

## *N*-(2-Chlorophenylsulfonyl)-2,2-dimethylpropanamide

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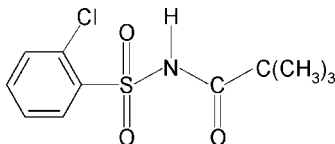
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 Key indicators: single-crystal X-ray study;  $T = 293$  K; mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å;  $R$  factor = 0.036;  $wR$  factor = 0.107; data-to-parameter ratio = 17.3.

In the title compound,  $\text{C}_{11}\text{H}_{14}\text{ClNO}_3\text{S}$ , the C—S—N—C torsion angle is  $-61.69$  (17)°. In the crystal, inversion dimers linked by pairs of N—H...O hydrogen bonds occur, generating  $R_2^2(8)$  loops.

### Related literature

For the sulfanilamide moiety in sulfonamide drugs, see: Maren (1976). For the ability of sulfonamides to form hydrogen bonds in the solid state, see: Yang & Guillory (1972). For hydrogen-bonding modes of sulfonamides, see: Adsmund & Grant (2001). For our study of the effect of substituents on the structures of *N*-(aryl)-methanesulfonamides, see: Gowda *et al.* (2007). For related structures, see: Gowda *et al.* (2008); Shakuntala *et al.* (2011a,b).



### Experimental

#### Crystal data

$\text{C}_{11}\text{H}_{14}\text{ClNO}_3\text{S}$   
 $M_r = 275.74$   
 Triclinic,  $P\bar{1}$   
 $a = 8.785$  (1) Å  
 $b = 8.914$  (1) Å  
 $c = 9.313$  (1) Å

$\alpha = 103.12$  (1)°  
 $\beta = 107.14$  (1)°  
 $\gamma = 94.20$  (1)°  
 $V = 670.98$  (13) Å<sup>3</sup>  
 $Z = 2$   
 Mo  $K\alpha$  radiation

$\mu = 0.44$  mm<sup>-1</sup>  
 $T = 293$  K

$0.44 \times 0.40 \times 0.38$  mm

#### Data collection

Oxford Diffraction Xcalibur diffractometer with a Sapphire CCD detector  
 Absorption correction: multi-scan (*CrysAlis RED*; Oxford)

Diffraction, 2009)  
 $T_{\min} = 0.831$ ,  $T_{\max} = 0.852$   
 4355 measured reflections  
 2730 independent reflections  
 2321 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.012$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.036$   
 $wR(F^2) = 0.107$   
 $S = 1.06$   
 2730 reflections  
 158 parameters  
 1 restraint

H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\text{max}} = 0.26$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.26$  e Å<sup>-3</sup>

**Table 1**

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1N}\cdots\text{O1}^i$	0.83 (2)	2.22 (2)	3.042 (2)	178 (2)

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

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2009); cell refinement: *CrysAlis RED* (Oxford Diffraction, 2009); data reduction: *CrysAlis RED*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: TK2742).

### References

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**supplementary materials**

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## ***N*-(2-Chlorophenylsulfonyl)-2,2-dimethylpropanamide**

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### **Comment**

The molecular structures of sulfonamide drugs contain the sulfanilamide moiety (Maren, 1976). The propensity for hydrogen bonding in the solid state, due to the presence of various hydrogen bond donors and acceptors, gives rise to polymorphism (Yang & Guillory, 1972). The hydrogen bonding preferences of sulfonamides has also been investigated (Admond & Grant, 2001). The nature and position of substituents in *N*-(aryl)sulfonamides play a significant role on their crystal structures. As a part of a study of the substituent effects upon their crystal structures (Gowda *et al.*, 2007, 2008; Shakuntala *et al.*, 2011*a,b*), in the present work, the crystal structure of *N*-(2-chlorophenylsulfonyl)-2,2,2-trimethylacetamide (I) has been determined. The N—H and C=O bonds in the SO<sub>2</sub>—NH—CO—C segment of (I) are *anti* to each other (Fig. 1), similar to that observed in *N*-(2-methylphenylsulfonyl)-2,2,2-trimethylacetamide (II) (Shakuntala *et al.*, 2011*a*), *N*-(phenylsulfonyl)-2,2,2-trimethylacetamide (III) (Gowda *et al.*, 2008) and *N*-(2-chlorophenylsulfonyl)-acetamide (IV) (Shakuntala *et al.*, 2011*b*). Further, the amide hydrogen is *syn* to the *ortho*-chloro group in the benzene ring.

