Acta Crystallographica Section E
Structure Reports Online
ISSN 1600-5368

N-Benzyl-2-(2,6-dichlorophenoxy)-acetamide

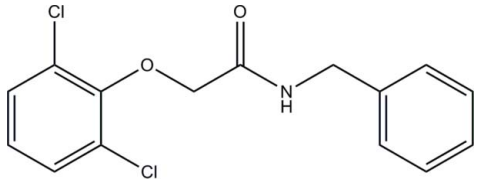
Zhu-Bo Li,[a]* Yong-Huang Luo,[a] Wen-Liang Dong,[b] Jing Li[a] and Hua Zuo[a]

[a]College of Pharmaceutical Sciences, Southwest University, Chongqing 400716, People's Republic of China, and [b]Shandong University of Traditional Chinese Medicine, Jinan 250355, People's Republic of China
Correspondence e-mail: lizhubo2007@163.com

Received 8 July 2008; accepted 18 July 2008

Key indicators: single-crystal X-ray study; T = 298 K; mean σ(C–C) = 0.003 Å; R factor = 0.045; wR factor = 0.132; data-to-parameter ratio = 18.9.

The structure determination of the title compound, $C_{15}H_{13}Cl_2NO_2$, was undertaken as part of a project on the interaction of small molecules with proteins. In the crystal structure, the dihedral angle between the two aryl rings is 40.71 (11)°. The molecules are connected *via* N—H···O hydrogen bonding into chains, which extend in the direction of the *b* axis.



Experimental

Crystal data

$C_{15}H_{13}Cl_2NO_2$
M_r = 310.16
Orthorhombic, *Pbca*
a = 14.8886 (10) Å
b = 8.6579 (6) Å
c = 22.9867 (14) Å
V = 2963.1 (3) Å^3
Z = 8
Mo $K\alpha$ radiation
μ = 0.44 mm^{-1}
T = 298 (2) K
0.20 × 0.20 × 0.10 mm

Data collection

Bruker APEXII CCD area-detector diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2005)
T_{min} = 0.918, T_{max} = 0.958
16445 measured reflections
3412 independent reflections
2103 reflections with $I > 2\sigma(I)$
R_{int} = 0.040

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.044
$wR(F^2)$ = 0.131
S = 1.02
3412 reflections
181 parameters
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.20 e Å^{-3}
$\Delta\rho_{min}$ = −0.34 e Å^{-3}

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

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1A···O1	0.86	2.23	2.644 (2)	109
N1—H1A···O2^{i}	0.86	2.31	2.970 (2)	133

Symmetry code: (i) $-x+\frac{3}{2}, y+\frac{1}{2}, z$.

Data collection: *APEX2* (Bruker, 2005); cell refinement: *APEX2*; data reduction: *APEX2*; program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

This study was supported by the Key Program Projects of the Municipal Natural Science Foundation of Chongqing, China (grant No. CSTC, 2007AC1042).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: NC2111).

References

Altomare, A., Burla, M. C., Camalli, M., Cascarano, G. L., Giacovazzo, C., Guagliardi, A., Moliterni, A. G. G., Polidori, G. & Spagna, R. (1999). *J. Appl. Cryst.* **32**, 115–119.
Bruker (2005). *SADABS* and *APEX2*. Bruker AXS Inc., Madison, Wisconsin, USA.
Farrugia, L. J. (1999). *J. Appl. Cryst.* **32**, 837–838.
Sheldrick, G. M. (2008). *Acta Cryst.* A**64**, 112–122.

supporting information

Acta Cryst. (2008). E**64**, o1610 [doi:10.1107/S1600536808022514]

N-Benzyl-2-(2,6-dichlorophenoxy)acetamide

Zhu-Bo Li, Yong-Huang Luo, Wen-Liang Dong, Jing Li and Hua Zuo

S1. Experimental

A solution of 2,6-dichlorophenol (1.0 mmol), *N*-benzyl-2-chloroacetamide (1.1 mmol), K_2CO_3 (1.1 mmol) in CH_3CN (20 ml) was refluxed for 3 h and afterwards cooled down to room temperature. The solvent was removed under reduced pressure and the residue was poured into water and adjusted to pH 6–7. with dilute hydrochloric acid (10%) and extracted with ethyl acetate, washed with brine and dried over anhydrous $MgSO_4$ to obtain the corresponding crude product. The product was obtained by column chromatography on silica gel using ethyl acetate as eluent. (yield 90%). Crystals suitable for X-ray diffraction were obtained by slow cooling of a solution of the solid in ethyl acetate/hexane at room temperature for 4 d.

S2. Refinement

All H atoms were placed in geometrically calculated positions and refined using a riding model with C—H = 0.97 Å (for CH_2 groups) and 0.96 Å (for CH_3 groups), their isotropic displacement parameters were set to 1.2 times (1.5 times for CH_3 groups) the equivalent displacement parameter of their parent atoms.

