Acta Crystallographica Section E
Structure Reports Online
ISSN 1600-5368

N-(2-Benzoyl-4-chlorophenyl)-4-chlorobenzenesulfonamide

Arto Valkonen,[a]* Ryszard Gawinecki,[b] Henryk Janota,[b] Borys Ośmiałowski[b] and Erkki Kolehmainen[a]

[a]Department of Chemistry, University of Jyväskylä, PO Box 35, FIN-40014 Jyväskylä, Finland, and [b]Department of Chemistry, University of Technology and Life Sciences, Seminaryjna 3, PL-85-326 Bydgoszcz, Poland
Correspondence e-mail: arto.valkonen@jyu.fi

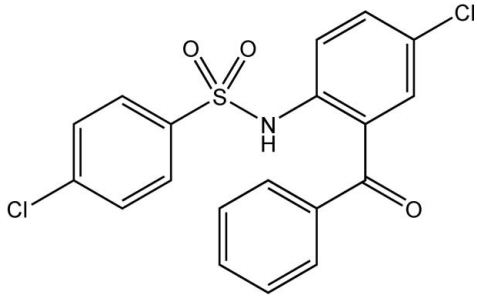
Received 28 February 2008; accepted 18 March 2008

Key indicators: single-crystal X-ray study; T = 173 K; mean σ(C–C) = 0.003 Å; R factor = 0.043; wR factor = 0.085; data-to-parameter ratio = 18.5.

The title compound, $C_{19}H_{13}Cl_2NO_3S$, is an *N*-arylsulfonyl derivative of 2-amino-5-chlorobenzophenone. The compound is biologically active and shows potential to be utilized as an inhibitor of CCR2 and CCR9 receptor functions. In the crystal structure, there is an intramolecular N—H···O hydrogen bond between the amide and carbonyl groups. The benzoyl and 4-chlorophenyl groups form intramolecular and intermolecular face-to-face contacts, with a dihedral angle of 10.6 (1)° between their mean planes in both cases, and centroid–centroid separations of 4.00 (1) and 4.25 (1) Å for the intra- and intermolecular interactions, respectively.

Related literature

For related literature, see: Basak *et al.* (2008); Fleming *et al.* (2003); Kolehmainen *et al.* (2003); Sternbach *et al.* (1962).



Experimental

Crystal data

$C_{19}H_{13}Cl_2NO_3S$
M_r = 406.26
Monoclinic, $P2_1/n$
a = 8.2307 (1) Å
b = 18.5014 (3) Å
c = 12.1364 (2) Å
β = 105.211 (1)°
V = 1783.38 (5) Å^3
Z = 4
Mo $K\alpha$ radiation
μ = 0.50 mm^{-1}
T = 173 (2) K
0.25 × 0.25 × 0.15 mm

Data collection

Bruker Kappa APEXII CCD diffractometer
Absorption correction: none
13987 measured reflections
4401 independent reflections
3534 reflections with $I > 2\sigma(I)$
R_{int} = 0.044

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.042
$wR(F^2)$ = 0.084
S = 1.05
4401 reflections
238 parameters
1 restraint
H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{max}$ = 0.32 e Å^{-3}
$\Delta\rho_{min}$ = −0.31 e Å^{-3}

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

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1···O1	0.865 (15)	2.20 (2)	2.798 (2)	126.1 (18)

Data collection: *COLLECT* (Bruker, 2004); cell refinement: *DENZO-SMN* (Otwinowski & Minor, 1997); data reduction: *DENZO-SMN*; program(s) used to solve structure: *SIR2002* (Burla *et al.*, 2003); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*, *PLATON* (Spek, 2003) and *Mercury* (Macrae *et al.*, 2006).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BI2286).