The N—C bond in the C—SO<sub>2</sub>—NH—C segment has *gauche* torsion with respect to the S=O bonds. The molecule is twisted at the *S*-atom with a C—S—N—C torsion angle of -61.7 (2)°, compared to the values of -65.4 (2)° in (II), -64.5 (3)° in (III), and -71.7 (3)° and 61.2 (3)° in the two independent molecules of (IV).

In the crystal structure, the pairs of intermolecular N—H...O hydrogen bonds (Table 1) link the molecules into inversion-related dimers; a view of the crystal packing is shown in Fig. 2.

### **Experimental**

The title compound was prepared by refluxing 2-chlorobenzenesulfonamide (0.10 mole) with an excess of pivalyl chloride (0.20 mole) for about an hour on a water bath. The reaction mixture was cooled and poured into ice cold water. The resulting solid was separated, washed thoroughly with water and dissolved in warm dilute sodium hydrogen carbonate solution. The title compound was re-precipitated by acidifying the filtered solution with glacial acetic acid. It was filtered, dried and recrystallized from ethanol. Colorless prisms of the title compound used in X-ray diffraction studies were obtained from a slow evaporation of an ethanolic solution of the compound.

### **Refinement**

The H atom of the NH group was located in a difference map and later restrained to the distance N—H = 0.86 (2) Å. The other H atoms were positioned with idealized geometry using a riding model with the aromatic C—H = 0.93 Å and methyl C—H = 0.96 Å. All H atoms were refined with isotropic displacement parameters (set to 1.2 times of the  $U_{eq}$  of the parent atom).

## Figures

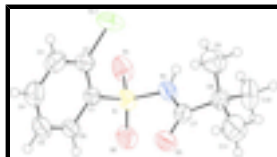


Fig. 1. Molecular structure of the title compound, showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level.

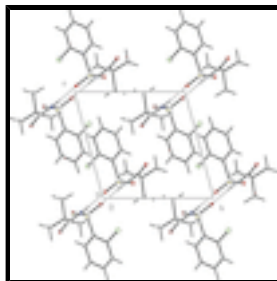


Fig. 2. Molecular packing in the title compound. Hydrogen bonds are shown as dashed lines.

## *N*-(2-Chlorophenylsulfonyl)-2,2-dimethylpropanamide

### Crystal data

$C_{11}H_{14}ClNO_3S$	$Z = 2$
$M_r = 275.74$	$F(000) = 288$
Triclinic, $P\bar{1}$	$D_x = 1.365 \text{ Mg m}^{-3}$
Hall symbol: $-P\ 1$	Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
$a = 8.785 (1) \text{ \AA}$	Cell parameters from 2522 reflections
$b = 8.914 (1) \text{ \AA}$	$\theta = 3.1\text{--}27.7^\circ$
$c = 9.313 (1) \text{ \AA}$	$\mu = 0.44 \text{ mm}^{-1}$
$\alpha = 103.12 (1)^\circ$	$T = 293 \text{ K}$
$\beta = 107.14 (1)^\circ$	Prism, colourless
$\gamma = 94.20 (1)^\circ$	$0.44 \times 0.40 \times 0.38 \text{ mm}$
$V = 670.98 (13) \text{ \AA}^3$	

### Data collection

Oxford Diffraction Xcalibur diffractometer with a Sapphire CCD detector	2730 independent reflections
Radiation source: fine-focus sealed tube graphite	2321 reflections with $I > 2\sigma(I)$
Rotation method data acquisition using $\omega$ and $\varphi$ scans	$R_{\text{int}} = 0.012$
Absorption correction: multi-scan ( <i>CrysAlis RED</i> ; Oxford Diffraction, 2009)	$\theta_{\text{max}} = 26.4^\circ$ , $\theta_{\text{min}} = 3.2^\circ$
$T_{\text{min}} = 0.831$ , $T_{\text{max}} = 0.852$	$h = -10 \rightarrow 10$
4355 measured reflections	$k = -11 \rightarrow 10$
	$l = -11 \rightarrow 7$

