

Oxidative Addition of Thioesters to Iron(0): Active-Site Models for Hmd, Nature's Third Hydrogenase

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Summary: The thioester $\text{Ph}_2\text{PC}_6\text{H}_4\text{-2-C(O)SPh}$ reacts with $\text{Fe}_2(\text{CO})_9$ to give $[\text{Ph}_2\text{PC}_6\text{H}_4\text{C(O)}]\text{Fe}(\text{SPh})(\text{CO})_3$, a model for the CO-inhibited active site of the enzyme Hmd. This species, which reversibly decarbonylates to give a diiron derivative, reacts with cyanide to give $[[\text{Ph}_2\text{PC}_6\text{H}_4\text{C(O)}]\text{Fe}(\text{SPh})(\text{CN})(\text{CO})_2]^-$.

One of the great surprises in bioinorganic chemistry was the discovery of the iron carbonyl sites in the [NiFe]- and the [FeFe]-hydrogenases.¹ A third hydrogenase, Hmd (methylenetetrahydromethanopterin dehydrogenase), formerly thought to be “metal-free,” was recently shown to also contain an iron carbonyl center.² Because it features only a single Fe center at its active site, this enzyme is also called the [Fe]-hydrogenase. Hmd represents possibly the last hydrogenase to be discovered: it is strongly expressed only under special conditions (Ni deficiency) and only by methanogenic archaea. The fact that three genetically independent hydrogenases evolved similar catalytic motifs underscores the versatility of the Fe–S–CO compounds for reactions involving hydrogen.

Synthetic modeling of the active sites of the hydrogenases, in parallel with biophysical studies, provides mechanistic insights. Good progress is being made in the biomimicry of the active sites of the [NiFe]- and [FeFe]-hydrogenases.^{3,4} The active site of Hmd consists of a $\text{Fe}(\text{CO})_2$ center bound to a thiolate and the pyridine-like nitrogen of the guanylylpyridinol (GP) cofactor. This ensemble, the Fe-GP cofactor, is extractable from the protein via transthioation using mercaptoethanol. Because it is bound to the protein via this

single exchangeable residue, this active site is particularly ripe for modeling, and indeed models have already been described.⁵ Recently, however, the structural assignment of the active site has been significantly revised.⁶ The new analysis indicates that Fe is bound also to an acyl ligand, provided by the GP cofactor (Figure 1). Acyl ligands are rarely encountered in bioinorganic chemistry,⁷ and their coexistence with thiolato ligands defines a novel platform from the perspective of homogeneous catalysis. The new structural information—i.e. $\text{Fe}(\text{SR})(\text{N-donor})(\text{CO})_2(\text{acyl})$ —provides sufficient information to enable the design of a first-generation model for this active site, which is described below.

In terms of retrosynthetic analysis, the structure of the Fe center at the active site suggests that thioesters would oxidatively add to Fe(0) reagents. The interaction of thioesters with metal complexes has been intermittently investigated,⁸ including studies suggesting that this interaction is of prebiotic significance.⁹ The oxidative addition of a thioester has been established for rhodium(I) complexes.¹⁰ Our approach focused on the use of a donor-functionalized thioester, which upon oxidative addition would simultaneously deliver the Lewis base, thiolate, and acyl groups. To simplify the analysis of the synthetic studies, we chose to use a phosphine in place of the nitrogen heterocycle, since ³¹P NMR analysis provides a convenient means to monitor reactive intermediates. The probe thioester, $\text{Ph}_2\text{PC}_6\text{H}_4\text{C(O)SPh}$, was efficiently generated by condensation of 2-diphenylphosphinobenzoic acid¹¹ and benzenethiol. Related phosphine thioesters are known¹² but have been only lightly studied.