References

Basak, A., Jin, J., Moore, J., Pennell, A. M. K., Punna, S., Ungashe, S. & Wei, Z. (2008). World Patent Application 2008008374.
Bruker (2004). *COLLECT*. Bruker AXS Inc., Madison, Wisconsin, USA.
Burla, M. C., Camalli, M., Carrozzini, B., Cascarano, G. L., Giacovazzo, C., Polidori, G. & Spagna, R. (2003). *J. Appl. Cryst.* **36**, 1103.
Farrugia, L. J. (1997). *J. Appl. Cryst.* **30**, 565.
Fleming, P., Harriman, G. C. B., Shi, Z. & Chen, S. (2003). World Patent Application 2003099773.
Kolehmainen, E., Nissinen, M., Janota, H., Gawinecki, R. & Ośmiałowski, B. (2003). *Pol. J. Chem.* **77**, 889–894.
Macrae, C. F., Edgington, P. R., McCabe, P., Pidcock, E., Shields, G. P., Taylor, R., Towler, M. & van de Streek, J. (2006). *J. Appl. Cryst.* **39**, 453–457.
Otwinowski, Z. & Minor, W. (1997). *Methods in Enzymology*, Vol. 276, *Macromolecular Crystallography*, Part A, edited by C. W. Carter Jr & R. M. Sweet, pp. 307–326. New York: Academic Press.
Sheldrick, G. M. (2008). *Acta Cryst.* A**64**, 112–122.
Spek, A. L. (2003). *J. Appl. Cryst.* **36**, 7–13.
Sternbach, L. H., Fryer, R. I., Metlesics, W., Sach, G. & Stempel, A. (1962). *J. Org. Chem.* **27**, 3781–3788.

supplementary materials

Acta Cryst. (2008). E**64**, o737 [**doi:10.1107/S1600536808007435**]

N-(2-Benzoyl-4-chlorophenyl)-4-chlorobenzenesulfonamide

A. Valkonen, R. Gawinecki, H. Janota, B. Osmialowski and E. Kolehmainen

Comment

The title compound was originally prepared to study its molecular structure by spectroscopy (Kolehmainen *et al.*, 2003). The compound has a sulfone group showing strong electron acceptor capability and two S=O double bonds with ineffective conjugation properties with other double bonds. The compound has also shown a potential to be utilized as inhibitor of CCR2 (Basak *et al.*, 2008) and CCR9 (Fleming *et al.*, 2003) receptor functions. This potent antagonist can possibly be utilized in pharmaceutical compositions for treatment of CCR2 and CCR9 related diseases.

In the crystal, there is an intramolecular N—H···O hydrogen bond between the amide and carbonyl groups (Fig. 1). The benzoyl and 4-chlorophenyl groups form intramolecular (Fig. 2) and intermolecular face-to-face contacts (Fig. 3), with a dihedral angle of 10.6 (1)° between their mean planes in both cases, and a centroid-centroid separation of 4.00 (1) and 4.25 (1)Å for the intra- and intermolecular interactions, respectively.

Experimental

The title compound was obtained by condensation of 2-amino-5-chlorobenzophenone and 4-chlorobenzenesulfonyl chloride according to a previously described method (Sternbach *et al.*, 1962). The reaction product was purified by crystallization from ethanol. The spectroscopic characterization (NMR, IR) has previously been reported by us (Kolehmainen *et al.*, 2003). The single-crystal suitable for X-ray determination was obtained by extremely slow evaporation of a $CDCl_3$ solution in a NMR tube.

Refinement

All H atoms were visible in electron density maps, but those bound to C were placed in idealized positions and allowed to ride on their parent atoms at C—H distances of 0.95 Å with $U_{iso}(H) = 1.2U_{eq}(C)$. The position of the N—H proton was refined with the N—H distance restrained to 0.91 (2) Å and with $U_{iso}(H) = 1.2U_{eq}(N)$.