### Refinement

Refinement on $F^2$	Secondary atom site location: difference Fourier map
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Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.036$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.107$	$w = 1/[\sigma^2(F_o^2) + (0.0533P)^2 + 0.2343P]$
$S = 1.06$	where $P = (F_o^2 + 2F_c^2)/3$
2730 reflections	$(\Delta/\sigma)_{\max} = 0.004$
158 parameters	$\Delta\rho_{\max} = 0.26 \text{ e } \text{\AA}^{-3}$
1 restraint	$\Delta\rho_{\min} = -0.26 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	Extinction correction: <i>SHELXL97</i> (Sheldrick, 2008), $F_c^* = kFc[1+0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$
	Extinction coefficient: 0.129 (8)

### Special details

**Experimental.** CrysAlis RED (Oxford Diffraction, 2009) Empirical absorption correction using spherical harmonics, implemented in SCALE3 ABSPACK scaling algorithm.

**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 ( $\text{\AA}^2$ )

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.1873 (2)	0.8220 (2)	0.6448 (2)	0.0421 (4)
C2	0.2567 (2)	0.9285 (2)	0.5857 (2)	0.0506 (5)
C3	0.1853 (3)	0.9346 (3)	0.4339 (3)	0.0635 (6)
H3	0.2316	1.0059	0.3940	0.076*
C4	0.0468 (3)	0.8358 (3)	0.3420 (3)	0.0666 (6)
H4	-0.0012	0.8417	0.2405	0.080*
C5	-0.0218 (2)	0.7287 (3)	0.3977 (3)	0.0603 (5)
H5	-0.1153	0.6614	0.3341	0.072*
C6	0.0485 (2)	0.7210 (2)	0.5489 (2)	0.0482 (4)
H6	0.0026	0.6476	0.5868	0.058*
C7	0.4718 (2)	0.6231 (2)	0.7753 (2)	0.0424 (4)
C8	0.6484 (2)	0.6029 (2)	0.8078 (2)	0.0473 (4)
C9	0.7309 (3)	0.7253 (3)	0.7535 (4)	0.0773 (7)
H9A	0.7220	0.8274	0.8089	0.093*
H9B	0.6802	0.7112	0.6439	0.093*
H9C	0.8427	0.7144	0.7735	0.093*
C10	0.7287 (3)	0.6256 (4)	0.9819 (3)	0.0817 (8)

## supplementary materials

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H10A	0.6750	0.5503	1.0166	0.098*
H10B	0.7221	0.7287	1.0368	0.098*
H10C	0.8398	0.6122	1.0018	0.098*
C11	0.6588 (3)	0.4422 (3)	0.7175 (4)	0.0894 (9)
H11A	0.6088	0.4308	0.6082	0.107*
H11B	0.6044	0.3650	0.7496	0.107*
H11C	0.7699	0.4287	0.7373	0.107*
N1	0.44486 (18)	0.76248 (19)	0.86001 (19)	0.0483 (4)
H1N	0.515 (2)	0.838 (2)	0.912 (2)	0.058*
O1	0.29358 (17)	0.96456 (19)	0.94151 (18)	0.0656 (4)
O2	0.16258 (17)	0.69277 (19)	0.86012 (18)	0.0635 (4)
O3	0.36032 (17)	0.52985 (17)	0.68368 (18)	0.0614 (4)
Cl1	0.43167 (8)	1.05545 (7)	0.69656 (9)	0.0820 (2)
S1	0.26460 (5)	0.81187 (6)	0.83971 (5)	0.04740 (17)

### Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C1	0.0325 (8)	0.0377 (8)	0.0535 (10)	0.0082 (6)	0.0111 (7)	0.0092 (7)
C2	0.0415 (9)	0.0402 (9)	0.0667 (12)	0.0038 (7)	0.0138 (9)	0.0127 (8)
C3	0.0661 (14)	0.0577 (12)	0.0771 (15)	0.0134 (10)	0.0264 (12)	0.0318 (11)
C4	0.0639 (13)	0.0745 (15)	0.0584 (13)	0.0198 (12)	0.0086 (10)	0.0229 (11)
C5	0.0428 (10)	0.0636 (13)	0.0600 (12)	0.0055 (9)	0.0030 (9)	0.0062 (10)
C6	0.0353 (9)	0.0434 (9)	0.0613 (11)	0.0039 (7)	0.0132 (8)	0.0082 (8)
C7	0.0408 (9)	0.0427 (9)	0.0450 (9)	0.0069 (7)	0.0136 (7)	0.0142 (7)
C8	0.0415 (9)	0.0493 (10)	0.0518 (10)	0.0140 (8)	0.0151 (8)	0.0119 (8)
C9	0.0606 (14)	0.0830 (17)	0.104 (2)	0.0109 (12)	0.0483 (14)	0.0255 (15)
C10	0.0679 (15)	0.109 (2)	0.0668 (15)	0.0413 (15)	0.0077 (12)	0.0302 (14)
C11	0.0721 (16)	0.0639 (15)	0.118 (2)	0.0272 (13)	0.0259 (16)	-0.0034 (15)
N1	0.0327 (8)	0.0500 (9)	0.0538 (9)	0.0090 (6)	0.0079 (6)	0.0042 (7)
O1	0.0498 (8)	0.0707 (10)	0.0623 (9)	0.0196 (7)	0.0132 (7)	-0.0068 (7)
O2	0.0460 (8)	0.0832 (11)	0.0725 (10)	0.0109 (7)	0.0249 (7)	0.0343 (8)
O3	0.0470 (8)	0.0519 (8)	0.0717 (9)	-0.0013 (6)	0.0117 (7)	0.0019 (7)
Cl1	0.0639 (4)	0.0658 (4)	0.1008 (5)	-0.0238 (3)	0.0154 (3)	0.0167 (3)
S1	0.0350 (2)	0.0546 (3)	0.0507 (3)	0.01154 (19)	0.01324 (19)	0.0089 (2)

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

C1—C6	1.389 (2)	C8—C9	1.524 (3)
C1—C2	1.388 (3)	C8—C10	1.524 (3)
C1—S1	1.7664 (19)	C9—H9A	0.9600
C2—C3	1.383 (3)	C9—H9B	0.9600
C2—Cl1	1.728 (2)	C9—H9C	0.9600
C3—C4	1.370 (3)	C10—H10A	0.9600
C3—H3	0.9300	C10—H10B	0.9600
C4—C5	1.368 (3)	C10—H10C	0.9600
C4—H4	0.9300	C11—H11A	0.9600
C5—C6	1.381 (3)	C11—H11B	0.9600
C5—H5	0.9300	C11—H11C	0.9600

