

## Rearrangement of Symmetrical Dicubane Clusters into Topological Analogues of the P Cluster of Nitrogenase: Nature's Choice?

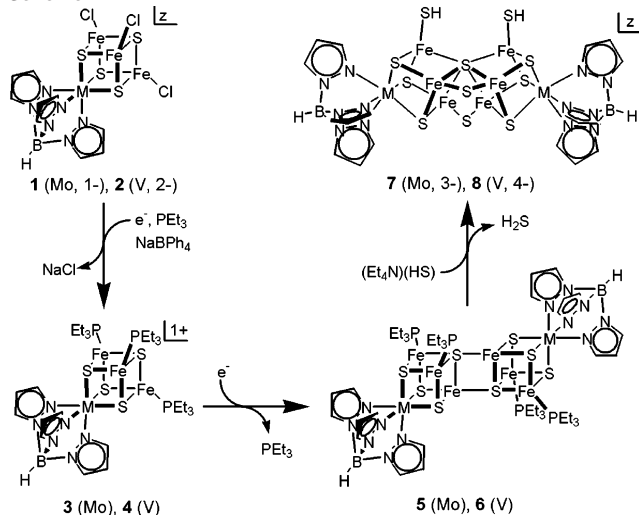
Yugen Zhang, Jing-Lin Zuo, Hong-Cai Zhou, and R. H. Holm\*

Department of Chemistry and Chemical Biology, Harvard University, Cambridge, Massachusetts 02138

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The electron-transfer P cluster ( $\text{Fe}_8\text{S}_7$ ) and the catalytic cofactor cluster ( $\text{MFe}_7\text{S}_9$ ;  $\text{M} = \text{Mo}, \text{V}$ ) of the MoFe protein of nitrogenase<sup>1–3</sup> have been assembled only by biosynthesis. Noting spectroscopic evidence that the protein as normally isolated contains substantially reduced clusters (majority of  $\text{Fe(II)}$ ),<sup>4–6</sup> we are investigating the reactivity of reduced  $\text{Fe–S}^7$  and  $\text{M–Fe–S}^{8–12}$  clusters, often with tertiary phosphine ligation as a device to stabilize low oxidation states. We describe here the use of reduced clusters in the synthesis of the initial molecular topological analogues of the P cluster in the crystallographically defined  $\text{P}^{\text{N}}$  state.<sup>13,14</sup> The synthetic procedure, outlined in Scheme 1, includes both molybdenum- and vanadium-containing clusters; heterometal  $\text{M}$  and charge  $z$  are specified. To ensure regioselective reactions and avoidance of potentially isomeric final products, tris(pyrazolyl)hydroborate(1–) ( $\text{Tp}$ ) at the heterometal site is a protecting group and in single cubanes a stipulator of trigonal symmetry. All structures have been proven by X-ray structure determinations.<sup>11,12,15</sup>