Figures

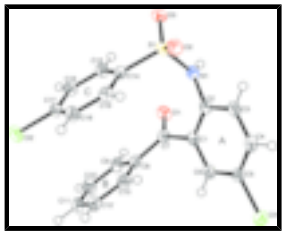


Fig. 1. The molecular structure of the title compound with displacement ellipsoids drawn at the 50% probability level. H atoms are represented by circles of arbitrary size

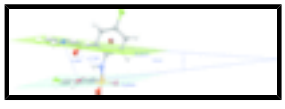


Fig. 2. Molecular conformation with selected geometric parameters

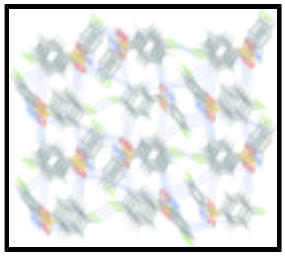


Fig. 3. Packing viewed along the *a*-axis showing the stack-like architecture

N-(2-Benzoyl-4-chlorophenyl)-4-chlorobenzenesulfonamide

Crystal data

$C_{19}H_{13}Cl_2NO_3S$	F_{000} = 832
M_r = 406.26	D_x = 1.513 Mg m^{-3}
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation λ = 0.71073 Å
Hall symbol: -P 2yn	Cell parameters from 31954 reflections
a = 8.2307 (1) Å	θ = 0.4–28.3º
b = 18.5014 (3) Å	μ = 0.50 mm^{-1}
c = 12.1364 (2) Å	T = 173 (2) K
β = 105.211 (1)º	Block, yellow
V = 1783.38 (5) Å^3	0.25 × 0.25 × 0.15 mm
Z = 4	

Data collection

Bruker Kappa APEXII CCD diffractometer	3534 reflections with $I > 2\sigma(I)$
Radiation source: fine-focus sealed tube	R_{int} = 0.044
Monochromator: graphite	θ_{max} = 28.3º
T = 173(2) K	θ_{min} = 2.1º
φ and ω scans	h = −10→10
Absorption correction: none	k = −24→24
13987 measured reflections	l = −14→16
4401 independent reflections	

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)]$ = 0.042	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2)$ = 0.084	$w = 1/[\sigma^2(F_o^2) + (0.0187P)^2 + 1.5712P]$ where $P = (F_o^2 + 2F_c^2)/3$
S = 1.05	$(\Delta/\sigma)_{max} < 0.001$
4401 reflections	$\Delta\rho_{max}$ = 0.32 e Å^{-3}

238 parameters	$\Delta\rho_{min} = -0.31$ e Å^{-3}
1 restraint	Extinction correction: none
Primary atom site location: structure-invariant direct methods	

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	0.44574 (7)	0.01345 (3)	0.80411 (4)	0.03185 (13)
S1	0.95358 (6)	0.13899 (3)	0.46317 (4)	0.02381 (11)
Cl2	0.64457 (7)	0.44781 (3)	0.45821 (5)	0.03514 (13)
O1	0.47941 (18)	0.13671 (8)	0.33815 (11)	0.0301 (3)
O3	1.00797 (18)	0.13033 (8)	0.36105 (13)	0.0331 (3)
O2	1.06708 (17)	0.12706 (8)	0.57308 (12)	0.0310 (3)
N1	0.7971 (2)	0.08200 (9)	0.44914 (14)	0.0239 (3)
H1	0.729 (2)	0.0856 (12)	0.3815 (14)	0.029*
C4	0.7144 (3)	0.01218 (10)	0.71469 (17)	0.0270 (4)
H4	0.7690	−0.0164	0.7787	0.032*
C15	0.7816 (2)	0.25730 (11)	0.35920 (17)	0.0261 (4)
H15	0.7689	0.2312	0.2900	0.031*
C7	0.4749 (2)	0.15092 (10)	0.43573 (16)	0.0220 (4)
C6	0.4742 (2)	0.08317 (10)	0.61488 (16)	0.0231 (4)
H6	0.3652	0.1018	0.6107	0.028*
C1	0.5564 (2)	0.10128 (10)	0.53194 (16)	0.0215 (4)
C14	0.8692 (2)	0.22691 (10)	0.46257 (16)	0.0227 (4)
C10	0.2063 (3)	0.31757 (12)	0.3887 (2)	0.0358 (5)
H10	0.1250	0.3397	0.3278	0.043*
C17	0.7328 (2)	0.36241 (10)	0.45946 (17)	0.0246 (4)
C18	0.8210 (3)	0.33297 (11)	0.56290 (17)	0.0279 (4)
H18	0.8345	0.3594	0.6318	0.033*
C16	0.7129 (3)	0.32578 (11)	0.35766 (17)	0.0268 (4)
H16	0.6530	0.3472	0.2876	0.032*
C2	0.7158 (2)	0.07170 (10)	0.53854 (16)	0.0220 (4)
C8	0.3942 (2)	0.21889 (10)	0.46066 (16)	0.0222 (4)
C19	0.8891 (2)	0.26433 (11)	0.56418 (17)	0.0261 (4)
H19	0.9491	0.2431	0.6343	0.031*
C9	0.2776 (3)	0.25261 (11)	0.37109 (17)	0.0280 (4)

