Characterization of the non-covalent docking motif in the isolated reactant complex of a double proton-coupled electron transfer reaction with cryogenic ion spectroscopy

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ABSTRACT

The solution kinetics of a proton-coupled electron transfer reaction involving two-electron oxidation of a Ru compound with concomitant transfer of two protons to a quinone derivative have been interpreted to indicate the formation of a long-lived intermediate between the reactants. We characterize the ionic reactants, products, and an entrance channel reaction complex in the gas phase using high-resolution mass spectrometry augmented by cryogenic ion IR photodissociation spectroscopy. Collisional activation of this trapped entrance channel complex does not drive the reaction to products but rather yields dissociation back to reactants. Electronic structure calculations indicate that there are four low-lying isomeric forms of the non-covalently bound complex. Comparison of their predicted vibrational spectra with the observed band pattern indicates that the C=O groups of the ortho-quinone attach to protons on two different $-NH_2$ groups of the reactant scaffold, exhibiting strong O–H–N contact motifs. Since collisional activation does not lead to the products observed in the liquid phase, these results indicate that the reaction most likely proceeds through reorientation of the H-atom donor ligand about the metal center.

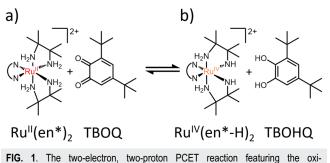
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I. INTRODUCTION

Recent advances in cold ion vibrational spectroscopy, coupled with ambient ionization methods, enable structural characterization of increasingly complex molecular ensembles.¹ An important class of species becoming accessible with these methods involves trapping non-covalently bound complexes between reactants,^{2–5} thus providing an unprecedented view of how the two species dock together and how this arrangement is related to the reaction coordinate that leads to products. This strategy was employed earlier to determine the docking arrangements at play in a biomimetic bromination catalyst,² as well as the degree of small molecule (CO₂, N₂, and O₂) activation by the active coordination site on an organometallic coordination compound.^{6,7} Here, we expand the scope of this approach to

address the geometry of a non-covalently bound complex formed by the reactants in a proton-coupled electron transfer reaction between a Ru^{II} coordination compound and the orthoquinone (hereafter denoted reaction I) depicted in Fig. 1. From the experimental standpoint, this system presents a more demanding scenario than those studied previously^{6,8–10} because of the +2 charge state of the catalyst under study and the many isotopologues arising from the Ru and C atoms. This challenge has very recently been met, however, by coupling cold ion spectroscopy with a commercial high-resolution mass spectrometer (Thermo Velos Pro), which we herein exploit for this experimental demonstration.^{11,12}

We chose this PCET system because it involves the interesting scenario where two hydrogen atoms are transferred to the orthoquinone along with two-electron oxidation of Ru^{II} to $Ru^{IV,13}$



dation of the ruthenium metal center from (a) +II $([Ru^{II}(en^*)_2]^{2*} = [Ru^{II}(bpy)(H_2NCMe_2CMe_2NH_2)_2]^{2*})$ to (b) +IV $([[Ru^{IV}(en^*-H)_2]^{2*} = [Ru^{IV}(bpy)(H_2NCMe_2CMe_2NH_2)_2]^{2*})$. The reactant ortho-quinone (TBOQ = 3,5-di-tert-butyl ortho-quinone) is converted to product ortho-hydroquinone (TBOHQ = 3,5-di-tert-butyl ortho-hydroquinone).