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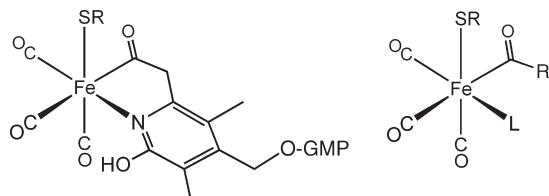


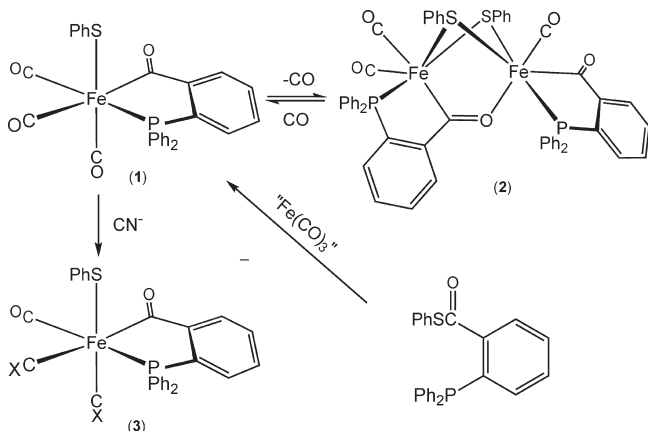
Figure 1. Structure of the CO-inhibited active site of Hmd (left) and a general motif of proposed models (right).

Table 1. Comparison of ν_{CO} and ν_{CN} Bands for Hmd Derivatives and Models

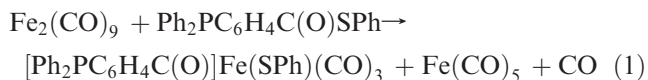
sample ^a	ν_{CO} (cm ⁻¹)
Hmd ^{CO}	2074, 2025, 2001
[Ph ₂ PC ₆ H ₄ C(O)]Fe(SPh)(CO) ₃ (1)	2075, 2020, 1981
Hmd ^{CN}	2090 (ν_{CN}), 2020, 1956
[Fe(SPh)(Ph ₂ PC ₆ H ₄ C(O)) ₂ (CN)(CO) ₂] ⁻ (3)	2094 (ν_{CN}), 2013, 1954
Hmd ¹³ CN	2047 (ν_{CN}), 2017, 1956
[Fe(SPh)(Ph ₂ PC ₆ H ₄ C(O)) ₂ (¹³ CN)(CO) ₂] ⁻ (3 ¹³ CN)	2050 (ν_{CN}), 2012, 1954

^a Enzyme data from ref 18.

Scheme 1. Oxidative Addition of Thioester–Phosphines to Fe(0) and Reactions of the Ferrous Acylthiolate



We found that Fe₂(CO)₉, a source of the reactive Fe(CO)₄ entity, reacted readily with the phosphine thioester to give a new derivative **1** (³¹P NMR: δ 72.5). Compound **1** is assumed to be the targeted acyliron(II) thiolate [Ph₂PC₆H₄C(O)]Fe(SPh)(CO)₃ (eq 1).



Compound **1** proved to be labile: it partially decarbonylates upon standing. It is known that Hmd also binds a third CO ligand only weakly.¹³ When solutions of **1** were allowed to decarbonylate under an inert gas, we obtained black, air-stable crystals of derivative **2**. Fresh solutions obtained from these crystals gave a simple ³¹P NMR spectrum consistent with a pair of uncoupled, nonequivalent phosphine ligands. As indicated crystallographically, **2** has the formula Fe₂(SPh)₂[Ph₂PC₆H₄C(O)]₂(CO)₃. Although it features ferrous centers, **2** is related to the well-studied diiron(I) dithiolates (Fe₂(SR)₂(CO)₆)¹⁴

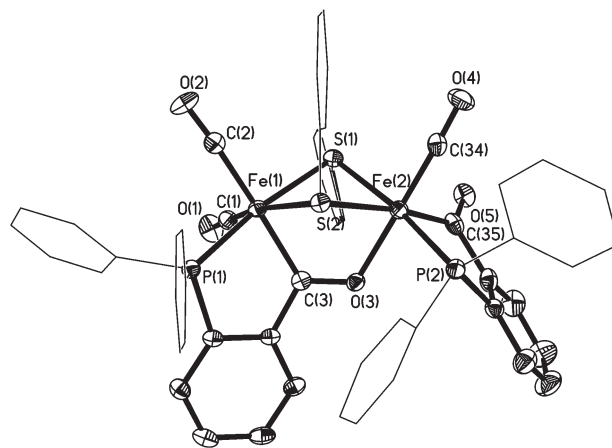
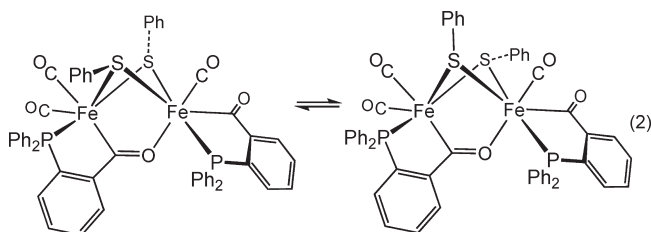


