

Dong-Sheng Ma

College of Chemistry and Materials Science,
Heilongjiang University, Harbin 150080,
People's Republic of China

Correspondence e-mail: hg1000@163.com

Key indicators

Single-crystal X-ray study
 $T = 291$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.049
 wR factor = 0.117
Data-to-parameter ratio = 17.6For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

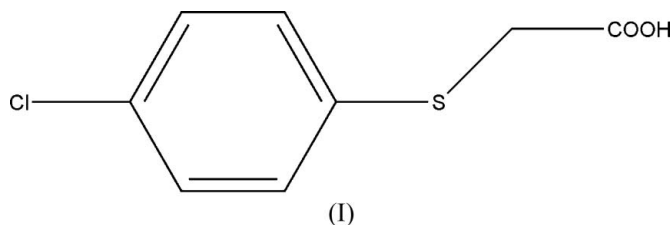
(4-Chlorophenyl)thioglycolic acid

In the title compound, $\text{C}_8\text{H}_7\text{ClO}_2\text{S}$, all bond lengths and angles are normal. Inversion-related molecules are linked by paired $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds into dimers containing an $R_2^2(8)$ motif.

Received 22 November 2006
Accepted 18 January 2007

Comment

(4-Chlorophenyl)thioglycolic acid, (I), is a flexible ligand with versatile binding abilities and the capability of participating in hydrogen bonds as both a donor and an acceptor, thus representing an excellent candidate for the construction of supramolecular complexes. The crystal structure of (4-chlorophenyl)thioglycolic acid was first reported by Shklover *et al.* (1983); it consists of a tris(2-hydroxyethyl)ammonium cation and a 4-chlorophenylthioacetate anion. Two crystal structures of (4-chlorophenyl)thioglycolic acid derivatives have also been reported previously, namely 2,4-dichloro-5-methylphenylthioacetic acid (Mak *et al.*, 1989) and 4-dichloro-2,5-methylphenylthioacetic acid (Kennard *et al.*, 1989).



Interestingly, in contrast to the previously reported structures, the molecular skeleton of (I) is almost planar, the largest deviation being 0.2759 (19) Å for atom O2; however, the sulfanylacetate group is slightly twisted with a C1—S1—C7—C8 torsion angle of 4.94 (18)°. In two of the previous structures, the acetate group is approximately perpendicular to the benzene ring (Shklover *et al.*, 1983; Kennard *et al.*, 1989), but it is approximately parallel to the benzene ring in the third (Mak *et al.*, 1989) and in (I) (Fig. 1).

A familiar carboxylic acid centrosymmetric dimer, containing an $R_2^2(8)$ motif, is built up by $\text{O}-\text{H}\cdots\text{O}$ hydrogen-bonding interactions between the carboxylic acid groups (Fig. 2 and Table 1).

Experimental

(4-Chlorophenyl)thioglycolic acid was prepared by nucleophilic reaction of chloroacetic acid with 4-chlorothiophenol under basic conditions (Nobles *et al.*, 1965). Crystals of (I) suitable for single-crystal X-ray diffraction were grown by slow evaporation of an ethanol solution.

Crystal data

$C_8H_7ClO_2S$
 $M_r = 202.65$
 Monoclinic, $P2_1/n$
 $a = 5.5743$ (11) Å
 $b = 5.3123$ (11) Å
 $c = 29.086$ (6) Å
 $\beta = 90.60$ (3)°
 $V = 861.2$ (3) Å³

$Z = 4$
 $D_x = 1.563$ Mg m⁻³
 Mo $K\alpha$ radiation
 $\mu = 0.64$ mm⁻¹
 $T = 291$ (2) K
 Block, colourless
 $0.33 \times 0.32 \times 0.16$ mm

Data collection

Rigaku R-AXIS RAPID
 diffractometer
 ω scans
 Absorption correction: multi-scan
 (ABSCOR; Higashi, 1995)
 $T_{\min} = 0.817$, $T_{\max} = 0.907$