C6—H6	0.9300	N1—S1	1.6434 (15)
C7—O3	1.202 (2)	N1—H1N	0.826 (16)
C7—N1	1.389 (2)	O1—S1	1.4274 (15)
C7—C8	1.523 (2)	O2—S1	1.4209 (16)
C8—C11	1.512 (3)		
C6—C1—C2	119.18 (18)	C8—C9—H9A	109.5
C6—C1—S1	117.88 (15)	C8—C9—H9B	109.5
C2—C1—S1	122.92 (14)	H9A—C9—H9B	109.5
C3—C2—C1	119.74 (18)	C8—C9—H9C	109.5
C3—C2—C11	118.34 (16)	H9A—C9—H9C	109.5
C1—C2—C11	121.92 (16)	H9B—C9—H9C	109.5
C4—C3—C2	120.2 (2)	C8—C10—H10A	109.5
C4—C3—H3	119.9	C8—C10—H10B	109.5
C2—C3—H3	119.9	H10A—C10—H10B	109.5
C5—C4—C3	120.8 (2)	C8—C10—H10C	109.5
C5—C4—H4	119.6	H10A—C10—H10C	109.5
C3—C4—H4	119.6	H10B—C10—H10C	109.5
C4—C5—C6	119.62 (19)	C8—C11—H11A	109.5
C4—C5—H5	120.2	C8—C11—H11B	109.5
C6—C5—H5	120.2	H11A—C11—H11B	109.5
C5—C6—C1	120.45 (19)	C8—C11—H11C	109.5
C5—C6—H6	119.8	H11A—C11—H11C	109.5
C1—C6—H6	119.8	H11B—C11—H11C	109.5
O3—C7—N1	120.27 (17)	C7—N1—S1	123.31 (13)
O3—C7—C8	124.89 (17)	C7—N1—H1N	125.0 (16)
N1—C7—C8	114.84 (15)	S1—N1—H1N	110.5 (16)
C11—C8—C7	109.01 (18)	O2—S1—O1	118.83 (10)
C11—C8—C9	109.4 (2)	O2—S1—N1	109.93 (9)
C7—C8—C9	108.39 (16)	O1—S1—N1	104.49 (9)
C11—C8—C10	110.8 (2)	O2—S1—C1	107.67 (9)
C7—C8—C10	109.76 (17)	O1—S1—C1	109.39 (9)
C9—C8—C10	109.4 (2)	N1—S1—C1	105.81 (8)
C6—C1—C2—C3	1.3 (3)	N1—C7—C8—C9	-65.3 (2)
S1—C1—C2—C3	-177.01 (16)	O3—C7—C8—C10	-126.5 (2)
C6—C1—C2—C11	-178.68 (14)	N1—C7—C8—C10	54.2 (2)
S1—C1—C2—C11	3.0 (2)	O3—C7—N1—S1	-1.5 (3)
C1—C2—C3—C4	-0.1 (3)	C8—C7—N1—S1	177.89 (13)
C11—C2—C3—C4	179.88 (18)	C7—N1—S1—O2	54.30 (18)
C2—C3—C4—C5	-0.9 (4)	C7—N1—S1—O1	-177.13 (15)
C3—C4—C5—C6	0.6 (3)	C7—N1—S1—C1	-61.69 (17)
C4—C5—C6—C1	0.6 (3)	C6—C1—S1—O2	0.55 (16)
C2—C1—C6—C5	-1.5 (3)	C2—C1—S1—O2	178.86 (15)
S1—C1—C6—C5	176.85 (15)	C6—C1—S1—O1	-129.89 (14)
O3—C7—C8—C11	-4.9 (3)	C2—C1—S1—O1	48.42 (17)
N1—C7—C8—C11	175.7 (2)	C6—C1—S1—N1	118.06 (15)
O3—C7—C8—C9	114.0 (2)	C2—C1—S1—N1	-63.63 (17)

## supplementary materials

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### Hydrogen-bond geometry (Å, °)

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N1-H1N\cdots O1^i$	0.83 (2)	2.22 (2)	3.042 (2)	178 (2)

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



Fig. 1

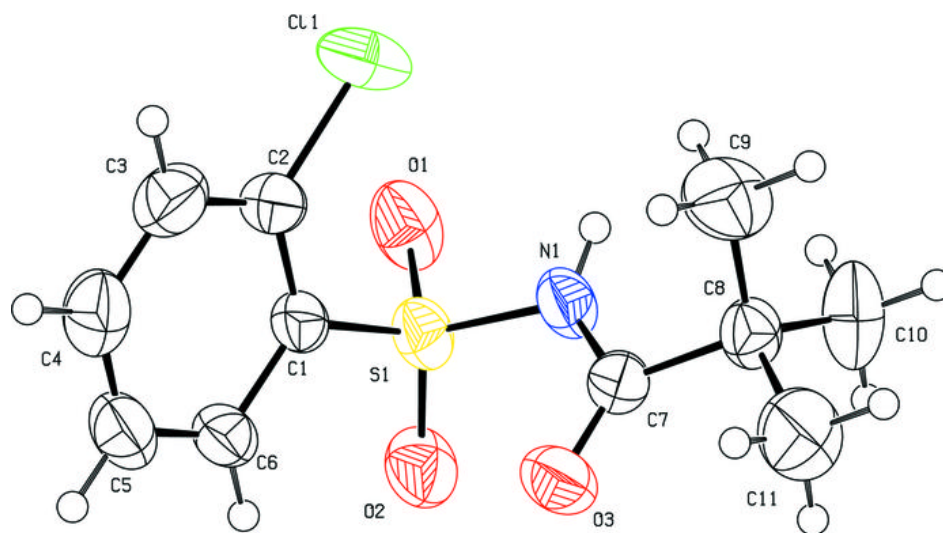


Fig. 2

