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Key indicators

Single-crystal X-ray study
 $T = 150\text{ K}$
 Mean $\sigma(\text{C-C}) = 0.002\text{ \AA}$
 R factor = 0.031
 wR factor = 0.084
 Data-to-parameter ratio = 12.3

For details of how these key indicators were automatically derived from the article, see
<http://journals.iucr.org/e>.

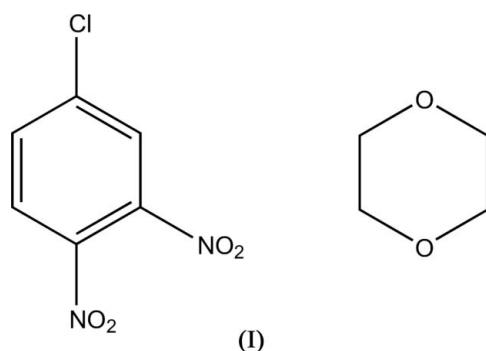
1-Chloro-3,4-dinitrobenzene–1,4-dioxane (1/1)

Received 21 March 2006
 Accepted 28 March 2006

The solvate structure of 1-chloro-3,4-dinitrobenzene with 1,4-dioxane, $\text{C}_6\text{H}_3\text{ClN}_2\text{O}_4\cdot\text{C}_4\text{H}_8\text{N}_2$, is reported. Alternating molecules of 3,4-dinitro-1-chlorobenzene and 1,4-dioxane are linked by $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds into a continuous two-dimensional sheet.

Comment

The title compound, (I), was produced during an experimental crystallization polymorph screen on 1-chloro-3,4-dinitrobenzene (3,4-DNCB). Compound (I) crystallizes in the space group $P\bar{1}$ with one molecule of 3,4-DNCB and one molecule of 1,4-dioxane in the asymmetric unit (Fig. 1).



The crystal structure of (I) is characterized by alternating molecules of 3,4-DNCB and 1,4-dioxane, linked by a series of $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds (Table 1) into a continuous two-dimensional sheet which lies parallel to the $(1\bar{1}\bar{1})$ plane (Fig. 2). Alternating 3,4-DNCB molecules and 1,4-dioxane are linked by pairwise $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds, forming a chain

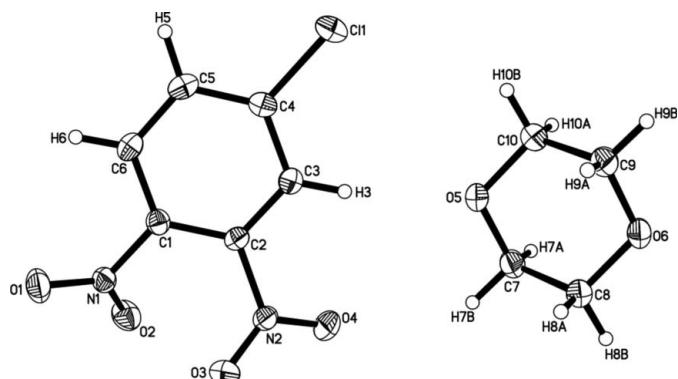
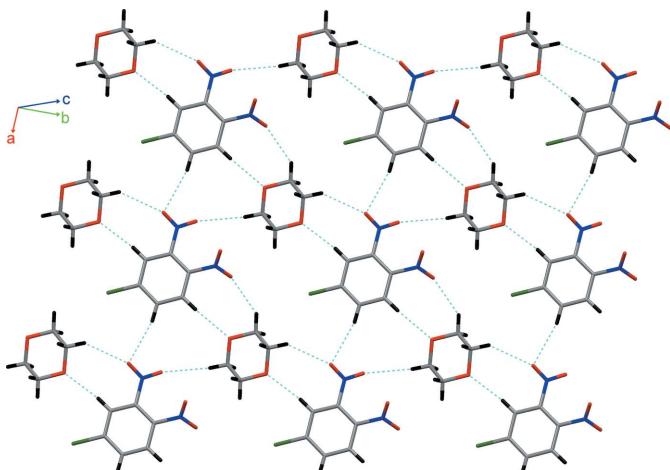


Figure 1

The structure of the asymmetric unit of (I), showing the numbering scheme used. Displacement ellipsoids are drawn at the 50% probability level.

