

bond dissociation enthalpy (BDE) estimate of 37 ± 3 kcal/mol. This is the highest^{3,37} Co-C BDE yet measured,³⁸ slightly above Toscano's 33 ± 2 kcal/mol BDE for Me-Co(DH)₂py in bromoform.³⁹

The activation parameters allow computation of a MeCbl Co-CH₃ homolysis rate constant at -30 °C of $k_{h,on} = 10^{-19\pm 4}$ s⁻¹. This is the highest temperature at which the rate for (MeCbl)^{•+} homolysis is sufficiently slow⁴⁰ to be measurable electrochemically (rate constant = 1200 s⁻¹ at -30 °C in DMF/1-propanol).¹⁶ Comparing these two rate constants quantifies the $10^{22\pm 4}$ homolysis rate enhancement at -30 °C due to the extra, Co-CH₃ antibonding electron in (MeCbl)^{•+}.

Informative rate comparisons at higher temperatures can be made if one compares MeCbl to methylcobinamide,⁹ MeCbi⁺ (the benzimidazole-base-free form of MeCbl; the lack of the axial base in MeCbi⁺/MeCbi^{•+} slows the Co-C cleavage rates enough to make them measurable electrochemically at 25 °C). Rigorously, the MeCbi⁺/MeCbi^{•+} electrochemical data¹⁶ serve as a *lower limit*⁴⁰ to the rates for MeCbl^{•+} Co-CH₃ cleavage at other temperatures. That is, the rate enhancements that follow are *lower limits* to the true values. (If desired, the Co-C cleavage rates from MeCbi^{•+} and MeCbl^{•+} can be taken as equivalent⁴⁰ within the estimated $\pm 10^{2-3}$ error bars, and given the large rate enhancements observed.)

The electrochemically derived,¹⁶ temperature-dependent MeCbi^{•+} Co-CH₃ homolysis rates, k_h ,⁴¹ provide the activation parameters $\Delta H^\ddagger = 19 (\pm 1)$ kcal/mol and $\Delta S^\ddagger = 21 (\pm 3)$ eu. Hence at 25 °C the MeCbi^{•+} k_h is 4400 s⁻¹, which, compared to our MeCbl $k_{h,on} = 10^{-12\pm 3}$ s⁻¹, demonstrates a *rate enhancement of* $> 10^{15\pm 3}$ at 25 °C. The rate enhancement is still $> 10^{13}$ or $> 10^{11}$ at even 90 or 135 °C, respectively.

Comparing activation parameters for reduced ($\sigma^2(\sigma^*)$) MeCbi^{•+} and (σ^2) MeCbl suggests that *an antibonding electron lowers the Co-C bond strength by more than half* (i.e., from 37 kcal mol⁻¹ down to approximately⁴²⁻⁴⁴ 12 kcal mol⁻¹). The effect of the M-C antibonding electron—the first such measurement for any M-C/M-C⁻ pair—is impressive.⁴²

(35) (a) An efficient cage ($F_c \approx 1$) and BDE $\approx \Delta H^\ddagger_{\text{obsd}}(\text{soln}) - F_c \Delta H^\ddagger_{\text{g}}$ are assumed.² (b) For this 0.96–1.5 cP viscosity³⁶ solvent (at 120–150 °C), the cage-escape diffusion barrier is approximated as $\Delta H^\ddagger_{\text{g}}$ (4.0 kcal mol⁻¹; calculated via the Frenkel form^{35c} of Guzman's "Andrade" equation²). (c) Frenkel, J. *Nature (London)* **1930**, *125*, 581–582.

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(40) (a) Axial-base on alkylcobalamins undergo Co-C homolysis faster than the corresponding base-off alkylcobalamins or benzimidazole-base-free alkylcobinamides.^{1,3,4,40b} For example, AdoCbl^{4,5} homolyzes 10² times faster than³ AdoCbi^{•+} at room temperature. Furthermore, MeCbl^{•+} homolyzes 440 times faster than MeCbi^{•+} at -30 °C (1200 s⁻¹ and 2.7 s⁻¹, respectively).¹⁶ (b) Schrauzer, G. N.; Grate, J. H. *J. Am. Chem. Soc.* **1981**, *103*, 541–546.

