organic compounds

2-Hydroxy-N-(4-methoxybenzyl)-4-nitro-anilinium chloride

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The crystal structure of the title compound, $C_{14}H_{15}N_2O_4^+ \cdot Cl^-$, can be described as being composed of layers containing both cations and anions that are staggered along [010]. Two types of the hydrogen bonds are observed, *viz*. cation–anion and cation–cation. The chloride anions are acceptors of the strong hydrogen bonds donated by the secondary amine and the hydroxy groups. The packing is also stabilized by weak C—H···O intermolecular hydrogen bonds. An intramolecular N—H···O interaction also occurs.

Related literature

For the preparation of amines, see: Apodaca & Xiao (2001); Baxter & Reitz (2002); Salvatore *et al.* (2002); Sato *et al.* (2004). For applications of amines, see: Bergeron *et al.* (1997); Seayad *et al.* (2002). For background to hydrogen bonding, see: Desiraju (2003); Dorn *et al.* (2005) and for hydrogen-bond motifs, see: Etter *et al.* (1990).
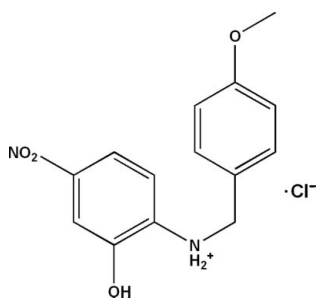
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Experimental

Crystal data

$C_{14}H_{15}N_2O_4^+ \cdot Cl^-$	V = 2983.09 (15) Å³
M_r = 310.73	Z = 8
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
a = 32.1166 (9) Å	μ = 0.27 mm⁻¹
b = 7.4888 (2) Å	T = 100 K
c = 13.0907 (4) Å	0.18 × 0.12 × 0.06 mm
β = 108.655 (2)°	

Data collection

Bruker APEXII CCD area-detector diffractometer	11129 measured reflections
	3359 independent reflections
Absorption correction: multi-scan (*SADABS*; Sheldrick, 2002)	2290 reflections with $I > 2\sigma(I)$
T_{min} = 0.784, T_{max} = 0.984	R_{int} = 0.049

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.045	191 parameters
$wR(F^2)$ = 0.079	H-atom parameters constrained
S = 1.35	$\Delta\rho_{max}$ = 0.32 e Å⁻³
3359 reflections	$\Delta\rho_{min}$ = −0.25 e Å⁻³

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
O1—H1···Cl1	0.84	2.16	2.9950 (13)	174
N10—H10A···O1	0.92	2.22	2.652 (2)	108
N10—H10A···Cl1ⁱ	0.92	2.30	3.1082 (17)	146
N10—H10B···Cl1ⁱⁱ	0.92	2.23	3.0518 (15)	149
C8—H8···O5ⁱⁱ	0.93	2.58	3.273 (2)	132
C14—H14···O6ⁱⁱⁱ	0.93	2.48	3.388 (3)	165

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

Data collection: *APEX2* (Bruker, 2003); cell refinement: *SAINT* (Bruker, 2003); data reduction: *SAINT*; program(s) used to solve structure: *SIR2002* (Burla *et al.*, 2003); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997) and *DIAMOND* (Brandenburg & Berndt, 2001); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FB2237).

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supplementary materials

2-Hydroxy-*N*-(4-methoxybenzyl)-4-nitroanilinium chloride

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Comment

Amines are one of the most important classes of biologically active compounds of natural origin. They are also widely used in the chemical industry as basic intermediates for preparation of *e.g.* fine chemicals, pharmaceuticals and agrochemicals (Seayad *et al.*, 2002).

Due to their biological properties, amines have played important role in chemotherapeutic treatment of different diseases (Bergeron *et al.*, 1997).

Alkylation of the secondary amines with alkyl halides is the most straightforward method for the synthesis of the tertiary amines (Salvatore *et al.*, 2002). Reductive amination of aldehydes andketones is a powerful tool for the synthesis of amines. This approach is extensively used for rapid access to diverse sets of amines (Sato *et al.*, 2004; Baxter *et al.*, 2002; Apodaca *et al.*, 2001).

