

Electronic Supplementary Information: Direct Simulation of Electron Transfer in the Cobalt Hexammine(II/III) Self-Exchange Reaction

Rachel L. Kenion[†] and Nandini Ananth^{*,‡}

[†]*Robert Frederick Smith School of Chemical and Biomolecular Engineering, Cornell University, Ithaca, New York 14853, USA*

[‡]*Department of Chemistry and Chemical Biology, Cornell University, Ithaca, New York 14853, USA*

E-mail: ananth@cornell.edu

Contents

I. Co(NH ₃) ₆ ³⁺ Optimization for Partial Charges	1
II. Force Field Parameters for Atomistic Simulations	2
III. Solvent Relaxation Timescales	3

I. Co(NH₃)₆³⁺ Optimization for Partial Charges

As described in the text, Mulliken analysis from gas phase density functional theory is used to assign partial charges to the atoms of the TMC for the atomistic simulations. Using the Gaussian 09¹ software package, the complex is optimized using the M06 hybrid functional² with the 6-31G(d,p) basis set³ for N and H atoms and the LANL2DZ⁴ basis set for Co.

The resulting optimized structure is provided in Table S1. The computed energy at this configuration is -483.282740937 hartrees. Partial charges on H atoms from the Mulliken analysis were summed into the N to which they are bonded, and those net partial charges are averaged over each of the N atoms in order to achieve a total charge of $+3\text{ e}$ on $\text{Co}(\text{NH}_3)_6^{3+}$. The resulting partial charges used in the simulation are $+0.41281\text{ e}$ on Co and $+0.43119\text{ e}$ on N.

Table S1: Optimized $\text{Co}(\text{NH}_3)_6^{3+}$ geometry

atom type	x	y	z
Co	0.000192	0.000106	-0.000008
N	1.275211	-0.743840	1.357761
N	-1.280279	0.657544	-1.397605
N	1.159948	1.624901	-0.193382
N	-1.035952	0.985046	1.406268
N	-1.150609	-1.617457	0.288628
N	1.031384	-0.906287	-1.461649
H	0.954795	-0.687125	2.330209
H	1.469827	-1.741134	1.217457
H	2.200171	-0.300657	1.367293
H	-1.562368	-0.075561	-2.057387
H	-2.164499	1.026081	-1.030941
H	-0.920427	1.407129	-1.997678
H	0.720576	2.417464	-0.673103
H	1.456396	2.012570	0.708929
H	2.035037	1.469386	-0.705342
H	-1.268573	1.945628	1.131574
H	-1.938671	0.562311	1.648358
H	-0.555137	1.092274	2.305603
H	-2.100953	-1.540449	-0.089489
H	-0.785633	-2.487385	-0.113283
H	-1.286385	-1.839099	1.280812
H	1.970246	-1.220781	-1.193406
H	0.585346	-1.741836	-1.854485
H	1.187132	-0.301032	-2.275043

II. Force Field Parameters for Atomistic Simulations

Table S2: Simulation Force Field Parameters

atom types	potential type ^a	parameter	value
Co-N	harmonic bond ^b	k_b^c	352.5
		r_0	1.94 Å
Co-N	rigid bond ^d	r^\ddagger	2.05 Å
N-H	rigid bond	r	1.03 Å
Co-N-H	harmonic angle	k_θ^e	30
		θ_0	114°
H-N-H	harmonic angle	k_θ^e	44
		θ_0	104°
O-N	12-6	A^f	8.78×10^5
		B^g	7.83×10^2
H_{SPC} -N	12-6	A^f	6.09×10^2
		B^g	15.3
N-N	12-6	A^f	1.32×10^6
		B^g	1.03×10^3

^a Functional forms of the potentials appropriate for the provided parameters are in accordance with the CHARMM⁵ convention. ^b Parameter adapted from Ref. 6; ^c harmonic bond force constant reported in units of kcal/mol Å⁻²; ^d observed simulated transition state bond length in agreement with previously reported value;^{7,8} ^e harmonic angle force constant reported in units of kcal/mol rad⁻²; ^f van der Waals 12-6 A parameter reported in units of kcal/mol Å¹²; ^g van der Waals 12-6 B parameter reported in units of kcal/mol Å⁶.

III. Solvent Relaxation Timescales

We demonstrate the ability of the approach outlined in this work to provide insight into solvent relaxation timescales. Defining a collective coordinate ΔU reminiscent of the solvent coordinate in Marcus theory,⁹ the energy difference between the diabatic electronic states within the tight binding approximation is expressed as:

$$\Delta U(\mathbf{Q}) = -\frac{e}{4\pi\varepsilon_0} \sum_{k=1}^N \left(\frac{q_k}{|\mathbf{Q}_D - \mathbf{Q}_k|} - \frac{q_k}{|\mathbf{Q}_A - \mathbf{Q}_k|} \right). \quad (1)$$

Here, q_k the charge on solvent atom k , and \mathbf{Q}_D , \mathbf{Q}_A , and \mathbf{Q}_k positions of the “donor” Co center, “acceptor” Co center, and solvent atom k , respectively. Time resolving the auto-

correlation of ΔU provides insight into the relaxation time of the solvent environment. In this study, an initial relaxation time of about 300 fs is indicated (Fig. S1), consistent with solvent relaxation on the same timescale of our electronic transition (shown by κ in Fig. 5 in the main text), followed by a second longer relaxation mechanism that extends beyond the simulation time.

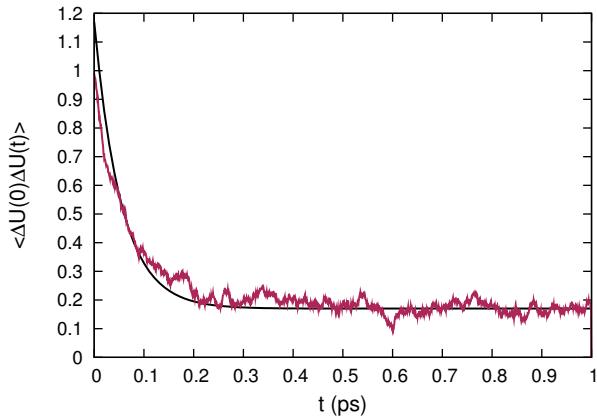


Figure S1: The normalized autocorrelation function of ΔU Eq. 1 along RPMD trajectory time.

References

- (1) Frisch, M. J. et al. Gaussian 09 Revision C.01. 2009.
- (2) Zhao, Y.; Truhlar, D. G. *Theor. Chem. Acc.* **2008**, *120*, 215–241.
- (3) Hehre, W. J.; Ditchfield, R.; Pople, J. A. *J. Chem. Phys.* **1972**, *56*, 2257–2261.
- (4) Hay, P. J.; Wadt, W. R. *J. Chem. Phys.* **1985**, *82*, 270–283.
- (5) MacKerell Jr., A. D. et al. *J. Phys. Chem. B* **1998**, *102*, 3586–3616.
- (6) Zheng, C.; McCammon, J. A.; Wolynes, P. G. *Proc. Natl. Acad. Sci. U.S.A.* **1989**, *86*, 6441–6444.
- (7) Newton, M. D. *J. Phys. Chem.* **1991**, *95*, 30–38.

- (8) Endres, R. G.; LaBute, M. X.; Cox, D. L. *J. Chem. Phys.* **2003**, *118*, 8706–8714.
- (9) Menzelev, A. R.; Ananth, N.; Miller III, T. F. *J. Chem. Phys.* **2011**, *135*, 074106–1–074106–17.