H9	0.2473	0.2307	0.2977	0.034*
C3	0.7952 (2)	0.02892 (10)	0.63138 (17)	0.0262 (4)
H3	0.9055	0.0111	0.6376	0.031*
C5	0.5527 (2)	0.03774 (10)	0.70344 (16)	0.0232 (4)
C13	0.4391 (3)	0.25198 (11)	0.56727 (17)	0.0266 (4)
H13	0.5174	0.2291	0.6290	0.032*
C11	0.2525 (3)	0.35055 (11)	0.4944 (2)	0.0362 (5)
H11	0.2036	0.3955	0.5058	0.043*
C12	0.3699 (3)	0.31836 (11)	0.58384 (19)	0.0334 (5)
H12	0.4030	0.3415	0.6562	0.040*

Atomic displacement parameters (Å^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.0399 (3)	0.0286 (3)	0.0302 (3)	−0.0016 (2)	0.0148 (2)	0.0021 (2)
S1	0.0222 (2)	0.0227 (2)	0.0275 (3)	0.00388 (18)	0.00828 (19)	0.00123 (19)
Cl2	0.0431 (3)	0.0224 (2)	0.0419 (3)	0.0072 (2)	0.0147 (2)	−0.0016 (2)
O1	0.0341 (8)	0.0324 (8)	0.0215 (7)	0.0044 (6)	0.0029 (6)	−0.0029 (6)
O3	0.0374 (8)	0.0304 (8)	0.0375 (8)	0.0066 (6)	0.0202 (7)	0.0006 (6)
O2	0.0221 (7)	0.0336 (8)	0.0349 (8)	0.0041 (6)	0.0031 (6)	0.0044 (6)
N1	0.0255 (8)	0.0219 (8)	0.0239 (9)	−0.0003 (7)	0.0058 (7)	−0.0030 (7)
C4	0.0304 (10)	0.0195 (9)	0.0283 (10)	0.0012 (8)	0.0029 (8)	0.0045 (8)
C15	0.0321 (11)	0.0250 (10)	0.0210 (10)	0.0016 (8)	0.0064 (8)	−0.0016 (8)
C7	0.0191 (9)	0.0223 (9)	0.0229 (10)	−0.0021 (7)	0.0024 (7)	−0.0010 (7)
C6	0.0217 (9)	0.0202 (9)	0.0267 (10)	0.0005 (7)	0.0050 (8)	−0.0016 (8)
C1	0.0227 (9)	0.0169 (9)	0.0223 (9)	−0.0004 (7)	0.0013 (7)	−0.0017 (7)
C14	0.0226 (9)	0.0223 (9)	0.0243 (10)	0.0010 (7)	0.0079 (8)	0.0013 (7)
C10	0.0364 (12)	0.0279 (11)	0.0399 (13)	0.0081 (9)	0.0042 (10)	0.0105 (10)
C17	0.0255 (10)	0.0184 (9)	0.0312 (11)	0.0003 (7)	0.0099 (8)	−0.0014 (8)
C18	0.0327 (11)	0.0268 (10)	0.0242 (10)	−0.0010 (8)	0.0075 (8)	−0.0051 (8)
C16	0.0299 (10)	0.0258 (10)	0.0234 (10)	0.0030 (8)	0.0047 (8)	0.0017 (8)
C2	0.0243 (9)	0.0179 (9)	0.0232 (9)	0.0003 (7)	0.0051 (7)	−0.0014 (7)
C8	0.0201 (9)	0.0205 (9)	0.0255 (10)	−0.0004 (7)	0.0051 (7)	0.0038 (7)
C19	0.0265 (10)	0.0276 (10)	0.0222 (10)	−0.0007 (8)	0.0031 (8)	0.0004 (8)
C9	0.0309 (11)	0.0253 (10)	0.0251 (10)	0.0009 (8)	0.0027 (8)	0.0052 (8)
C3	0.0235 (10)	0.0200 (9)	0.0336 (11)	0.0025 (7)	0.0046 (8)	0.0010 (8)
C5	0.0292 (10)	0.0181 (9)	0.0223 (9)	−0.0034 (8)	0.0067 (8)	−0.0016 (7)
C13	0.0303 (10)	0.0253 (10)	0.0230 (10)	−0.0003 (8)	0.0051 (8)	0.0017 (8)
C11	0.0416 (13)	0.0198 (10)	0.0519 (14)	0.0049 (9)	0.0205 (11)	0.0053 (10)
C12	0.0439 (13)	0.0251 (10)	0.0350 (12)	−0.0047 (9)	0.0168 (10)	−0.0043 (9)

