Intermolecular Electron-Transfer Catalyzed on Nanoparticle Surfaces

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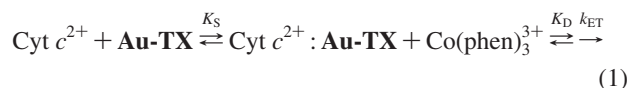
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Interprotein electron transfer (ET) is crucial to energy transduction in photosynthesis and respiration and is a direct example of protein recognition coupled with chemistry. The effects of distance and driving force are well understood and can be manipulated in synthetic and natural systems to enhance ET kinetics.[1,2] In contrast, the effects of surface binding and dynamics on ET rates are harder to interpret,[3−6] making new model systems crucial for understanding the role of dynamics in ET reactivity.

Nanoparticles are capable of catalyzing selected reactions, by acting as artificial receptors for substrates.[7,8] For example, the rates of peptide ligation[7] and phosphodiester cleavage were enhanced by up to 10^3-fold on functionalized nanoparticle surfaces.[9] Here, we show a 10^5-fold enhancement of an intermolecular ET rate through use of surface-functionalized nanoparticles (**Au-TX**) as artificial receptors (Figure 1). This catalysis arises through bringing
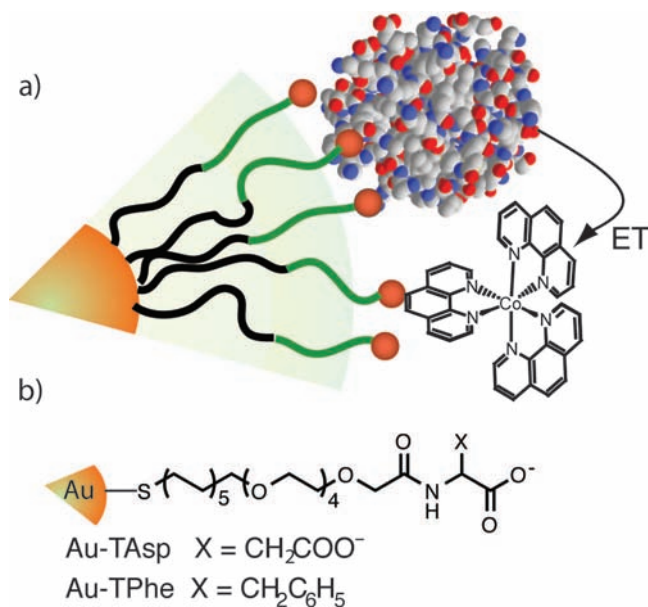

Figure 1. (a) Encounter complex for electron transfer from Cyt c to $Co(phen)_3^{3+}$ on the surface of **Au-TX**; (b) structure of **TX** ligands.

an ET donor and acceptor into close proximity, demonstrating the ability to modulate protein ET at surfaces, and provides a tool for observing and controlling the interplay between binding, dynamics, and reactivity.

Bimolecular ET kinetics follow Marcus theory,[2] with the driving force, reorganization, and steric factor determining rate. Cytochrome c (Cyt c) undergoes rapid ET with other proteins, as well as with small molecules. The dominant ET pathway is over the exposed
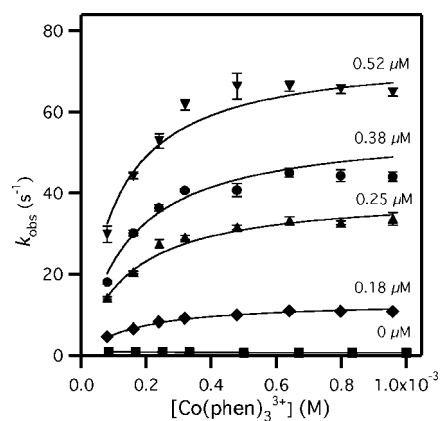
heme edge; however heme access is limited by surface Lys residues,[10] making ET selective for partners that have the appropriate binding surface.

