## Acta Crystallographica Section E

## Structure Reports

Online
ISSN 1600-5368

## 2-[(Methoxycarbonothioyl)sulfanyl]acetic acid

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Received 8 December 2010; accepted 31 January 2011
Key indicators: single-crystal X-ray study; $T=150 \mathrm{~K}$; mean $\sigma(\mathrm{C}-\mathrm{C})=0.002 \AA$;
$R$ factor $=0.021 ; w R$ factor $=0.056 ;$ data-to-parameter ratio $=20.5$.

The title compound, $\mathrm{C}_{4} \mathrm{H}_{6} \mathrm{O}_{3} \mathrm{~S}_{2}$, features a characteristic xanthate group; the $\mathrm{C}=\mathrm{S}$ double bond is shorter than the $\mathrm{C}-\mathrm{S}$ single bond, and the methyl group is coplanar with the xanthate group. In the crystal pairs of molecules form dimers through intermolecular $\mathrm{O}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonding.

## Related literature

For a related structure, see: Xiao \& Charpentier (2010). For the design and applications of the title compound, see: Moad et al. (2005, 2008); Stenzel et al. (2003); Coote \& Radom (2004); Coote et al. (2006).


## Experimental

## Crystal data

$\mathrm{C}_{4} \mathrm{H}_{6} \mathrm{O}_{3} \mathrm{~S}_{2}$
$M_{r}=166.21$
Monoclinic, $P 2_{1} / c$
$a=7.1009$ (3) А
$b=10.6485(5) \AA$
$c=9.2022$ (4) $\AA$
$\beta=93.370(1)^{\circ}$

## Data collection

## Bruker APEXII CCD

diffractometer
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
$T_{\text {min }}=0.931, T_{\text {max }}=0.963$

## Refinement

$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.021 \quad 84$ parameters
$w R\left(F^{2}\right)=0.056 \quad \mathrm{H}$-atom parameters constrained
$S=1.05$
1723 reflections

33976 measured reflections 1723 independent reflections 1517 reflections with $I>2 \sigma(I)$ $R_{\text {int }}=0.038$

Table 1
Hydrogen-bond geometry ( $\AA,^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :---: | :---: | :--- | :--- |
| $\mathrm{O} 2-\mathrm{H} 2 \cdots \mathrm{O}^{\mathrm{i}}$ | 0.84 | 1.82 | $2.6540(12)$ | 175 |
| Symmetry code: (i) $-x+1,-y,-z+2$. |  |  |  |  |

Data collection: APEX2 (Bruker, 2009); cell refinement: SAINT (Bruker, 2009); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

This work was supported by the Canadian Natural Sciences and Engineering Research Council (NSERC) Idea to Innovation (I2I) Program. The authors are grateful to Dr Guerman Popov of the Department of Chemistry, the University of Western Ontario, for the XRD data acquisition and interpretation.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: NG5085).

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## supporting information

Acta Cryst. (2011). E67, o575 [doi:10.1107/S1600536811003941]

## 2-[(Methoxycarbonothioyl)sulfanyl]acetic acid

## Shude Xiao and Paul A. Charpentier

## S1. Comment

Carbonothioylthio ( $\mathrm{S}=\mathrm{C}-\mathrm{S}$ ) compounds are used as chain transfer agents (CTA) in addition-fragmentation chain-transfer (RAFT) polymerization. In the addition-fragmentation equilibria, addition of the propagating radicals to the $\mathrm{S}=\mathrm{C}$ group followed by fragmentation of the intermediate radical at the C - S bond generates a new radical and a polymeric carbonothioylthio compound (Moad et al., 2005, 2008). $O$-alkyl xanthates show low reactivity in RAFT equilibria due to the conjugation of the $O$ lone pair electrons and the $\mathrm{C}=\mathrm{S}$ bond which is favorable to the zwitterionic canonical forms of xanthates (Moad et al., 2005; Coote et al., 2006). However, xanthates can promote fragmentation of unstable radicals, such as vinyl acetate radicals that undergo fast addition and slow fragmentation (Coote et al., 2006). Though studies have been done on RAFT polymerization of vinyl acetate with methyl 2-(methoxycarbonothioylthio)acetate (Stenzel et al., 2003; Coote \& Radom, 2004), 2-(methoxycarbonothioylthio)acetic acid has not been used in RAFT polymerization. Therefore, efforts were made to use 2-(methoxycarbonothioylthio)acetic acid as the CTA in RAFT polymerization, and poly(vinyl acetate)s containing carboxylic acid end groups were successfully prepared. A similar compound, 2-(isopropoxycarbonothioylthio)acetic acid, has been reported for the same application (Xiao \& Charpentier, 2010).

