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Key indicators

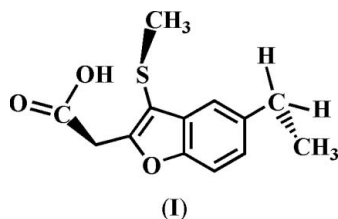
Single-crystal X-ray study
 $T = 173$ K
Mean $\sigma(C-C) = 0.002$ Å
 R factor = 0.035
 wR factor = 0.096
Data-to-parameter ratio = 14.9For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.2-(5-Ethyl-3-methylsulfanyl-1-benzofuran-2-yl)-
acetic acid

The title compound, $C_{13}H_{14}O_3S$, was prepared by alkaline hydrolysis of ethyl 2-(5-ethyl-3-methylsulfanyl-1-benzofuran-2-yl)acetate. The crystal structure is stabilized by aromatic π - π stacking interactions and inversion-related intermolecular $O-H\cdots O$ hydrogen bonds between adjacent carboxyl groups.

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Comment

This work is related to our previous communications on (3-methylsulfanyl-1-benzofuran-5-yl)acetic acid derivatives (Choi *et al.*, 2006, 2007), with the purpose of understanding how the molecular conformation and crystal structure are affected by acetic acid substituents on the 1-benzofuran ring system. The title compound, (I) (Fig. 1), was obtained by hydrolysis of ethyl 2-(5-ethyl-3-methylsulfanyl-1-benzofuran-2-yl)acetate with potassium hydroxide.



In (I), bond lengths and angles are as expected for this type of compound (Choi *et al.*, 2006, 2007). In the crystal structure (Fig. 2), π - π stacking interactions are observed between the furan and benzene rings of adjacent benzofuran systems, with $Cg1\cdots Cg2^i = 3.801(2)$ Å ($Cg1$ and $Cg2$ are the centroids of the $O1/C8/C1/C2/C7$ and $C2-C7$ rings; symmetry code as in Fig. 2). Classical inversion-related $O2-H14\cdots O3^{iii}$ hydrogen bonds link the carboxyl groups of adjacent molecules (Table 1 and Fig. 2).

Experimental

Ethyl 2-(5-ethyl-3-methylsulfanyl-1-benzofuran-2-yl)acetate (834 mg, 3 mmol) was added to a solution of potassium hydroxide (561 mg, 10.0 mmol) in water (10 ml) and methanol (10 ml); the mixture was heated at 333 K for 4 h and then cooled. Water (20 ml) was added, and the solution was washed with dichloromethane. The aqueous layer was acidified to $pH = 1.0$ with concentrated hydrochloric acid and then extracted with chloroform, dried over magnesium sulfate, filtered and concentrated under vacuum. The residue was purified by column chromatography (hexane-ethyl acetate, 1:2 *v/v*) to give (I) as a white solid. Crystals suitable for X-ray analysis were grown by slow evaporation of a benzene solution [yield 92%, m.p. 396–397 K; $R_F = 0.46$ (hexane-ethyl acetate, 1:2 *v/v*)].

Crystal data

$C_{13}H_{14}O_3S$
 $M_r = 250.30$
 Triclinic, $P\bar{1}$
 $a = 5.0746$ (4) Å
 $b = 8.0636$ (6) Å
 $c = 15.058$ (1) Å
 $\alpha = 94.711$ (1)°
 $\beta = 93.153$ (1)°

$\gamma = 90.538$ (1)°
 $V = 613.10$ (8) Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 0.26$ mm⁻¹
 $T = 173$ (2) K
 $0.32 \times 0.25 \times 0.20$ mm

Data collection

Bruker SMART CCD
 diffractometer
 Absorption correction: none
 3461 measured reflections

2368 independent reflections
 2145 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.031$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.035$
 $wR(F^2) = 0.096$
 $S = 1.10$
 2368 reflections
 159 parameters

H atoms treated by a mixture of
 independent and constrained
 refinement
 $\Delta\rho_{max} = 0.28$ e Å⁻³
 $\Delta\rho_{min} = -0.22$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$O2-H14\cdots O3^i$	0.82 (3)	1.87 (3)	2.684 (2)	175 (3)

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

Atom H14 of the carboxyl group was found in a difference Fourier map and refined freely with an isotropic displacement parameter. The other H atoms were positioned geometrically and refined using a riding model, with $C-H = 0.95$ Å for aromatic H atoms, 0.98 Å for methyl H atoms and 0.99 Å for methylene H atoms, and with $U_{iso}(H) = 1.2U_{eq}(C)$ for aromatic and methylene H atoms and $1.5U_{eq}(C)$ for methyl H atoms.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1997); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *DIAMOND* (Brandenburg, 1998); software used to prepare material for publication: *SHELXL97*.

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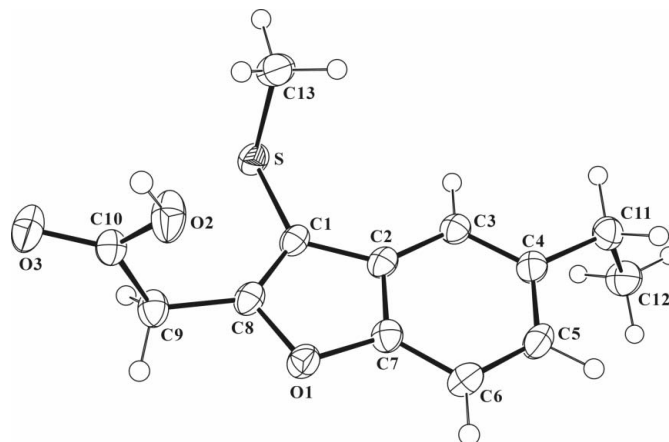


Figure 1

The molecular structure of (I), with displacement ellipsoids drawn at the 50% probability level for non-H atoms.

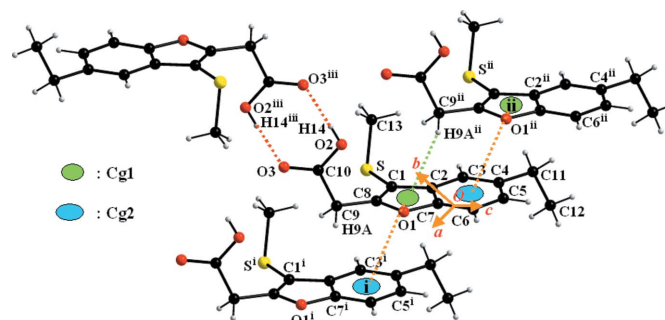


Figure 2

Hydrogen-bond and π - π interactions (dotted lines) in (I). [Symmetry codes: (i) $1 + x, y, z$; (ii) $x - 1, y, z$; (iii) $2 - x, 2 - y, 1 - z$.]

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