

3-(1*H*-1,2,4-Triazol-3-ylsulfanylmethyl)-benzoic acid

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Key indicators

Single-crystal X-ray study

 $T = 293\text{ K}$ Mean $\sigma(\text{C}-\text{C}) = 0.007\text{ \AA}$ R factor = 0.058 wR factor = 0.133

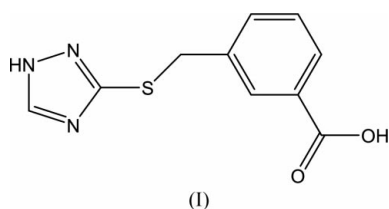
Data-to-parameter ratio = 12.6

For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

The title compound, $\text{C}_{10}\text{H}_9\text{N}_3\text{O}_2\text{S}$, crystallizes in the orthorhombic non-centrosymmetric space group $P2_12_12_1$. The molecules are linked into a three-dimensional network through $\text{O}-\text{H}\cdots\text{N}$ and $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds and weak $\text{S}\cdots\text{O}$ interactions.

Comment

The construction of chiral coordination polymers combining asymmetric catalysis and optical activity is a rapidly developing area of supramolecular chemistry (Seo *et al.*, 2000; Evans & Lin, 2002). Generally, chiral compounds can be obtained either by stereoselective synthesis using chiral organic ligands or by spontaneous resolution upon crystallization without any chiral sources, and the chirally discriminative interactions may arise from coordination bonds and/or hydrogen bonds. We rationalize that the twisted binding sites of asymmetric bridging ligands containing sulfide moieties can induce the formation of chiral structures when linked by suitable metal nodes (Han *et al.*, 2003). Additionally, the incorporation of S atoms in the related ligands will lead to weak interactions between the chalcogen centers, which play an important role in the crystal structures. We report here the synthesis and crystal structure of the organic ligand 3-(1*H*-1,2,4-triazol-3-ylsulfanylmethyl)benzoic acid, (I) (Fig. 1).



Compound (I), prepared from 1*H*-1,2,4-triazole-3-thiol and 3-(chloromethyl)benzoic acid, crystallizes in the orthorhombic non-centrosymmetric space group $P2_12_12_1$. Intriguing features of the compound arise not only from its flexibility but also from the presence of triazole and benzenecarboxylate

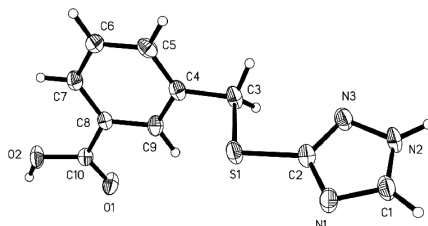


Figure 1

Perspective view of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.

moieties connected by a $-\text{CH}_2\text{S}-$ spacer. The sp^3 configurations of the C and S atoms of the $-\text{CH}_2\text{S}-$ spacer force the molecule to be twisted. Actually, the C—C—S and C—S—C angles are 105.4 (3) and 101.3 (2) $^\circ$, respectively. The dihedral angle between the benzene and triazole rings is 70.1 (2) $^\circ$. The molecules are linked into a three-dimensional network through O—H \cdots N and N—H \cdots O hydrogen bonds (Table 2) and weak S \cdots O (3.251 Å) interactions (Fig. 2).

Experimental

1*H*-1,2,4-Triazole-3-thiol (1.01 g, 10 mmol) was added to a stirred solution of NaOH (0.4 g, 10 mmol) in H₂O (20 ml). The mixture was warmed to reflux, then 3-(chloromethyl)benzoic acid (1.70 g, 10 mmol) was added and the mixture was refluxed for about 10 h with vigorous stirring. After cooling, the mixture was left to stand overnight. The precipitate was filtered off and washed with water, giving a fine white powder in 80% yield. Colorless needle-shaped crystals suitable for X-ray diffraction were isolated after recrystallization from a chloroform–ether solution.

Crystal data

C ₁₀ H ₉ N ₃ O ₂ S	Mo <i>K</i> α radiation
<i>M_r</i> = 235.26	Cell parameters from 3393 reflections
Orthorhombic, <i>P</i> 2 ₁ 2 ₁ 2 ₁	θ = 2.1–25.1 $^\circ$
<i>a</i> = 5.5079 (1) Å	μ = 0.30 mm ^{−1}
<i>b</i> = 13.1233 (6) Å	<i>T</i> = 293 (2) K
<i>c</i> = 14.3414 (7) Å	Needle, colorless
<i>V</i> = 1036.62 (7) Å ³	0.20 × 0.10 × 0.08 mm
<i>Z</i> = 4	
<i>D_x</i> = 1.507 Mg m ^{−3}	

