

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

2,2-Dichloro-1-[(2*R*,5*S*)-5-ethyl-2-methyl-2-propyl-1,3-oxazolidin-3-yl]ethanone

Ying Fu, Fei Ye* and Xiaotian Wen

College of Science, Northeast Agricultural University, Harbin 150030, People's Republic of China

Correspondence e-mail: yefei@neau.edu.cn

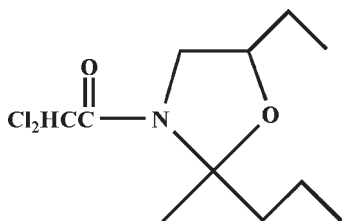
Received 23 April 2010; accepted 1 May 2010

 Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.009$ Å; R factor = 0.084; wR factor = 0.258; data-to-parameter ratio = 23.6.

In the title compound, $\text{C}_{11}\text{H}_{19}\text{Cl}_2\text{NO}_2$, the oxazolidine ring is in an envelope conformation with the O atom forming the flap. In the crystal structure, molecules are linked by weak intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds, forming chains.

Related literature

For general background to *N*-dichloroacetyl oxazolidine, see: Agami & Couty (2004); Abu-Qare & Duncan (2002); Guirado *et al.* (2003); Davies & Caseley (1999). For the bioactivity of related compounds, see: Del Buono *et al.* (2007); Hatzios & Burgos (2004). For details of the synthesis, see: Fu *et al.* (2009).



Experimental

Crystal data

 $\text{C}_{11}\text{H}_{19}\text{Cl}_2\text{NO}_2$
 $M_r = 268.17$

 Orthorhombic, $P2_12_12_1$
 $a = 6.4834$ (12) Å

 $b = 10.795$ (2) Å

 $c = 20.030$ (4) Å

 $V = 1401.8$ (5) Å³
 $Z = 4$

 Mo $K\alpha$ radiation

 $\mu = 0.45$ mm⁻¹
 $T = 293$ K

 $0.32 \times 0.24 \times 0.20$ mm

Data collection

Bruker SMART CCD

diffractometer

Absorption correction: multi-scan

 (*SADABS*; Sheldrick, 1996)

 $T_{\min} = 0.869$, $T_{\max} = 0.915$

14134 measured reflections

3499 independent reflections

 2117 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.058$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.084$
 $wR(F^2) = 0.258$
 $S = 1.02$

3499 reflections

148 parameters

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.58$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.39$ e Å⁻³

Absolute structure: Flack (1983)

1468 Friedels

Flack parameter: 0.02 (15)

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C}1-\text{H}1\cdots\text{O}2^i$	0.98	2.38	3.327 (5)	163

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

Data collection: *SMART* (Bruker, 1998); cell refinement: *S SAINT* (Bruker, 1998); data reduction: *S SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

We thank the Heilongjiang Province Foundation for Young Scholars (QC2009C44), the Innovation Foundation for Young Scholars of Harbin (No.2007RFQXN017), the China Postdoctoral Science Foundation (20080430951), the Heilongjiang Province Postdoctoral Science Foundation and the Northeast Agricultural University Doctoral Foundation for generously supporting this study.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH5034).

References

- Abu-Qare, A. W. & Duncan, H. J. (2002). *Chemosphere*, **48**, 965–974.
- Agami, C. & Couty, F. (2004). *Eur. J. Org. Chem.* **69**, 677–685.
- Bruker (1998). *SMART* and *S SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Davies, J. & Caseley, J. C. (1999). *Pestic. Sci.* **55**, 1043–1046.
- Del Buono, D., Scarponi, L. & Espen, L. (2007). *Phytochemistry*, **68**, 2614–2618.
- Flack, H. D. (1983). *Acta Cryst.* **A39**, 876–881.
- Fu, Y., Fu, H. G., Ye, F., Mao, J. D. & Wen, X. T. (2009). *Synth. Commun.* **39**, 2454–2463.
- Guirado, A., Andreu, R. & Galvez, J. (2003). *Tetrahedron Lett.* **44**, 3809–3841.
- Hatzios, K. K. & Burgos, N. (2004). *Weed Sci.* **52**, 454–467.
- Sheldrick, G. M. (1996). *SADABS*. University of Göttingen, Germany.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.

supporting information

Acta Cryst. (2010). E66, o1293 [https://doi.org/10.1107/S1600536810016090]

2,2-Dichloro-1-[(2*R*,5*S*)-5-ethyl-2-methyl-2-propyl-1,3-oxazolidin-3-yl]ethanone

