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#### Key indicators

Single-crystal X-ray study T = 153 K Mean  $\sigma$ (Cu–S) = 0.001 Å R factor = 0.023 wR factor = 0.054 Data-to-parameter ratio = 17.7

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

# $RbHo_2Cu_3S_5$

Rubidium diholmium tricopper pentasulfide, RbHo<sub>2</sub>Cu<sub>3</sub>S<sub>5</sub>, crystallizes in the orthorhombic space group *Cmcm* and is isostructural with RbSm<sub>2</sub>Ag<sub>3</sub>Se<sub>5</sub>. In the asymmetric unit, the site symmetries of atoms Rb, Cu1, and S2 are *mm* and those of the other atoms are *m*. The structure has a three-dimensional tunnel framework, with tunnels built from HoS<sub>6</sub> octahedra and CuS<sub>4</sub> tetrahedra. The tunnels are filled with Rb atoms.

### Comment

Ternary and quaternary rare-earth chalcogenides containing a combination of d- and f-elements have been reviewed recently (Mitchell & Ibers, 2002). We report here the structure of RbHo<sub>2</sub>Cu<sub>3</sub>S<sub>5</sub>, a new member of this large family. RbHo<sub>2</sub>Cu<sub>3</sub>S<sub>5</sub>, which has the RbSm<sub>2</sub>Ag<sub>3</sub>Se<sub>5</sub> structure type (Huang & Ibers, 2000), crystallizes in space group Cmcm of the orthorhombic system. In the asymmetric unit (Fig. 1), the site symmetries of atoms Rb, Cu1, and S2 are mm and those of the other atoms are m. The structure of RbHo<sub>2</sub>Cu<sub>3</sub>S<sub>5</sub> (Fig. 2) possesses a threedimensional tunnel framework built from HoS<sub>6</sub> octahedra and CuS<sub>4</sub> tetrahedra. The tunnel, comprising ten-membered rings of six Cu-S bonds and four Ho-S bonds, is only large enough in cross-section to accommodate one Rb atom. Each Rb atom is surrounded by a bicapped trigonal prism of eight S atoms, with Rb-S separations ranging from 3.293 (1) to 3.460(1) Å, comparable with those of 3.247(2)-3.7951(4) Å in RbNd<sub>2</sub>CuS<sub>4</sub> (Huang & Ibers, 2000). The Ho-S bond distances range from 2.6497 (8) to 2.787 (1) Å, consistent with those of 2.672 (2)–2.8009 (3) Å in  $K_2Ho_4Cu_4S_9$  (Yao *et al.*, 2003), and the Cu–S bond distances range from 2.321 (1) to 2.547 (2) Å, comparable with those of 2.3448 (9) to 2.534 (2) Å in K<sub>2</sub>Ho<sub>4</sub>Cu<sub>4</sub>S<sub>9</sub> (Yao *et al.*, 2003).



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# Experimental

RbHo<sub>2</sub>Cu<sub>3</sub>S<sub>5</sub> was obtained as yellow needles from a solid-state reaction of Rb<sub>2</sub>S<sub>3</sub> (1.2 mmol), Ho (Aldrich, 99%, 1.0 mmol), Cu (Aldrich, 99.999%, 0.5 mmol), and S (Aldrich, 99.5%, 2.0 mmol). The Rb<sub>2</sub>S<sub>3</sub> reactive flux (Sunshine *et al.*, 1987) was prepared by the stoichiometric reaction of Rb (Aldrich, 98+%) and S in liquid NH<sub>3</sub>. The reactants were loaded into a fused-silica tube under an argon atmosphere in a glove-box. The tube was sealed under a  $10^{-4}$  Torr atmosphere and then placed in a computer-controlled furnace. The sample was heated to 1173 K over a period of 25 h, kept at 1173 K for 3 d, slowly cooled at a rate of 10 K h<sup>-1</sup> to 473 K, and then cooled rapidly to room temperature.