Data collection

Bruker SMART CCD diffractometer	1827 independent reflections
ω scans	1460 reflections with <i>I</i> > 2σ(<i>I</i>)
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	<i>R</i> _{int} = 0.044
<i>T</i> _{min} = 0.972, <i>T</i> _{max} = 0.978	θ _{max} = 25.1 $^\circ$
3393 measured reflections	<i>h</i> = −6 → 6
	<i>k</i> = −10 → 15
	<i>l</i> = −17 → 8

Refinement

Refinement on <i>F</i> ²	$w = 1/[\sigma^2(F_o^2) + (0.0275P)^2 + 1.1042P]$
$R[F^2 > 2\sigma(F^2)] = 0.058$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.133$	(Δ/σ) _{max} = 0.001
<i>S</i> = 1.26	Δρ _{max} = 0.22 e Å ^{−3}
1827 reflections	Δρ _{min} = −0.27 e Å ^{−3}
145 parameters	Absolute structure: Flack (1983),
H atoms treated by a mixture of independent and constrained refinement	728 Friedel pairs
	Flack parameter = 0.0 (2)

Table 1

Selected geometric parameters (Å, $^\circ$).

S1—C2	1.760 (4)	C3—C4	1.524 (6)
S1—C3	1.818 (5)	C4—C5	1.384 (7)
O1—C10	1.220 (5)	C4—C9	1.390 (7)
O2—C10	1.317 (5)	C5—C6	1.396 (7)
N1—C1	1.338 (6)	C6—C7	1.381 (7)
N1—C2	1.352 (6)	C7—C8	1.383 (7)
N2—C1	1.314 (7)	C8—C9	1.403 (6)
N2—N3	1.373 (5)	C8—C10	1.486 (7)
N3—C2	1.318 (6)		
C2—S1—C3	101.3 (2)	C4—C3—S1	105.4 (3)

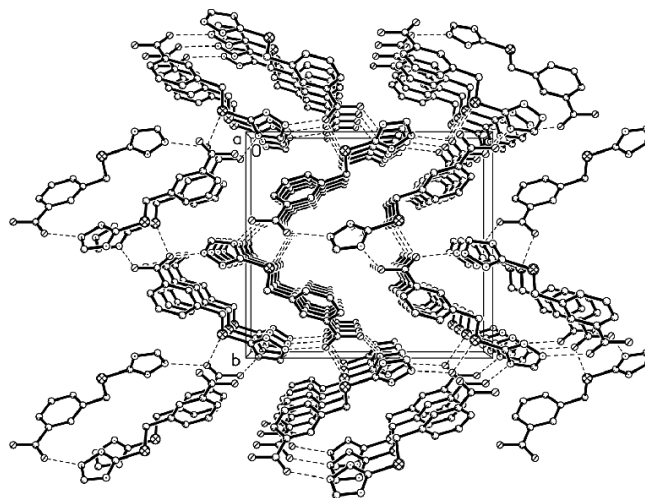


Figure 2

Packing diagram, viewed along the *a* axis. Hydrogen bonds and weak S \cdots O interactions are indicated by dashed lines. H atoms not involved in hydrogen bonding have been omitted for clarity.

Table 2

Hydrogen-bonding geometry (Å, $^\circ$).

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
O2—H2B \cdots N1 ⁱ	0.84	1.89	2.707 (6)	163
N2—H2A \cdots O1 ⁱⁱ	0.88	1.90	2.767 (5)	170

Symmetry codes: (i) 1 − *x*, *y* − ½, ½ − *z*; (ii) *x* − ½, ½ − *y*, 1 − *z*.

H atoms were positioned geometrically and allowed to ride on their respective parent atoms with *U*_{iso} = 1.5 (carboxyl H) or 1.2 (other H atoms) times *U*_{eq} of the parent atoms

Data collection: *SMART* (Siemens, 1994); cell refinement: *SMART* and *SAINT* (Siemens, 1994); data reduction: *SAINT* and *SHELXTL* (Sheldrick, 1997*b*); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997*a*); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997*a*); molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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References

- Evans, O. R. & Lin, W. (2002). *Acc. Chem. Res.* **35**, 511–522.
 Flack, H. D. (1983). *Acta Cryst.* **A39**, 876–881.
 Han, L., Hong, M., Wang, R., Luo, J., Lin, Z. & Yuan, D. (2003). *Chem. Commun.* pp. 2580–2581.
 Seo, J. S., Whang, D., Lee, H., Jun, S. I., Oh, J., Jeon, Y. J. & Kim, K. (2000). *Nature (London)*, **404**, 982–986.
 Sheldrick, G. M. (1996). *SADABS*. University of Göttingen, Germany.
 Sheldrick, G. M. (1997*a*). *SHELXS97* and *SHELXL97*. Siemens Analytical X-ray Instruments Inc., Madison, Wisconsin, USA.
 Sheldrick, G. M. (1997*b*). *SHELXTL*. Bruker AXS Inc., Madison, Wisconsin, USA.
 Siemens (1994). *SMART* and *SAINT*. Siemens Analytical X-ray Instruments Inc., Madison, Wisconsin, USA.