The synthetic route we envisioned for preparation of the title compounds consists of a one-step reductive amination of aromatic aldehydes with primary amine at acid conditions (pH = 4-5). We found that this could be efficiently conducted in methanol at room temperature using the excess of reductive agent (NaBH$_3$CN). Under these conditions 2-(4-methoxyben-zylamino)-5-nitrophenol was cleanly obtained in very good chemical yield (85%).

Fig. 1 shows the title molecule. The two benzene rings contain the interplanar angle equal to 35.35 (6)°. In the crystal packing, the important role play the hydrogen bonds. In the title structure, two types of hydrogen bonds are present, inter-connecting the cations with the anions as well as mutually the cations. The chloride anions are involved as acceptors in the strong hydrogen bonds (Desiraju, 2003; Dorn *et al.* 2005) with the secondary amine and the hydroxy group stemming from the cation (Tab. 1), *i.e.* in [O—H···Cl⁻ and N—H···Cl⁻] hydrogen bonds interactions.

The layers staggered along the *b* axis can be discerned in the crystal structure (Fig. 2). Each layer contains dimers composed of the cations and Cl⁻. These dimers are situated about the crystallographic two-fold axes. The dimers form the motifs R$^2_4$(14) (Etter *et al.*, 1990) with a pair of chains O1-H1···Cl1···H10*a*-N10-C1-C9 (Fig. 3). Moreover, the dimers are interconnected with those in the adjacent layer by another pair of the hydrogen bonds Cl1···H10*b*-N10-H10*a* with the graph set motif R$^2_4$(8) (Fig. 3). The latter motifs are situated about the crystallographic inversion centres. The packing is also stabilized by weak N—H···O (intramolecular) and C—H···O (intermolecular) interactions (Fig. 4, Tab. 1). Fig. 5 shows the projection of the structure along the *a* axis.

Experimental

To the solution of 4-methoxybenzaldehyde (2 mmol) in dry methanol (10 ml) 5-nitro-2-aminophenol (2 mmol) was added and acidified by concentrated HCl until pH = 6. After vigorous stirring for 2 h at room temperature, NaBH$_3$CN (6

mmol) was added. On completion of the reaction, as indicated by thin layer chromatography (ethyl acetate/hexane: 1/3 as eluent), the excess of the hydride was carefully destroyed by slow addition of 20 ml of cold water. The mixture was left for several hours and the resulting precipitate was filtered off, washed with water, then with ethanol and with hexane. 2-(4-methoxybenzylamino)-5-nitrophenol was identified by IR, ^{1}H and ^{13}C NMR spectroscopies. Colourless prismatic crystals (0.06×0.12×0.18 mm) of the title structure were obtained by slow crystallization from the aqueous solution with pH = 5.5.

Refinement

Approximate positions for all the H atoms were first obtained from the difference electron density map. However, the H atoms were situated into idealized positions and the H-atoms have been refined within the riding atom approximation. The applied constraints were as follow: C_{aryl}—H_{aryl} = 0.95 Å; C_{methyl}—H_{methyl} = 0.98 Å; $C_{methylene}$—$H_{methylene}$ = 0.99Å and $N_{sec.amine}$—$H_{sec.amine}$ = 0.84 Å. The idealized methyl group was allowed to rotate about the C—C bond during the refinement by application of the command AFIX 137 in *SHELXL97* (Sheldrick, 2008). $U_{iso}(H_{methyl}/_{hydroxy}) = 1.5U_{eq}(C_{methyl}/O_{hydroxy})$ or $U_{iso}(H_{aryl}/_{methylene}/_{sec. amine}) = 1.2U_{eq}(C_{aryl}/C_{methylene}/N_{sec. amine})$.
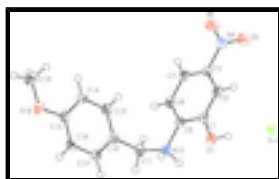
Figures