## Crystal data

RbHo<sub>2</sub>Cu<sub>3</sub>S<sub>5</sub>  $M_r = 766.25$ Orthorhombic, *Cmcm*  a = 3.9451 (11) Å b = 13.915 (4) Å c = 16.408 (5) Å  $V = 900.8 (4) Å^3$  Z = 4  $D_x = 5.650 \text{ Mg m}^{-3}$  *Data collection* Bruker SMART 1000 CCD diffractometer  $\omega$  scans Absorption correction: numerical face indexed

 $T_{\min} = 0.047, T_{\max} = 0.405$ 5433 measured reflections

## Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.023$   $wR(F^2) = 0.054$  S = 1.39673 reflections 38 parameters Cell parameters from 4507 reflections  $\theta = 2.5-29.0^{\circ}$  $\mu = 30.77 \text{ mm}^{-1}$ T = 153 (2) KNeedle, yellow  $0.42 \times 0.068 \times 0.030 \text{ mm}$ 

Mo  $K\alpha$  radiation

673 independent reflections 661 reflections with  $I > 2\sigma(I)$   $R_{int} = 0.050$   $\theta_{max} = 29.0^{\circ}$   $h = -5 \rightarrow 5$   $k = -18 \rightarrow 18$  $l = -21 \rightarrow 21$ 

$$\begin{split} &w = 1/[\sigma^2(F_o^2) + (0.03P)^2] \\ &where \ P = (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{max} = 0.001 \\ \Delta\rho_{max} = 2.37 \ e \ \text{\AA}^{-3} \\ \Delta\rho_{min} = -2.55 \ e \ \text{\AA}^{-3} \\ &\text{Extinction correction: SHELXL97} \\ &\text{Extinction coefficient: 0.00404 (19)} \end{split}$$

# Table 1

Selected geometric parameters (Å, °).

Rb-S3	3.2929 (15)	Ho-S3 <sup>vi</sup>	2.7865 (10)
Rb-S2 <sup>i</sup>	3.3410 (15)	Cu1-S2 <sup>vii</sup>	2.3212 (10)
Rb-S1 <sup>ii</sup>	3.4604 (12)	Cu1-S1 <sup>iv</sup>	2.5438 (14)
Ho-S2 <sup>iii</sup>	2.6497 (8)	Cu2-S1 <sup>viii</sup>	2.3641 (14)
Ho-S3 <sup>iv</sup>	2.6957 (14)	Cu2-S3 <sup>v</sup>	2.3674 (8)
Ho-S1 <sup>v</sup>	2.6995 (9)	Cu2-S1 <sup>iv</sup>	2.5473 (15)
S2 <sup>iii</sup> -Ho-S3 <sup>iv</sup>	172.13 (4)	S3 <sup>vi</sup> -Ho-S3 <sup>v</sup>	90.13 (4)
S2 <sup>iii</sup> -Ho-S1 <sup>v</sup>	92.47 (3)	$S2^{vii}$ -Cu1-S2 <sup>i</sup>	116.38 (7)
S3 <sup>iv</sup> -Ho-S1 <sup>v</sup>	92.89 (3)	S2 <sup>i</sup> -Cu1-S1 <sup>iv</sup>	105.05 (2)
S1 <sup>v</sup> -Ho-S1 <sup>vi</sup>	93.89 (4)	S1 <sup>iv</sup> -Cu1-S1	120.96 (6)
S2 <sup>iii</sup> -Ho-S3 <sup>vi</sup>	87.95 (4)	S1 <sup>viii</sup> -Cu2-S3 <sup>v</sup>	108.94 (4)
S3 <sup>iv</sup> -Ho-S3 <sup>vi</sup>	86.50 (3)	S3 <sup>v</sup> -Cu2-S3 <sup>vi</sup>	112.86 (5)
S1 <sup>v</sup> -Ho-S3 <sup>vi</sup>	178.05 (3)	S1 <sup>viii</sup> -Cu2-S1 <sup>iv</sup>	115.43 (4)
$S1^{vi}$ -Ho- $S3^{vi}$	87.99 (3)	S3 <sup>v</sup> -Cu2-S1 <sup>iv</sup>	105.36 (4)

Symmetry codes: (i)  $x - \frac{1}{2}, y - \frac{1}{2}, z$ ; (ii)  $\frac{1}{2} + x, \frac{1}{2} + y, \frac{1}{2} - z$ ; (iii) -x, 1 - y, 1 - z; (iv)  $x, y, \frac{1}{2} - z$ ; (v)  $\frac{1}{2} - x, \frac{1}{2} - y, \frac{1}{2} + z$ ; (vi)  $-\frac{1}{2} - x, \frac{1}{2} - y, \frac{1}{2} + z$ ; (vii)  $\frac{1}{2} + x, y - \frac{1}{2}, z$ ; (viii)  $-x, -y, \frac{1}{2} + z$ .



The structure of  $RbHo_2Cu_3S_5$ , viewed down [100].

The highest residual electron density is 0.87 Å from the Ho site and the deepest hole is 0.70 Å from this same site.

Data collection: *SMART* (Bruker, 2003); cell refinement: *SAINT* (Bruker, 2003); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2003); program(s) used to refine structure: *SHELXTL*; molecular graphics: *XP* in *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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