## S2. Experimental

Potassium hydroxide $5.6 \mathrm{~g}(50 \mathrm{mmol})$ was dissolved in methanol 30 ml at room temperature. The solution was cooled with an ice bath when carbon disulfide 20 ml was charged into the flask dropwise. After 1 day reaction at room temperature, a solution of 2-bromoacetic acid $6.9 \mathrm{~g}(50 \mathrm{mmol}) /$ methanol 20 ml was added into the flask dropwise in an ice bath. The precipitates were removed by filtration after 2 days reaction at room temperature, and the solvent was evaporated with a rotary evaporator. The crude product was run through a silica gel column with a mixture of ethyl ether / hexanes (5:1). Colorless crystals were obtained from crystalization in hexanes/ cyclohexane (4:1). m.p.: $112.6^{\circ} \mathrm{C}$ (DSC). MS: 165.9764.

## S3. Refinement

The structure was solved and refined using the Bruker SHELXTL Software Package, using the space group P 1 21/c 1, with $Z=4$ for the formula unit, $\mathrm{C}_{4} \mathrm{H}_{6} \mathrm{O}_{3} \mathrm{~S}_{2}$. All of the non-hydrogen atoms were refined with anisotropic thermal parameters. The hydrogen atom positions were calculated geometrically and were included as riding on their respective carbon/oxygen atoms. The final anisotropic full-matrix least-squares refinement on $\mathrm{F}^{2}$ with 84 variables converged at R 1 $=2.13 \%$, for the observed data and wR2 $=5.55 \%$ for all data. The goodness-of-fit was 1.047. The largest peak in the final difference electron density synthesis was $0.288 \mathrm{e}^{-} / \AA^{3}$ and the largest hole was $-0.195 \mathrm{e}^{-} / \AA^{3}$ with an RMS deviation of $0.040 \mathrm{e}^{-} / \AA^{3}$. On the basis of the final model, the calculated density was $1.589 \mathrm{~g} / \mathrm{cm}^{3}$ and $\mathrm{F}(000), 344 \mathrm{e}^{-}$.


Figure 1
View of the title compound ( $50 \%$ probability displacement ellipsoids).


Figure 2
Packing diagram of the structure with H -bonds.

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## Crystal data

$\mathrm{C}_{4} \mathrm{H}_{6} \mathrm{O}_{3} \mathrm{~S}_{2}$

$$
M_{r}=166.21
$$

$$
\begin{aligned}
& a=7.1009(3) \AA \\
& b=10.6485(5) \AA \\
& c=9.2022(4) \AA \\
& \beta=93.370(1)^{\circ}
\end{aligned}
$$

Monoclinic, $P 2_{1} / c$
Hall symbol: - P 2 ybc
$V=694.61(5) \AA^{3}$
$Z=4$
$F(000)=344$
$D_{\mathrm{x}}=1.589 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation, $\lambda=0.71073 \AA$
Cell parameters from 9941 reflections

## Data collection

## Bruker APEXII CCD <br> diffractometer

Radiation source: fine-focus sealed tube
Graphite monochromator
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
$T_{\text {min }}=0.931, T_{\text {max }}=0.963$

$$
\begin{aligned}
& \theta=2.9-30.2^{\circ} \\
& \mu=0.70 \mathrm{~mm}^{-1} \\
& T=150 \mathrm{~K} \\
& \text { Block, colourless } \\
& 0.10 \times 0.07 \times 0.06 \mathrm{~mm} \\
& \\
& \\
& 33976 \text { measured reflections } \\
& 1723 \text { independent reflections } \\
& 1517 \text { reflections with } I>2 \sigma(I) \\
& R_{\text {int }}=0.038 \\
& \theta_{\max }=28.3^{\circ}, \theta_{\min }=2.9^{\circ} \\
& h=-9 \rightarrow 9 \\
& k=-14 \rightarrow 13 \\
& l=-12 \rightarrow 12
\end{aligned}
$$