Ying Fu, Fei Ye and Xiaotian Wen

S1. Comment

N-dichloroacetyl oxazolidines are becoming increasingly important with their excellent biological activity (Agami & Couty, 2004; Abu-Qare & Duncan, 2002; Guirado *et al.*, 2003; Davies & Caseley, 1999). The discovery of N-dichloroacetyl oxazolidine as a herbicide safener has drawn widespread attention in agricultural biochemistry (Del Buono *et al.*, 2007; Hatzios & Burgos, 2004). As a part of our ongoing investigation of oxazolidine derivatives (Fu *et al.*, 2009) we prepared the title compound.

The molecular structure of the title compound is shown in Fig. 1. In the crystal structure, molecules are linked by weak intermolecular C—H \cdots O hydrogen bonds to form one-dimensional chains (Fig. 2).

S2. Experimental

The title compound was prepared according to the literature procedure (Fu *et al.*, 2009). The single crystal suitable for X-ray structural analysis was obtained by slow evaporation in petroleum ether and ethyl acetate at room temperature. The title enantiomer spontaneously resolved from a racemic mixture during the crystallization.

S3. Refinement

All H atoms were initially located in a different Fourier map. The C—H atoms were then constrained to an ideal geometry, with C—H = 0.96–0.98 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $1.5U_{\text{eq}}(\text{C})$ for methyl H atoms.

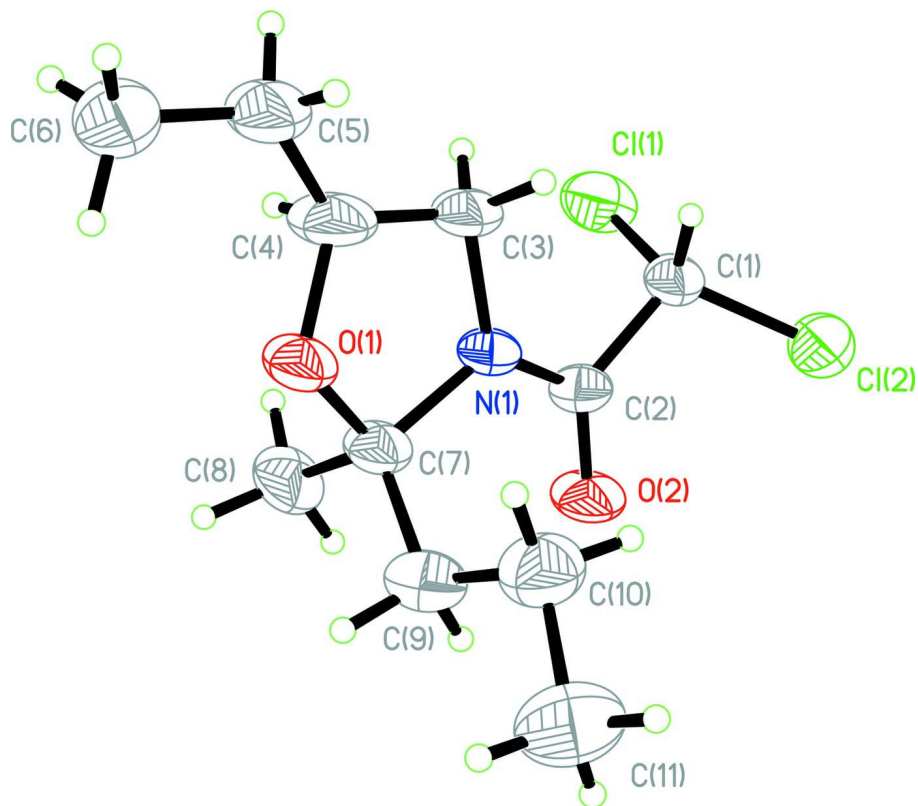


Figure 1

The molecular structure of the title compound, with the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level.

