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N-[2-(2-Chlorophenyl)-2-hydroxyethyl]-propan-2-aminium benzoate

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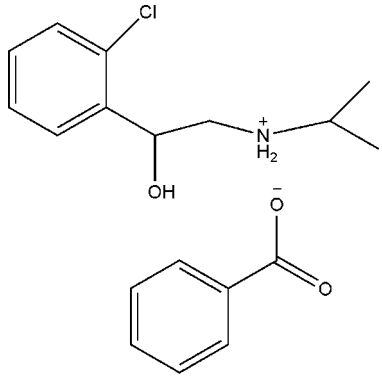
Key indicators: single-crystal X-ray study; T = 296 K; mean σ(C–C) = 0.003 Å; R factor = 0.042; wR factor = 0.110; data-to-parameter ratio = 19.6.

In the title compound, $C_{11}H_{17}ClNO^{+}\cdot C_7H_5O_2^{-}$, obtained by the reaction of chlorprenaline {or 1-(2-chlorophenyl)-2-[(1-methylethyl)amino]ethanol} and benzoic acid, the chlorprenaline is twisted moderately [C—C—C—C torsion angle = −76.00 (17)°] compared with related compounds. The molecules as usual form dimers. In the crystal structure, the two components are connected by classical O—H···O and N—H···O hydrogen bonds.

Related literature

For related structures, see: Feld *et al.* (1981); Feng *et al.* (2010); Tang *et al.* (2009*a*,*b*).



Experimental

Crystal data

$C_{11}H_{17}ClNO^{+}\cdot C_7H_5O_2^{-}$ M_r = 335.82
Monoclinic, $P2_1/n$
a = 7.8343 (3) Å
b = 13.1260 (5) Å
c = 17.7308 (7) Å
β = 94.330 (1)°
V = 1818.11 (12) Å^3
Z = 4
Mo $K\alpha$ radiation
μ = 0.22 mm^{-1}
T = 296 K
0.53 × 0.48 × 0.46 mm

Data collection

Rigaku R-AXIS RAPID/ZJUG diffractometer
Absorption correction: multi-scan (*ABSCOR*; Higashi, 1995)
T_{min} = 0.871, T_{max} = 0.904
17400 measured reflections
4123 independent reflections
3186 reflections with $I > 2\sigma(I)$
R_{int} = 0.023

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.042
$wR(F^2)$ = 0.110
S = 1.00
4123 reflections
210 parameters
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.31 e Å^{-3}
$\Delta\rho_{min}$ = −0.32 e Å^{-3}

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

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1A···O2	0.90	1.85	2.7231 (15)	162
O1—H1···O3	0.82	1.93	2.7219 (15)	162
N1—H1B···O3^{i}	0.90	1.88	2.7710 (15)	169

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

Data collection: *PROCESS-AUTO* (Rigaku/MSC, 2006); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/MSC, 2007); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

This work was supported by the Key Scientific and Technological Research Project of the Science and Technology Department of Zhejiang Province of China (grant No. 2009 C32078).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: RK2237).

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supporting information

Acta Cryst. (2010). E**66**, o2910 [https://doi.org/10.1107/S1600536810040274]

N-[2-(2-Chlorophenyl)-2-hydroxyethyl]propan-2-aminium benzoate

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S1. Comment

A recent study reports the structure of bis{*N*-[2-(2-chlorophenyl)-2-hydroxyethyl]propan-2-aminium} oxalate (Tang *et al.*, 2009*b*), which was synthesized by oxalic acid and chlorprenaline (Tang *et al.*, 2009*a*). Here using benzoic acid instead of oxalic acid and following a similar synthetic procedure yields the title compound, **I**.

