

## L-Glutamic acid hydrochloride at 153 K

Yi Jian Zhang,<sup>a</sup> Zhan Shu,<sup>b</sup> Wei Xu,<sup>b</sup> Gang Chen<sup>b</sup> and Zu-Guang Li<sup>c\*</sup>

<sup>a</sup>College of Materials Chemistry and Chemical Engineering, Hangzhou Normal University, Hangzhou 310036, People's Republic of China, <sup>b</sup>College of Pharmaceutical Sciences, Zhejiang University of Technology, Hangzhou 310014, People's Republic of China, and <sup>c</sup>College of Chemical Engineering and Materials Science, Zhejiang University of Technology, Hangzhou 310014, People's Republic of China  
Correspondence e-mail: zgdlzg@sina.com

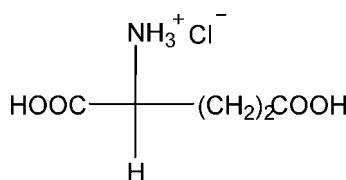
Received 16 November 2007; accepted 27 December 2007

Key indicators: single-crystal X-ray study;  $T = 153$  K; mean  $\sigma(C-C) = 0.002$  Å;  $R$  factor = 0.020;  $wR$  factor = 0.056; data-to-parameter ratio = 17.8.

The title compound [systematic name: (*S*)-1,3-dicarboxypropanaminium chloride],  $C_5H_{10}NO_4^+\cdot Cl^-$ , has been investigated previously by Dawson [*Acta Cryst.* (1953). **6**, 81–83], with  $R = 0.106$  and without the location of H atoms, and then by Sequeira, Rajagopal & Chidambaram [*Acta Cryst.* (1972). **B28**, 2514–2519] using neutron diffraction with  $R = 0.043$ . The present determination at 153 K has  $R = 0.017$  and all the H atoms are located. There are obvious differences in some C–C bond lengths between the present and previous studies. In the present structure, L-glutamic acid is protonated and is linked to the  $Cl^-$  anion by an O–H···Cl hydrogen bond. The crystal structure is established by a three-dimensional network of O–H···O, N–H···O and N–H···Cl hydrogen bonds.

### Related literature

For related literature, see: Delfino *et al.* (1978); Kirfel & Wallrafen (1985). For previous structure determinations, see: Dawson (1953) [Cambridge Structural Database (CSD; Version 5.26; Allen, 2002) refcode LGLUTA01]; Sequeira *et al.* (1972) (refcode LGLUTA).



### Experimental

#### Crystal data

$C_5H_{10}NO_4^+\cdot Cl^-$	$V = 786.73 (4)$ Å <sup>3</sup>
$M_r = 183.59$	$Z = 4$
Orthorhombic, $P2_12_12_1$	Mo $K\alpha$ radiation
$a = 5.1016 (1)$ Å	$\mu = 0.45$ mm <sup>-1</sup>
$b = 11.6386 (4)$ Å	$T = 153 (2)$ K
$c = 13.2500 (3)$ Å	$0.49 \times 0.48 \times 0.35$ mm

#### Data collection

Siemens P4 diffractometer	7742 measured reflections
Absorption correction: $\psi$ scan ( <i>SHELXTL</i> ; Bruker, 1998)	1798 independent reflections
$T_{\min} = 0.808$ , $T_{\max} = 0.858$	1775 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.016$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.019$	$\Delta\rho_{\max} = 0.29$ e Å <sup>-3</sup>
$wR(F^2) = 0.055$	$\Delta\rho_{\min} = -0.25$ e Å <sup>-3</sup>
$S = 1.15$	Absolute structure: Flack (1983),
1798 reflections	725 Friedel pairs
101 parameters	Flack parameter: 0.02 (4)
H-atom parameters constrained	

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1–H1A···Cl1	0.91	2.29	3.1406 (10)	157
N1–H1B···Cl1 <sup>i</sup>	0.91	2.34	3.1890 (10)	155
N1–H1C···O2 <sup>ii</sup>	0.91	2.04	2.8700 (13)	151
O1–H1···O3 <sup>iii</sup>	0.84	1.79	2.6208 (13)	170
O4–H4···Cl1 <sup>iv</sup>	0.84	2.21	3.0389 (9)	168

Symmetry codes: (i)  $x - 1, y, z$ ; (ii)  $x + \frac{1}{2}, -y + \frac{3}{2}, -z$ ; (iii)  $-x + \frac{1}{2}, -y + 1, z - \frac{1}{2}$ ; (iv)  $x - \frac{1}{2}, -y + \frac{3}{2}, -z + 1$ .

