Acta Crystallographica Section E
Structure Reports Online
ISSN 1600-5368

(*E*)-(2,4-Dichlorobenzylidene)amino cyclopropanecarboxylate

Mei-Yi Wang* and Ya Zhang

College of Chemistry and Chemical Engineering, Beifang University of Nationalities, Yinchuan 750021, People's Republic of China
Correspondence e-mail: aseonliu@gmail.com

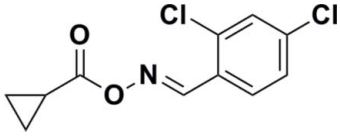
Received 10 April 2012; accepted 23 April 2012

Key indicators: single-crystal X-ray study; T = 113 K; mean σ(C–C) = 0.002 Å; R factor = 0.027; wR factor = 0.076; data-to-parameter ratio = 13.4.

In the title compound $C_{11}H_9Cl_2NO_2$, the dihedral angle between the benzene and cyclopropane ring planes is 89.95 (13)°. The carbonyl–oxime grouping is almost coplanar with the benzene ring [dihedral angle = 4.08 (6)°]. In the crystal, molecules are linked by C—H···O interactions into [100] chains.

Related literature

For further synthetic details, see: Liu *et al.* (2011*b*, 2012). For related structures, see: Liu & Liu (2011) Liu *et al.* (2011*d*). For the biological activity of related compounds, see: Liu *et al.* (2010, 2011*a*,*c*).



Experimental

Crystal data

$C_{11}H_9Cl_2NO_2$
M_r = 258.09
Triclinic, $P\overline{1}$
a = 6.4381 (13) Å
b = 7.6030 (15) Å
c = 11.956 (2) Å
α = 94.90 (3)°
β = 100.42 (3)°
γ = 102.70 (3)°
V = 556.71 (19) Å^3
Z = 2
Mo $K\alpha$ radiation
μ = 0.57 mm^{-1}
T = 113 K
0.24 × 0.20 × 0.10 mm

Data collection

Rigaku Saturn CCD diffractometer
Absorption correction: multi-scan (*CrystalClear*; Rigaku/MSC, 2005)
T_{min} = 0.876, T_{max} = 0.946
3737 measured reflections
1937 independent reflections
1563 reflections with $I > 2\sigma(I)$
R_{int} = 0.030

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.027
$wR(F^2)$ = 0.076
S = 1.03
1937 reflections
145 parameters
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.24 e Å^{-3}
$\Delta\rho_{min}$ = −0.21 e Å^{-3}

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

D—H···A	D—H	H···A	D···A	D—H···A
C1—H1B···O1^{i}	0.97	2.57	3.5008 (19)	161

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

Data collection: *CrystalClear* (Rigaku/MSC, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; 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: *SHELXL97*.

This research work was supported by the Research Program of Beifang University of Nationalities (2010Y047) and the Key Laboratory of Chemical Technology of the SEAC (2010SY16).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB6735).

References

Liu, X.-H., Chen, P.-Q., Wang, B.-L., Dong, W.-L., Li, Y.-H., Xie, X.-Q. & Li, Z.-M. (2010). *Chem. Biol. Drug Des.* **75**, 228–232.
Liu, X. H., Jian, Q. W. & Tan, C. X. (2011*a*). *Asian J. Chem.* **23**, 4064–4066.
Liu, X.-F. & Liu, X.-H. (2011). *Acta Cryst.* E**67**, o202.
Liu, X.-H., Pan, L., Tan, C.-X., Weng, J.-Q., Wang, B.-L. & Li, Z.-M. (2011*b*). *Pest. Biochem. Physiol.* **101**, 143–147.
Liu, X. H., Tan, C. X. & Jian, Q. W. (2011*c*). *Phosphorus Sulfur Silicon*, **186**, 552–557.
Liu, X.-H., Tan, C.-X., Weng, J.-Q. & Liu, H.-J. (2012). *Acta Cryst.* E**68**, o493.
Liu, H.-J., Weng, J.-Q., Tan, C.-X. & Liu, X.-H. (2011*d*). *Acta Cryst.* E**67**, o1940.
Rigaku/MSC (2005). *CrystalClear*. Rigaku/MSC Inc. The Woodlands, Texas, USA.
Sheldrick, G. M. (2008). *Acta Cryst.* A**64**, 112–122.