(41) (a) The Co-CH₃ cleavage mechanism we expect for reduced alkylcobalamins differs from that presented in the electrochemical literature¹⁴⁻¹⁶ by incorporating *reversible* Co(II)-CH₃ cleavage^{41b} followed by CH₃[•] trapping, Me[Co(II)corrin]^{•+} \rightleftharpoons Co(I)^{•+} + CH₃[•], then CH₃[•] + trap \rightarrow CH₃-trap, $k_{\text{obsd}} = k_{\text{h,apparent}}$ = a composite (with the reverse of the first step probably favored by the preferred, base-off form^{41c} of Co(I)). Fortunately, the solvent mixture DMF/1-propanol is apparently serving as a trap (a H[•] source as previously noted),¹⁶ thereby preventing Co(I) + Me[•] recombination (and thus $k_{\text{h,apparent}} \approx k_{\text{h,true}} = k_h$ in DMF/1-propanol, but not in H₂O¹⁴). This mechanism, the evidence for it, and its implications will be discussed in a full paper.^{8a} (b) The trapping of a R[•] by a diamagnetic metal has precedent: Collman, J. P.; Hegedus, L. S.; Norton, J. R.; Finke, R. G. *Principles and Applications of Organotransition Metal Chemistry*; University Science Books: Mill Valley, CA, 1987; pp 314–315. Finke, R. G.; Keenan, S. R.; Watson, P. L. *Organometallics* **1989**, *8*, 263–277, especially p 269 and footnote 26. (c) Lexa, D.; Savéant, J.-M. *J. Am. Chem. Soc.* **1976**, *98*, 2652.

(42) Radical-cage effects,² although undoubtedly present in $k_{h,on}$ (F_c assumed ≈ 1)³⁵ and possibly $k_{h,apparent}$ (F_c assumed⁴³ ≈ 1 , but arguably as small as ≈ 0),^{41a,43} will not influence the conclusions in this paper (that are based on $> 10^{15}$ rate differences) and are most likely negligible compared to the indicated $\pm 10^{2-3}$ error bars.

It is of interest to consider the possible biological relevance of this mechanism for greatly enhancing M-C cleavage. Extremely labile M-alkyls are hereby predicted for systems isoelectronic to d⁷ Co(II)-CH₃, notably any d⁷ Ni(III)-alkyls related to cofactor F₄₃₀.⁴⁵ On the other hand, rather stable Co-methyl bonds (BDE = 37 kcal/mol) that are *not reducible by biological reductants*⁴⁶ are the apparent rule for d⁶ Co-CH₃ corrinoids. This latter statement is supported by the work of Ragsdale and co-workers, who have recently tested for, but found no evidence of, reductive cleavage of a d⁶ Co(III)-CH₃ bond in the corrinoid/4Fe-4S-containing protein which serves as the methyl carrier protein in the acetyl-CoA pathway of *Clostridium thermoaceticum*.⁴⁷ Perhaps it is the enormous stability difference between a d⁷ Ni(III)-CH₃ and a d⁶ Co(III)-CH₃ that Nature is exploiting.

Consistent with the above, the mechanism responsible for the observed enzymatic rate enhancement¹ of Co-C homolysis in AdoCbl probably does *not* involve (AdoCbl)^{•+}.^{6,46} Our reasoning behind this statement, and a parallel analysis of the rate enhancement following AdoCbl reduction, is presented elsewhere.⁴⁶

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Registry No. MeCbl, 13422-55-4; MeCbl^{•+}, 67087-21-2; Co¹¹B₁₂, 14463-33-3; TEMPO, 2564-83-2; TEMPO-Me, 34672-84-9.

(43) $\Delta H^\ddagger_{\text{g}} = 18.9$ kcal mol⁻¹ for (MeCbi^{•+})^{•+} homolysis. Subtracting both 4.5 kcal mol⁻¹ for the axial-base contribution^{3,4} and $\Delta H^\ddagger_{\text{g}}$ (< 2.3 kcal mol⁻¹; i.e., assuming $F_c \approx 1$)³⁵ yields an *estimated* BDE for (MeCbl)^{•+} of 12 kcal mol⁻¹.