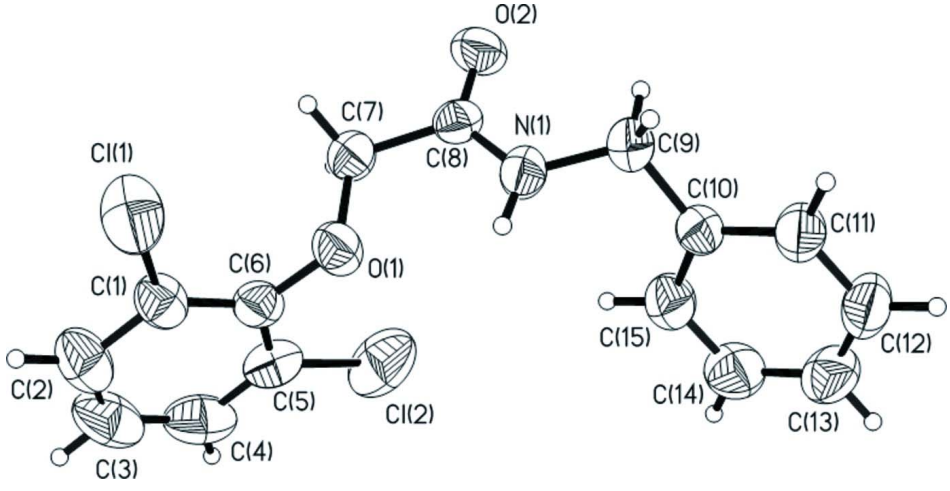


Figure 1

The molecular structure of the title compound with labelling and displacement ellipsoids drawn at the 50% probability level.

N-Benzyl-2-(2,6-dichlorophenoxy)acetamide

Crystal data

$C_{15}H_{13}Cl_2NO_2$
$M_r = 310.16$
Orthorhombic, *Pbca*
$a = 14.8886$ (10) Å
$b = 8.6579$ (6) Å
$c = 22.9867$ (14) Å
$V = 2963.1$ (3) Å^3
$Z = 8$
$F(000) = 1280$
$D_x = 1.391$ Mg m^{-3}
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 2733 reflections
$\theta = 2.2$–$21.8°$
$\mu = 0.44$ mm^{-1}
$T = 298$ K
Block, colorless
0.20 × 0.20 × 0.10 mm

Data collection

Bruker SMART CCD area-detector diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
φ and ω scans
Absorption correction: multi-scan (*SADABS*; Bruker, 2005)
$T_{min} = 0.918$, $T_{max} = 0.958$
16445 measured reflections
3412 independent reflections
2103 reflections with $I > 2\sigma(I)$
$R_{int} = 0.040$
$\theta_{max} = 27.5°$, $\theta_{min} = 1.8°$
$h = -19 \rightarrow 18$
$k = -10 \rightarrow 11$
$l = -20 \rightarrow 29$

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)] = 0.045$
$wR(F^2) = 0.131$
$S = 1.02$
3412 reflections
181 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
$w = 1/[\sigma^2(F_o^2) + (0.0631P)^2 + 0.3666P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} = 0.001$
$\Delta\rho_{max} = 0.20$ e Å^{-3}
$\Delta\rho_{min} = -0.34$ 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}
N1	0.74097 (11)	0.0194 (2)	0.64026 (7)	0.0521 (4)
H1A	0.7004	0.0734	0.6230	0.063*
Cl1	0.74730 (5)	0.13003 (7)	0.41952 (3)	0.0787 (2)
Cl2	0.55214 (5)	−0.24316 (9)	0.55609 (4)	0.0985 (3)
O1	0.68835 (9)	−0.00499 (15)	0.53078 (6)	0.0526 (4)