Fig. 1. The title molecule (Farrugia, 1997) with the atomic labelling scheme. The displacement parameters are drawn at the 50% probability level.
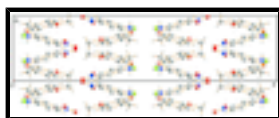

Fig. 2. The packing of the title structure viewed down the *c* axis (Brandenburg & Berndt, 2001). Cl is shown in green, N in blue and C in grey.
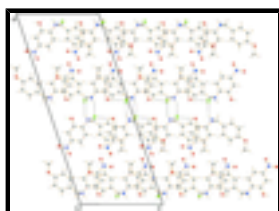
Fig. 3. A section of the title structure showing the hydrogen bonds N—H···Cl and O—H···Cl as dashed lines (Brandenburg & Berndt, 2001). Cl is shown in green, N in blue and C in grey.
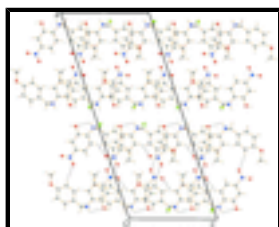
Fig. 4. A section of the title structure showing intermolecular hydrogen bond N—H···O and weak C—H···O interactions as dashed lines (Brandenburg & Berndt, 2001). Cl is shown in green, N in blue and C in grey.

Fig. 5. The packing of the title structure viewed down the *a* axis (Brandenburg & Berndt, 2001). Cl is shown in green, N in blue and C in grey.
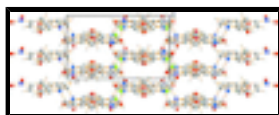
2-Hydroxy-*N*-(4-methoxybenzyl)-4-nitroanilinium chloride

Crystal data

$C_{14}H_{15}N_2O_4{}^+\cdot Cl^-$	$F(000) = 1296$
$M_r = 310.73$	$D_x = 1.384$ Mg m^{-3}
Monoclinic, $C2/c$	Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Hall symbol: -C 2yc	Cell parameters from 1996 reflections
$a = 32.1166$ (9) Å	$\theta = 2.7–25.9°$
$b = 7.4888$ (2) Å	$\mu = 0.27$ mm^{-1}
$c = 13.0907$ (4) Å	$T = 100$ K
$\beta = 108.655$ (2)°	Prism, colourless
$V = 2983.09$ (15) Å^3	$0.18 \times 0.12 \times 0.06$ mm
$Z = 8$	

Data collection

Bruker APEXII CCD area-detector diffractometer	2290 reflections with $I > 2\sigma(I)$
graphite	$R_{int} = 0.049$
φ and ω scans	$\theta_{max} = 27.5°$, $\theta_{min} = 3.1°$
Absorption correction: multi-scan (*SADABS*; Sheldrick, 2002)	$h = -41 \rightarrow 36$
$T_{min} = 0.784$, $T_{max} = 0.984$	$k = -9 \rightarrow 8$
11129 measured reflections	$l = -16 \rightarrow 13$
3359 independent reflections	

Refinement

Refinement on F^2	Primary atom site location: structure-invariant direct methods
Least-squares matrix: full	Secondary atom site location: difference Fourier map
$R[F^2 > 2\sigma(F^2)] = 0.045$	Hydrogen site location: difference Fourier map
$wR(F^2) = 0.079$	H-atom parameters constrained
$S = 1.35$	$w = 1/[\sigma^2(F_o{}^2) + (0.010P)^2]$ where $P = (F_o{}^2 + 2F_c{}^2)/3$
3359 reflections	$(\Delta/\sigma)_{max} = 0.001$
191 parameters	$\Delta\rho_{max} = 0.32$ e Å^{-3}
0 restraints	$\Delta\rho_{min} = -0.25$ e Å^{-3}
59 constraints	