As such, this "double-PCET" process raises the question of whether the entrance channel complex can be captured in a configuration involving C=O interactions with the two reactive protons on the quinone. The motivation for addressing this question in part derives from an earlier investigation¹³ of the mechanism for this reaction using traditional tools of chemical analysis. UV-Vis spectroscopy of the reaction mixture, for example, revealed an isosbestic point, indicating production of Ru^{IV} without buildup of an intermediate Ru^{III} species.¹³ In addition, features in the ¹H-NMR spectra were interpreted in the context of the formation of a long-lived association complex between the reactants. In this report, we examine the intrinsic hydrogen-bonding interaction of the isolated Ru^{II}amine dication $([Ru^{II}(bpy)(H_2NCMe_2CMe_2NH_2)_2]^{2+}$ (bpy = 2,2'bipyridine, H₂NCMe₂CMe₂NH₂ = 2,3-dimethyl-2,3-butanediamine = en^*), denoted $[Ru^{II}(en^*)_2]^{2+}$ with the quinone (3,5-di-tert-butyl ortho-quinone, denoted TBOQ), where both are displayed schematically in Fig. 1(a) This characterization was accomplished by extracting the ions from solution using an electrospray ambient ion source and cryogenically cooling them to ~20 K in a radio frequency ion trap. Their vibrational spectra were then obtained in a linear action mode by IR photodissociation of a weakly bound D₂ molecule.¹, Finally, the nature of the binding pocket was assessed by comparing the observed band patterns with those calculated for various local minima and consideration of their relative computed energies.¹¹

II. EXPERIMENTAL

 $[Ru^{II}(en^*)_2(PF_6)_2]$ and $[Ru^{IV}(en^*-H)_2(PF_6)_2]$ were synthesized according to protocols reported previously.¹³ 3,5-di-tertbutyl ortho-quinone (TBOQ), 3,5-di-tert-butyl ortho-hydroquinone (TBOHQ), and acetonitrile (HPLC grade) were purchased from Sigma-Aldrich and used without further purification. The $[Ru^{II}(en^*)$ TBOQ]²⁺ complex ion was generated via electrospray ionization using a (~1 mM), 1:1 solution of $[Ru^{II}(en^*)_2(PF_6)_2]$ and TBOQ in acetonitrile. Mass spectra and photofragmentation infrared spectra were collected using a hybrid instrument that combines a commercial Thermo Fisher Scientific LTQ Orbitrap Velos Pro with a custom-built triple-focusing time-of-flight photofragmentation mass spectrometer, shown in Fig. S1 and described in detail previously.¹¹ In brief, mass spectra were collected using the instrument's Orbitrap capability at maximum resolution (Δ m/m = 100 000) before transmitting the ions to the photofragmentation spectrometer for infrared structural characterization. The vibrational spectra were obtained by photodissociating weakly bound D₂ molecules using the "messenger tagging" approach,²¹ where the D₂ molecules were condensed onto target ions in a radio frequency ion trap held at ~20 K.

Density functional theory (DFT) was used to calculate the normal modes of the target ions at the M062X/6-311G* level of theory (C, H, N, O, P, F) and def2-TZVP with an effective core potential for the Ru center.^{22,23} All calculations were conducted using the Gaussian 09²⁴ suite of electronic structure software. Crystal structures of $[Ru^{II}(en^*)_2(PF_6)_2]$ and $[Ru^{IV}(en^*-H)_2(PF_6)_2]$ were used as initial structures for DFT optimization after omitting the PF₆⁻ anion.²⁵ A conformational survey was performed in which TBOQ was placed at different positions around the NH moieties in the $[Ru^{II}(en^*)_2]^{2+}$ compound groups and then optimizing the initial structure into the local energetic minima. See Table S1 of the supplementary material for relative energies and BSSE corrected binding energies. Additionally, all possible isomers for [Ru^{IV}(en*-H)₂] after double proton abstraction from the en^* groups were identified. The relative energies of the isolated reactants and products, as well as their intermediates, are displayed in Fig. S2.