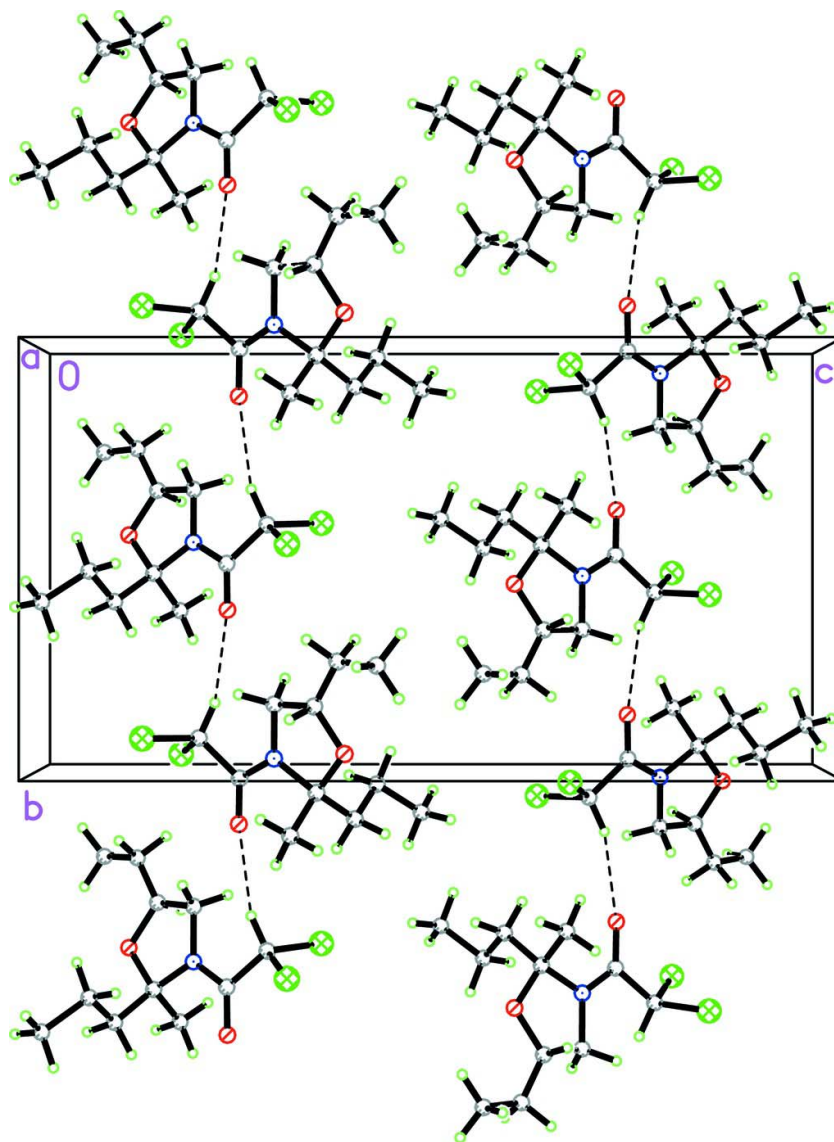


Figure 2

A packing diagram for (I), showing weak hydrogen bonds as dashed lines.

2,2-Dichloro-1-[(2R,5S)-5-ethyl-2-methyl-2-propyl-1,3-oxazolidin-3-yl]ethanone

Crystal data

$C_{11}H_{19}Cl_2NO_2$

$M_r = 268.17$

Orthorhombic, $P2_12_12_1$

Hall symbol: P 2ac 2ab

$a = 6.4834$ (12) Å

$b = 10.795$ (2) Å

$c = 20.030$ (4) Å

$V = 1401.8$ (5) Å³

$Z = 4$

$F(000) = 568.0$

$D_x = 1.271$ Mg m⁻³

$D_m = 1.271$ Mg m⁻³

D_m measured by not measured

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

Cell parameters from 3142 reflections

$\theta = 2.8$ – 20.6°

$\mu = 0.45$ mm⁻¹

$T = 293$ K

Block, colourless

$0.32 \times 0.24 \times 0.20$ mm

Data collection

Bruker SMART CCD diffractometer	14134 measured reflections
Radiation source: fine-focus sealed tube	3499 independent reflections
Graphite monochromator	2117 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\text{int}} = 0.058$
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	$\theta_{\text{max}} = 28.4^\circ$, $\theta_{\text{min}} = 2.8^\circ$
$T_{\text{min}} = 0.869$, $T_{\text{max}} = 0.915$	$h = -8 \rightarrow 8$
	$k = -14 \rightarrow 14$
	$l = -25 \rightarrow 26$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.084$	H-atom parameters constrained
$wR(F^2) = 0.258$	$w = 1/[\sigma^2(F_o^2) + (0.165P)^2]$
$S = 1.02$	where $P = (F_o^2 + 2F_c^2)/3$
3499 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
148 parameters	$\Delta\rho_{\text{max}} = 0.58 \text{ e } \text{\AA}^{-3}$
0 restraints	$\Delta\rho_{\text{min}} = -0.39 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	Absolute structure: Flack (1983) 1468 Friedels
	Absolute structure parameter: 0.02 (15)