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Long-Range Electron Transfer in Ruthenium-Modified Cytochrome c: Evaluation of Porphyrin-Ruthenium Electronic Couplings in the *Candida krusei* and Horse Heart Proteins

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Experiments in several laboratories have shown that electron transfer (ET) can take place at appreciable rates over long distances (> 10 Å) in organic and inorganic molecules¹⁻⁶ and in

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Table I. Rate Constants and Activation Parameters for Ru(His-33)-Zn-cyt *c* and Ru(His-39)-Zn-cyt *c* ET Reactions

reactn ^b	Ru(His-33)-Zn-cyt <i>c</i> ^d				Ru(His-39)-Zn-cyt <i>c</i> ^d		
	$-\Delta G^\circ$, eV	k_{ET} , s ⁻¹	ΔH^\ddagger , kcal mol ⁻¹	ΔS^\ddagger , eu	k_{ET} , s ⁻¹	ΔH^\ddagger , kcal mol ⁻¹	ΔS^\ddagger , eu
Ru ₄ (isn)(His) ²⁺ → ZnP ⁺ (ET ^b)	0.66	2.0 × 10 ⁵	<0.5	-35	6.5 × 10 ⁵	-1.7	-39
ZnP* → Ru ₄ (His) ³⁺ (ET*)	0.70	7.7 × 10 ⁵	1.7	-27	1.5 × 10 ⁶	1.3	-27
Ru ₄ (py)(His) ²⁺ → ZnP ⁺ (ET ^b)	0.74	4.2 × 10 ⁵	<0.5	-34	1.5 × 10 ⁶	-1.8	-37
ZnP* → Ru ₄ (py)(His) ³⁺ (ET*)	0.97	3.3 × 10 ⁶	2.2	-22	8.9 × 10 ⁶	0.2	-27
Ru ₄ (His) ²⁺ → ZnP ⁺ (ET ^b)	1.01	2.2 × 10 ⁶	-0.4	-31	5.7 × 10 ⁶	-0.2	-29
ZnP* → Ru ₄ (isn)(His) ³⁺ (ET*)	1.05	2.9 × 10 ⁶	<0.5	-30	1.0 × 10 ⁷	0.2	-27

^a From ref 21. ^b a = NH₃. ^c k_{ET} values are reported for 22 °C (μ 0.1 M sodium phosphate; pH 7.0); thermodynamic parameters were calculated from rate data obtained in the range 5–35 °C.

proteins.^{6–15} In non-protein systems, the evidence suggests that ET rates depend upon the number of covalent bonds separating the donor and the acceptor, rather than upon their direct separation distance.^{2,3} There is a bewildering array of potential ET pathways in proteins;^{16–19} interestingly, the through-peptide routes (if there are any!) generally involve so many bonds that they cannot possibly account for the observed rates.^{20,21} Pathways that include ionic contacts (e.g., hydrogen bonds) or small through-space jumps often can be found, and it has been postulated that such shortcuts greatly enhance the donor–acceptor electronic coupling.^{16,22} In searching for good pathways through cytochrome *c*, we discovered a relatively short route from His-39 to the heme in the *Candida krusei* (*C.k.*) protein.²³ Along this route, the crucial shortcut is a hydrogen bond that bridges Gly-41 and the heme. Since experimental information relevant to protein pathway models is lacking, we have extracted donor–acceptor electronic coupling constants from an analysis of the driving-force dependence of ET rates in Ru(His-39)-modified *C.k.* zinc cytochrome *c*.

Ruthenium [Ru(NH₃)₄L(His-39)], with L = NH₃, pyridine, isonicotinamide] derivatives of *C.k.* Zn-substituted cyt *c* [Ru-

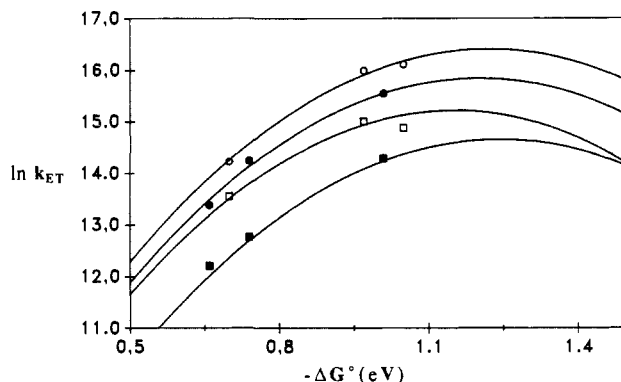


Figure 1. Plots of $\ln k_{ET}$ vs $-\Delta G^\circ$ for the ruthenated Zn-cytochrome *c* ET reactions (see Table I): ET* (○) and ET^b (●), *C.k.* protein; ET* (□) and ET^b (■), horse heart protein. Solid lines represent best fits to eq 1. *C.k.* protein: ET*, $\lambda = 1.22$ eV, $H_{AB} = 0.24$ cm⁻¹; ET^b, $\lambda = 1.20$ eV, $H_{AB} = 0.18$ cm⁻¹. Horse heart protein: ET*, $\lambda = 1.15$ eV, $H_{AB} = 0.13$ cm⁻¹; ET^b, $\lambda = 1.24$ eV, $H_{AB} = 0.10$ cm⁻¹.