In **I** (Fig. 1), the chlorprenaline molecule and the benzoic acid molecule are linked to each other by the N1—H1A···O2 hydogen bond (2.7231 (15)Å) and the O1—H1···O3 hydogen bond (2.7219 (15)Å) (Fig. 2 & Table 1). The chlorprenaline in **I** are twisted moderately as compared with those of other compounds. The C12—O2 distance of 1.2456 (18)Å is much shorter than the similar distance of 1.2675 (15)Å (Feld *et al.*, 1981). The C1—C6—C7—C8 torsion angle of -76.00 (17)° (104.0 (17)°) is larger than the value of the similar torsion angle of 91.9 (2)° (Tang *et al.*, 2009*a*). The C9—N1 distance of 1.5096 (17)Å is longer than the value of the similar bond distance of 1.473 (4)Å (Tang *et al.*, 2009*b*), as similar as the value of the similar bond distance of 1.503 (2)Å (Feng *et al.*, 2010).

Classical hydrogen bonds (O—H···O and N—H···O) are found in the cystal structure (Fig. 2 & Table 1) are essential forces in crystal formation.

S2. Experimental

Racemic chlorprenaline was prepared by chlorprenaline hydrochloride purchased from ShangHai Shengxin Medicine & Chemical Co., Ltd. ShangHai, China. chlorprenaline hydrochloride and NaOH in a molar ratio of 1:1 were mixed and dissolved in a methanol–water solution (1:1 *v*/*v*). The precipitate formed was filtered off, washed with water and dried. It was used without further purification. Racemic chlorprenaline (0.5 g, 0.0023 mol) was dissolved in methanol (5 ml) and then Benzoic acid (0.3 g, 0.0023 mol) was added. The mixture was dissovled by heating to 323 K where a clear solution resulted. The resulting solution was concentrated at ambient temperature. Colourless crystals of **I** separated from the solution in about 68% yield after one day.

S3. Refinement

All of the H atoms were placed in calculated positions and allowed to ride on their parent atoms at distances of 0.93Å (aromatic), 0.98Å (methine), 0.97Å (methylene), 0.96Å (methyl) 0.82Å (hydroxyl) and N—H = 0.90Å, with U_{iso}(H) = 1.2–1.5 U_{eq}(C,O,N).

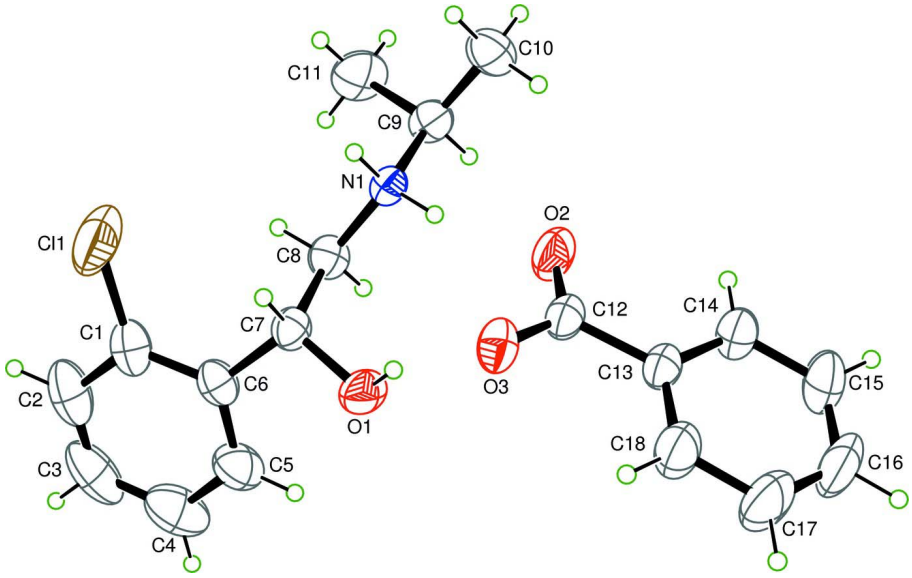

Figure 1

The asymmetric unit of **I** with atom numbering scheme. Displacement ellipsoids are drawn at 50% probability level. H atoms are presented as a small spheres of arbitrary radius.