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/MSC, 2004); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Bruker, 1998); software used to prepare material for publication: *SHELXTL*.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SI2055).

### References

- Allen, F. H. (2002). *Acta Cryst.* **B58**, 380–388.
- Bruker (1998). *SHELXTL*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Dawson, B. (1953). *Acta Cryst.* **6**, 81–87.
- Delfino, M., Loiacono, G. M. & Nicolosi, J. A. (1978). *J. Solid State Chem.* **23**, 289–296.
- Flack, H. D. (1983). *Acta Cryst.* **A39**, 876–881.
- Kirfel, A. & Wallrafen, F. (1985). *Z. Kristallogr.* **171**, 121–126.
- Rigaku (1998). *PROCESS-AUTO*. Rigaku Corporation, Tokyo, Japan.
- Rigaku/MSC (2004). *CrystalStructure*. Version 3.6.0. Rigaku/MSC, The Woodlands, Texas, USA.
- Sequeira, A., Rajagopal, H. & Chidambaram, R. (1972). *Acta Cryst.* **B28**, 2514–2519.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.

# supporting information

*Acta Cryst.* (2008). E64, o446 [doi:10.1107/S1600536807068560]

## L-Glutamic acid hydrochloride at 153 K

**Yi Jian Zhang, Zhan Shu, Wei Xu, Gang Chen and Zu-Guang Li**

### S1. Comment

The crystal structure of the title compound has been investigated by Dawson (1953) using the multiple-film technique for  $h0l$  and  $0kl$  intensity data collection, then by Sequeira *et al.* (1972) using neutron diffraction. Besides, the L-glutamic acid (LGA) halogen acid salts of LGA hydroiodide (Kirfel & Wallrafen, 1985), LGA HBr and LGA HCl HBr (Delfino *et al.*, 1978) have been reported before. Recently we have made a structure determination of the title compound at 153 K and make comparisons with Dawson's and Sequeira's studies, hereby.

The unit-cell parameters and space group of the title compound (see crystal data) and the respective values in Dawson's structure [ $a = 5.16$  (1),  $b = 11.80$  (2),  $c = 13.30$  (2) Å] and Sequeira's structure [ $a = 5.151$  (6),  $b = 11.789$  (19),  $c = 13.347$  (20) Å] demonstrate the high quality of modern X-ray diffractometer and software. The final  $R$  value of the present case [0.0171] is much smaller than that of Dawson's [0.106] and Sequeira's [0.043]. The hydrogen atoms were not determined in Dawson's study, due to the incompleteness of the estimated intensity data.

The C4—C5 [1.5066 (15) Å] bond length intermediates Dawson's [1.54 Å] and Sequeira's [1.475 (7) Å] (Table 1). The C2—C3 distance [1.5312 (15) Å] differs from that of Dawson's [1.55 Å], the C1—C2 [1.5156 (16) Å] deviates from that of Sequeira's [1.535 (7) Å]. The C—O, C—N and the other C—C bond lengths of these three cases have little distinctions respectively. Between the present and Dawson's study, the major distinctions of bond angles lie in C1—C2—N1 and C3—C2—N1 [108.02 (9) and 111.94 (9)° for present, 111 and 110° for Dawson's case, respectively]. The torsion angles of the main skeleton are finely coherent with those in Sequeira's study, respectively (Table 1). But, some torsion angles related to H atoms that are involved in hydrogen bonds are deviated greatly from those in Sequeira's study, respectively [H1C—N1—C2—C1 (-60.0, -48.0°), H1B—N1—C2—C1 (60.0, 73.2°) and H1A—N1—C2—C1 (-180.0, -165.1°); H1—O1—C1—C2 (179.8, 168.9°), the former and latter values for the present (Table 1) and Sequeira's study, respectively].