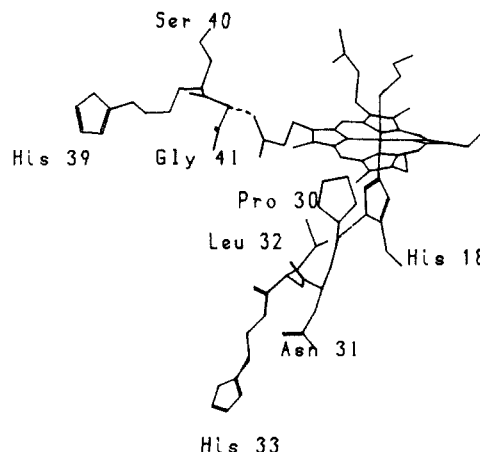


Figure 2. Possible pathways for electron transfer from histidines 33 and 39 to the heme in cytochrome *c*.²¹ Edge–edge distances are as follows:²⁹ His-39 to heme, 13.0; His-33 to His-18, 11.7; His-33 to heme, 13.2 Å.

(His-39)-Zn-cyt *c*] were prepared by standard procedures.^{9,23,24} Intramolecular ET can be initiated in these protein derivatives by photoexcitation of the Zn porphyrin (ZnP) to its strongly reducing triplet excited state.²¹ In addition to its intrinsic radiative and nonradiative decay pathways, this triplet can decay by ET to a histidine-bound Ru(III)-ammine complex (ET*). The metastable product of the ET* reaction, Ru(II)-ZnP⁺, relaxes via a thermal ET process (ET^b) to reform the ground-state complex. The kinetics of these ET reactions were measured with the three *C.k.* Ru(His-39)-Zn-cyt *c* derivatives by laser flash photolysis.⁹ ET rates and activation parameters for both *C.k.* and horse heart proteins²¹ are set out in Table I.

A nonadiabatic expression (eq 1) can be utilized to analyze long-range ET rates in derivatized proteins.²⁵ The term H_{AB} in

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$$k_{ET} = \frac{2\pi(H_{AB})^2}{\hbar(4\pi\lambda kT)^{1/2}} \exp\left[\frac{-(\Delta G^\circ + \lambda)^2}{4\lambda kT}\right] \quad (1)$$

eq 1 is the electronic coupling matrix element for the ET reaction, and λ is the nuclear reorganization parameter. Analyses of the Ru(His-33)-Zn-cyt *c* and Ru(His-39)-Zn-cyt *c* data are shown in Figure 1. Only an increase in H_{AB} , as opposed to small variations in λ or ΔG° , can account for the fact that the Ru(His-39)-Zn-cyt *c* ET rates are 3 times the Ru(His-33)-Zn-cyt *c* rates over the 0.66–1.05 eV driving-force range.²⁶ The invariance of λ found in the ET^a and ET^b reactions of the two proteins is consistent with theoretical considerations.^{25,27,28}

The H_{AB} values for Ru(His-39)-Zn-cyt *c* (ET^a, 0.24 cm⁻¹; ET^b, 0.18 cm⁻¹) are almost twice as large as those for Ru(His-33)-Zn-cyt *c* (ET^a, 0.13 cm⁻¹; ET^b, 0.10 cm⁻¹).²¹ It is likely that the electronic couplings in both proteins involve a superexchange mechanism in which electronic states of the intervening medium mix with localized donor states to produce a nonzero H_{AB} .^{16,29,30} Calculations indicate that there are two relatively good pathways for ET from His-33 to the metalloporphyrin:³¹ a 16-bond route to the Zn atom through His-18 that includes a 1.85-Å H bond between the Pro-30 carboxyl oxygen and the proton on the His-18 nitrogen and a 13-bond route ending with a 3.6-Å through-space contact between the δ -carbon of Pro-30 and a pyrrole carbon of the porphyrin. The shortest pathway from His-39 is a 13-bond route that includes a 2.4-Å H bond between the α -amino hydrogen atom of Gly-41 and the carboxyl oxygen of a propionate side chain on the porphyrin (Figure 2). Because a hydrogen bond is predicted¹⁶ to be a better shortcut than a through-space jump, the 13-bond route from His-39 to the porphyrin should lead to stronger electronic coupling than the 13-bond pathway from His-33. The 16-bond bridge from His-33 to the Zn that includes an H bond may also provide better coupling than the 13-bond (Pro-30 through space to porphyrin) pathway. It should be only slightly less effective in coupling the ruthenium and porphyrin centers than the 13-bond pathway in Ru(His-39)-Zn-cyt *c*.³²