O2	0.84330 (10)	−0.17088 (18)	0.62980 (6)	0.0591 (4)
C1	0.65927 (14)	0.0023 (2)	0.42821 (9)	0.0515 (5)
C2	0.60907 (17)	−0.0417 (3)	0.38036 (11)	0.0679 (7)
H2A	0.6217	−0.0014	0.3438	0.081*
C3	0.54023 (18)	−0.1457 (3)	0.38759 (14)	0.0802 (8)
H3A	0.5057	−0.1746	0.3557	0.096*
C4	0.52178 (17)	−0.2071 (3)	0.44101 (13)	0.0761 (8)
H4A	0.4755	−0.2783	0.4454	0.091*
C5	0.57263 (15)	−0.1625 (3)	0.48853 (11)	0.0603 (6)
C6	0.64111 (13)	−0.0555 (2)	0.48288 (9)	0.0466 (5)
C7	0.76294 (14)	−0.1008 (3)	0.54620 (9)	0.0522 (5)
H7A	0.8148	−0.0717	0.5231	0.063*
H7B	0.7487	−0.2076	0.5375	0.063*
C8	0.78542 (13)	−0.0856 (2)	0.60971 (9)	0.0433 (5)
C9	0.75781 (14)	0.0471 (3)	0.70161 (9)	0.0597 (6)
H9A	0.7848	0.1484	0.7060	0.072*
H9B	0.8007	−0.0286	0.7155	0.072*
C10	0.67475 (13)	0.0393 (2)	0.73885 (8)	0.0471 (5)
C11	0.67143 (16)	0.1244 (3)	0.78942 (9)	0.0586 (6)
H11A	0.7190	0.1894	0.7988	0.070*
C12	0.59859 (17)	0.1148 (3)	0.82645 (10)	0.0696 (7)
H12A	0.5979	0.1718	0.8607	0.084*
C13	0.52742 (17)	0.0216 (3)	0.81283 (11)	0.0693 (7)
H13A	0.4780	0.0165	0.8375	0.083*
C14	0.52928 (16)	−0.0638 (3)	0.76294 (11)	0.0697 (7)
H14A	0.4814	−0.1283	0.7538	0.084*
C15	0.60248 (15)	−0.0545 (3)	0.72586 (10)	0.0609 (6)
H15A	0.6029	−0.1123	0.6918	0.073*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0540 (10)	0.0620 (11)	0.0404 (9)	0.0117 (8)	0.0022 (8)	−0.0039 (8)
Cl1	0.0982 (5)	0.0733 (4)	0.0644 (4)	−0.0303 (4)	−0.0038 (3)	0.0045 (3)
Cl2	0.0904 (5)	0.1033 (6)	0.1019 (6)	−0.0081 (4)	0.0394 (4)	0.0184 (4)
O1	0.0594 (9)	0.0521 (8)	0.0463 (8)	0.0140 (7)	−0.0084 (7)	−0.0094 (6)
O2	0.0484 (8)	0.0657 (9)	0.0633 (10)	0.0121 (7)	−0.0112 (7)	−0.0085 (7)
C1	0.0582 (12)	0.0460 (12)	0.0503 (12)	0.0008 (10)	−0.0087 (10)	−0.0080 (9)
C2	0.0867 (18)	0.0606 (14)	0.0565 (14)	0.0067 (13)	−0.0203 (12)	−0.0093 (11)
C3	0.0734 (18)	0.0740 (18)	0.093 (2)	0.0044 (14)	−0.0355 (16)	−0.0234 (15)
C4	0.0485 (13)	0.0675 (17)	0.112 (2)	−0.0034 (12)	−0.0089 (14)	−0.0132 (15)
C5	0.0466 (12)	0.0599 (13)	0.0745 (16)	0.0051 (11)	0.0084 (11)	−0.0035 (11)
C6	0.0443 (11)	0.0458 (11)	0.0497 (12)	0.0084 (9)	−0.0027 (9)	−0.0087 (9)
C7	0.0507 (12)	0.0603 (13)	0.0456 (12)	0.0127 (10)	0.0006 (9)	−0.0098 (9)
C8	0.0362 (10)	0.0448 (11)	0.0488 (12)	−0.0023 (9)	0.0028 (9)	−0.0019 (9)
C9	0.0522 (12)	0.0828 (17)	0.0441 (12)	−0.0021 (11)	0.0004 (10)	−0.0138 (11)
C10	0.0481 (11)	0.0523 (12)	0.0409 (11)	0.0042 (9)	−0.0021 (9)	0.0014 (9)
C11	0.0599 (13)	0.0665 (15)	0.0495 (12)	−0.0068 (11)	0.0043 (11)	−0.0085 (10)

C12	0.0749 (16)	0.0836 (17)	0.0503 (13)	−0.0039 (14)	0.0138 (12)	−0.0113 (12)
C13	0.0610 (14)	0.0832 (18)	0.0636 (16)	−0.0015 (13)	0.0164 (12)	0.0084 (13)
C14	0.0580 (14)	0.0770 (16)	0.0740 (17)	−0.0139 (13)	0.0017 (12)	0.0038 (13)
C15	0.0621 (14)	0.0654 (14)	0.0552 (14)	−0.0045 (12)	−0.0020 (11)	−0.0112 (11)

Geometric parameters (Å, º)