Geometric parameters (Å, °)

Cl1—C5	1.742 (2)	C14—C19	1.386 (3)
S1—O2	1.4307 (15)	C10—C9	1.379 (3)
S1—O3	1.4329 (15)	C10—C11	1.381 (3)
S1—N1	1.6383 (17)	C10—H10	0.950
S1—C14	1.7679 (19)	C17—C16	1.381 (3)
Cl2—C17	1.7374 (19)	C17—C18	1.386 (3)

O1—C7	1.223 (2)	C18—C19	1.387 (3)
N1—C2	1.429 (2)	C18—H18	0.950
N1—H1	0.865 (15)	C16—H16	0.950
C4—C3	1.384 (3)	C2—C3	1.392 (3)
C4—C5	1.385 (3)	C8—C13	1.391 (3)
C4—H4	0.950	C8—C9	1.394 (3)
C15—C16	1.386 (3)	C19—H19	0.950
C15—C14	1.391 (3)	C9—H9	0.950
C15—H15	0.950	C3—H3	0.950
C7—C8	1.490 (3)	C13—C12	1.391 (3)
C7—C1	1.499 (3)	C13—H13	0.950
C6—C5	1.384 (3)	C11—C12	1.384 (3)
C6—C1	1.393 (3)	C11—H11	0.950
C6—H6	0.950	C12—H12	0.950
C1—C2	1.404 (3)		
O2—S1—O3	120.89 (9)	C17—C18—C19	118.95 (18)
O2—S1—N1	107.51 (9)	C17—C18—H18	120.5
O3—S1—N1	104.68 (9)	C19—C18—H18	120.5
O2—S1—C14	107.79 (9)	C17—C16—C15	118.98 (18)
O3—S1—C14	108.11 (9)	C17—C16—H16	120.5
N1—S1—C14	107.13 (9)	C15—C16—H16	120.5
C2—N1—S1	121.25 (13)	C3—C2—C1	120.09 (18)
C2—N1—H1	114.7 (15)	C3—C2—N1	118.44 (17)
S1—N1—H1	110.0 (15)	C1—C2—N1	121.43 (17)
C3—C4—C5	119.05 (18)	C13—C8—C9	119.28 (18)
C3—C4—H4	120.5	C13—C8—C7	122.45 (17)
C5—C4—H4	120.5	C9—C8—C7	118.12 (17)
C16—C15—C14	119.66 (18)	C14—C19—C18	119.64 (18)
C16—C15—H15	120.2	C14—C19—H19	120.2
C14—C15—H15	120.2	C18—C19—H19	120.2
O1—C7—C8	120.44 (17)	C10—C9—C8	120.21 (19)
O1—C7—C1	120.10 (17)	C10—C9—H9	119.9
C8—C7—C1	119.43 (16)	C8—C9—H9	119.9
C5—C6—C1	119.39 (17)	C4—C3—C2	120.33 (18)
C5—C6—H6	120.3	C4—C3—H3	119.8
C1—C6—H6	120.3	C2—C3—H3	119.8
C6—C1—C2	119.26 (17)	C6—C5—C4	121.67 (18)
C6—C1—C7	120.47 (17)	C6—C5—Cl1	118.85 (15)
C2—C1—C7	120.26 (17)	C4—C5—Cl1	119.48 (15)
C19—C14—C15	120.85 (18)	C12—C13—C8	120.28 (19)
C19—C14—S1	120.09 (15)	C12—C13—H13	119.9
C15—C14—S1	119.06 (15)	C8—C13—H13	119.9
C9—C10—C11	120.3 (2)	C10—C11—C12	120.3 (2)
C9—C10—H10	119.9	C10—C11—H11	119.9
C11—C10—H10	119.9	C12—C11—H11	119.9
C16—C17—C18	121.92 (18)	C11—C12—C13	119.7 (2)
C16—C17—Cl2	119.17 (15)	C11—C12—H12	120.2
C18—C17—Cl2	118.92 (15)	C13—C12—H12	120.2