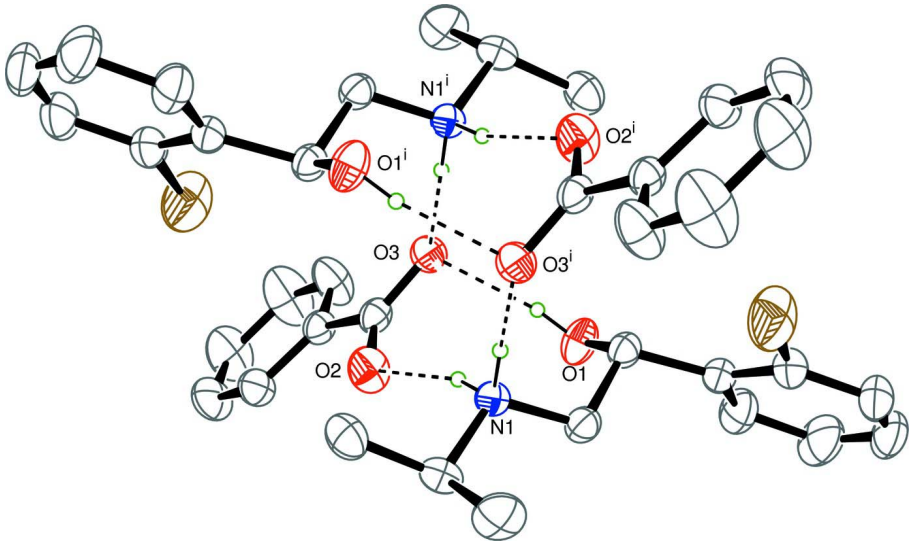

Figure 2

The hydrogen bonds (dashed lines) system in **I**.

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Crystal data

$C_{11}H_{17}ClNO^{+}\cdot C_7H_5O_2^{-}$
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Monoclinic, $P2_1/n$
Hall symbol: -P 2yn
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c = 17.7308 (7) Å
β = 94.330 (1)°
V = 1818.11 (12) Å^3
Z = 4
F(000) = 712
D_x = 1.227 Mg m^{-3}
Mo $K\alpha$ radiation, λ = 0.71073 Å
Cell parameters from 12983 reflections

θ = 3.0–27.4°
μ = 0.22 mm^{-1}
T = 296 K

Chunk, colourless
0.53 × 0.48 × 0.46 mm

Data collection

Rigaku R-AXIS RAPID/ZJUG diffractometer
Radiation source: rolling anode
Graphite monochromator
Detector resolution: 10.00 pixels mm^{-1}
ω scans
Absorption correction: multi-scan (*ABSCOR*; Higashi, 1995)
T_{min} = 0.871, T_{max} = 0.904

17400 measured reflections
4123 independent reflections
3186 reflections with $I > 2\sigma(I)$
R_{int} = 0.023
θ_{max} = 27.4°, θ_{min} = 3.0°
h = −9→10
k = −15→17
l = −22→22

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)]$ = 0.042
$wR(F^2)$ = 0.110
S = 1.00
4123 reflections
210 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
$w = 1/[\sigma^2(F_o^2) + (0.0437P)^2 + 0.6461P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} < 0.001$
$\Delta\rho_{max}$ = 0.31 e Å^{-3}
$\Delta\rho_{min}$ = −0.32 e Å^{-3}
Extinction correction: *SHELXL97* (Sheldrick, 2008), $Fc^*=kFc[1+0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$
Extinction coefficient: 0.0125 (12)

Special details

Geometry. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'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}
Cl1	0.45832 (8)	0.73075 (5)	0.69565 (4)	0.0898 (2)
N1	0.27625 (14)	0.59370 (8)	0.48693 (6)	0.0326 (2)
H1A	0.2765	0.5272	0.4744	0.039*
H1B	0.3858	0.6149	0.4912	0.039*
O3	0.40196 (12)	0.31773 (8)	0.50725 (6)	0.0456 (3)
O2	0.23389 (16)	0.40526 (8)	0.42406 (7)	0.0543 (3)
O1	0.25272 (15)	0.43338 (8)	0.61142 (6)	0.0497 (3)
H1	0.3143	0.4081	0.5810	0.075*
C13	0.21403 (17)	0.22582 (10)	0.42001 (8)	0.0369 (3)
C8	0.20481 (18)	0.60417 (11)	0.56199 (8)	0.0392 (3)
H8A	0.0844	0.5862	0.5571	0.047*