N1—C8	1.326 (2)	C7—C8	1.504 (3)
N1—C9	1.452 (3)	C7—H7A	0.9700
N1—H1A	0.8600	C7—H7B	0.9700
Cl1—C1	1.726 (2)	C9—C10	1.506 (3)
Cl2—C5	1.730 (3)	C9—H9A	0.9700
O1—C6	1.378 (2)	C9—H9B	0.9700
O1—C7	1.431 (2)	C10—C11	1.378 (3)
O2—C8	1.225 (2)	C10—C15	1.381 (3)
C1—C6	1.379 (3)	C11—C12	1.381 (3)
C1—C2	1.383 (3)	C11—H11A	0.9300
C2—C3	1.374 (4)	C12—C13	1.368 (3)
C2—H2A	0.9300	C12—H12A	0.9300
C3—C4	1.366 (4)	C13—C14	1.365 (3)
C3—H3A	0.9300	C13—H13A	0.9300
C4—C5	1.384 (3)	C14—C15	1.386 (3)
C4—H4A	0.9300	C14—H14A	0.9300
C5—C6	1.384 (3)	C15—H15A	0.9300
C8—N1—C9	122.78 (18)	O2—C8—N1	124.39 (19)
C8—N1—H1A	118.6	O2—C8—C7	118.03 (18)
C9—N1—H1A	118.6	N1—C8—C7	117.58 (17)
C6—O1—C7	114.23 (14)	N1—C9—C10	113.75 (17)
C6—C1—C2	121.2 (2)	N1—C9—H9A	108.8
C6—C1—Cl1	119.12 (15)	C10—C9—H9A	108.8
C2—C1—Cl1	119.65 (18)	N1—C9—H9B	108.8
C3—C2—C1	119.2 (2)	C10—C9—H9B	108.8
C3—C2—H2A	120.4	H9A—C9—H9B	107.7
C1—C2—H2A	120.4	C11—C10—C15	118.0 (2)
C4—C3—C2	120.9 (2)	C11—C10—C9	119.02 (19)
C4—C3—H3A	119.5	C15—C10—C9	122.96 (19)
C2—C3—H3A	119.5	C10—C11—C12	121.1 (2)
C3—C4—C5	119.4 (2)	C10—C11—H11A	119.5
C3—C4—H4A	120.3	C12—C11—H11A	119.5
C5—C4—H4A	120.3	C13—C12—C11	120.2 (2)
C6—C5—C4	121.0 (2)	C13—C12—H12A	119.9
C6—C5—Cl2	118.98 (18)	C11—C12—H12A	119.9
C4—C5—Cl2	120.0 (2)	C14—C13—C12	119.7 (2)
O1—C6—C1	120.84 (19)	C14—C13—H13A	120.1
O1—C6—C5	120.90 (19)	C12—C13—H13A	120.1
C1—C6—C5	118.23 (19)	C13—C14—C15	120.1 (2)
O1—C7—C8	111.28 (15)	C13—C14—H14A	120.0

O1—C7—H7A	109.4	C15—C14—H14A	120.0
C8—C7—H7A	109.4	C10—C15—C14	120.9 (2)
O1—C7—H7B	109.4	C10—C15—H15A	119.5
C8—C7—H7B	109.4	C14—C15—H15A	119.5
H7A—C7—H7B	108.0		

C6—C1—C2—C3	−0.4 (3)	C6—O1—C7—C8	−154.72 (17)
Cl1—C1—C2—C3	178.76 (18)	C9—N1—C8—O2	1.3 (3)
C1—C2—C3—C4	−0.9 (4)	C9—N1—C8—C7	−178.76 (19)
C2—C3—C4—C5	0.7 (4)	O1—C7—C8—O2	174.13 (18)
C3—C4—C5—C6	0.7 (4)	O1—C7—C8—N1	−5.8 (3)
C3—C4—C5—Cl2	−178.2 (2)	C8—N1—C9—C10	−126.9 (2)
C7—O1—C6—C1	−95.4 (2)	N1—C9—C10—C11	−151.9 (2)
C7—O1—C6—C5	86.3 (2)	N1—C9—C10—C15	30.6 (3)
C2—C1—C6—O1	−176.57 (18)	C15—C10—C11—C12	0.9 (3)
Cl1—C1—C6—O1	4.3 (3)	C9—C10—C11—C12	−176.7 (2)
C2—C1—C6—C5	1.7 (3)	C10—C11—C12—C13	−1.1 (4)
Cl1—C1—C6—C5	−177.41 (15)	C11—C12—C13—C14	1.1 (4)
C4—C5—C6—O1	176.40 (19)	C12—C13—C14—C15	−0.9 (4)
Cl2—C5—C6—O1	−4.6 (3)	C11—C10—C15—C14	−0.7 (3)
C4—C5—C6—C1	−1.9 (3)	C9—C10—C15—C14	176.8 (2)
Cl2—C5—C6—C1	177.05 (16)	C13—C14—C15—C10	0.7 (4)

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
N1—H1*A*···O1	0.86	2.23	2.644 (2)	109
N1—H1*A*···O2^{i}	0.86	2.31	2.970 (2)	133

Symmetry code: (i) −*x*+3/2, *y*+1/2, *z*.