O2—S1—N1—C2	46.32 (17)	C7—C1—C2—N1	−6.2 (3)
O3—S1—N1—C2	176.05 (14)	S1—N1—C2—C3	−78.1 (2)
C14—S1—N1—C2	−69.30 (16)	S1—N1—C2—C1	104.42 (19)
C5—C6—C1—C2	1.4 (3)	O1—C7—C8—C13	−154.24 (19)
C5—C6—C1—C7	−179.43 (17)	C1—C7—C8—C13	23.5 (3)
O1—C7—C1—C6	−136.12 (19)	O1—C7—C8—C9	21.4 (3)
C8—C7—C1—C6	46.2 (2)	C1—C7—C8—C9	−160.91 (17)
O1—C7—C1—C2	43.0 (3)	C15—C14—C19—C18	0.0 (3)
C8—C7—C1—C2	−134.72 (18)	S1—C14—C19—C18	−179.36 (15)
C16—C15—C14—C19	−0.2 (3)	C17—C18—C19—C14	0.6 (3)
C16—C15—C14—S1	179.15 (15)	C11—C10—C9—C8	1.5 (3)
O2—S1—C14—C19	−13.39 (18)	C13—C8—C9—C10	−0.8 (3)
O3—S1—C14—C19	−145.63 (16)	C7—C8—C9—C10	−176.61 (19)
N1—S1—C14—C19	102.04 (17)	C5—C4—C3—C2	0.9 (3)
O2—S1—C14—C15	167.28 (15)	C1—C2—C3—C4	3.3 (3)
O3—S1—C14—C15	35.04 (18)	N1—C2—C3—C4	−174.15 (17)
N1—S1—C14—C15	−77.28 (17)	C1—C6—C5—C4	2.9 (3)
C16—C17—C18—C19	−0.9 (3)	C1—C6—C5—Cl1	−178.11 (14)
Cl2—C17—C18—C19	178.95 (15)	C3—C4—C5—C6	−4.0 (3)
C18—C17—C16—C15	0.7 (3)	C3—C4—C5—Cl1	176.93 (15)
Cl2—C17—C16—C15	−179.17 (15)	C9—C8—C13—C12	−0.7 (3)
C14—C15—C16—C17	−0.1 (3)	C7—C8—C13—C12	174.83 (18)
C6—C1—C2—C3	−4.5 (3)	C9—C10—C11—C12	−0.5 (3)
C7—C1—C2—C3	176.37 (17)	C10—C11—C12—C13	−1.1 (3)
C6—C1—C2—N1	172.92 (17)	C8—C13—C12—C11	1.7 (3)

Hydrogen-bond geometry (Å, °)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
N1—H1···Cl1^{i}	0.865 (15)	2.967 (19)	3.6519 (17)	137.5 (18)
N1—H1···O1	0.865 (15)	2.20 (2)	2.798 (2)	126.1 (18)

Symmetry codes: (i) −*x*+1, −*y*, −*z*+1.

