N20—C21—H21	119.8
C1—O1—H1	116 (2)	C22—C21—H21	119.8
O1—C1—C6	116.0 (2)	C21—C22—C23	121.0 (3)
O1—C1—C2	122.3 (2)	C21—C22—H22	119.5
C6—C1—C2	121.6 (2)	C23—C22—H22	119.5
C3—C2—C1	117.7 (2)	C24—C23—C22	116.4 (2)
C3—C2—C7	120.8 (2)	C24—C23—C26	123.2 (2)
C1—C2—C7	121.5 (2)	C22—C23—C26	120.5 (2)
C4—C3—C2	122.9 (2)	C25—C24—C23	120.8 (3)
C4—C3—H3	118.6	C25—C24—H24	119.6
C2—C3—H3	118.6	C23—C24—H24	119.6
C5—C4—C3	117.2 (2)	N20—C25—C24	120.0 (3)
C5—C4—C9	123.9 (2)	N20—C25—H25	120.0
C3—C4—C9	118.9 (2)	C24—C25—H25	120.0
C4—C5—C6	122.5 (3)	C31—C26—C29	126.9 (6)
C4—C5—H5	118.8	C31—C26—C23	116.9 (5)
C6—C5—H5	118.8	C29—C26—C23	114.7 (3)
C1—C6—C5	118.1 (2)	C29—C26—C27	111.2 (5)
C1—C6—C8	120.7 (2)	C23—C26—C27	109.2 (3)
C5—C6—C8	121.2 (3)	C31—C26—C30	115.0 (8)
C2—C7—H7A	109.5	C29—C26—C30	55.7 (5)
C2—C7—H7B	109.5	C23—C26—C30	109.1 (4)

H7A—C7—H7B	109.5	C27—C26—C30	141.3 (5)
C2—C7—H7C	109.5	C31—C26—C28	67.4 (8)
H7A—C7—H7C	109.5	C29—C26—C28	109.0 (4)
H7B—C7—H7C	109.5	C23—C26—C28	106.1 (3)
C6—C8—H8A	109.5	C27—C26—C28	106.2 (4)
C6—C8—H8B	109.5	C30—C26—C28	57.2 (5)
H8A—C8—H8B	109.5	C31—C26—C32	108.5 (8)
C6—C8—H8C	109.5	C29—C26—C32	46.9 (4)
H8A—C8—H8C	109.5	C23—C26—C32	103.1 (4)
H8B—C8—H8C	109.5	C27—C26—C32	73.8 (5)
C12—C9—C4	110.1 (2)	C30—C26—C32	102.6 (6)
C12—C9—C11	107.2 (2)	C28—C26—C32	148.7 (4)
C4—C9—C11	112.3 (2)	C26—C27—H27A	109.5
C12—C9—C10	111.8 (2)	C26—C27—H27B	109.5
C4—C9—C10	107.6 (2)	C26—C27—H27C	109.5
C11—C9—C10	107.9 (2)	C26—C28—H28A	109.5
C9—C10—H10A	109.5	C26—C28—H28B	109.5
C9—C10—H10B	109.5	C26—C28—H28C	109.5
H10A—C10—H10B	109.5	C26—C29—H29A	109.5
C9—C10—H10C	109.5	C26—C29—H29B	109.5
H10A—C10—H10C	109.5	C26—C29—H29C	109.5
H10B—C10—H10C	109.5	C26—C30—H30A	109.5
C9—C11—H11A	109.5	C26—C30—H30B	109.5
C9—C11—H11B	109.5	H30A—C30—H30B	109.5
H11A—C11—H11B	109.5	C26—C30—H30C	109.5
C9—C11—H11C	109.5	H30A—C30—H30C	109.5
H11A—C11—H11C	109.5	H30B—C30—H30C	109.5
H11B—C11—H11C	109.5	C26—C31—H31A	109.5
C17—C12—C13	117.0 (2)	C26—C31—H31B	109.5
C17—C12—C9	123.5 (2)	H31A—C31—H31B	109.5
C13—C12—C9	119.4 (2)	C26—C31—H31C	109.5
C14—C13—C12	122.7 (2)	H31A—C31—H31C	109.5
C14—C13—H13	118.6	H31B—C31—H31C	109.5
C12—C13—H13	118.6	C26—C32—H32A	109.5
C13—C14—C15	117.9 (2)	C26—C32—H32B	109.5
C13—C14—C18	120.8 (2)	H32A—C32—H32B	109.5
C15—C14—C18	121.3 (2)	C26—C32—H32C	109.5
O2—C15—C14	122.1 (2)	H32A—C32—H32C	109.5
O2—C15—C16	116.3 (2)	H32B—C32—H32C	109.5
C14—C15—C16	121.6 (2)	C42—C41—H41A	109.5
C17—C16—C15	118.2 (2)	C42—C41—H41B	109.5
C17—C16—C19	121.0 (2)	H41A—C41—H41B	109.5
C15—C16—C19	120.7 (2)	C42—C41—H41C	109.5
C12—C17—C16	122.4 (2)	H41A—C41—H41C	109.5
C12—C17—H17	118.8	H41B—C41—H41C	109.5
C16—C17—H17	118.8	C44—C42—C41	119.3 (4)
C14—C18—H18A	109.5	C44—C42—C43	119.4 (4)
C14—C18—H18B	109.5	C41—C42—C43	121.3 (5)
H18A—C18—H18B	109.5	C44 ⁱ —C43—C42	120.8 (4)

C14—C18—H18C	109.5	C44 ⁱ —C43—H43	119.6
H18A—C18—H18C	109.5	C42—C43—H43	119.6
H18B—C18—H18C	109.5	C43 ⁱ —C44—C42	119.9 (4)
C16—C19—H19A	109.5	C43 ⁱ —C44—H44	120.1
C16—C19—H19B	109.5	C42—C44—H44	120.1
H19A—C19—H19B	109.5		

Symmetry code: (i) $-x+1, -y+2, -z+2$.

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
N20—H20···Cl	1.06 (4)	1.98 (4)	3.035 (3)	173 (3)
O2—H2···Cl ⁱⁱ	0.85 (3)	2.30 (3)	3.079 (2)	153 (3)
O1—H1···Cl	0.90 (4)	2.27 (4)	3.118 (2)	157 (3)

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