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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=273 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.007 \AA$
$R$ factor $=0.054$
$w R$ factor $=0.125$
Data-to-parameter ratio $=14.4$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## Diaquabis[(pyrimidin-2-yIsulfanyl)acetato]copper(II) dihydrate

In the title compound, $\left[\mathrm{Cu}\left(\mathrm{C}_{6} \mathrm{H}_{5} \mathrm{~N}_{2} \mathrm{O}_{2} \mathrm{~S}\right)_{2}\left(\mathrm{H}_{2} \mathrm{O}\right)_{2}\right] \cdot 2 \mathrm{H}_{2} \mathrm{O}$, the Cu atom lies on an inversion centre in a distorted square coordination geometry that consists of two O atoms of two (pyrimidin-2-ylsulfanyl)acetate ligands $[\mathrm{Cu}-\mathrm{O}=1.953$ (2) $\AA$ A $]$ and two O atoms of two water molecules $[\mathrm{Cu}-\mathrm{O}=$ 1.942 (3) $\AA]$.

## Comment

Following our studies of complexes of 2-pyrimidylthioacetic acid (Ng et al., 1993; Ma et al., 2004; Hao et al., 2005), we report the structure of the centrosymmetric title compound, (I). The four-coordinate Cu atom is in a square coordination geometry that is made up of two O atoms of two carboxylate groups and two O atoms of two water molecules (Fig. 1). Hydrogen bonds connect the molecules and the solvent water molecules into a three-dimensional network structure.

(I)

## Experimental

Cupric nitrate ( $120.8 \mathrm{mg}, 0.5 \mathrm{mmol}$ ) was dissolved in water ( 10 ml ). Aqueous ammonium hydroxide ( $6 M$ ) was added until the solution turned blue. 2-Pyrimidylthioacetic acid ( $170.2 \mathrm{mg}, 1 \mathrm{mmol}$ ) was suspended in a small volume of water-ethanol ( $1: 1 \mathrm{v} / \mathrm{v}$ ); aqueous ammonium hydroxide ( $6 M$ ) was added until the compound dissolved completely. The two solutions were then mixed. After three weeks, dark-blue crystals were obtained in $65 \%$ yield. Analysis found: C 30.41, H 3.83, N $11.82 \%$; calculated for $\mathrm{C}_{12} \mathrm{H}_{18} \mathrm{CuN}_{4} \mathrm{O}_{8} \mathrm{~S}_{2}$ : C 30.20, H $3.91, \mathrm{~N} 11.78 \%$. IR $\left(\mathrm{cm}^{-1}\right)$ : $3423\left(v_{\mathrm{OH}}\right.$ for $\left.\mathrm{H}_{2} \mathrm{O}\right), 1605,1384\left(v_{\text {as }}\right.$ and $v_{\mathrm{s}}$ for $\mathrm{COO}^{-}$), 1551, 1309, 1280.


Figure 1
ORTEPII (Johnson, 1976) plot of (I). Displacement ellipsoids are drawn at the $30 \%$ probability level. [Symmetry code: (i) $1-x, 1-y, 1-z$.]

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- $2 \mathrm{H}_{2} \mathrm{O}$


## Crystal data

$\left[\mathrm{Cu}\left(\mathrm{C}_{6} \mathrm{H}_{5} \mathrm{~N}_{2} \mathrm{O}_{2} \mathrm{~S}\right)_{2}\left(\mathrm{H}_{2} \mathrm{O}\right)_{2}\right] \cdot 2 \mathrm{H}_{2} \mathrm{O}$
$M_{r}=473.96$
Monoclinic, $P 2_{1} / c$
$a=17.160$ (5) А
$b=5.1577$ (16) A
$c=10.568$ (3) $\AA$
$\beta=99.942(5)^{\circ}$
$V=921.3(5) \AA^{3}$
$Z=2$
$D_{x}=1.709 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
Cell parameters from 3246 reflections
$\theta=3.6-26.9^{\circ}$
$\mu=1.46 \mathrm{~mm}^{-1}$
$T=273$ (2) K
Plate, blue
$0.27 \times 0.16 \times 0.07 \mathrm{~mm}$

## Data collection

Bruker APEX 2000 area-detector diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
$T_{\text {min }}=0.694, T_{\text {max }}=0.905$
9709 measured reflections

## Refinement

Refinement on $F^{2}$

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.0443 P)^{2}\right. \\
& \quad+1.6524 P] \\
& \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
& (\Delta / \sigma)_{\max }=0.00 \\
& \Delta \rho_{\max }=0.54 \mathrm{e} \AA^{-3} \\
& \Delta \rho_{\min }=-0.66 \mathrm{e}^{-3}
\end{aligned}
$$

Table 1
Selected geometric parameters ( $\left(\AA,{ }^{\circ}\right)$.

| $\mathrm{Cu} 1-\mathrm{O} 1 W$ | $1.942(3)$ | $\mathrm{O} 2-\mathrm{C} 6$ | $1.252(4)$ |
| :--- | :--- | :--- | :--- |
| $\mathrm{Cu} 1-\mathrm{O} 2$ | $1.953(2)$ | $\mathrm{O} 1-\mathrm{C} 6$ | $1.202(4)$ |
|  |  |  |  |
| $\mathrm{O} 1 W-\mathrm{Cu} 1-\mathrm{O} 2$ | $89.34(12)$ | $\mathrm{O} 1-\mathrm{C} 6-\mathrm{O} 2$ | $124.4(3)$ |
| $\mathrm{O} 1 W^{\mathrm{i}}-\mathrm{Cu} 1-\mathrm{O} 2$ | $90.66(12)$ |  |  |
| Symmetry code: $(\mathrm{i})-x+1,-y+1,-z+1$ |  |  |  |

Table 2
Hydrogen-bond geometry $\left(\AA,^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{O} 2 W-\mathrm{H} 2 W A \cdots \mathrm{O} 1$ | $0.85(1)$ | $1.86(2)$ | $2.695(4)$ | $169(5)$ |

The water H atoms were located in difference Fourier maps and were refined with distance restraints of $\mathrm{O}-\mathrm{H}=0.85$ (1) $\AA$ and $\mathrm{H} \cdots \mathrm{H}=1.39(1) \AA\left[U_{\text {iso }}(\mathrm{H})=0.85 \AA^{2}\right]$. The aromatic and aliphatic H atoms were placed at calculated positions ( $\mathrm{C}-\mathrm{H}=0.93$ and $0.97 \AA$ ) and refined using the riding-model approximation, with $U_{\mathrm{iso}}(\mathrm{H})=$ $1.2 U_{\text {eq }}(\mathrm{C})$.

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPII (Johnson, 1976); software used to prepare material for publication: SHELXL97.

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