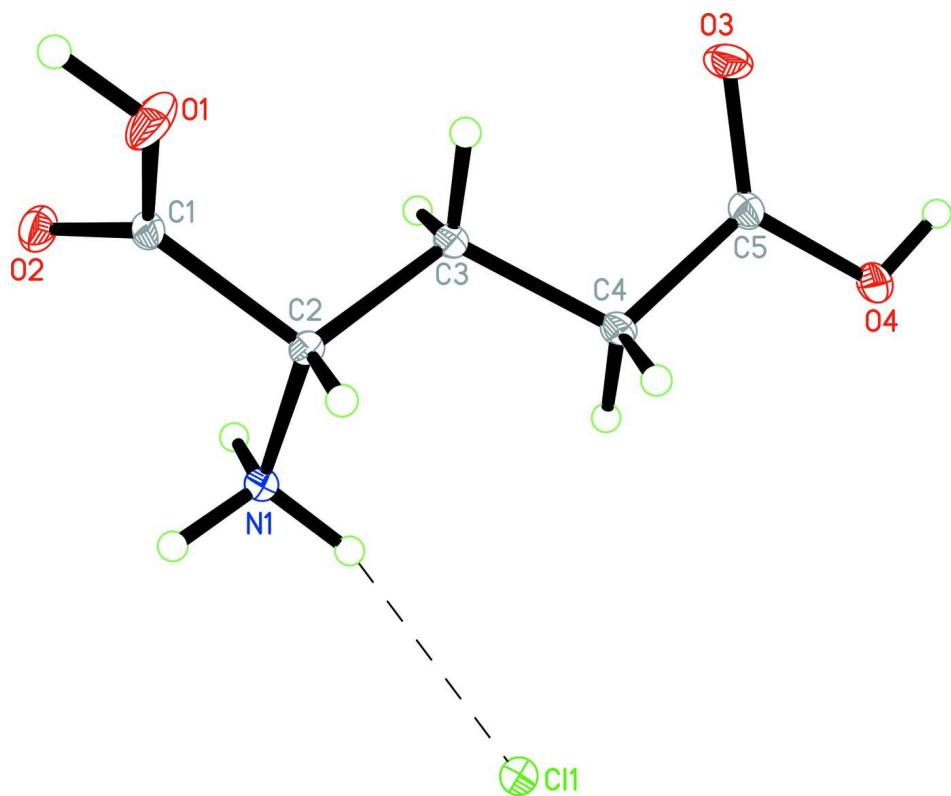
In the title compound, L-glutamic acid is protonated and is linked with the Cl<sup>-</sup> anion by a O—H···Cl hydrogen bond (Fig. 1, Table 2). The crystal structure is established by a three-dimensional network of O—H···O, N—H···O and N—H···Cl classic hydrogen bonds (Fig. 2, Table 2).

### S2. Experimental

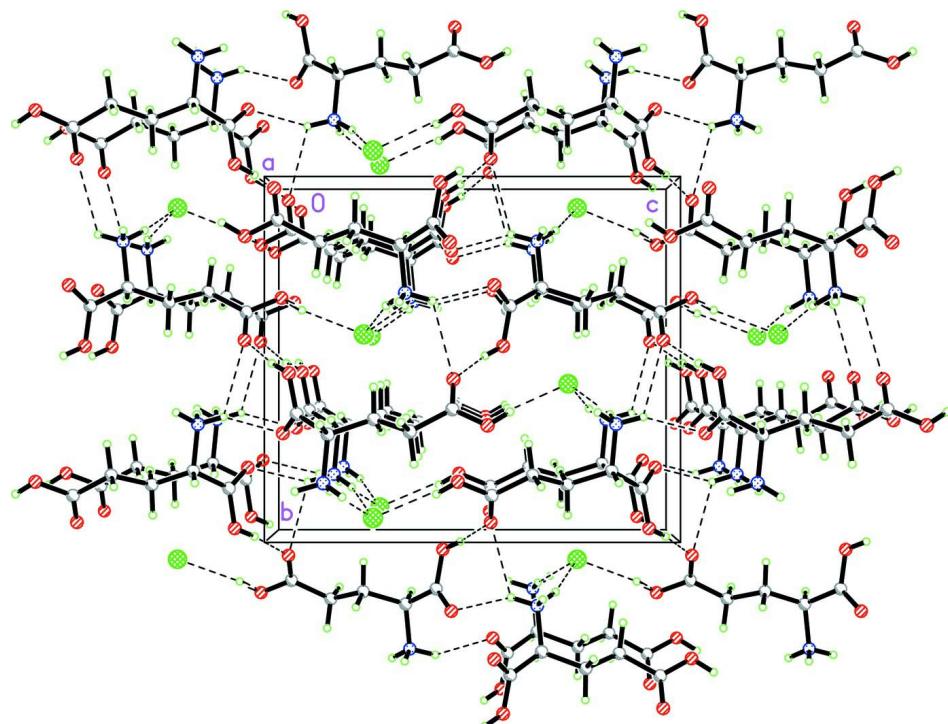
L-Glutamic acid and hydrochloric acid in equal molar ratio were mixed together with enough water, and heated to a temperature where clear solution was obtained. The solution were layed in a ventilative place. Colorless single crystals of the title compound were obtained from the solution by slow volatilization of water at room temperature for 7 days.