**Figure 2**

The two-dimensional network formed by compound (I), showing the intermolecular interactions as thin pale-blue lines. Key: C grey, N blue, O red, Cl green and H black.

which runs parallel to the body diagonal (111). These chains are then hydrogen-bonded together, forming a sheet *via* two C—H···O interactions between two 3,4-DNCB molecules and one C—H···O interaction between the 3,4-DNCB molecule and a 1,4-dioxane molecule of the adjacent chain. Viewing the crystal structure down the *a* axis reveals that there are alternating layers of 3,4-DNCB and 1,4-dioxane (Fig. 3).

Experimental

The title compound was recrystallized from 1,4-dioxane solution by slow evaporation at 298 K.

Crystal data

$C_6H_3ClN_2O_4 \cdot C_4H_8O_2$	$Z = 2$
$M_r = 290.66$	$D_x = 1.616 \text{ Mg m}^{-3}$
Triclinic, $P\bar{1}$	Mo $K\alpha$ radiation
$a = 8.2976 (12) \text{ \AA}$	Cell parameters from 3669
$b = 8.7112 (13) \text{ \AA}$	reflections
$c = 8.8015 (13) \text{ \AA}$	$\theta = 2.5\text{--}28.2^\circ$
$\alpha = 103.661 (2)^\circ$	$\mu = 0.35 \text{ mm}^{-1}$
$\beta = 103.909 (2)^\circ$	$T = 150 (2) \text{ K}$
$\gamma = 91.718 (2)^\circ$	Block, yellow
$V = 597.52 (15) \text{ \AA}^3$	$0.86 \times 0.67 \times 0.45 \text{ mm}$

Data collection

Bruker SMART APEX diffractometer	2650 independent reflections
Narrow-frame ω scans	2529 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2001)	$R_{\text{int}} = 0.014$
$T_{\min} = 0.738$, $T_{\max} = 0.856$	$\theta_{\max} = 28.2^\circ$
4424 measured reflections	$h = -10 \rightarrow 10$
	$k = -11 \rightarrow 11$
	$l = -11 \rightarrow 11$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0478P)^2 + 0.1587P]$
$R[F^2 > 2\sigma(F^2)] = 0.031$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.084$	$(\Delta/\sigma)_{\max} < 0.001$
$S = 1.07$	$\Delta\rho_{\max} = 0.36 \text{ e \AA}^{-3}$
2647 reflections	$\Delta\rho_{\min} = -0.32 \text{ e \AA}^{-3}$
216 parameters	
All H-atom parameters refined	

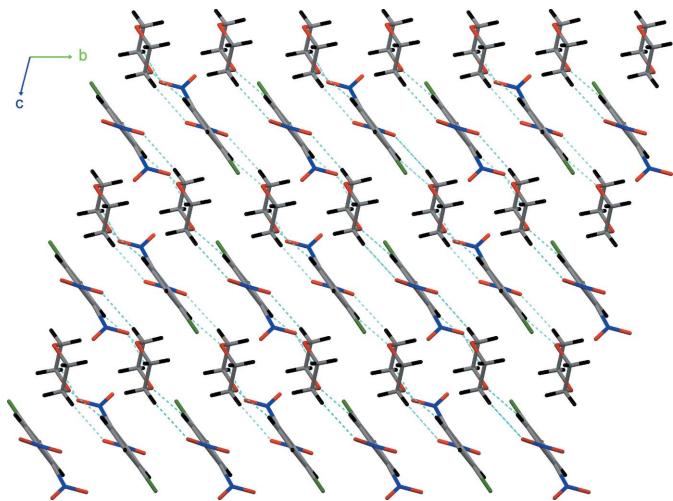
$$w = 1/[\sigma^2(F_o^2) + (0.0478P)^2 + 0.1587P]$$

where $P = (F_o^2 + 2F_c^2)/3$

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

$$\Delta\rho_{\max} = 0.36 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.32 \text{ e \AA}^{-3}$$

**Figure 3**

Packing diagram, showing the stacking of the sheets. C—H···O interactions are shown as thin pale-blue lines. Key: C grey, N blue, O red, Cl green and H black.