Figure 2. Structure of Fe₂(SPh)₂[Ph₂PC₆H₄C(O)]₂(CO)₃ (**2**). Selected distances (Å): Fe1–C1, 1.760(3); Fe1–C2, 1.828(3); Fe1–C3, 1.963(3); Fe1–P1, 2.2115(8); Fe1–S2, 2.3258(8); Fe1–S1, 2.3581(8); Fe2–C34, 1.748(3); Fe2–C35, 1.937(3); Fe2–O3, 2.0104(18); Fe2–P2, 2.2094(8); Fe2–S1, 2.3102(8); Fe2–S2, 2.3972(8).

and thiolato acyls (Fe₂(SR)(C(O)R')(CO)₆).¹⁵ Two acyl–phosphine ligands are chelating, one on each Fe. One acyl group is bridging, indicative of the basicity inherent in acyl oxygen.^{15,16} In solution, **2** was found to isomerize over the course of several hours, as indicated by the appearance of new pairs of singlets in the ³¹P NMR spectrum. The isomerization is attributed to the reorientation of the μ -SPh groups, a well-known phenomenon (eq 2).^{14,17}



Treatment of solutions of **2** with 1600 psi of CO (24 h, 25 °C) resulted in its conversion back to **1**. The carbonylation of **2** provided us with highly pure samples of **1**, allowing its more definitive characterization. The IR spectrum of **1** resembles that for Hmd^{CO}, the CO-inhibited state of the active site (Table 1).¹⁸

The presence of three ν_{CO} bands in the IR spectrum established the facial geometry for **1**, analogous to the case for CO-inhibited Hmd.¹⁸ Exposure of a solution of **1** to an atmosphere of ¹³CO resulted in rapid exchange of all sites to give Fe(SPh)[Ph₂PC₆H₄C(O)](¹³CO)₃. The ³¹P NMR spectrum of this species confirmed the arrangement of the CO ligands, since three separate ¹³C–³¹P couplings are observed, (J = 58, 21, and 16 Hz). Treatment of Hmd with ¹³CO results in stereoselective substitution,¹⁹ in contrast to our model. The differing CO exchange pathways for the protein vs the

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model may be due to the phosphine vs pyridine or may arise from the role of the oxygen functionality on the 2-position of the pyridine ring in the GP cofactor (see Figure 1).

Analogous to the behavior of the enzyme,¹⁸ cyanide forms a stable adduct from **1**, giving $[\text{Fe}(\text{SPh})(\text{Ph}_2\text{PC}_6\text{H}_4\text{C}(\text{O}))_2(\text{CN})(\text{CO})_2]^-$ (**3**) (Scheme 1 and Figure 2). This yellowish species, which is highly sensitive to air, was initially characterized by ESI-MS. The ^{31}P NMR spectrum indicated a single isomer. The IR data for the cyanide again match well with the data for the cofactor. Using $^{13}\text{CN}^-$, we generated the corresponding labeled derivative. The close match of the IR data for **3** and Hmd^{CN} is consistent with a low-spin iron(II) center in Hmd (Table 1). From the ^{31}P NMR spectrum of ^{13}CN -labeled **3**, we obtained $J(^{31}\text{P}, ^{13}\text{C})$ of

24 Hz, indicating that CN is cis to the phosphine.²⁰ Unlike **1**, **3** is stable with respect to loss of CO.

In summary, we have shown that donor-functionalized thioesters oxidatively add to Fe(0) to afford acyl ferrous thiolates, which provide models for the inhibited active site of Hmd. In view of the assembly strategy described here, functional modeling of this active site appears highly feasible. More broadly, the work shows that thioesters are precursors to versatile organometallic platforms.^{8–10,21}

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Supporting Information Available: Text, figures, and a CIF file giving preparative, spectroscopic, and crystallographic details. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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