III. RESULTS AND DISCUSSION

A. Mass analysis of complex formation with ESI

The kinetics and energetics of $[Ru^{II}(en^*)_2]^{2+}$ reactions with various quinones have been characterized extensively by Cattaneo *et al.*¹³ Of these, we selected the endothermic reaction with TBOQ $(\Delta G^0 = 1.5 \text{ kcal/mol in acetone})$ on the basis that both reactants should be present at significant concentrations in equilibrium solutions of acetonitrile prepared with equimolar (~1 mM) initial concentrations of both reactants. Figure 2(a) presents the electrospray ionization (ESI) mass spectrum of the [Ru^{II}(en^{*})₂]²⁺ dication extracted from a 1 mM solution of the $[Ru^{II}(en^*)_2(PF_6)_2]$ salt. The stoichiometry of this species was verified using the high-resolution capability of the mass analyzer, with the resulting mass spectrum displayed in Fig. S3. Introduction of the TBOQ neutral reactant into the solution yields the mass spectrum shown in Fig. 2(b), confirming formation of a binary complex between [Ru^{II}(en^{*})₂]²⁺ and TBOQ with m/z of 355.19. Furthermore, we mixed solutions of each ruthenium complex ([Ru^{II}(en*)₂]²⁺ and [Ru^{IV}(en*-H)₂]²⁺) with solutions of TBOQ or TBOHQ. In general, we might expect to capture all four binary combinations of reactants and products, but of these, we only observed TBOQ complexes with $[Ru^{II}(en^*)_2]^{2+}$ and [Ru^{IV}(en*-H)₂]²⁺; attachment of TBOHQ was never observed. This is consistent with the calculated binding energies plotted in Fig. S2, which predict that the binding energies of the various isomeric forms of the exit channel complexes are about half those of the entrance channel. Although these ions $([Ru^{II}(en^*)_2]^{2+})$ and $[Ru^{IV}(en^*-H)_2]^{2+})$ display interleaving isotopic distributions, the high-resolution mode of the Orbitrap enables clear differentiation to establish the presence of both species, as shown in Fig. S3. [Ru^{IV}(en*-H)₂TBOQ]²⁺ and [Ru^{IV}(en*-H)₂]²⁺ were observed to build up over time as the reactant [Ru^{II}(en*)₂]²⁺ degrades and

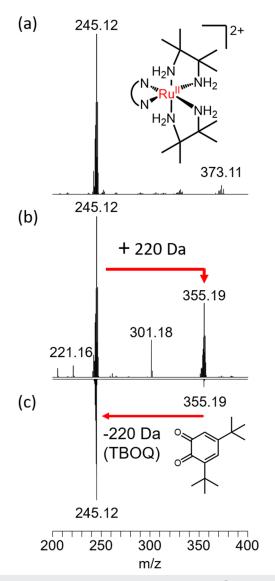


FIG. 2. (a) Mass spectrum of the solution containing the Ru^{II}(en*)₂(PF₆)₂ salt displaying the [Ru^{II}(en*)₂]²⁺ ion at 245 m/z. (b) Addition of TBOQ to the Ru^{II}(en*)₂(PF₆)₂ solution leads to the appearance of a mass at 355 m/z, which is assigned to the [Ru^{II}(en*)₂TBOQ]²⁺ ion. (c) Collision Induced Dissociation (CID) of the [Ru^{II}(en*)₂TBOQ]²⁺ ion displaying loss of TBOQ without hydrogen transfer. See the supplementary material for conformation of stoichiometry by high resolution.

were isolated and characterized without interference by spraying a solution containing only $[Ru^{IV}(en^*-H)_2]^{2+}$ and TBOQ.

To evaluate the chemical nature of the $[Ru^{II}(en^*)_2]^{2+}$ and $[Ru^{IV}(en^*-H)_2]^{2+}$ dications with masses that nominally correspond to the addition of TBOQ, we carried out collisional induced dissociation (CID) in the linear quadrupole ion trap (LTQ) section of the Velos Pro. In both cases, we exclusively observed the ejection of neutral TBOQ, with the CID mass spectrum of the $[Ru^{II}(en^*)_2 TBOQ]^{2+}$ complex displayed in Fig. 2(c). As such, we conclude that the