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

$D\text{--H}\cdots A$	$D\text{--H}$	$H\cdots A$	$D\cdots A$	$D\text{--H}\cdots A$
$C3\text{--H}3\cdots O5$	0.938 (15)	2.331 (15)	3.2553 (14)	168.3 (13)
$C5\text{--H}5\cdots O4^i$	0.969 (17)	2.697 (18)	3.6166 (16)	158.7 (13)
$C6\text{--H}6\cdots O6^{ii}$	0.949 (15)	2.468 (15)	3.4017 (14)	168.1 (12)
$C7\text{--H}7B\cdots O4$	0.990 (17)	2.624 (17)	3.4618 (16)	142.5 (13)
$C8\text{--H}8B\cdots O1^{iii}$	0.996 (19)	2.446 (18)	3.2604 (16)	138.6 (13)
$C9\text{--H}9B\cdots O3^{iv}$	0.951 (18)	2.649 (18)	3.5930 (16)	172.3 (14)

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

H atoms were located in a difference Fourier map and refined freely [$\text{C--H} = 0.938 (15)\text{--}0.996 (19) \text{ \AA}$]. The three reflections with the greatest discrepancies were omitted from the refinement.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2000) and *MERCURY* (Bruno *et al.*, 2002); software used to prepare material for publication: *SHELXL97* and *PLATON* (Spek, 2003).

The authors acknowledge the Research Councils UK Basic Technology Programme for supporting ‘Control and Prediction of the Organic Solid State’ (URL: www.cpos.org.uk).

References

- Bruker (2000). *SHELXTL*. Version 6.10. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruker (2001). *SMART* (Version 5.625), *SAINT* (Version 6.22) and *SADABS* (Version 2.03). Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruno, I. J., Cole, J. C., Edgington, P. R., Kessler, M. K., Macrae, C. F., McCabe, P., Pearson, J. & Taylor, R. (2002). *Acta Cryst.* **B58**, 389–397.
- Sheldrick, G. M. (1997). *SHELXS97 & SHELXL97*. University of Göttingen, Germany.
- Spek, A. L. (2003). *J. Appl. Cryst.* **36**, 7–13.

supporting information

Acta Cryst. (2006). E62, o1662–o1663 [https://doi.org/10.1107/S1600536806011226]

1-Chloro-3,4-dinitrobenzene–1,4-dioxane (1/1)

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4-Chloro-1,2-dinitrobenzene–1,4-dioxane (1/1)

Crystal data



$M_r = 290.66$

Triclinic, $P\bar{1}$

$a = 8.2976 (12) \text{ \AA}$

$b = 8.7112 (13) \text{ \AA}$

$c = 8.8015 (13) \text{ \AA}$

$\alpha = 103.661 (2)^\circ$

$\beta = 103.909 (2)^\circ$

$\gamma = 91.718 (2)^\circ$

$V = 597.52 (15) \text{ \AA}^3$

$Z = 2$

$F(000) = 300$

$D_x = 1.616 \text{ Mg m}^{-3}$

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

Cell parameters from 3669 reflections

$\theta = 2.5\text{--}28.2^\circ$

$\mu = 0.35 \text{ mm}^{-1}$

$T = 150 \text{ K}$

Block, yellow

$0.86 \times 0.67 \times 0.45 \text{ mm}$

Data collection

Bruker SMART APEX

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Narrow-frame ω scans

Absorption correction: multi-scan
(SADABS; Bruker, 2001)