H8B	0.2135	0.6748	0.5779	0.047*
C9	0.18157 (18)	0.65197 (11)	0.42338 (9)	0.0418 (3)
H9	0.0642	0.6258	0.4164	0.050*
C1	0.3003 (2)	0.65860 (13)	0.73480 (9)	0.0522 (4)
C14	0.1011 (2)	0.22632 (12)	0.35609 (9)	0.0463 (4)
H14	0.0760	0.2873	0.3309	0.056*
C12	0.28937 (17)	0.32356 (10)	0.45198 (8)	0.0373 (3)
C6	0.23953 (18)	0.57020 (11)	0.69877 (8)	0.0410 (3)
C7	0.29652 (18)	0.53722 (10)	0.62261 (8)	0.0369 (3)
H7	0.4206	0.5458	0.6218	0.044*
C10	0.2707 (2)	0.63197 (14)	0.35160 (9)	0.0545 (4)
H10A	0.2723	0.5600	0.3419	0.082*
H10B	0.3860	0.6571	0.3577	0.082*
H10C	0.2100	0.6661	0.3098	0.082*
C11	0.1747 (3)	0.76461 (13)	0.44186 (11)	0.0617 (5)
H11A	0.1177	0.7740	0.4874	0.093*
H11B	0.1130	0.8000	0.4010	0.093*
H11C	0.2890	0.7911	0.4488	0.093*
C16	0.0598 (3)	0.04706 (15)	0.36633 (13)	0.0800 (7)
H16	0.0070	−0.0128	0.3489	0.096*
C18	0.2509 (3)	0.13432 (13)	0.45566 (11)	0.0644 (5)
H18	0.3292	0.1323	0.4978	0.077*
C15	0.0252 (2)	0.13674 (14)	0.32929 (11)	0.0625 (5)
H15	−0.0495	0.1377	0.2860	0.075*
C5	0.1165 (2)	0.51537 (15)	0.73318 (9)	0.0572 (4)
H5	0.0756	0.4549	0.7113	0.069*
C3	0.1146 (3)	0.63805 (19)	0.83306 (11)	0.0776 (7)
H3	0.0712	0.6611	0.8774	0.093*
C17	0.1724 (4)	0.04543 (14)	0.42930 (14)	0.0901 (8)
H17	0.1961	−0.0157	0.4545	0.108*
C2	0.2388 (3)	0.69251 (17)	0.80124 (10)	0.0688 (6)
H2	0.2816	0.7519	0.8241	0.083*
C4	0.0528 (3)	0.5488 (2)	0.79991 (11)	0.0783 (6)
H4	−0.0309	0.5113	0.8221	0.094*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.0863 (4)	0.0823 (4)	0.1027 (4)	−0.0357 (3)	0.0192 (3)	−0.0458 (3)
N1	0.0330 (5)	0.0275 (5)	0.0370 (6)	−0.0016 (4)	0.0007 (4)	−0.0014 (4)
O3	0.0384 (5)	0.0479 (6)	0.0497 (6)	−0.0055 (5)	−0.0019 (4)	−0.0129 (5)
O2	0.0723 (8)	0.0301 (5)	0.0588 (7)	−0.0024 (5)	−0.0053 (6)	−0.0076 (5)
O1	0.0659 (7)	0.0344 (5)	0.0512 (6)	−0.0035 (5)	0.0199 (5)	−0.0064 (5)
C13	0.0375 (7)	0.0323 (7)	0.0412 (7)	−0.0023 (6)	0.0046 (6)	−0.0077 (6)
C8	0.0381 (7)	0.0382 (7)	0.0416 (8)	0.0030 (6)	0.0053 (6)	−0.0045 (6)
C9	0.0370 (7)	0.0397 (8)	0.0473 (8)	−0.0004 (6)	−0.0054 (6)	0.0072 (6)
C1	0.0531 (9)	0.0565 (10)	0.0458 (9)	0.0081 (8)	−0.0039 (7)	−0.0146 (7)
C14	0.0491 (8)	0.0384 (8)	0.0502 (9)	0.0006 (7)	−0.0040 (7)	−0.0073 (6)