355.19 m/z species is indeed the non-covalently bound, entrance channel complex of reaction I. We speculate that the absence of the non-covalent adducts with the TBOHQ product is due to the fact that they are too fragile to survive the ion transfer optics in the commercial ion source. On the other hand, an extended survey of CID conditions of the mass isolated [Ru^{II}(en*)₂TBOQ]²⁺ did not yield formation of $[Ru^{IV}(en^*-H)_2]^{2+}$ products by collisional activation, raising the question of whether this non-covalently bound complex is indeed arranged along the entrance channel reaction coordinate. In contrast, we note that for the bare $[Ru^{II}(en^*)_2]^{2+}$ reactant ion, CID yields elimination of singly charged fragments in addition to simple ejection of the neutral ligand en* (Table S2 and Figs. S3 and S4). The production of two singly charged fragments necessarily involves proton migration in which one of the two en* ligands abstracts a proton, presumably from the other en* ligand, to produce $[en^* + H]^+$ and $[Ru^{II}(en^*)(en^* - H)]^+$.

B. Cryogenic vibrational spectroscopy

1. [Ru^{II}(en^{*})₂]²⁺ and [Ru^{IV}(en^{*}-H)₂]²⁺

The observed vibrational spectrum of the $[Ru^{II}(en^*)_2]^{2+} \bullet D_2$ ion in the NH stretching region (black trace) is presented in Fig. 3(a), along with the calculated harmonic spectrum [inverted bars in Fig. 3(b)] for the structure presented in Fig. S5. As indicated in the schematic structure at the upper right of Fig. 3, there are four NH₂ groups coordinated to the metal center in the *C*₂-symmetry

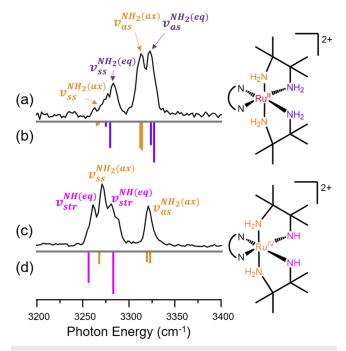


FIG. 3. NH₂ stretching region of the photofragmentation IR spectrum of $[Ru^{II}(en^*)_2]^{2^*} \bullet D_2$ (a) and $[Ru^{IV}(en^*-H)_2]^{2^*} \bullet D_2$ (c). DFT calculated frequencies [(b) and (d)] were scaled by 0.9425:1 to match the broadened line shape to the experiment maxima. Equatorial NH₂ groups (and NH groups) are highlighted in purple and pink, respectively, while axial NH₂ groups are labeled in orange.

 $[Ru^{II}(en^*)_2]^{2+}$ compound, resulting in eight vibrational fundamentals. These are divided into four symmetric (v_{ss}) and four asymmetric (v_{as}) normal modes derived from the NH stretches. The local environments of these NH₂ moieties differ according to their positions relative to the aromatic bipyridine ligand. Specifically, the two equatorial NH₂ groups [same plane as the bipyridine ligand, purple in the upper structure inset of Fig. 3, NH₂(eq)] are arranged to promote electronic interaction with the bipyridine ligand, a process that is suppressed in the axial NH₂ groups [orange in the upper inset of Fig. 3, NH₂(ax)]. The v_{as} band arising from the NH₂(eq) groups lies just above those assigned to v_{as} from the NH₂(ax) groups, which accounts for the doublet structure of the observed v_{as} band centered at ~3325 cm⁻¹. The calculated patterns [inverted trace in Fig. 3(b)] underlying this doublet are further split according to the collective in- and out-of-phase nature of the weakly coupled groups in each class, but this fine structure is not observed at the present experimental resolution (~5 cm⁻¹). The observed asymmetric peak near 3275 cm⁻¹ in Fig. 3(a) can then be assigned to the expected $v_{ss}^{\text{NH}_2(eq)}$ and $v_{ss}^{\text{NH}_2(ax)}$ bands, which are calculated to be evenly distributed in this region [orange and purple in Fig. 3(b)]. We note that the splitting between the v_{ss} and v_{as} bands (40 cm⁻¹) in the [Ru^{II}(en^{*})₂] complex is smaller than that observed in methylamine (66 cm⁻¹) and ethylenediamine (61 cm⁻¹), although it is found to be reduced in complexes involving dicationic metal ions: zinc, cadmium, and mercury (53 cm⁻¹, 55 cm⁻¹, and 55 cm⁻¹, respectively).²