$T_{\min} = 0.738$, $T_{\max} = 0.856$

4424 measured reflections

2650 independent reflections

2529 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.014$

$\theta_{\max} = 28.2^\circ$, $\theta_{\min} = 2.4^\circ$

$h = -10 \rightarrow 10$

$k = -11 \rightarrow 11$

$l = -11 \rightarrow 11$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.031$

$wR(F^2) = 0.084$

$S = 1.07$

2647 reflections

216 parameters

0 restraints

Primary atom site location: Direct methods

Secondary atom site location: Difference

Fourier synthesis

Hydrogen site location: Found from delta-F

All H-atom parameters refined

$w = 1/[\sigma^2(F_o^2) + (0.0478P)^2 + 0.1587P]$

where $P = (F_o^2 + 2F_c^2)/3$

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

$\Delta\rho_{\max} = 0.36 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.32 \text{ e \AA}^{-3}$

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.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.01692 (14)	0.14991 (13)	0.38735 (13)	0.0189 (2)
C2	0.13820 (13)	0.25986 (13)	0.49998 (13)	0.0182 (2)
C3	0.09882 (14)	0.37166 (13)	0.62157 (13)	0.0198 (2)
H3	0.1791 (19)	0.4482 (18)	0.6968 (18)	0.024 (4)*
C4	-0.06714 (14)	0.36928 (13)	0.62793 (13)	0.0205 (2)
C5	-0.18948 (14)	0.25951 (15)	0.51873 (14)	0.0233 (2)
H5	-0.304 (2)	0.262 (2)	0.526 (2)	0.035 (4)*
C6	-0.14693 (14)	0.14773 (14)	0.39675 (14)	0.0218 (2)
H6	-0.2275 (19)	0.0712 (18)	0.3186 (18)	0.022 (3)*
Cl1	-0.11770 (4)	0.50731 (4)	0.78178 (4)	0.02978 (11)
N1	0.05511 (12)	0.04236 (11)	0.24729 (11)	0.0204 (2)
O1	-0.00297 (12)	-0.09537 (10)	0.21122 (11)	0.0300 (2)
O2	0.13920 (12)	0.09876 (11)	0.17386 (10)	0.0295 (2)
N2	0.31546 (12)	0.25335 (12)	0.50183 (11)	0.0217 (2)
O3	0.36588 (11)	0.12279 (11)	0.46586 (11)	0.0301 (2)
O4	0.40217 (11)	0.37924 (11)	0.54379 (12)	0.0318 (2)
C7	0.50812 (15)	0.71687 (15)	0.86037 (14)	0.0265 (3)
H7A	0.512 (2)	0.821 (2)	0.8349 (19)	0.034 (4)*
H7B	0.533 (2)	0.635 (2)	0.772 (2)	0.036 (4)*
C8	0.62967 (15)	0.72245 (16)	1.01944 (15)	0.0264 (3)
H8A	0.6304 (19)	0.6196 (19)	1.0408 (18)	0.026 (4)*
H8B	0.746 (2)	0.756 (2)	1.020 (2)	0.040 (4)*
C9	0.42101 (16)	0.79512 (16)	1.15303 (14)	0.0273 (3)
H9A	0.414 (2)	0.695 (2)	1.1806 (18)	0.029 (4)*
H9B	0.398 (2)	0.875 (2)	1.237 (2)	0.039 (4)*
C10	0.29983 (16)	0.78846 (16)	0.99319 (15)	0.0272 (3)
H10A	0.297 (2)	0.891 (2)	0.970 (2)	0.037 (4)*
H10B	0.186 (2)	0.757 (2)	0.991 (2)	0.035 (4)*
O5	0.34241 (11)	0.67501 (10)	0.86581 (10)	0.0258 (2)
O6	0.58807 (11)	0.83433 (11)	1.14873 (10)	0.0273 (2)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0205 (5)	0.0180 (5)	0.0177 (5)	0.0016 (4)	0.0039 (4)	0.0044 (4)
C2	0.0155 (5)	0.0194 (5)	0.0203 (5)	0.0011 (4)	0.0046 (4)	0.0058 (4)
C3	0.0193 (5)	0.0190 (5)	0.0204 (5)	0.0002 (4)	0.0045 (4)	0.0046 (4)
C4	0.0223 (5)	0.0211 (5)	0.0207 (5)	0.0051 (4)	0.0083 (4)	0.0068 (4)
C5	0.0170 (5)	0.0298 (6)	0.0248 (5)	0.0025 (4)	0.0055 (4)	0.0099 (5)
C6	0.0179 (5)	0.0246 (6)	0.0211 (5)	-0.0012 (4)	0.0014 (4)	0.0067 (4)

