

Octakis(μ_3 -*tert*-butylthiolato)bis(*tert*-butylthiolato)hexakis(1-methylpyrrolidin-2-one)- μ_6 -sulfido-hexabarium(II)

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Key indicators

Single-crystal X-ray study
 T = 296 K
 Mean $\sigma(\text{C}-\text{C}) = 0.017 \text{ \AA}$
 Disorder in main residue
 R factor = 0.036
 wR factor = 0.082
 Data-to-parameter ratio = 27.7

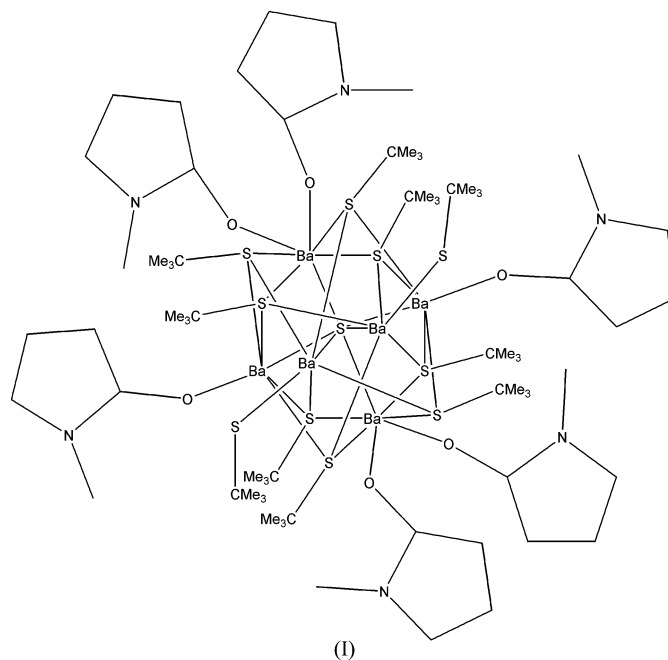
For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title compound, $[\text{Ba}_6(\text{C}_4\text{H}_9\text{S})_{10}\text{S}(\text{C}_5\text{H}_9\text{NO})_6]$, possessing twofold symmetry, contains a central μ_6 -sulfido atom surrounded by six Ba atoms in an almost perfect octahedron. Of the six Ba atoms, two are seven-coordinate with an S_5O_2 donor set, while four are six-coordinate. Two of these have S_5O and two have S_6 donor sets. There are six 1-methyl-2-pyrrolidone donor molecules, two each coordinated to two Ba atoms and one each coordinated to two other Ba atoms.

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Comment

There have been numerous examples of high nuclearity transition metal clusters encapsulating S (Allen, 2002). Such SM_6 clusters have been reported for Fe (Zhang *et al.*, 2002, 2003, 2004; Zuo *et al.*, 2003; Osterloh *et al.*, 1999; Ohki *et al.*, 2003), Cu (Wu, *et al.*, 1988; Marsh, 1997; Hu, 2001; Lin *et al.*, 2000a; Chen *et al.*, 2003; Tang *et al.*, 1993), Co (Shu *et al.*, 1991), Zr (Fenske *et al.*, 1991) and Ag (Chen *et al.*, 2004; Lin *et al.*, 1997; Matsumoto *et al.*, 2000; Lin *et al.*, 2000b; Jin *et al.*, 1996), while SM_8 clusters have been reported for Rh (Vidal *et al.*, 1978) and Cu (Lin *et al.*, 1999; Huang *et al.*, 1991; Dehnen & Fenske, 1996; Liu *et al.*, 1995; Fenske *et al.*, 2004). With the exception of one report of an SLi_8 cluster (Banister *et al.*, 1988), there has been no report of an SM_x cluster involving a non-transition metal.



The title compound, (I), crystallized out of a reaction medium containing barium *tert*-butylsulfide, 1-methyl-2-

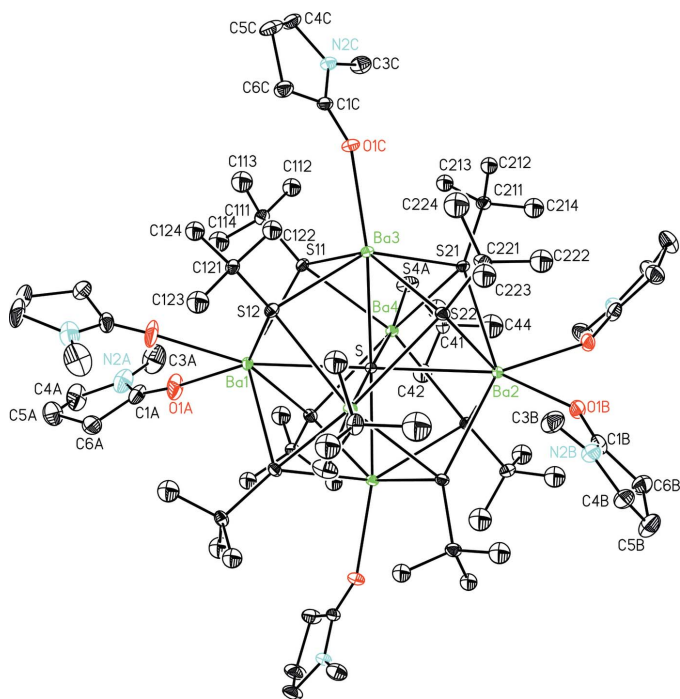


Figure 1
View of the title compound. Displacement ellipsoids are drawn at the 20% probability level. H atoms have been omitted for clarity.