Figure 3(c) presents the vibrational spectrum of the $[Ru^{IV}(en^*$ $-H_{2}^{2+} \bullet D_{2}$ product ion from reaction I, which explores how the conversion of two of the NH₂(eq) groups to NHs impacts the spectrum. Consistent with the calculated pattern [Fig. 3(d)], this modification inverts the relative intensities of the feature in the v_{as} region and the absorption just below it. The strong band at 3275 cm⁻¹ is predicted to drive from the $v_{str}^{NH(eq)}$ fundamentals associated with the NH(eq) groups in the product ion [pink bars in Fig. 3(d)]. We note that all combinations (eq and ax) of the two amido NH locations were calculated (Fig. S6), and the lowest energy structure was that with the amido groups both occupying the equatorial (eq-eq) positions with respect to the bipyridine, as shown in Fig. S6(b), in agreement with the reported crystallographic structure.²⁵ The (eq-ax and ax-ax) conformers are predicted to lie significantly higher in energy (by 15.0 kcal/mol and 25.4 kcal/mol, respectively) and display vibrational band patterns that are significantly different than what is observed experimentally [see Figs. S6(c) and S6(d)]. We can rationalize the preference for the (eq-eq) isomer by the formation of a π bond between the symmetric combination of filled amide p orbitals and the Ru d_{xy} orbital (considering the bpy ligand to lie in the xy plane). This interaction, which involves only one of the σ nonbonding t_{2g} -like d orbitals, accounts for the diamagnetism of the $d^4 [Ru^{IV}(en^*-H)_2]^{2+}$ compound.²⁵

2. The $[Ru^{II}(en^*)_2 TBOQ]^{2+}$ entrance channel complex

Figure 4 compares the observed D_2 -tagged spectrum of the isolated $[Ru^{II}(en^*)_2]^{2+}$ dication with that of the putative entrance channel complex, $[Ru^{II}(en^*)_2 TBOQ]^{2+}$, in the high frequency NH stretching region as well as in the lower frequency range of the NH bending and C=O stretching fundamentals. Although the highest energy NH feature of the $[Ru^{II}(en^*)_2]^{2+}$ band near

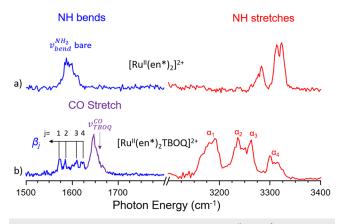


FIG. 4. Vibrational predissociation spectra of (a) $[Ru^{II}(en^*)_2]^{2^*} \bullet D_2$ and (b) $[Ru^{II}(en^*)_2 TBOQ]^{2^*} \bullet D_2$. The purple arrow in b) indicates the solution phase FTIR value for the C=O stretching mode in neutral TBOQ.²⁷