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

supplementary materials

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 ($Å^2$)

	x	*y*	*z*	$U_{iso}*/U_{eq}$
O1	0.05089 (4)	0.24108 (16)	0.25570 (10)	0.0282 (4)
H1	0.0481	0.2513	0.317	0.042*
O5	0.19661 (4)	0.3946 (2)	0.52794 (12)	0.0450 (4)
O6	0.23649 (5)	0.4782 (2)	0.43081 (12)	0.0596 (5)
O18	0.16637 (4)	0.13654 (17)	−0.26241 (11)	0.0333 (4)
N4	0.20173 (6)	0.4259 (2)	0.44038 (15)	0.0365 (5)
N10	0.05796 (5)	0.31371 (19)	0.06366 (12)	0.0219 (4)
H10A	0.0329	0.3166	0.0834	0.026*
H10B	0.0562	0.4062	0.0162	0.026*
C1	0.09069 (6)	0.3047 (2)	0.25869 (16)	0.0223 (5)
C2	0.12573 (6)	0.3321 (2)	0.35182 (16)	0.0250 (5)
H2	0.1233	0.3074	0.4194	0.03*
C3	0.16445 (6)	0.3973 (3)	0.34105 (16)	0.0259 (5)
C7	0.16978 (6)	0.4387 (3)	0.24313 (16)	0.0289 (5)
H7	0.1963	0.4833	0.2395	0.035*
C8	0.13458 (6)	0.4118 (2)	0.15093 (16)	0.0247 (5)
H8	0.1369	0.439	0.0837	0.03*
C9	0.09576 (6)	0.3441 (2)	0.15932 (15)	0.0207 (5)
C11	0.05879 (6)	0.1389 (2)	0.00555 (16)	0.0273 (5)
H11A	0.068	0.044	0.0584	0.033*
H11B	0.0293	0.1116	−0.0407	0.033*
C12	0.08902 (6)	0.1425 (2)	−0.06154 (16)	0.0232 (5)
C13	0.13234 (6)	0.0854 (2)	−0.02110 (16)	0.0267 (5)
H13	0.1432	0.0459	0.0499	0.032*
C14	0.15987 (6)	0.0858 (2)	−0.08398 (16)	0.0264 (5)
H14	0.1891	0.0507	−0.0548	0.032*
C15	0.14329 (6)	0.1393 (2)	−0.19111 (16)	0.0236 (5)
C16	0.10009 (6)	0.1979 (2)	−0.23310 (15)	0.0245 (5)
H16	0.089	0.2353	−0.3045	0.029*
C17	0.07379 (6)	0.2001 (2)	−0.16807 (16)	0.0248 (5)
H17	0.045	0.2412	−0.1962	0.03*
C19	0.21172 (6)	0.0849 (3)	−0.22251 (18)	0.0447 (6)
H19A	0.2139	−0.0351	−0.1955	0.067*
H19B	0.2273	0.1641	−0.1654	0.067*
H19C	0.2243	0.0912	−0.2799	0.067*
Cl1	0.042616 (15)	0.30709 (6)	0.47404 (4)	0.02477 (14)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0289 (8)	0.0383 (9)	0.0212 (8)	−0.0058 (6)	0.0133 (7)	0.0010 (6)
O5	0.0345 (9)	0.0751 (12)	0.0242 (9)	0.0038 (8)	0.0079 (7)	−0.0032 (8)
O6	0.0226 (9)	0.1194 (16)	0.0372 (10)	−0.0117 (9)	0.0102 (8)	−0.0126 (10)
O18	0.0293 (8)	0.0455 (10)	0.0306 (9)	0.0039 (7)	0.0174 (7)	−0.0011 (7)
N4	0.0260 (11)	0.0543 (13)	0.0292 (12)	0.0056 (9)	0.0087 (9)	−0.0083 (10)
N10	0.0229 (9)	0.0259 (10)	0.0201 (9)	−0.0018 (7)	0.0112 (7)	−0.0006 (8)
C1	0.0221 (11)	0.0214 (11)	0.0265 (12)	0.0034 (9)	0.0120 (9)	0.0004 (9)
C2	0.0275 (12)	0.0290 (13)	0.0200 (11)	0.0072 (9)	0.0096 (9)	0.0017 (9)
C3	0.0215 (11)	0.0328 (13)	0.0230 (12)	0.0054 (9)	0.0066 (9)	−0.0049 (10)
C7	0.0209 (11)	0.0378 (14)	0.0309 (13)	0.0003 (9)	0.0125 (10)	−0.0048 (10)
C8	0.0268 (11)	0.0309 (13)	0.0211 (11)	0.0013 (9)	0.0140 (9)	−0.0004 (9)
C9	0.0205 (10)	0.0206 (12)	0.0212 (11)	0.0015 (8)	0.0072 (9)	−0.0018 (9)
C11	0.0337 (12)	0.0237 (12)	0.0263 (12)	−0.0035 (9)	0.0121 (10)	−0.0044 (9)
C12	0.0285 (12)	0.0216 (12)	0.0217 (11)	−0.0027 (9)	0.0112 (9)	−0.0035 (9)
C13	0.0321 (12)	0.0284 (13)	0.0192 (11)	0.0032 (10)	0.0073 (9)	−0.0007 (9)
C14	0.0220 (11)	0.0317 (13)	0.0246 (12)	0.0039 (9)	0.0060 (9)	−0.0026 (10)
C15	0.0254 (11)	0.0257 (12)	0.0228 (12)	−0.0031 (9)	0.0120 (9)	−0.0046 (9)
C16	0.0276 (11)	0.0276 (12)	0.0171 (11)	0.0006 (9)	0.0053 (9)	0.0017 (9)
C17	0.0214 (11)	0.0267 (12)	0.0253 (12)	−0.0014 (9)	0.0064 (9)	−0.0019 (10)
C19	0.0297 (13)	0.0644 (17)	0.0467 (16)	0.0054 (12)	0.0218 (12)	−0.0007 (13)
Cl1	0.0245 (3)	0.0286 (3)	0.0224 (3)	0.0021 (2)	0.0091 (2)	−0.0001 (2)