C11	0.02999 (17)	0.03041 (18)	0.02956 (17)	0.00620 (12)	0.01481 (12)	0.00096 (12)
N1	0.0198 (4)	0.0199 (5)	0.0190 (4)	0.0023 (3)	0.0018 (3)	0.0031 (4)
O1	0.0339 (5)	0.0195 (4)	0.0312 (4)	-0.0023 (3)	0.0049 (4)	0.0000 (3)
O2	0.0367 (5)	0.0284 (5)	0.0260 (4)	0.0016 (4)	0.0151 (4)	0.0047 (3)
N2	0.0168 (4)	0.0261 (5)	0.0198 (4)	0.0005 (4)	0.0047 (3)	0.0017 (4)
O3	0.0229 (4)	0.0300 (5)	0.0334 (5)	0.0083 (3)	0.0067 (4)	0.0005 (4)
O4	0.0217 (4)	0.0307 (5)	0.0390 (5)	-0.0074 (4)	0.0089 (4)	0.0010 (4)
C7	0.0261 (6)	0.0276 (6)	0.0228 (5)	-0.0011 (5)	0.0077 (5)	-0.0002 (5)
C8	0.0224 (6)	0.0268 (6)	0.0258 (6)	0.0001 (5)	0.0058 (4)	-0.0009 (5)
C9	0.0252 (6)	0.0327 (7)	0.0218 (5)	-0.0001 (5)	0.0070 (5)	0.0017 (5)
C10	0.0240 (6)	0.0301 (6)	0.0242 (6)	0.0034 (5)	0.0061 (5)	0.0002 (5)
O5	0.0224 (4)	0.0267 (4)	0.0223 (4)	-0.0012 (3)	0.0034 (3)	-0.0027 (3)
O6	0.0239 (4)	0.0292 (5)	0.0225 (4)	-0.0033 (3)	0.0045 (3)	-0.0036 (3)

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

C1—C6	1.3816 (16)	N2—O3	1.2210 (14)
C1—C2	1.3881 (15)	C7—O5	1.4267 (14)
C1—N1	1.4691 (14)	C7—C8	1.5052 (18)
C2—C3	1.3768 (15)	C7—H7A	0.982 (17)
C2—N2	1.4700 (14)	C7—H7B	0.990 (17)
C3—C4	1.3913 (15)	C8—O6	1.4314 (14)
C3—H3	0.938 (15)	C8—H8A	0.958 (16)
C4—C5	1.3811 (16)	C8—H8B	0.996 (19)
C4—C11	1.7291 (11)	C9—O6	1.4292 (15)
C5—C6	1.3908 (17)	C9—C10	1.5071 (17)
C5—H5	0.969 (17)	C9—H9A	0.962 (17)
C6—H6	0.949 (15)	C9—H9B	0.951 (18)
N1—O1	1.2184 (13)	C10—O5	1.4306 (14)
N1—O2	1.2215 (13)	C10—H10A	0.964 (18)
N2—O4	1.2208 (13)	C10—H10B	0.973 (17)
C6—C1—C2	120.56 (10)	O5—C7—H7A	108.9 (10)
C6—C1—N1	117.89 (10)	C8—C7—H7A	110.4 (10)
C2—C1—N1	121.33 (10)	O5—C7—H7B	107.0 (10)
C3—C2—C1	121.33 (10)	C8—C7—H7B	110.9 (10)
C3—C2—N2	117.30 (9)	H7A—C7—H7B	109.0 (13)
C1—C2—N2	121.19 (10)	O6—C8—C7	110.89 (10)
C2—C3—C4	117.37 (10)	O6—C8—H8A	109.6 (9)
C2—C3—H3	122.1 (9)	C7—C8—H8A	110.3 (9)
C4—C3—H3	120.5 (9)	O6—C8—H8B	106.8 (10)
C5—C4—C3	122.32 (10)	C7—C8—H8B	111.9 (10)
C5—C4—C11	119.85 (9)	H8A—C8—H8B	107.3 (14)
C3—C4—C11	117.82 (9)	O6—C9—C10	111.09 (10)
C4—C5—C6	119.34 (10)	O6—C9—H9A	109.7 (10)
C4—C5—H5	120.1 (10)	C10—C9—H9A	110.3 (9)
C6—C5—H5	120.6 (10)	O6—C9—H9B	106.5 (11)
C1—C6—C5	119.06 (10)	C10—C9—H9B	110.7 (11)