3325 cm⁻¹ (α_4) persists with some distortion in the line shape upon complexation, the dominant activity in the NH stretching region is dramatically red-shifted. Since two of the -NH₂ groups are not involved in the interaction in all calculated structures, the highest energy feature (a4) arises from the associated NH2 asymmetric stretch. The red-shifted features (α_{1-3}) , on the other hand, are expected for the TBOQ adduct and are therefore traced to the stretching bands of the NH groups involved in hydrogen bonds to the quinone. In the lower energy fingerprint region, a very strong band (purple) near 1650 cm⁻¹ appears upon complexation with the quinone, which lies just below the collective symmetric stretch of the two C=O groups of the isolated quinone in solution (1660 cm^{-1}) ,²¹ indicated by the downward arrow denoted v_{TBOQ}^{CO} in Fig. 4(b). This feature establishes that the C=O groups are intact in the complex because formation of C-OH single bonds in the neutral TBOHQ product after PCET would yield much lower frequency fundamentals.²⁸ We also note that the NH bending modes $(v_{bend}^{\rm NH_2})$, which appear as an unresolved broad feature in the bare $[Ru^{II}(en^*)_2]^{2+}$ spectrum, are split apart upon formation of the complex to form a set of distinct features [β_{1-4} in Fig. 4(b)]. The fact that two of these $(\beta_{3,4})$ lie above the bends in the bare reactant ion is consistent with the formation of NH-O linkages, which are known to blue shift the bending fundamentals.²⁹ We remark that we have focused on the NH bending and C=O stretching bands in this region because the C-C stretching fundamentals, which also occur in this frequency range, are calculated to be much weaker as we describe further below.

We identified many local minima for the entrance channel complex that all feature two H-bond docking sites in which the TBOQ C=O groups attach to one of the two NH bonds in separate $-NH_2$ groups. Interestingly, there are configurations in which the quinone binds to either both equatorial NH groups (eq-eq), which would nominally correspond to the two H atoms that are transferred in the 2xPCET process, or structures where one C=O binds to an equatorial NH and the other one to an axial group (eq-ax). Because the quinone is itself asymmetrical with respect to the locations of the

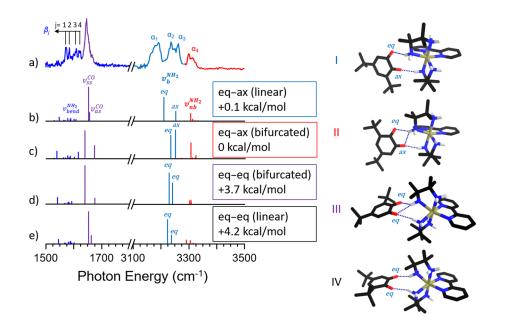


FIG. 5. Vibrational predissociation spectrum of (a) [Ru^{II}(en*)₂TBOQ]²⁺•D₂ with those calculated in (b) through (e) for the four isomers lying lowest in energy (relative values in boxes, after correction for vibrational ZPE). Calculated spectra are scaled by 0.9425 in order to match the CO and α_4 bands of the experimental spectrum. Isomers III and IV [spectra (d) and (e), respectively] correspond to a hydrogen-bonding interaction with the two equatorial NH₂ groups (eq-eq), whereas isomers I and II [spectra (b) and (c), respectively] feature attachment to one equatorial and one axial NH2 (eqax). Linear and bifurcated monikers indicate the nature of the hydrogen bonds between the two C=O groups and the two N-H groups. 3-D structures can be seen in supplementary material. Fig. S7.

two t-butyl groups, each of these forms occurs in two isomeric forms that differ by roughly 180° rotation of the TBOQ in the docking site. This results in four low-lying isomers (hereafter denoted I, II, III, and IV) with the structures indicated at the right of Fig. 5. Rotatable pdf structures are included in Fig. S7 to better display the local interactions. Note that isomers II and III feature a bifurcated binding motif in which an NH group lies between two C=O groups, while I and IV are based on more local quasilinear (C=O-HN) hydrogen bond linkages. The two eq-ax isomers (I and II) are calculated to be nearly degenerate and lie about 4 kcal/mol lower in energy than those (III and IV) with the eq-eq binding motif.