Geometric parameters (Å, °)

O1—C1	1.353 (2)	C8—C9	1.382 (2)
O1—H1	0.84	C8—H8	0.93
O5—N4	1.231 (2)	C11—C12	1.503 (2)
O6—N4	1.2268 (19)	C11—H11A	0.97
O18—C15	1.365 (2)	C11—H11B	0.97
O18—C19	1.434 (2)	C12—C13	1.389 (2)
N4—C3	1.475 (2)	C12—C17	1.390 (3)
N10—C9	1.456 (2)	C13—C14	1.387 (2)
N10—C11	1.519 (2)	C13—H13	0.93
N10—H10A	0.9201	C14—C15	1.390 (3)
N10—H10B	0.9201	C14—H14	0.93
C1—C2	1.385 (2)	C15—C16	1.390 (2)
C1—C9	1.393 (2)	C16—C17	1.378 (2)
C2—C3	1.385 (2)	C16—H16	0.93
C2—H2	0.93	C17—H17	0.93
C3—C7	1.381 (3)	C19—H19A	0.96
C7—C8	1.379 (2)	C19—H19B	0.96
C7—H7	0.93	C19—H19C	0.96
C1—O1—H1	109.3	C12—C11—H11A	108.9
C15—O18—C19	117.73 (15)	N10—C11—H11A	108.9