C12	0.0355 (7)	0.0351 (7)	0.0419 (7)	−0.0045 (6)	0.0074 (6)	−0.0090 (6)
C6	0.0425 (7)	0.0454 (8)	0.0348 (7)	0.0094 (6)	0.0005 (6)	−0.0031 (6)
C7	0.0381 (7)	0.0342 (7)	0.0386 (7)	0.0006 (6)	0.0050 (6)	−0.0052 (5)
C10	0.0610 (10)	0.0605 (10)	0.0410 (8)	0.0011 (8)	−0.0027 (7)	0.0093 (7)
C11	0.0728 (12)	0.0393 (9)	0.0729 (12)	0.0131 (8)	0.0032 (9)	0.0116 (8)
C16	0.1048 (17)	0.0444 (10)	0.0876 (15)	−0.0270 (11)	−0.0145 (13)	−0.0166 (10)
C18	0.0916 (14)	0.0393 (8)	0.0581 (11)	−0.0045 (9)	−0.0216 (10)	−0.0008 (8)
C15	0.0630 (11)	0.0572 (11)	0.0642 (11)	−0.0086 (9)	−0.0147 (9)	−0.0186 (9)
C5	0.0658 (11)	0.0623 (11)	0.0453 (9)	0.0028 (9)	0.0154 (8)	0.0000 (8)
C3	0.0973 (16)	0.1001 (17)	0.0361 (9)	0.0424 (14)	0.0090 (10)	−0.0085 (10)
C17	0.144 (2)	0.0329 (9)	0.0878 (16)	−0.0147 (11)	−0.0269 (15)	0.0046 (9)
C2	0.0803 (13)	0.0757 (13)	0.0483 (10)	0.0247 (11)	−0.0089 (9)	−0.0237 (9)
C4	0.0884 (15)	0.0955 (17)	0.0549 (11)	0.0159 (13)	0.0311 (11)	0.0092 (11)

Geometric parameters (Å, º)

Cl1—C1	1.7438 (19)	C6—C7	1.5169 (19)
N1—C8	1.4884 (17)	C7—H7	0.9800
N1—C9	1.5096 (17)	C10—H10A	0.9600
N1—H1A	0.9000	C10—H10B	0.9600
N1—H1B	0.9000	C10—H10C	0.9600
O3—C12	1.2704 (17)	C11—H11A	0.9600
O2—C12	1.2456 (18)	C11—H11B	0.9600
O1—C7	1.4158 (16)	C11—H11C	0.9600
O1—H1	0.8200	C16—C15	1.365 (3)
C13—C18	1.378 (2)	C16—C17	1.370 (3)
C13—C14	1.384 (2)	C16—H16	0.9300
C13—C12	1.5050 (18)	C18—C17	1.384 (3)
C8—C7	1.526 (2)	C18—H18	0.9300
C8—H8A	0.9700	C15—H15	0.9300
C8—H8B	0.9700	C5—C4	1.390 (3)
C9—C11	1.516 (2)	C5—H5	0.9300
C9—C10	1.520 (2)	C3—C2	1.363 (3)
C9—H9	0.9800	C3—C4	1.382 (3)
C1—C2	1.380 (2)	C3—H3	0.9300
C1—C6	1.391 (2)	C17—H17	0.9300
C14—C15	1.386 (2)	C2—H2	0.9300
C14—H14	0.9300	C4—H4	0.9300
C6—C5	1.381 (2)		

C8—N1—C9	115.08 (11)	C6—C7—H7	109.8
C8—N1—H1A	108.5	C8—C7—H7	109.8
C9—N1—H1A	108.5	C9—C10—H10A	109.5
C8—N1—H1B	108.5	C9—C10—H10B	109.5
C9—N1—H1B	108.5	H10A—C10—H10B	109.5
H1A—N1—H1B	107.5	C9—C10—H10C	109.5
C7—O1—H1	109.5	H10A—C10—H10C	109.5
C18—C13—C14	118.61 (14)	H10B—C10—H10C	109.5