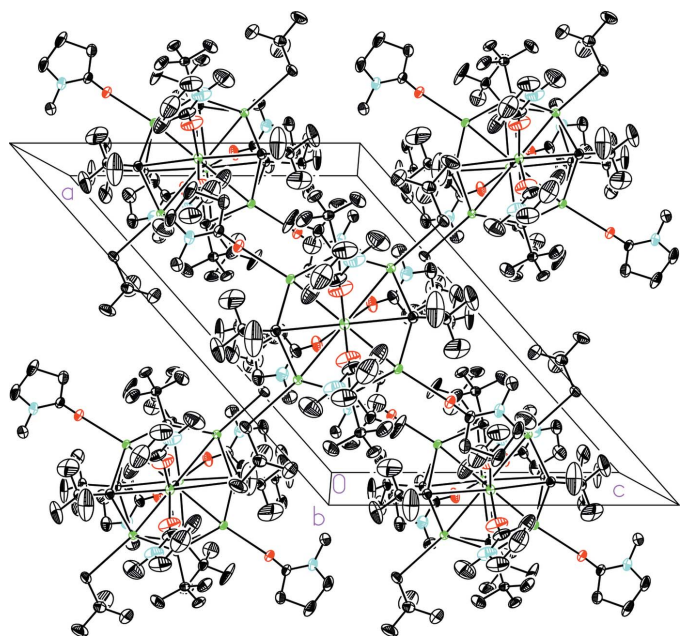


Figure 2
The molecular packing of the title compound viewed along the *b* axis. H atoms have been omitted for clarity.

pyrrolidone (NMP) and C_6D_6 over a period of 6.5 years. The molecule has crystallographically imposed twofold symmetry with Ba1, Ba2 and the central S on a twofold axis. The coordination spheres for both Ba1 and Ba2 are similar, both being seven-coordinate with an S_5O_2 donor set, while Ba3 and Ba4

are both six-coordinate with S_5O and S_6 donor sets, respectively. Of the ten *t*-butylsulfide ligands, eight contain triply bridging S donors, while two are terminally coordinated. There are six 1-methyl-2-pyrrolidone donor molecules coordinated to four of the six Ba atoms. For the central S atom, the six Ba—S distances range from 3.1011 (3) to 3.279 (4) Å with an average of 3.20 (7) Å. The Ba—S distances for the triply bridging S donors range from 3.1389 (18) to 3.2901 (15) Å with an average of 3.209 Å. The terminal Ba—S distance is 3.1258 (13) Å. For the 1-methyl-2-pyrrolidone ligands, the Ba—O distances range from 2.693 (4) to 2.732 (2) Å with an average of 2.707 (18) Å. The six Ba atoms and the central S form an almost perfect octahedron, the maximum deviation of the Ba—S—Ba angles being 0.65 (5).

Experimental

$Ba(SCMe_3)_2$ (Purdy *et al.*, 1997) was dissolved in 1-methyl-2-pyrrolidone containing about 10% C_6D_6 and a drop of trimethylsilane and sealed in an NMR tube. After 6.5 years, a mass of colorless crystals was discovered in the NMR tube. The crystals were sealed in thin-walled glass capillaries.

Crystal data

$[Ba_6(C_4H_9S)_{10}S(C_5H_9NO)_6]$
 $M_r = 2340.60$
 Monoclinic, $C2$
 $a = 21.0316$ (11) Å
 $b = 21.7686$ (11) Å
 $c = 15.2669$ (8) Å
 $\beta = 131.391$ (1)°
 $V = 5243.7$ (5) Å³
 $Z = 2$

$D_x = 1.482$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 7746 reflections
 $\theta = 2.6$ – 29.2 °
 $\mu = 2.48$ mm⁻¹
 $T = 296$ (2) K
 Chunk, colourless
 $0.4 \times 0.3 \times 0.2$ mm

Data collection

Bruker SMART 1K CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{min} = 0.498$, $T_{max} = 0.609$
 21162 measured reflections

11503 independent reflections
 9871 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.023$
 $\theta_{max} = 29.3$ °
 $h = -27 \rightarrow 25$
 $k = -29 \rightarrow 29$
 $l = -20 \rightarrow 20$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.082$
 $S = 1.05$
 11503 reflections
 416 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.029P)^2 + 14.7453P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} = 0.006$
 $\Delta\rho_{max} = 0.65$ e Å⁻³
 $\Delta\rho_{min} = -0.63$ e Å⁻³
 Absolute structure: Flack (1983), with 4126 Friedel pairs
 Flack parameter: 0.45 (2)

All H atoms were initially located in a difference Fourier map. The methyl H atoms were then constrained to an ideal geometry with C—H distances of 0.96 Å and $U_{iso}(H) = 1.5U_{eq}(C)$, but each group was allowed to rotate freely about its C—C bond. All other H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C—H distances in the range 0.96–0.97 Å and $U_{iso}(H) = 1.2U_{eq}(C)$. There are several disordered components in this structure: (a) S4 is disordered over two positions (S4A and S4B) with occupancies of 0.81 (2) and 0.19 (2); (b) the *tert*-butyl group attached to S11 is disordered over two conformations with occupancies of

0.64 (1) and 0.36 (1); (c) the *tert*-butyl group attached to S21 is disordered over two conformations with occupancies of 0.55 (2) and 0.45 (2). Both *tert*-butyl groups were restrained to be tetrahedral. The value of the Flack parameter (Flack, 1983) suggests inversion twinning.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1999); software used to prepare material for publication: *SHELXTL*.

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