Au nanoparticles functionalized with thiol ligands containing the free-carboxylate form of amino acids (**Au-TX**)[8] bind to Cyt c (pI = 10) with $K_S \sim 10^7$ M^{-1}.[11,12] Moreover **Au-TX** binds selectively to the surface of Cyt c near the heme edge.[11] As the charges of both $Co(phen)_3^{3+}$ and Cyt c are complementary to that of **Au-TX**, we tested whether concurrent binding could catalyze ET. The kinetic scheme accounted for the pre-equilibrium binding of Cyt c to **Au-TX**, followed by $Co(phen)_3^{3+}$ binding (K_D) and ET (k_{ET}):

$$\text{Cyt } c^{2+} + \textbf{Au-TX} \underset{}{\overset{K_S}{\rightleftharpoons}} \text{Cyt } c^{2+} : \textbf{Au-TX} + Co(phen)_3^{3+} \overset{K_D \ k_{ET}}{\rightleftharpoons} \rightarrow \tag{1}$$

Cyt c was reduced with dithionite followed by gel filtration and $Co(phen)_3^{3+}$ synthesized as per the literature. Cyt c oxidation was monitored as a single exponential decrease in A_{550} following the mixing of Cyt c (5 μM after mixing, 10 mM Tris, pH 7.40, $I = 20$ mM, 12 mM NaCl) with $Co(phen)_3^{3+}$ (0.083−1.00 mM after mixing, in same buffer) in a stopped-flow spectrometer. Cyt c oxidation by $Co(phen)_3^{3+}$ was nearly independent of [Co^{3+}], with $k_{ET} = 0.66 \ (\pm 0.02)$ s^{-1} (Figure 2),[13] in good agreement with prior

Figure 2. Cyt c oxidation (5 μM) by $Co(phen)_3^{3+}$ in the presence of varied **Au-TAsp** (0−0.52 μM); buffer is 10 mM Tris, pH 7.40, 12 mM NaCl, $I = 20$ mM, 25.0 °C.

reports.[14] This has been attributed to a rate-limiting conformational change at the heme edge of Cyt c; however, under moderate ionic strength the bimolecular ET rate is $k_{ET}/K_D = 10^3$ M^{-1} s^{-1}.[15]

The Cyt c/**Au-TAsp** adduct was formed by preincubating Cyt c (10 μM) with **Au-TAsp** (0.16−1.04 μM; higher concentrations were not used due to high optical density) and then mixing with $Co(phen)_3^{3+}$ as above. Cyt c/**Au-TAsp** oxidation was evident as single-exponential decays in A_{550}, indicating that Cyt c binding to

Au-TAsp was in a rapid pre-equilibrium (eq 1). As the position of this equilibrium favored unbound Cyt c, the ET process may be considered as coupled.[16] The observed rates for Cyt c/**Au-TAsp** exhibited saturation due to Co(phen)$_3^{3+}$ binding, with a k_{obs} as high as 65 s^{-1} (Figure 2).

Analogous data sets for Cyt c/**Au-TPhe** exhibited similar saturation kinetics.[13] The second-order kinetics were fitted to obtain apparent rate constants for Cyt c oxidation: $k_{obs} = k_{ET(app)}[Co]_0/([Co]_0 + K_D)$.

The apparent maximum ET rate, $(k_{ET})_{app}$, was a linear function of nanoparticle concentration, indicating that the binding of Cyt c to **Au-TX** (K_S of eq 1) was far from equilibrium (Figure 3). Under
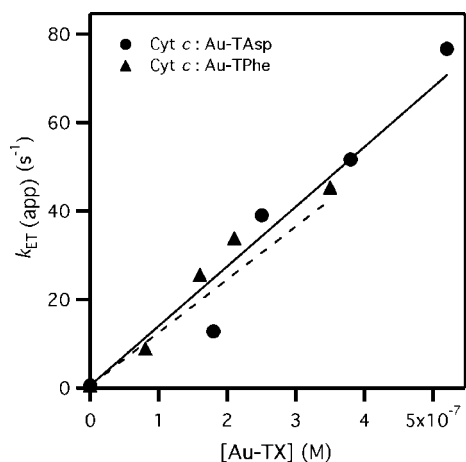