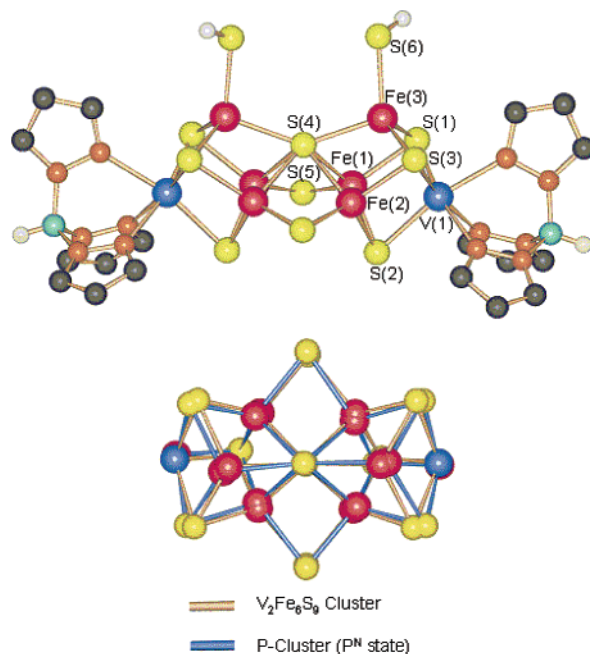
## Scheme 1



Treatment of  $[(\text{Tp})\text{MoFe}_3\text{S}_4\text{Cl}_3]^{1-}$  (**1**,  $[\text{MoFe}_3\text{S}_4]^{3+}$ ) with 5 equiv each of  $\text{PEt}_3$  and  $\text{NaBPh}_4$  for 12 h results in reductive substitution and formation of  $[(\text{Tp})\text{MoFe}_3\text{S}_4(\text{PEt}_3)_3]^{1+}$  (**3**,  $[\text{MoFe}_3\text{S}_4]^{2+}$ ), isolated as the  $\text{BPh}_4^-$  salt<sup>15</sup> (80%). Reduction of **3** with 1.3 equiv of  $(\text{Bu}_4\text{N})(\text{BH}_4)$  for 16 h gives the neutral edge-bridged double cubane  $[(\text{Tp})_2\text{Mo}_2\text{Fe}_6\text{S}_8(\text{PEt}_3)_4]$  (**5**, 70%). Its structure<sup>15</sup> resembles that of  $[(\text{Cl}_4\text{cat})_2(\text{Et}_3\text{P})_2\text{Mo}_2\text{Fe}_6\text{S}_8(\text{PEt}_3)_4]$ <sup>8,9</sup> (**9**) but is more reduced, being the first example of an isolated  $[\text{MoFe}_3\text{S}_4]^{1+}$  cluster. Reaction of a slurry of **5** with 4 equiv of  $(\text{Et}_3\text{N})(\text{HS})$  causes the solid to dissolve immediately with formation of  $[(\text{Tp})_2\text{Mo}_2\text{Fe}_6\text{S}_9(\text{SH})_2]^{3-}$

(**7**). Slow diffusion of ether into the reaction mixture leads to separation of  $(\text{Et}_3\text{N})_3[\text{7}] \cdot 9\text{MeCN}$  (73%). The reaction sequence  $2 \rightarrow 4 \rightarrow 6$  of vanadium-containing clusters has been demonstrated recently.<sup>12</sup> The double cubane  $[(\text{Tp})_2\text{V}_2\text{Fe}_6\text{S}_8(\text{PEt}_3)_4]$  (**6**) is isostructural with **5** and is also strongly reduced ( $2[\text{VFe}_3\text{S}_4]^{1+}$ ). It reacts similarly with 4 equiv of  $(\text{Et}_3\text{N})(\text{HS})$  to form  $[(\text{Tp})_2\text{V}_2\text{Fe}_6\text{S}_9(\text{SH})_2]^{4-}$  [**8**], isolated as  $(\text{Et}_3\text{N})_4[\text{8}] \cdot 6\text{MeCN}$  (70%).

Clusters **7** and **8** are isostructural and nearly isometric.<sup>15</sup> The structure of vanadium cluster **8**, which has crystallographically imposed  $\text{C}_2$  symmetry, is set out in Figure 1. The core exhibits the



**Figure 1.** (Upper) Structure of cluster **8** in which a crystallographic  $\text{C}_2$  axis passes through  $\mu_6\text{-S}(4)$  and the center of square  $\text{Fe}(1,1')(2,2')$  and relates primed and unprimed atoms. (Lower) Best-fit superposition of the  $\text{V}_2\text{Fe}_6\text{S}_9$  core of **8** and the  $\text{Fe}_8\text{S}_7(\mu_2\text{-S}_{\text{Cys}})_2$  core of the *Klebsiella pneumoniae*  $\text{P}^{\text{N}}$  cluster,<sup>14</sup> which in its entirety is  $\text{Fe}_8\text{S}_7(\mu_2\text{-S}_{\text{Cys}})_2(\text{S}_{\text{Cys}})_4$ . The bridging cysteine atoms are simulated by  $\text{S}(5,5')$  and two terminal cysteinates by  $\text{S}(6,6')$  of **8**.

bridging pattern  $[\text{V}_2\text{Fe}_6(\mu_2\text{-S})_2(\mu_3\text{-S})_6(\mu_6\text{-S})]$ , which reduces to two distorted cuboidal fragments  $\text{VFe}_3(\mu_3\text{-S})_3$  sharing the common bridge atom  $\mu_6\text{-S}(4)$  and externally bridged by  $\mu_2\text{-S}(5,5')$ . Each iron atom has distorted tetrahedral stereochemistry, with terminal hydrosulfide ligands at  $\text{Fe}(3,3')$  and trigonally distorted octahedral coordination at the vanadium atoms with parameters typical of  $\text{Tp}$  binding.<sup>11,12</sup> The most conspicuous individual feature of **7** and **8** is the infrequently encountered  $\mu_6\text{-S}$  atom, at which there are nine independent  $\text{Fe–S–Fe}$  angles. Three of these are notably obtuse,  $\text{Fe}(3)\text{–S}(4)\text{–Fe}(1',2',3') = 136\text{–}141^\circ$ , and convey the open nature