O6—N4—O5	123.56 (18)	C12—C11—H11B	108.9
O6—N4—C3	117.71 (18)	N10—C11—H11B	108.9
O5—N4—C3	118.73 (17)	H11A—C11—H11B	107.7
C9—N10—C11	114.97 (13)	C13—C12—C17	117.77 (18)
C9—N10—H10A	108.5	C13—C12—C11	121.83 (18)
C11—N10—H10A	108.5	C17—C12—C11	120.38 (17)
C9—N10—H10B	108.5	C14—C13—C12	121.61 (18)
C11—N10—H10B	108.5	C14—C13—H13	119.2
H10A—N10—H10B	107.5	C12—C13—H13	119.2
O1—C1—C2	124.88 (17)	C13—C14—C15	119.21 (18)
O1—C1—C9	116.05 (17)	C13—C14—H14	120.4
C2—C1—C9	119.07 (17)	C15—C14—H14	120.4
C3—C2—C1	117.78 (18)	O18—C15—C16	115.21 (17)
C3—C2—H2	121.1	O18—C15—C14	124.67 (17)
C1—C2—H2	121.1	C16—C15—C14	120.11 (18)
C7—C3—C2	123.74 (19)	C17—C16—C15	119.42 (18)
C7—C3—N4	118.62 (17)	C17—C16—H16	120.3
C2—C3—N4	117.63 (18)	C15—C16—H16	120.3
C3—C7—C8	117.97 (18)	C16—C17—C12	121.83 (18)
C3—C7—H7	121	C16—C17—H17	119.1
C8—C7—H7	121	C12—C17—H17	119.1
C9—C8—C7	119.44 (18)	O18—C19—H19A	109.5
C9—C8—H8	120.3	O18—C19—H19B	109.5
C7—C8—H8	120.3	H19A—C19—H19B	109.5
C8—C9—C1	121.97 (18)	O18—C19—H19C	109.5
C8—C9—N10	120.92 (17)	H19A—C19—H19C	109.5
C1—C9—N10	117.10 (16)	H19B—C19—H19C	109.5
C12—C11—N10	113.27 (14)		
O1—C1—C2—C3	179.67 (17)	C11—N10—C9—C8	82.5 (2)
C9—C1—C2—C3	−0.2 (3)	C11—N10—C9—C1	−98.47 (19)
C1—C2—C3—C7	0.9 (3)	C9—N10—C11—C12	−77.3 (2)
C1—C2—C3—N4	−179.87 (16)	N10—C11—C12—C13	93.1 (2)
O6—N4—C3—C7	−3.1 (3)	N10—C11—C12—C17	−88.4 (2)
O5—N4—C3—C7	177.68 (18)	C17—C12—C13—C14	0.3 (3)
O6—N4—C3—C2	177.70 (18)	C11—C12—C13—C14	178.75 (17)
O5—N4—C3—C2	−1.6 (3)	C12—C13—C14—C15	−2.1 (3)
C2—C3—C7—C8	−0.6 (3)	C19—O18—C15—C16	177.25 (17)
N4—C3—C7—C8	−179.78 (17)	C19—O18—C15—C14	−3.9 (3)
C3—C7—C8—C9	−0.5 (3)	C13—C14—C15—O18	−176.41 (17)
C7—C8—C9—C1	1.2 (3)	C13—C14—C15—C16	2.4 (3)
C7—C8—C9—N10	−179.87 (16)	O18—C15—C16—C17	178.03 (16)
O1—C1—C9—C8	179.28 (16)	C14—C15—C16—C17	−0.9 (3)
C2—C1—C9—C8	−0.8 (3)	C15—C16—C17—C12	−1.0 (3)
O1—C1—C9—N10	0.3 (2)	C13—C12—C17—C16	1.3 (3)
C2—C1—C9—N10	−179.82 (15)	C11—C12—C17—C16	−177.19 (17)

Hydrogen-bond geometry (Å, °)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*

O1—H1···Cl1	0.84	2.16	2.9950 (13)	174
N10—H10A···O1	0.92	2.22	2.652 (2)	108
N10—H10A···Cl1[i]	0.92	2.30	3.1082 (17)	146
N10—H10B···Cl1[ii]	0.92	2.23	3.0518 (15)	149
C8—H8···O5[ii]	0.93	2.58	3.273 (2)	132
C14—H14···O6[iii]	0.93	2.48	3.388 (3)	165

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

Fig. 1

Fig. 2

Fig. 3

Fig. 4

Fig. 5