supporting information

Acta Cryst. (2012). E**68**, o1594 [doi:10.1107/S1600536812018016]

(*E*)-(2,4-Dichlorobenzylidene)amino cyclopropanecarboxylate

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S1. Experimental

Dropwised the cyclopropanecarbonyl chloride to 2,4-dichlorobenzaldehyde oxime (7.50 mmol in 25 ml THF) and 7.5 mmol Et3N, then vigorously stirred at ambient temperature for overnight. The corresponding product precipitated immediately. Compound was dissolved in hot alcohol and the resulting solution was allowed to stand in air at room temperature to give colourless blocks of the title compound.

S2. Refinement

All the H atoms were positioned geometrically (C—H = 0.93–0.97 Å) and refined as riding with U_{iso}(H) = 1.2U_{eq}(C) or 1.5U_{eq}(methyl C).

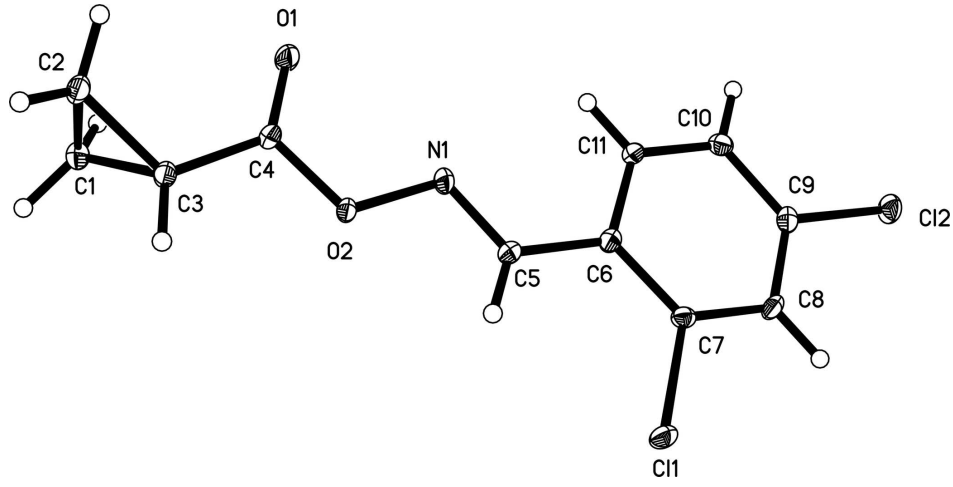

Figure 1

The molecular structure of (I). Displacement ellipsoids are drawn at the 30% probability level.

(*E*)-(2,4-Dichlorobenzylidene)amino cyclopropanecarboxylate

Crystal data

$C_{11}H_9Cl_2NO_2$
M_r = 258.09
Triclinic, $P\overline{1}$
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γ = 102.70 (3)°
V = 556.71 (19) Å^3
Z = 2
F(000) = 264
D_x = 1.540 Mg m^{-3}
Mo $K\alpha$ radiation, λ = 0.71073 Å
Cell parameters from 1887 reflections
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μ = 0.57 mm^{-1}
T = 113 K

Block, coloress
0.24 × 0.20 × 0.10 mm

Data collection

Rigaku Saturn CCD diffractometer
Radiation source: rotating anode
Confocal monochromator
ω scans
Absorption correction: multi-scan (*CrystalClear*; Rigaku/MSC, 2005)
T_{min} = 0.876, T_{max} = 0.946

3737 measured reflections
1937 independent reflections
1563 reflections with $I > 2\sigma(I)$
R_{int} = 0.030
θ_{max} = 25.0°, θ_{min} = 1.8°
h = −7→7
k = −9→6
l = −14→14