Fig. 1

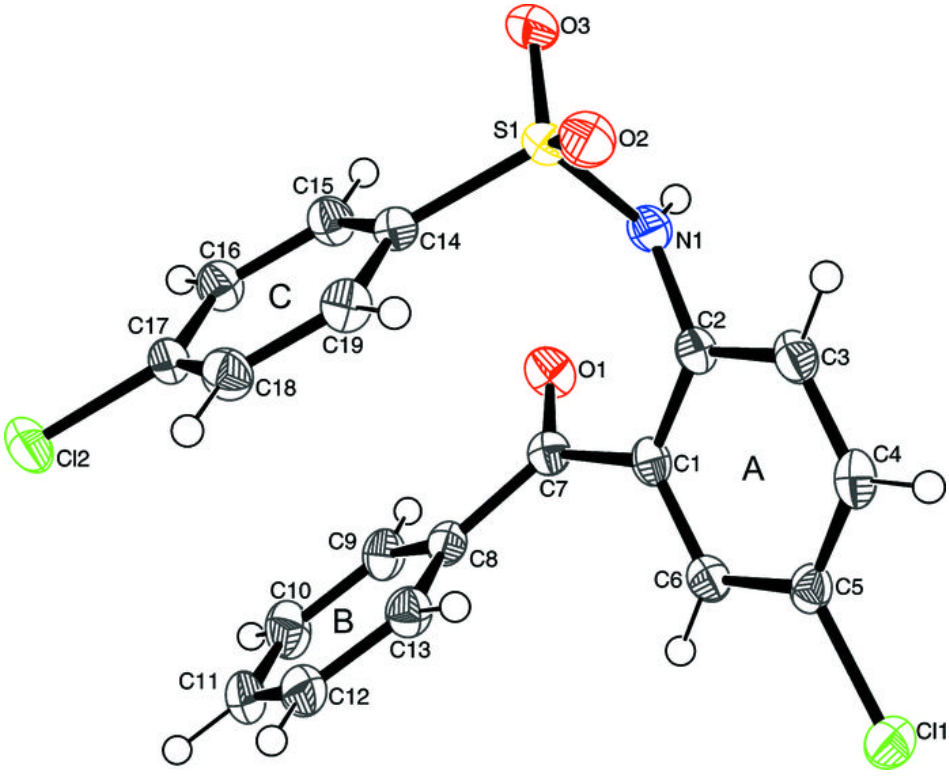

O3
S1
O2
C15
C14
N1
C16
C
C17
C19
C2
C3
O1
C18
Cl2
C7
C1
A
C4
C9
C8
C6
C5
C10
B
C13
C11
C12
Cl1

Fig. 2

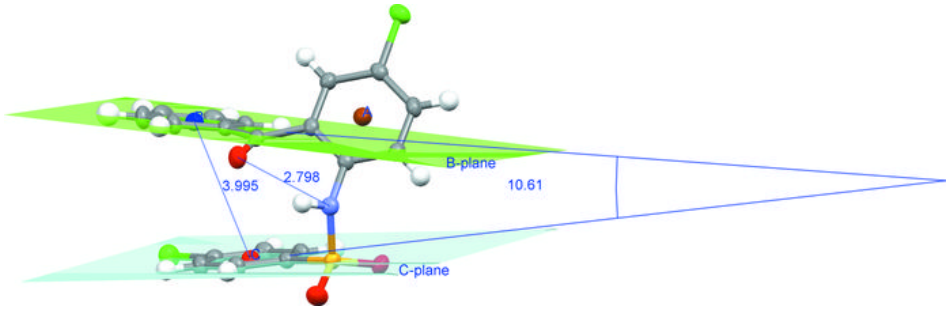


Fig. 3