\* To whom correspondence should be addressed. E-mail: holm@chemistry.harvard.edu.

of the structure along the Fe(3,3')S(4) edge at which the angle is 140.8(1)°. In **7** the range is similar (135–141°), and the edge angle is 141.0(1)°. Other parameters in both structures occur in the intervals normally found for cubane-type Fe<sub>4</sub>S<sub>4</sub> and MFe<sub>3</sub>S<sub>4</sub> structures.<sup>7,11,12,16</sup> However, the most important result is that the core shapes of **7** and **8** are clearly similar to that of the P<sup>N</sup> cluster as determined in two different enzymes.<sup>13,14</sup> Shown in Figure 1 is a best-fit superposition of the 17 atoms of the V<sub>2</sub>Fe<sub>6</sub>S<sub>9</sub> core of **8** and the Fe<sub>8</sub>S<sub>7</sub>(μ<sub>2</sub>-S<sub>Cys</sub>)<sub>2</sub> core of the P<sup>N</sup> cluster, described by a weighted rms deviation of 0.33 Å. For the Mo<sub>2</sub>Fe<sub>6</sub>S<sub>9</sub> core of **7** and the P<sup>N</sup> cluster, the corresponding value is 0.38 Å. Notwithstanding the less precise protein data, obtained at resolutions of 2.0 and 1.6 Å for the proteins from *Azotobacter vinelandii*<sup>13</sup> and *Klebsiella pneumoniae*,<sup>14</sup> respectively, we conclude that *the synthetic and native cluster cores approach congruency*. Other than the presence of six-coordinate heterometals, a number of apparent differences contribute to the structural deviations. Among these is a much larger edge-bridge angle (158° in the P cluster) and its attendant effect on atom positions. Of lesser influence is the presence of two Fe–(μ<sub>2</sub>-S)–Fe bridges at the periphery of **7** and **8** (74.6–77.6°) instead of two Fe–(μ<sub>2</sub>-S<sub>Cys</sub>)–Fe bridges (73°, 76°) as in the protein.

The core structure of **7** has been observed before as sulfide-bridged fragments in the high-nuclearity clusters [(Cl<sub>4</sub>cat)<sub>6</sub>(Et<sub>3</sub>P)<sub>6</sub>Mo<sub>6</sub>Fe<sub>20</sub>S<sub>30</sub>]<sup>8–17</sup> and [(Cl<sub>4</sub>cat)<sub>2</sub>(Et<sub>3</sub>P)<sub>4</sub>Mo<sub>4</sub>Fe<sub>12</sub>S<sub>20</sub>K<sub>3</sub>(DMF)]<sup>5–10</sup>. These clusters were prepared by the reaction of edge-bridged **9** with (Et<sub>4</sub>N)(SH) under specific reaction and workup conditions. Their formation also illustrates the value of reduced double cubanes as precursors to new structures. However, because of the complicated and tight sulfide-bridged cluster structures, properties intrinsic to the fragments of interest, such as oxidation level, charge distribution, and redox states, cannot be deconvoluted from those of the whole clusters. *Molecular* clusters **7** and **8** provide that opportunity. Oxidation states of metal atoms in heterometal clusters are often difficult to assess. Terminal Fe–SR bond lengths increase as the oxidation level of Fe<sub>4</sub>S<sub>4</sub> clusters decreases. The Fe–SH distances in **7** (mean 2.291 Å) and **8** (2.327(3) Å), when compared with the values in [Fe<sub>4</sub>S<sub>4</sub>(SH)<sub>4</sub>]<sup>2–</sup> (mean 2.262 Å)<sup>18</sup> and [Fe<sub>4</sub>S<sub>4</sub>(SH)<sub>4</sub>]<sup>3–</sup> (2.317(2) Å),<sup>16</sup> are consistent with substantial Fe(II) character at these sites. The Mössbauer spectrum<sup>15</sup> of **8** at 4.2 K consists of two overlapping doublets fitted in an intensity ratio of 3:1. The majority/minority doublet has δ = 0.52/0.59 mm/s (mean 0.54 mm/s) and ΔE<sub>Q</sub> = 1.23/0.65 mm/s. The spectrum of **7** is a broadened quadrupole doublet with δ = 0.55 mm/s and ΔE<sub>Q</sub> = 0.62 mm/s. While comparison with isomer shifts of other reduced clusters<sup>7,9–12</sup> is not precise because of differences in structure and terminal ligands, we do note the values for [Fe<sub>4</sub>S<sub>4</sub>(SEt)<sub>4</sub>]<sup>3–</sup> (0.59 mm/s), double cubane {(Tp)<sub>2</sub>Mo<sub>2</sub>Fe<sub>6</sub>S<sub>8</sub>(SPh)<sub>4</sub>]<sup>4–</sup> (0.64 mm/s), and the P<sup>N</sup> cluster (~0.64 mm/s)<sup>4</sup> and conclude that **7** and **8** are substantially reduced clusters. In keeping with its reduced nature, **7** shows three quasireversible oxidations at E<sub>1/2</sub> = –1.09, –0.71, and –0.43 V; redox steps of **8** appear to be irreversible. Electronic structural properties of these clusters will be described subsequently.