Figure 3. Bimolecular rate plot for Cyt c oxidation (5 μM, 10 mM Tris, pH 7.4, I = 20 mM) by Co(phen)$_3^{3+}$ in the presence of **Au-TX** (X = Asp, Phe). Linear fitting to $(k_{ET}/K_S)[\textbf{Au-TX}] = k_{ET(app)}$ yielded k_{ET}/K_S = 1.35 ($\pm$0.03) $\times$ 10^8 (X = Asp) and 1.20 ($\pm$0.03) $\times$ 10^8 M^{-1} s^{-1} (X = Phe). Error bars are approximately the size of each data point.

rapid pre-equilibrium conditions, $(k_{ET}/K_S)[\textbf{Au-TX}] = (k_{ET})_{app}$; the slope of Figure 3 is the absolute bimolecular rate constant for Co(phen)$_3^{3+}$ reacting with Cyt c on the **Au-TX** surface. Linear least-squares fitting for Cyt c/**Au-TX** (X = Asp, Phe) yielded slopes that were nearly identical: k_{ET}/K_S = 1.35 ($\pm$0.03) $\times$ 10^8 and 1.20 ($\pm$0.03) $\times$ 10^8 M^{-1} s^{-1}, respectively. This is significantly larger than the ET rate in the absence of **Au-TX** (1 $\times$ 10^3 M^{-1} s^{-1}),[15] indicating that complexation by **Au-TX** catalyzed the ET reaction by 10^5.

This catalysis can be understood within the context of Marcus theory,[2] which states that the rate of bimolecular ET depends on the collision rate (Z $\sim$1 $\times$ 10^{11} M^{-1} s^{-1}) and the activation energy (ΔG^*): $k = Z \exp(-\Delta G^*/RT)$. In the limit of low driving force ($\Delta G^0 \ll \lambda$), $\Delta G^* = \lambda/4 + w$, where w is the work to bring the reactants together. The reaction of Cyt c(Fe^{2+}) with Co(phen)$_3^{3+}$ (k = 1 $\times$ 10^3 M^{-1} s^{-1}) can be attributed to λ = 44 kcal/mol, assuming that w = 0. This λ is consistent with the self-exchange reaction rate for each reagent.[2] The faster rate in the presence of **Au-TX** may arise from changes in either λ or w; should λ remain unchanged, the data could be accommodated with w = -7 kcal/mol. This value for work is attributed to the electrostatic attraction of Co(phen)$_3^{3+}$ and Cyt c to **Au-TX**.

This electrostatic attraction increases the local concentration of each reagent. Calorimetric and kinetic data indicate that the Co(phen)$_3^{3+}$/**Au-TX** binding equilibrium was saturated at 1 mM

[Co(phen)$_3^{3+}$], leading to the plateau in k_{obs} (Figure 2); in contrast, Cyt c binding to **Au-TX** was subsaturated, leading to the linear dependence of k_{ET} on [**Au-TX**] (Figure 3). Entropic factors (i.e., the "Circe effect"[17]) presumably dominate intermolecular ET catalysis in the Cyt c/**Au-TX** system; however enthalpic contributions cannot be excluded.

Kinetic complexity masked the rate of electron flow within encounter complexes in previous studies of the relationship between conformational dynamics and interprotein ET. In some cases, the overall rate was limited by a conformational change, leading to gated ET.[16,18,19] In other cases, ET was rate-limiting but occurred following a disfavored conformational change with concomitant coupled ET.[3,4] Although amide H/D exchange showed that **Au-TPhe** bound to a smaller Cyt c face than **Au-TAsp**,[11] which could increase the rate of conformational dynamics, this did not translate into faster ET kinetics for Cyt c/**Au-TPhe**. This suggests that conformational dynamics within the {Cyt c/**Au-TX**/Co^{3+}} encounter complex are much faster than ET and that dynamics may be uncoupled from ET under these conditions.

In summary, we have demonstrated the use of nanoparticles as highly efficient catalysts for intermolecular ET. These catalysts function by reversibly binding the ET donor and acceptor, thus increasing the local concentration of the redox partners. This process may provide a model for understanding intermolecular ET, in which binding is coupled to reactivity.

Supporting Information Available: Additional kinetics and calorimetric data. This material is available free of charge via the Internet at http://pubs.acs.org.

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