7327 measured reflections
 1935 independent reflections
 1324 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.048$
 $\theta_{\text{max}} = 27.5^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.049$
 $wR(F^2) = 0.118$
 $S = 1.07$
 1935 reflections
 110 parameters

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0538P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.32$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.22$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$O2-H7\cdots O1^i$	0.82	1.88	2.689 (3)	171

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

All H atoms were placed in calculated positions and treated as riding on their parent atoms, with $C-H = 0.93$ (aromatic) or 0.97 Å (methylene), $O-H = 0.82$ Å (carboxyl), and with $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C)$ or $1.5U_{\text{eq}}(O)$.

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997a); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997a); molecular graphics: *SHELXTL* (Sheldrick, 1997b); software used to prepare material for publication: *SHELXL97*.

The author thanks Heilongjiang University for supporting this study.

References

Higashi, T. (1995). *ABSCOR*. Rigaku Corporation, Tokyo, Japan.

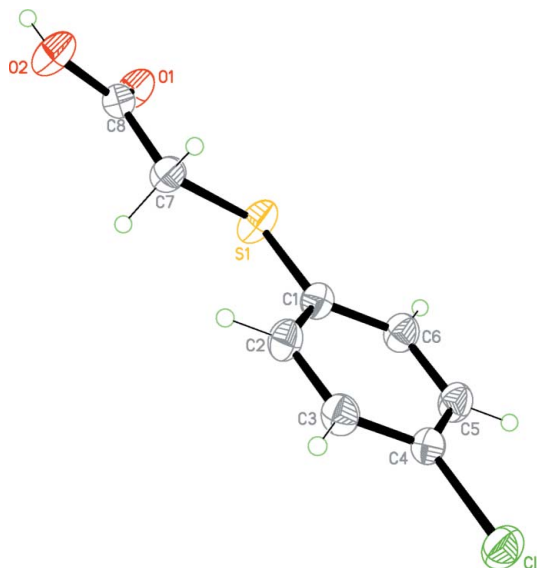


Figure 1

The molecular structure of the title compound, showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are represented as spheres of arbitrary radius.

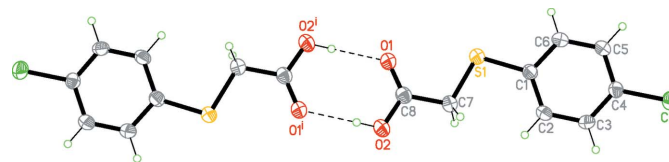


Figure 2

A view of the hydrogen-bonded (dashed lines) dimer [symmetry code: (i) $-x + 1, -y - 1, -z$].

- Kennard, C. H. L., Smith, G., Mak, T. C. W., Yip, W. H. & Chan, W. H. (1989). *Polyhedron*, **8**, 2147–2150.
- Mak, T. C. W., Yip, W. H., Chan, W. H., Kennard, C. H. L., Smith, G. & O'Reilly, E. J. (1989). *Polyhedron*, **8**, 793–800.
- Nobles, W. L. & Thompson, B. B. (1965). *J. Pharm. Sci.* **54**, 709–713.
- Rigaku (1998). *RAPID-AUTO*. Rigaku Corporation, Tokyo, Japan.
- Rigaku/MS (2002). *CrystalStructure*. Rigaku/MS Inc., The Woodlands, Texas, USA.
- Sheldrick, G. M. (1997a). *SHELXL97* and *SHELXS97*. University of Göttingen, Germany.
- Sheldrick, G. M. (1997b). *SHELXTL*. Version 5.10. Bruker AXS Inc., Madison, Wisconsin, USA.
- Shklover, V. E., Gridunova, G. V., Struchkov, Y. T., Voronkov, M. G., Kryukova, Y. I. & Mirskova, A. N. (1983). *Dokl. Akad. Nauk SSSR*, **269**, 387–393. (In Russian.)