The apparent means of formation of **7** and **8** can be summarized by the reaction [(Tp)<sub>2</sub>M<sub>2</sub>Fe<sub>6</sub>S<sub>8</sub>(PEt<sub>3</sub>)<sub>4</sub>] + 4HS<sup>–</sup> → [(Tp)<sub>2</sub>M<sub>2</sub>Fe<sub>6</sub>S<sub>9</sub>(SH)<sub>4</sub>]<sup>4–</sup> + H<sub>2</sub>S + 4PEt<sub>3</sub>. In the case of **7** (for which E<sub>1/2</sub> (4–/3–)

< –1.1 V), the cluster is isolated in a one-electron oxidized form; the oxidant is currently unknown. While clusters **5** and **6** have nonphysiological Tp ligands and a core structure yet to be found in a protein-bound condition, this reaction does raise a provocative question in biosynthesis. Can two closely juxtaposed Fe<sub>4</sub>S<sub>4</sub> clusters be induced to coalesce into the P<sup>N</sup> topology by reaction with hydrosulfide? Further, is the P cluster a precursor to the all-iron nitrogenase cofactor, and are one Fe<sub>4</sub>S<sub>4</sub> cluster and one MFe<sub>3</sub>S<sub>4</sub> cluster component building blocks to the FeMo- and FeV-cofactor clusters of nitrogenase? Last, we note the contributions of others in the synthesis of clusters relevant to nitrogenase clusters by use of reduced Mo–Fe–S double cubanes.<sup>19,20</sup>

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**Supporting Information Available:** Crystallographic data in CIF format of the compounds in footnote 15. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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- Experimental Details:* All reactions were carried out under strictly anaerobic conditions in acetonitrile; products were isolated as black, air-sensitive solids. X-ray diffraction data (θ = 1.2–22.5°) were collected at 213 K on a Siemens SMART CCD-based diffractometer; structures were solved and refined by standard methods. [2](BPh<sub>4</sub>): rhombohedral (R3); a = 15.4332(8) Å, c = 44.421(5) Å; Z = 6, R1(wR2) = 0.066(0.198). **5**: monoclinic (P2<sub>1</sub>/n); a = 11.2863(9) Å, b = 25.282(2) Å, c = 12.8320(9) Å; β = 112.743(1)°, Z = 4, R1(wR2) = 0.049(0.120). (Et<sub>4</sub>N)<sub>3</sub>[7]·9MeCN: monoclinic (P2<sub>1</sub>/n); a = 18.358(1) Å, b = 17.534(1) Å, c = 29.424(2) Å; β = 96.619(1)°, Z = 4, R1(wR2) = 0.066(0.151). (Et<sub>4</sub>N)<sub>4</sub>[8]·6MeCN: monoclinic (I2/a); a = 22.981(7) Å, b = 16.858(5) Å, c = 24.753(7) Å; β = 96.441(5)°, Z = 4, R1(wR2) = 0.074(0.160). The <sup>1</sup>H NMR spectra of **1–4** and **5–8** in CD<sub>3</sub>CN are consistent with three-fold and two-fold symmetry, respectively. Isomer shifts δ of <sup>57</sup>Fe are referenced to Fe metal at room temperature; all quoted values are at 4.2 K.
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