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)]$ = 0.027
$wR(F^2)$ = 0.076
S = 1.03
1937 reflections
145 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
$w = 1/[\sigma^2(F_o^2) + (0.0407P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max}$ = 0.001
$\Delta\rho_{max}$ = 0.24 e Å^{-3}
$\Delta\rho_{min}$ = −0.21 e Å^{-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 (Å^2)

	x	y	z	U_{iso}*/U_{eq}
Cl1	0.32359 (6)	0.70035 (5)	0.47089 (3)	0.02317 (14)
Cl2	1.15092 (6)	0.98054 (6)	0.68823 (4)	0.02531 (14)
O1	0.08882 (17)	0.30779 (16)	0.94813 (9)	0.0263 (3)
O2	0.00692 (16)	0.37138 (15)	0.76492 (9)	0.0186 (3)
N1	0.2306 (2)	0.46280 (18)	0.78101 (11)	0.0202 (3)
C1	−0.3868 (2)	0.2548 (2)	0.94150 (14)	0.0217 (4)
H1A	−0.2971	0.3293	1.0106	0.026*
H1B	−0.5348	0.2689	0.9234	0.026*
C2	−0.3504 (3)	0.0724 (2)	0.91336 (14)	0.0208 (4)
H2A	−0.4764	−0.0249	0.8783	0.025*
H2B	−0.2388	0.0354	0.9655	0.025*
C3	−0.2777 (2)	0.2198 (2)	0.84222 (14)	0.0198 (4)
H3	−0.3629	0.2113	0.7644	0.024*
C4	−0.0426 (2)	0.3004 (2)	0.86199 (13)	0.0177 (4)

C5	0.2637 (2)	0.5304 (2)	0.69054 (14)	0.0177 (4)
H5	0.1511	0.5129	0.6268	0.021*
C6	0.4823 (2)	0.6365 (2)	0.68810 (14)	0.0164 (4)
C7	0.5265 (3)	0.7218 (2)	0.59348 (13)	0.0168 (4)
C8	0.7305 (2)	0.8271 (2)	0.59131 (13)	0.0188 (4)
H8	0.7566	0.8836	0.5275	0.023*
C9	0.8947 (2)	0.8453 (2)	0.68750 (14)	0.0187 (4)
C10	0.8595 (3)	0.7612 (2)	0.78232 (14)	0.0196 (4)
H10	0.9725	0.7736	0.8453	0.024*
C11	0.6539 (2)	0.6579 (2)	0.78256 (14)	0.0177 (4)
H11	0.6293	0.6016	0.8466	0.021*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.0227 (2)	0.0314 (3)	0.0143 (2)	0.00420 (18)	0.00140 (17)	0.00786 (18)
Cl2	0.0192 (2)	0.0275 (2)	0.0273 (3)	−0.00120 (17)	0.00716 (17)	0.00581 (18)
O1	0.0182 (6)	0.0425 (8)	0.0166 (7)	0.0021 (5)	0.0021 (5)	0.0121 (6)
O2	0.0123 (5)	0.0267 (6)	0.0165 (6)	0.0012 (5)	0.0038 (4)	0.0082 (5)
N1	0.0117 (6)	0.0267 (8)	0.0213 (8)	0.0005 (6)	0.0039 (5)	0.0072 (6)
C1	0.0179 (8)	0.0266 (9)	0.0221 (10)	0.0051 (7)	0.0073 (7)	0.0061 (7)
C2	0.0191 (8)	0.0226 (9)	0.0198 (9)	0.0004 (7)	0.0059 (7)	0.0054 (7)
C3	0.0151 (8)	0.0264 (9)	0.0160 (9)	0.0002 (7)	0.0027 (7)	0.0059 (7)
C4	0.0187 (8)	0.0201 (8)	0.0162 (9)	0.0045 (7)	0.0070 (7)	0.0059 (7)
C5	0.0178 (8)	0.0218 (9)	0.0145 (9)	0.0048 (7)	0.0041 (7)	0.0056 (7)
C6	0.0184 (8)	0.0158 (8)	0.0162 (9)	0.0042 (6)	0.0061 (7)	0.0031 (6)
C7	0.0185 (8)	0.0193 (8)	0.0134 (9)	0.0070 (6)	0.0019 (6)	0.0024 (6)
C8	0.0244 (9)	0.0189 (9)	0.0165 (10)	0.0057 (7)	0.0100 (7)	0.0071 (7)
C9	0.0181 (8)	0.0170 (8)	0.0223 (9)	0.0038 (7)	0.0082 (7)	0.0024 (7)
C10	0.0189 (8)	0.0226 (9)	0.0183 (9)	0.0077 (7)	0.0022 (7)	0.0040 (7)
C11	0.0194 (8)	0.0206 (9)	0.0166 (9)	0.0079 (7)	0.0065 (7)	0.0073 (7)