### S3. Refinement

All hydrogen atoms were placed in calculated positions and allowed to ride on their parent atoms at distances of 0.99 (methylene), 1.00 (methine), 0.91 Å (N—H), 0.84 Å (O—H), with  $U_{\text{iso}}(\text{H})$  values 1.2–1.5 times  $U_{\text{eq}}$  of the parent atoms.

**Figure 1**

A view of the title compound showing with 40% probability displacement ellipsoids. Hydrogen bonds are indicated by thin lines.

**Figure 2**

The packing diagram of the title compound viewed down along the  $a$  axis. Hydrogen bonds are indicated by thin lines.

### (S)-1,3-dicarboxypropanaminium chloride

#### Crystal data



$M_r = 183.59$

Orthorhombic,  $P2_12_12_1$

Hall symbol: P 2ac 2ab

$a = 5.1016(1)$  Å

$b = 11.6386(4)$  Å

$c = 13.2500(3)$  Å

$V = 786.73(4)$  Å<sup>3</sup>

$Z = 4$

$F(000) = 384$

$D_x = 1.550$  Mg m<sup>-3</sup>

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

Cell parameters from 7746 reflections

$\theta = 3.1\text{--}27.5^\circ$

$\mu = 0.45$  mm<sup>-1</sup>

$T = 153$  K

Block, colorless

$0.49 \times 0.48 \times 0.35$  mm

#### Data collection

Siemens P4

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

$\omega$  scans

Absorption correction:  $\psi$  scan  
(*SHELXTL*; Bruker, 1998)

$T_{\min} = 0.809$ ,  $T_{\max} = 0.858$

7742 measured reflections

1798 independent reflections

1775 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.016$

$\theta_{\max} = 27.5^\circ$ ,  $\theta_{\min} = 3.1^\circ$

$h = -6 \rightarrow 6$

$k = -15 \rightarrow 15$

$l = -15 \rightarrow 16$

*Refinement*Refinement on  $F^2$ 

Least-squares matrix: full

$$R[F^2 > 2\sigma(F^2)] = 0.019$$

$$wR(F^2) = 0.055$$

$$S = 1.15$$

1798 reflections

101 parameters

0 restraints

Primary atom site location: structure-invariant  
direct methodsSecondary atom site location: difference Fourier  
mapHydrogen site location: inferred from  
neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0313P)^2 + 0.1681P]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

$$(\Delta/\sigma)_{\max} = 0.001$$

$$\Delta\rho_{\max} = 0.29 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.25 \text{ e \AA}^{-3}$$

Extinction correction: *SHELXL97* (Sheldrick,  
1997),  $F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$ 

Extinction coefficient: 0.058 (4)

Absolute structure: Flack (1983), 725 Friedel  
pairs

Absolute structure parameter: 0.02 (4)

*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 ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C11	1.05471 (5)	0.93130 (2)	0.26203 (2)	0.01395 (9)
O1	0.4387 (2)	0.53632 (7)	0.07337 (7)	0.0205 (2)
H1	0.3341	0.5102	0.0301	0.031*
O2	0.20701 (18)	0.69846 (7)	0.05162 (7)	0.0161 (2)
O3	0.4214 (2)	0.55251 (8)	0.45585 (6)	0.0204 (2)
O4	0.70041 (18)	0.67059 (8)	0.53439 (6)	0.0167 (2)
H4	0.6454	0.6361	0.5859	0.025*
N1	0.5606 (2)	0.82410 (8)	0.15658 (7)	0.0120 (2)
H1A	0.6701	0.8569	0.2025	0.018*
H1B	0.3939	0.8490	0.1676	0.018*
H1C	0.6117	0.8442	0.0932	0.018*
C1	0.3864 (2)	0.64491 (10)	0.08985 (9)	0.0116 (2)
C2	0.5705 (2)	0.69659 (9)	0.16728 (8)	0.0102 (2)
H2	0.7532	0.6689	0.1547	0.012*
C3	0.4798 (2)	0.65720 (10)	0.27187 (8)	0.0115 (2)
H3A	0.3074	0.6926	0.2863	0.014*
H3B	0.4550	0.5728	0.2705	0.014*
C4	0.6668 (2)	0.68690 (10)	0.35714 (8)	0.0124 (2)
H4A	0.8458	0.6610	0.3395	0.015*
H4B	0.6712	0.7713	0.3666	0.015*
C5	0.5815 (3)	0.63012 (9)	0.45392 (8)	0.0119 (2)