## Refinement

Refinement on $F^{2}$
Least-squares matrix: full
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.021$
$w R\left(F^{2}\right)=0.056$
$S=1.05$
1723 reflections
84 parameters
0 restraints
Primary atom site location: structure-invariant direct methods

> Secondary atom site location: difference Fourier map
> Hydrogen site location: inferred from neighbouring sites
> H -atom parameters constrained
> $w=1 /\left[\sigma^{2}\left(F_{0}{ }^{2}\right)+(0.0234 P)^{2}+0.2541 P\right]$
> where $P=\left(F_{\mathrm{o}}{ }^{2}+2 F_{\mathrm{c}}{ }^{2}\right) / 3$
> $(\Delta / \sigma)_{\max }=0.001$
> $\Delta \rho_{\text {max }}=0.29$ e $\AA^{-3}$
> $\Delta \rho_{\text {min }}=-0.20$ e $\AA^{-3}$

## Special details

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 $w R$ 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 $F^{2}>\sigma\left(F^{2}\right)$ 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 ( $A^{2}$ )

|  | $x$ | $y$ | $z$ | $U_{\mathrm{iso}} * / U_{\mathrm{eq}}$ |
| :--- | :--- | :--- | :--- | :--- |
| S1 | $0.15918(5)$ | $0.31498(3)$ | $0.78725(4)$ | $0.02980(9)$ |
| S2 | $-0.11784(5)$ | $0.12843(3)$ | $0.90015(4)$ | $0.03122(9)$ |
| O1 | $-0.10815(13)$ | $0.37609(8)$ | $0.93394(10)$ | $0.0311(2)$ |
| O2 | $0.41851(14)$ | $-0.01297(9)$ | $0.81022(10)$ | $0.0319(2)$ |
| H2 | 0.4754 | -0.0511 | 0.8797 | $0.048^{*}$ |
| O3 | $0.38409(12)$ | $0.13435(8)$ | $0.98001(9)$ | $0.02570(19)$ |
| C1 | $-0.2814(2)$ | $0.36782(14)$ | $1.00825(15)$ | $0.0354(3)$ |
| H1A | -0.3839 | 0.3404 | 0.9395 | $0.053^{*}$ |
| H1B | -0.3120 | 0.4504 | 1.0476 | $0.053^{*}$ |
| H1C | -0.2659 | 0.3070 | 1.0880 | $0.053^{*}$ |
| C2 | $-0.03785(16)$ | $0.27021(11)$ | $0.88181(13)$ | $0.0233(2)$ |


|  |  |  |  |  |
| :--- | :--- | :--- | :--- | :--- |
| C3 | $0.25018(18)$ | $0.16615(12)$ | $0.73538(13)$ | $0.0279(3)$ |
| H3A | 0.1438 | 0.1134 | 0.6969 | $0.034^{*}$ |
| H3B | 0.3354 | 0.1794 | 0.6554 | $0.034^{*}$ |
| C4 | $0.35645(16)$ | $0.09552(11)$ | $0.85644(13)$ | $0.0224(2)$ |