C1—C6—H6	119.0 (9)	H9A—C9—H9B	108.3 (14)
C5—C6—H6	121.9 (9)	O5—C10—C9	110.50 (10)
O1—N1—O2	125.13 (10)	O5—C10—H10A	110.2 (10)
O1—N1—C1	117.35 (9)	C9—C10—H10A	110.8 (11)
O2—N1—C1	117.50 (9)	O5—C10—H10B	106.6 (10)
O4—N2—O3	124.94 (10)	C9—C10—H10B	113.0 (10)
O4—N2—C2	117.27 (10)	H10A—C10—H10B	105.5 (14)
O3—N2—C2	117.76 (9)	C7—O5—C10	109.34 (9)
O5—C7—C8	110.53 (10)	C9—O6—C8	109.74 (9)
C6—C1—C2—C3	-1.45 (17)	C2—C1—N1—O1	137.79 (11)
N1—C1—C2—C3	173.08 (10)	C6—C1—N1—O2	130.77 (11)
C6—C1—C2—N2	173.60 (10)	C2—C1—N1—O2	-43.91 (15)
N1—C1—C2—N2	-11.86 (16)	C3—C2—N2—O4	-37.51 (14)
C1—C2—C3—C4	0.51 (17)	C1—C2—N2—O4	147.24 (11)
N2—C2—C3—C4	-174.73 (10)	C3—C2—N2—O3	140.47 (11)
C2—C3—C4—C5	0.50 (17)	C1—C2—N2—O3	-34.78 (15)
C2—C3—C4—Cl1	179.11 (8)	O5—C7—C8—O6	-58.54 (13)
C3—C4—C5—C6	-0.58 (18)	O6—C9—C10—O5	57.95 (14)
Cl1—C4—C5—C6	-179.16 (9)	C8—C7—O5—C10	58.67 (13)
C2—C1—C6—C5	1.35 (17)	C9—C10—O5—C7	-58.31 (13)
N1—C1—C6—C5	-173.37 (10)	C10—C9—O6—C8	-56.48 (14)
C4—C5—C6—C1	-0.36 (17)	C7—C8—O6—C9	56.69 (13)
C6—C1—N1—O1	-47.53 (14)		

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
C3—H3···O5	0.938 (15)	2.331 (15)	3.2553 (14)	168.3 (13)
C5—H5···O4 ⁱ	0.969 (17)	2.697 (18)	3.6166 (16)	158.7 (13)
C6—H6···O6 ⁱⁱ	0.949 (15)	2.468 (15)	3.4017 (14)	168.1 (12)
C7—H7B···O4	0.990 (17)	2.624 (17)	3.4618 (16)	142.5 (13)
C8—H8B···O1 ⁱⁱⁱ	0.996 (19)	2.446 (18)	3.2604 (16)	138.6 (13)
C9—H9B···O3 ^{iv}	0.951 (18)	2.649 (18)	3.5930 (16)	172.3 (14)

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