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.5074 (7)	0.9192 (4)	0.2071 (2)	0.0675 (11)
H1	0.4966	0.8365	0.2270	0.081*
C2	0.5889 (7)	1.0103 (3)	0.2598 (2)	0.0638 (10)
C3	0.7886 (9)	0.8326 (4)	0.3065 (3)	0.0789 (13)
H3A	0.6784	0.7758	0.3181	0.095*
H3B	0.8454	0.8089	0.2636	0.095*
C4	0.9470 (13)	0.8334 (4)	0.3575 (3)	0.105 (2)
H4	1.0627	0.8594	0.3293	0.126*
C5	1.0407 (13)	0.7301 (5)	0.3866 (4)	0.122 (3)
H5A	1.0922	0.6801	0.3500	0.147*
H5B	0.9306	0.6828	0.4072	0.147*
C6	1.2052 (13)	0.7357 (6)	0.4353 (3)	0.111 (2)
H6A	1.3358	0.7362	0.4127	0.166*
H6B	1.1976	0.6647	0.4640	0.166*
H6C	1.1915	0.8098	0.4614	0.166*

C7	0.8170 (7)	1.0342 (4)	0.3589 (2)	0.0680 (10)
C8	0.9703 (10)	1.1251 (5)	0.3287 (3)	0.0925 (16)
H8A	1.0635	1.1533	0.3627	0.139*
H8B	0.8973	1.1946	0.3104	0.139*
H8C	1.0470	1.0847	0.2939	0.139*
C9	0.6750 (10)	1.1009 (5)	0.4051 (3)	0.0875 (14)
H9A	0.7550	1.1310	0.4427	0.105*
H9B	0.6205	1.1727	0.3819	0.105*
C10	0.4982 (11)	1.0277 (6)	0.4315 (3)	0.1013 (18)
H10A	0.5481	0.9494	0.4488	0.122*
H10B	0.4023	1.0103	0.3955	0.122*
C11	0.3855 (15)	1.0989 (8)	0.4874 (4)	0.127 (3)
H11A	0.4781	1.1119	0.5242	0.190*
H11B	0.2688	1.0516	0.5023	0.190*
H11C	0.3396	1.1775	0.4707	0.190*
Cl1	0.6885 (3)	0.91462 (14)	0.14073 (7)	0.0996 (5)
Cl2	0.2642 (2)	0.96549 (13)	0.17873 (9)	0.1008 (5)
N1	0.7143 (5)	0.9597 (3)	0.30507 (18)	0.0629 (8)
O1	0.9265 (6)	0.9417 (3)	0.39428 (17)	0.0793 (9)
O2	0.5407 (6)	1.1193 (3)	0.2580 (2)	0.0859 (10)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.072 (2)	0.0401 (18)	0.090 (3)	0.0031 (17)	-0.013 (2)	0.0002 (18)
C2	0.071 (2)	0.0396 (17)	0.081 (3)	0.0077 (17)	0.000 (2)	-0.0023 (17)
C3	0.093 (3)	0.0354 (18)	0.108 (3)	0.014 (2)	-0.021 (3)	-0.0090 (18)
C4	0.154 (5)	0.046 (2)	0.114 (4)	0.033 (3)	-0.043 (4)	-0.013 (3)
C5	0.149 (6)	0.059 (3)	0.160 (6)	0.031 (4)	-0.071 (5)	-0.020 (3)
C6	0.127 (5)	0.070 (3)	0.134 (5)	0.010 (4)	-0.031 (4)	0.013 (3)
C7	0.084 (3)	0.0379 (17)	0.082 (3)	0.0039 (18)	-0.010 (2)	-0.0029 (16)
C8	0.101 (4)	0.052 (2)	0.124 (4)	-0.020 (3)	-0.013 (3)	0.003 (3)
C9	0.109 (4)	0.060 (3)	0.093 (3)	0.006 (3)	0.003 (3)	-0.010 (2)
C10	0.105 (4)	0.075 (3)	0.124 (5)	0.010 (3)	0.006 (3)	0.000 (3)
C11	0.149 (6)	0.124 (6)	0.107 (4)	0.021 (5)	0.035 (4)	-0.005 (4)
Cl1	0.1169 (11)	0.0780 (8)	0.1038 (9)	-0.0063 (8)	0.0152 (9)	-0.0185 (7)
Cl2	0.0899 (8)	0.0761 (8)	0.1365 (12)	0.0154 (7)	-0.0350 (8)	-0.0093 (7)
N1	0.0715 (19)	0.0298 (13)	0.087 (2)	0.0036 (14)	-0.0060 (17)	-0.0042 (13)
O1	0.102 (2)	0.0413 (14)	0.095 (2)	0.0055 (15)	-0.0232 (19)	-0.0079 (14)
O2	0.114 (3)	0.0338 (13)	0.110 (2)	0.0152 (16)	-0.023 (2)	-0.0013 (14)