Atomic displacement parameters $\left(\hat{A}^{2}\right)$

|  | $U^{11}$ | $U^{22}$ | $U^{33}$ | $U^{12}$ | $U^{13}$ | $U^{23}$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| S1 | $0.03072(17)$ | $0.02100(15)$ | $0.03800(18)$ | $0.00050(12)$ | $0.00457(13)$ | $0.00871(12)$ |
| S2 | $0.03306(17)$ | $0.02151(16)$ | $0.03927(19)$ | $-0.00432(12)$ | $0.00377(13)$ | $0.00243(12)$ |
| O1 | $0.0293(5)$ | $0.0230(4)$ | $0.0407(5)$ | $0.0011(3)$ | $0.0005(4)$ | $-0.0062(4)$ |
| O2 | $0.0404(5)$ | $0.0291(5)$ | $0.0252(4)$ | $0.0120(4)$ | $-0.0050(4)$ | $-0.0049(4)$ |
| O3 | $0.0264(4)$ | $0.0255(4)$ | $0.0248(4)$ | $0.0046(3)$ | $-0.0011(3)$ | $-0.0034(3)$ |
| C1 | $0.0337(7)$ | $0.0385(7)$ | $0.0339(7)$ | $0.0061(6)$ | $0.0026(6)$ | $-0.0058(6)$ |
| C2 | $0.0245(6)$ | $0.0223(6)$ | $0.0224(5)$ | $0.0007(4)$ | $-0.0060(4)$ | $0.0009(4)$ |
| C3 | $0.0317(6)$ | $0.0281(6)$ | $0.0242(6)$ | $0.0041(5)$ | $0.0040(5)$ | $0.0041(5)$ |
| C4 | $0.0195(5)$ | $0.0226(5)$ | $0.0254(6)$ | $-0.0001(4)$ | $0.0040(4)$ | $0.0005(4)$ |

Geometric parameters ( $A,{ }^{\circ}$ )

| S1-C2 | 1.7564 (13) | O3-C4 | 1.2150 (14) |
| :---: | :---: | :---: | :---: |
| S1-C3 | 1.7870 (13) | $\mathrm{C} 1-\mathrm{H} 1 \mathrm{~A}$ | 0.9800 |
| S2-C2 | 1.6253 (12) | C1-H1B | 0.9800 |
| $\mathrm{O} 1-\mathrm{C} 2$ | 1.3336 (15) | $\mathrm{C} 1-\mathrm{H} 1 \mathrm{C}$ | 0.9800 |
| $\mathrm{O} 1-\mathrm{C} 1$ | 1.4451 (17) | C3-C4 | 1.5091 (16) |
| O2-C4 | 1.3159 (14) | C3-H3A | 0.9900 |
| $\mathrm{O} 2-\mathrm{H} 2$ | 0.8400 | C3-H3B | 0.9900 |
| C2-S1-C3 | 101.69 (6) | S2-C2-S1 | 126.65 (7) |
| $\mathrm{C} 2-\mathrm{O} 1-\mathrm{C} 1$ | 117.76 (10) | C4-C3-S1 | 114.70 (9) |
| C4-O2-H2 | 109.5 | C4-C3-H3A | 108.6 |
| $\mathrm{O} 1-\mathrm{C} 1-\mathrm{H} 1 \mathrm{~A}$ | 109.5 | S1-C3-H3A | 108.6 |
| $\mathrm{O} 1-\mathrm{C} 1-\mathrm{H} 1 \mathrm{~B}$ | 109.5 | C4-C3-H3B | 108.6 |
| $\mathrm{H} 1 \mathrm{~A}-\mathrm{C} 1-\mathrm{H} 1 \mathrm{~B}$ | 109.5 | S1-C3-H3B | 108.6 |
| $\mathrm{O} 1-\mathrm{C} 1-\mathrm{H} 1 \mathrm{C}$ | 109.5 | H3A-C3-H3B | 107.6 |
| $\mathrm{H} 1 \mathrm{~A}-\mathrm{C} 1-\mathrm{H} 1 \mathrm{C}$ | 109.5 | $\mathrm{O} 3-\mathrm{C} 4-\mathrm{O} 2$ | 124.23 (11) |
| H1B-C1-H1C | 109.5 | $\mathrm{O} 3-\mathrm{C} 4-\mathrm{C} 3$ | 124.58 (11) |
| $\mathrm{O} 1-\mathrm{C} 2-\mathrm{S} 2$ | 127.40 (10) | $\mathrm{O} 2-\mathrm{C} 4-\mathrm{C} 3$ | 111.18 (10) |
| $\mathrm{O} 1-\mathrm{C} 2-\mathrm{S} 1$ | 105.94 (8) |  |  |

Hydrogen-bond geometry ( $A,{ }^{\circ}$ )

| $D — \mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{O} 2 — \mathrm{H} 2 \cdots \mathrm{O} 3^{\mathrm{i}}$ | 0.84 | 1.82 | $2.6540(12)$ | 175 |

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