Geometric parameters (Å, º)

Cl1—C7	1.7477 (16)	C3—C4	1.470 (2)
Cl2—C9	1.7394 (16)	C3—H3	0.9800
O1—C4	1.1976 (18)	C5—C6	1.467 (2)
O2—C4	1.3800 (19)	C5—H5	0.9300
O2—N1	1.4253 (16)	C6—C7	1.394 (2)
N1—C5	1.269 (2)	C6—C11	1.401 (2)
C1—C2	1.478 (2)	C7—C8	1.385 (2)
C1—C3	1.517 (2)	C8—C9	1.389 (2)
C1—H1A	0.9700	C8—H8	0.9300
C1—H1B	0.9700	C9—C10	1.378 (2)
C2—C3	1.508 (2)	C10—C11	1.383 (2)
C2—H2A	0.9700	C10—H10	0.9300
C2—H2B	0.9700	C11—H11	0.9300

C4—O2—N1	112.18 (11)	O2—C4—C3	109.31 (12)
C5—N1—O2	109.05 (12)	N1—C5—C6	118.94 (14)
C2—C1—C3	60.45 (11)	N1—C5—H5	120.5
C2—C1—H1A	117.7	C6—C5—H5	120.5
C3—C1—H1A	117.7	C7—C6—C11	117.60 (14)
C2—C1—H1B	117.7	C7—C6—C5	121.71 (14)
C3—C1—H1B	117.7	C11—C6—C5	120.69 (15)
H1A—C1—H1B	114.8	C8—C7—C6	122.37 (14)
C1—C2—C3	61.08 (11)	C8—C7—Cl1	116.74 (13)
C1—C2—H2A	117.7	C6—C7—Cl1	120.89 (12)
C3—C2—H2A	117.7	C7—C8—C9	117.79 (15)
C1—C2—H2B	117.7	C7—C8—H8	121.1
C3—C2—H2B	117.7	C9—C8—H8	121.1
H2A—C2—H2B	114.8	C10—C9—C8	121.89 (15)
C4—C3—C2	116.62 (14)	C10—C9—Cl2	119.36 (12)
C4—C3—C1	115.92 (13)	C8—C9—Cl2	118.75 (13)
C2—C3—C1	58.47 (11)	C9—C10—C11	119.14 (14)
C4—C3—H3	117.5	C9—C10—H10	120.4
C2—C3—H3	117.5	C11—C10—H10	120.4
C1—C3—H3	117.5	C10—C11—C6	121.20 (15)
O1—C4—O2	123.97 (14)	C10—C11—H11	119.4
O1—C4—C3	126.72 (15)	C6—C11—H11	119.4

C4—O2—N1—C5	−177.60 (13)	C5—C6—C7—C8	177.96 (16)
C1—C2—C3—C4	105.44 (16)	C11—C6—C7—Cl1	179.11 (13)
C2—C1—C3—C4	−106.64 (16)	C5—C6—C7—Cl1	−2.0 (2)
N1—O2—C4—O1	−3.1 (2)	C6—C7—C8—C9	0.4 (3)
N1—O2—C4—C3	176.04 (13)	Cl1—C7—C8—C9	−179.65 (12)
C2—C3—C4—O1	−25.6 (3)	C7—C8—C9—C10	0.6 (3)
C1—C3—C4—O1	40.4 (2)	C7—C8—C9—Cl2	−178.69 (13)
C2—C3—C4—O2	155.33 (15)	C8—C9—C10—C11	−1.1 (3)
C1—C3—C4—O2	−138.66 (14)	Cl2—C9—C10—C11	178.24 (13)
O2—N1—C5—C6	178.12 (13)	C9—C10—C11—C6	0.5 (3)
N1—C5—C6—C7	−176.72 (16)	C7—C6—C11—C10	0.5 (2)
N1—C5—C6—C11	2.2 (3)	C5—C6—C11—C10	−178.44 (16)
C11—C6—C7—C8	−1.0 (3)		

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
C1—H1*B*···O1^{i}	0.97	2.57	3.5008 (19)	161

Symmetry code: (i) *x*−1, *y*, *z*.