Figure 5 compares the (scaled) predicted harmonic spectra [(b)-(e)] of the four lowest lying isomers with the observed spectrum of [Ru^{II}(en^{*})₂TBOQ]²⁺•D₂ in Fig. 5(a). Band positions and assignments are included in Table S3. It is of interest to note that, in all cases, the NH stretching fundamental associated with the eq NH group $(v_h^{\rm NH_2} eq)$ is calculated to be lowest in energy, which is expected to be the proton transferred in the PCET reaction. Of these, the spectrum calculated for isomer I is in overall best agreement with the observed bands in the NH and C=O stretching regions. The situation regarding the splitting between the collective symmetric and asymmetric C=O stretches is particularly useful for structure identification. Specifically, the two isomers (II and III) that feature bifurcated binding motifs to the NH groups are calculated to yield a clear splitting $(37 \text{ cm}^{-1} \text{ and } 39 \text{ cm}^{-1}, \text{ respectively})$, while I and IV with more linear binding arrangements are more in line with the single, asymmetric C=O stretching band in the observed spectrum. Based on the fact that the calculated pattern for isomer I in the NH stretching region is closest to the observed red-shifted bands, we propose that isomer I is the most likely candidate for the docking geometry of the entrance channel complex. We note, however, that although it is calculated to lie much higher in energy, we cannot rule out a contribution from isomer IV based on the

spectrum alone since both isomers yield similar patterns for the C=O stretches. Unfortunately, detailed comparisons in the NH stretching region are not straightforward as it is well known³⁰ that multiplet structures are often observed in this region due to strong Fermi-resonance interactions between the red-shifted NH fundamentals and the v = 2 level of the NH bends.^{29–31} We illustrate this in Fig. S8, which establishes that incorporation of the Fermi resonance interactions between the predicted NH stretch and bending levels with a typical matrix element (37 cm⁻¹) recovers the multiplet character of the observed spectrum. Having identified the most likely docking arrangement in the captured reactant complex, we note that only one of the two equatorial NH2 groups that are transformed into the product (which has two equatorial NHs) is in direct contact with the carbonyl. As such, although it is an appealing scenario that the system docks in a natural position to execute a double protonelectron transfer with minimal participation of the surrounding network, this study indicates that the reaction pathway might well be more complex, involving substantial rearrangement to access the transition state structure.

IV. CONCLUSION

Using a combination of high-resolution mass spectrometry and cryogenic ion vibrational spectroscopy techniques, we have isolated a non-covalently bound complex between an ortho-quinone and a $[Ru^{II}(en^*)_2]^{2+}$ coordination compound that is known to undergo a double proton-coupled electron transfer reaction in solution. Vibrational characterization of both the isolated reactant ion and $[Ru^{II}(en^*)_2 TBOQ]^{2+}$ reveals a double complexation motif in which two of the C=O groups attach to two of the NH₂ groups with strong hydrogen bonds. This is evidenced by the appearance of intense, red-shifted bands in the NH stretching region of the vibrational

spectrum upon complexation and a strong C=O fundamental in the lower energy region of the spectrum. Calculations indicate that the binary complex most likely corresponds to TBOQ attachment to two different classes of the NH₂ groups: one axial and one equatorial with respect to the bipyridine ligand. Because the $[Ru^{IV}(en^*-H)_2]^{2+}$ product is formed with loss of H atoms from two equatorial NH₂ groups, it appears that the transition state for the 2 × PCET reaction requires substantial rearrangement from the geometry of the isolated complex in the gas phase. This is consistent with the observation that the reaction could not be promoted by collisional activation of the entrance channel complex. As such, this study establishes that emergent methods in cryogenic ion spectroscopy can indeed engage structural aspects of complex systems under study in contemporary synthesis, but also points out limitations in their ability to directly unravel complex reaction pathways.

SUPPLEMENTARY MATERIAL

See the supplementary material for supporting experimental and computational data, including sample preparation, mass spectrometric data (MS and MS/MS), vibrational predissociation spectra, experimental and calculated frequencies, Fermi resonance analysis, as well as rotatable structures (PDF).

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DATA AVAILABILITY

The data that support the findings of this study are available from the corresponding author upon reasonable request.

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