Geometric parameters (Å, °)

C1—C2	1.536 (6)	C6—H6C	0.9600
C1—Cl2	1.750 (4)	C7—O1	1.416 (5)
C1—Cl1	1.775 (5)	C7—C9	1.491 (7)
C1—H1	0.9800	C7—N1	1.501 (5)
C2—O2	1.218 (5)	C7—C8	1.523 (7)

C2—N1	1.335 (6)	C8—H8A	0.9600
C3—C4	1.448 (8)	C8—H8B	0.9600
C3—N1	1.454 (5)	C8—H8C	0.9600
C3—H3A	0.9700	C9—C10	1.490 (9)
C3—H3B	0.9700	C9—H9A	0.9700
C4—O1	1.388 (6)	C9—H9B	0.9700
C4—C5	1.397 (7)	C10—C11	1.542 (9)
C4—H4	0.9800	C10—H10A	0.9700
C5—C6	1.446 (10)	C10—H10B	0.9700
C5—H5A	0.9700	C11—H11A	0.9600
C5—H5B	0.9700	C11—H11B	0.9600
C6—H6A	0.9600	C11—H11C	0.9600
C6—H6B	0.9600		
C2—C1—C12	110.5 (3)	O1—C7—N1	101.8 (3)
C2—C1—C11	107.8 (3)	C9—C7—N1	115.5 (4)
C12—C1—C11	111.1 (3)	O1—C7—C8	109.0 (4)
C2—C1—H1	109.1	C9—C7—C8	109.8 (4)
C12—C1—H1	109.1	N1—C7—C8	110.5 (4)
C11—C1—H1	109.1	C7—C8—H8A	109.5
O2—C2—N1	124.9 (4)	C7—C8—H8B	109.5
O2—C2—C1	120.7 (4)	H8A—C8—H8B	109.5
N1—C2—C1	114.5 (3)	C7—C8—H8C	109.5
C4—C3—N1	104.1 (4)	H8A—C8—H8C	109.5
C4—C3—H3A	110.9	H8B—C8—H8C	109.5
N1—C3—H3A	110.9	C10—C9—C7	116.0 (5)
C4—C3—H3B	110.9	C10—C9—H9A	108.3
N1—C3—H3B	110.9	C7—C9—H9A	108.3
H3A—C3—H3B	108.9	C10—C9—H9B	108.3
O1—C4—C5	119.5 (5)	C7—C9—H9B	108.3
O1—C4—C3	108.1 (4)	H9A—C9—H9B	107.4
C5—C4—C3	126.8 (5)	C9—C10—C11	111.0 (6)
O1—C4—H4	97.9	C9—C10—H10A	109.4
C5—C4—H4	97.9	C11—C10—H10A	109.4
C3—C4—H4	97.9	C9—C10—H10B	109.4
C4—C5—C6	124.7 (6)	C11—C10—H10B	109.4
C4—C5—H5A	106.2	H10A—C10—H10B	108.0
C6—C5—H5A	106.2	C10—C11—H11A	109.5
C4—C5—H5B	106.2	C10—C11—H11B	109.5
C6—C5—H5B	106.2	H11A—C11—H11B	109.5
H5A—C5—H5B	106.3	C10—C11—H11C	109.5
C5—C6—H6A	109.5	H11A—C11—H11C	109.5
C5—C6—H6B	109.5	H11B—C11—H11C	109.5
H6A—C6—H6B	109.5	C2—N1—C3	127.0 (3)
C5—C6—H6C	109.5	C2—N1—C7	122.6 (3)
H6A—C6—H6C	109.5	C3—N1—C7	110.1 (3)
H6B—C6—H6C	109.5	C4—O1—C7	112.1 (3)
O1—C7—C9	109.9 (4)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H··· <i>A</i>	<i>D</i> —H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> —H··· <i>A</i>
C1—H1···O2 ⁱ	0.98	2.38	3.327 (5)	163

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