C18—C13—C12	120.50 (13)	C9—C11—H11A	109.5
C14—C13—C12	120.84 (13)	C9—C11—H11B	109.5
N1—C8—C7	112.82 (11)	H11A—C11—H11B	109.5
N1—C8—H8A	109.0	C9—C11—H11C	109.5
C7—C8—H8A	109.0	H11A—C11—H11C	109.5
N1—C8—H8B	109.0	H11B—C11—H11C	109.5
C7—C8—H8B	109.0	C15—C16—C17	119.85 (16)
H8A—C8—H8B	107.8	C15—C16—H16	120.1
N1—C9—C11	110.87 (13)	C17—C16—H16	120.1
N1—C9—C10	107.86 (12)	C13—C18—C17	120.49 (17)
C11—C9—C10	112.06 (14)	C13—C18—H18	119.8
N1—C9—H9	108.7	C17—C18—H18	119.8
C11—C9—H9	108.7	C16—C15—C14	120.15 (17)
C10—C9—H9	108.7	C16—C15—H15	119.9
C2—C1—C6	122.12 (18)	C14—C15—H15	119.9
C2—C1—Cl1	117.95 (15)	C6—C5—C4	121.29 (19)
C6—C1—Cl1	119.92 (12)	C6—C5—H5	119.4
C13—C14—C15	120.55 (15)	C4—C5—H5	119.4
C13—C14—H14	119.7	C2—C3—C4	120.44 (18)
C15—C14—H14	119.7	C2—C3—H3	119.8
O2—C12—O3	124.03 (13)	C4—C3—H3	119.8
O2—C12—C13	117.97 (12)	C16—C17—C18	120.32 (19)
O3—C12—C13	117.97 (13)	C16—C17—H17	119.8
C5—C6—C1	117.24 (15)	C18—C17—H17	119.8
C5—C6—C7	120.44 (14)	C3—C2—C1	119.36 (19)
C1—C6—C7	122.23 (14)	C3—C2—H2	120.3
O1—C7—C6	108.36 (12)	C1—C2—H2	120.3
O1—C7—C8	111.07 (12)	C3—C4—C5	119.5 (2)
C6—C7—C8	107.95 (11)	C3—C4—H4	120.2
O1—C7—H7	109.8	C5—C4—H4	120.2

C9—N1—C8—C7	175.43 (11)	C1—C6—C7—C8	−76.00 (17)
C8—N1—C9—C11	57.38 (16)	N1—C8—C7—O1	−73.09 (15)
C8—N1—C9—C10	−179.59 (12)	N1—C8—C7—C6	168.25 (11)
C18—C13—C14—C15	0.9 (3)	C14—C13—C18—C17	−2.0 (3)
C12—C13—C14—C15	−176.53 (15)	C12—C13—C18—C17	175.4 (2)
C18—C13—C12—O2	−170.12 (16)	C17—C16—C15—C14	−1.3 (4)
C14—C13—C12—O2	7.3 (2)	C13—C14—C15—C16	0.7 (3)
C18—C13—C12—O3	7.8 (2)	C1—C6—C5—C4	1.8 (3)
C14—C13—C12—O3	−174.79 (13)	C7—C6—C5—C4	−174.81 (17)
C2—C1—C6—C5	−1.5 (2)	C15—C16—C17—C18	0.2 (4)
Cl1—C1—C6—C5	178.90 (13)	C13—C18—C17—C16	1.5 (4)
C2—C1—C6—C7	175.02 (15)	C4—C3—C2—C1	1.1 (3)
Cl1—C1—C6—C7	−4.6 (2)	C6—C1—C2—C3	0.1 (3)
C5—C6—C7—O1	−19.97 (19)	Cl1—C1—C2—C3	179.68 (15)
C1—C6—C7—O1	163.62 (14)	C2—C3—C4—C5	−0.8 (3)
C5—C6—C7—C8	100.41 (16)	C6—C5—C4—C3	−0.7 (3)

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
N1—H1*A*···O2	0.90	1.85	2.7231 (15)	162
O1—H1···O3	0.82	1.93	2.7219 (15)	162
N1—H1*B*···O3^{i}	0.90	1.88	2.7710 (15)	169

Symmetry code: (i) −*x*+1, −*y*+1, −*z*+1.