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(28) Dielectric continuum models of solvent reorganization predict that the outer-sphere contribution to λ (λ_o) will increase with donor–acceptor separation.^{25,27} Modeling the Ru-Zn-cyt *c* systems as single spheres suggests that λ_o for the Ru(His-33)-Zn-cyt *c* reactions should be nearly the same as that for the Ru(His-39)-Zn-cyt *c* reactions (0.58 and 0.59 eV, respectively). The cyt *c* molecule was taken as a 17-Å sphere and the Ru–ammine group as a 6-Å sphere. These two interpenetrating spheres were enclosed by a third sphere of radius 17.6 Å for Ru(His-33)-Zn-cyt *c* and 18.2 Å for Ru(His-39)-Zn-cyt *c*. The Zn and Ru redox centers were taken as 6.0 and 14.6 Å from the center of the sphere, respectively, and separated from one another by 18.6 Å in Ru(His-33)-Zn-cyt *c*. The corresponding distances for the Ru(His-39)-Zn-cyt *c* model were 6.3, 15.2, and 19.3 Å. The dielectric constant of the sphere was taken as 1.8; the solvent was assigned a static dielectric constant of 78.54 and an optical dielectric constant of 1.78.

(29) The shortest direct distances between porphyrin carbon atoms and imidazole carbon atoms of His-33 (13.2 Å) and His-39 (13.0 Å) are much too long for any direct donor–acceptor interaction.¹⁶ Calculations were made using BIOGRAF/III version 1.34 (BIOGRAF was designed and written by S. L. Mayo, B. D. Olafson, and W. A. Goddard III). The structures of horse heart cytochrome *c* and its Ru(His-33) derivatives were built from the structure of the tuna protein by side-chain substitution and molecular mechanics energy minimization.^{7a,7b,21} The structure of *C.k.* cytochrome *c* was generated from the structure of the tuna protein by side-chain substitution.²³ In both *C.k.* and horse heart proteins, an imidazole carbon on His-33 is 11.7 Å from an imidazole carbon of His-18, an axial ligand of the metalloporphyrin. This value has been used as the edge-to-edge distance in previous studies.^{9,21} His-18 is not likely to be as strongly coupled to the porphyrin-localized donor and acceptor states as are carbon atoms of the porphyrin ring.³⁰ Hence, in comparing donor–acceptor coupling in Ru(His-33)-Zn-cyt *c* and Ru(His-39)-Zn-cyt *c*, distances to porphyrin carbon atoms have been used.

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Our work highlights the need for in-depth theoretical and experimental investigations of the possible role of hydrogen bonds in the pathways for long-range electron transfer through proteins. Systematic studies of electronic couplings in donor–(spacer)–acceptor molecules with variable H-bond connectors could be particularly valuable.

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(32) The H bonds in the His-33 (horse) and His-39 (*C.k.*) pathways are assumed to be the same as in the tuna protein. This assumption is reasonable, because the amino acids involved in these interactions (His-18, Pro-30, Gly-41) are conserved in the three proteins.²³

Novel Pentacoordinate Anionic Silicate, [o-C₆H₄(SiPhF₂)₂F]⁻, K⁺-18-Crown-6, Containing a Bent Fluoride Bridge between Two Silicon Atoms

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Pentacoordinate anionic silicates have recently received much attention from structural and mechanistic points of view,¹ including the nature of bonding,² intramolecular ligand exchange,³ intermolecular ligand exchange with tetracoordinate silanes,⁴ enhanced reactivity toward nucleophiles,⁵ and activation of the silicon–carbon bonds.⁶ New aspects should further be accumulated.

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