

2-Chloro-4-(2-iodobenzenesulfonamido)-benzoic acid

Muhammad Nadeem Arshad,^{a*}‡ Islam Ullah Khan,^b H. M. Rafique,^a Abdullah M. Asiri^c and Muhammad Shafiq^b

^aX-ray Diffraction and Crystallography Laboratory, Department of Physics, School of Physical Sciences, University of the Punjab, Quaid-e-Azam Campus, Lahore 54590, Pakistan, ^bMaterials Chemistry Laboratory, Department of Chemistry, GC University, Lahore 54000, Pakistan, and ^cThe Center of Excellence for Advanced Materials Research, King Abdul Aziz University, Jeddah, PO Box 80203, Saudi Arabia
Correspondence e-mail: mnachemist@hotmail.com

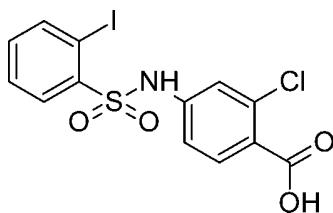
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Key indicators: single-crystal X-ray study; $T = 296\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.010\text{ \AA}$; R factor = 0.054; wR factor = 0.129; data-to-parameter ratio = 19.2.

In the title compound, $\text{C}_{13}\text{H}_9\text{ClINO}_4\text{S}$, the dihedral angle between the aromatic rings is $81.04(17)^\circ$. The disposition of the I and Cl atoms attached to the two rings is *anti*. In the crystal, molecules are connected via $\text{O}-\text{H}\cdots\text{O}$ and $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

For background to thiazine heterocycles, see: Arshad *et al.* (2008, 2011). For their biological activity, see: Medina *et al.* (1999). For related structures, see: Arshad *et al.* (2009a,b,c).



Experimental

Crystal data

$\text{C}_{13}\text{H}_9\text{ClINO}_4\text{S}$	$V = 1473.68(14)\text{ \AA}^3$
$M_r = 437.62$	$Z = 4$
Monoclinic, $P2_1/n$	$\text{Mo } K\alpha$ radiation
$a = 14.1522(8)\text{ \AA}$	$\mu = 2.51\text{ mm}^{-1}$
$b = 7.3203(4)\text{ \AA}$	$T = 296\text{ K}$
$c = 14.7193(8)\text{ \AA}$	$0.18 \times 0.15 \times 0.09\text{ mm}$
$\beta = 104.892(2)^\circ$	

Data collection

Bruker Kappa APEXII CCD diffractometer	16645 measured reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2007)	3668 independent reflections
$T_{\min} = 0.661$, $T_{\max} = 0.806$	1876 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.058$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.054$	191 parameters
$wR(F^2) = 0.129$	H-atom parameters constrained
$S = 1.03$	$\Delta\rho_{\max} = 1.24\text{ e \AA}^{-3}$
3668 reflections	$\Delta\rho_{\min} = -1.30\text{ e \AA}^{-3}$

Table 1

Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1—H1 \cdots O2 ⁱ	0.86	2.38	3.214 (6)	162
O1—H1O \cdots O2 ⁱⁱ	0.82	2.09	2.771 (6)	140

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

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *PLATON* (Spek, 2009); software used to prepare material for publication: *WinGX* (Farrugia, 1999) and *PLATON*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5865).

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‡ Materials Chemistry Laboratory, Department of Chemistry, GC University, Lahore 54000, Pakistan.