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

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C11	0.01329 (14)	0.01559 (13)	0.01296 (14)	-0.00183 (11)	-0.00097 (10)	0.00137 (10)
O1	0.0232 (4)	0.0145 (4)	0.0238 (4)	0.0019 (4)	-0.0111 (4)	-0.0082 (3)
O2	0.0166 (4)	0.0164 (4)	0.0153 (4)	0.0004 (3)	-0.0052 (3)	0.0000 (3)
O3	0.0284 (5)	0.0194 (5)	0.0134 (4)	-0.0119 (4)	0.0021 (4)	0.0018 (3)
O4	0.0214 (5)	0.0196 (4)	0.0089 (4)	-0.0053 (4)	-0.0014 (4)	0.0029 (3)
N1	0.0134 (4)	0.0119 (4)	0.0105 (4)	-0.0028 (4)	-0.0012 (4)	0.0014 (3)
C1	0.0119 (6)	0.0148 (5)	0.0081 (5)	-0.0022 (4)	0.0004 (4)	0.0006 (4)
C2	0.0107 (5)	0.0100 (5)	0.0101 (5)	-0.0009 (5)	-0.0006 (5)	-0.0006 (4)
C3	0.0121 (5)	0.0128 (5)	0.0096 (5)	-0.0025 (4)	0.0006 (4)	0.0016 (4)
C4	0.0138 (5)	0.0146 (5)	0.0089 (5)	-0.0039 (5)	0.0011 (4)	0.0014 (4)
C5	0.0139 (6)	0.0114 (5)	0.0105 (5)	0.0002 (4)	0.0007 (5)	0.0005 (4)

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

O1—C1	1.3101 (15)	C1—C2	1.5156 (16)
O1—H1	0.8400	C2—C3	1.5312 (15)
O2—C1	1.2176 (15)	C2—H2	1.0000
O3—C5	1.2183 (15)	C3—C4	1.5184 (16)
O4—C5	1.3139 (14)	C3—H3A	0.9900
O4—H4	0.8400	C3—H3B	0.9900
N1—C2	1.4916 (14)	C4—C5	1.5066 (15)
N1—H1A	0.9100	C4—H4A	0.9900
N1—H1B	0.9100	C4—H4B	0.9900
N1—H1C	0.9100		
C1—O1—H1	109.5	C3—C2—H2	109.7
C5—O4—H4	109.5	C4—C3—C2	114.55 (9)
C2—N1—H1A	109.5	C4—C3—H3A	108.6
C2—N1—H1B	109.5	C2—C3—H3A	108.6
H1A—N1—H1B	109.5	C4—C3—H3B	108.6
C2—N1—H1C	109.5	C2—C3—H3B	108.6
H1A—N1—H1C	109.5	H3A—C3—H3B	107.6
H1B—N1—H1C	109.5	C5—C4—C3	110.63 (10)
O2—C1—O1	125.29 (11)	C5—C4—H4A	109.5
O2—C1—C2	122.99 (10)	C3—C4—H4A	109.5
O1—C1—C2	111.67 (10)	C5—C4—H4B	109.5
N1—C2—C1	108.02 (9)	C3—C4—H4B	109.5
N1—C2—C3	111.94 (9)	H4A—C4—H4B	108.1
C1—C2—C3	107.85 (9)	O3—C5—O4	123.93 (10)
N1—C2—H2	109.7	O3—C5—C4	122.46 (10)
C1—C2—H2	109.7	O4—C5—C4	113.59 (10)
O2—C1—C2—N1	-21.37 (15)	C3—C4—C5—O3	14.84 (16)
O1—C1—C2—N1	161.21 (10)	C3—C4—C5—O4	-166.61 (10)
O2—C1—C2—C3	99.80 (13)	H1C—N1—C2—C1	-60.0

O1—C1—C2—C3	−77.62 (12)	H1B—N1—C2—C1	60.0
N1—C2—C3—C4	−69.92 (13)	H1A—N1—C2—C1	−180.0
C1—C2—C3—C4	171.39 (9)	H1—O1—C1—C2	179.8
C2—C3—C4—C5	−171.71 (9)		

*Hydrogen-bond geometry (Å, °)*

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1A···Cl1	0.91	2.29	3.1406 (10)	157
N1—H1B···Cl1 <sup>i</sup>	0.91	2.34	3.1890 (10)	155
N1—H1C···O2 <sup>ii</sup>	0.91	2.04	2.8700 (13)	151
O1—H1···O3 <sup>iii</sup>	0.84	1.79	2.6208 (13)	170
O4—H4···Cl1 <sup>iv</sup>	0.84	2.21	3.0389 (9)	168

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