DFT Provides Insight into the Additive-Free Conversion of Aqueous Methanol to Dihydrogen Catalyzed by [Ru(trop2dad)]: Importance of the (Electronic) Flexibility of the Diazadiene Moiety

Sinha, V.; Trincado, M.; Grutzmacher, H.; de Bruin, B.

DOI 10.1021/jacs.8b09011
Publication date 2018
Document Version Final published version
Published in Journal of the American Chemical Society
License CC BY-NC-ND

Citation for published version (APA):

General rights
It is not permitted to download or to forward/distribute the text or part of it without the consent of the author(s) and/or copyright holder(s), other than for strictly personal, individual use, unless the work is under an open content license (like Creative Commons).

Disclaimer/Complaints regulations
If you believe that digital publication of certain material infringes any of your rights or (privacy) interests, please let the Library know, stating your reasons. In case of a legitimate complaint, the Library will make the material inaccessible and/or remove it from the website. Please Ask the Library: https://uba.uva.nl/en/contact, or a letter to: Library of the University of Amsterdam, Secretariat, Singel 425, 1012 WP Amsterdam, The Netherlands. You will be contacted as soon as possible.

Download date: 13 Jul 2023
DFT Provides Insight into the Additive-Free Conversion of Aqueous Methanol to Dihydrogen Catalyzed by [Ru(trop2dad)]: Importance of the (Electronic) Flexibility of the Diazadiene Moiety

Vivek Sinha, Monica Trincado, Hansjörg Grützmacher, Bas de Bruin

ABSTRACT: The mechanism for complete dehydrogenation of aqueous methanol to CO$_2$ and three equivalents of H$_2$ catalyzed by [Ru(trop$_2$dad)] was investigated with DFT (trop$_2$dad = 1,4-bis(5H-dibenzo[a,d]cyclohepten-5-yl)-1,4-diazabuta-1,3-diene). To date, this is the only catalyst that promotes the acceptorless dehydrogenation of aqueous methanol in homogeneous phase under mild conditions without the addition of an additive (base, acid, or a secondary catalyst). A detailed understanding of the mechanism of this transformation may therefore be of significant importance for the conversion of liquid organic fuels. Previous computational studies using simplified models of the catalyst suggested entirely ligand-centered reaction pathways with rather high-energy barriers for complete dehydrogenation of aqueous methanol. These, however, are not consistent with the experimental data. In the present paper, we reveal a different reaction mechanism for aqueous methanol dehydrogenation that involves metal–ligand cooperativity involving the diazadiene (dad) ligand and has substantially lower barriers, in good agreement with the experimental data. The dad moiety of the ligand actively participates in the alcohol activation mechanism. In the first step of the reaction, the dad ligand rearranges from a $\sigma$- to a $\pi$-bound coordination mode. This adjusts the electronic structure of both the metal and the ligand, leading to an enhanced Brønsted basicity of the nitrogen centers and higher Lewis acidity of the ruthenium center. As a result, concerted proton-hydride transfer to/from metal-hydride and N-protonated dad-ligand moieties becomes possible, leading to low-barrier metal–ligand cooperative elementary steps for alcohol activation and H$_2$ elimination.

INTRODUCTION

Hydrogen is an attractive energy carrier for the clean and efficient generation of electricity using fuel cell technologies. In spite of the advantages that fuel cells entail, hydrogen storage is one of the main limitations to enable a hydrogen-based energy system. Consequently, the reversible chemical fixation of hydrogen in a carrier molecule, especially a stable liquid compound, may offer a practical alternative for hydrogen handling. Methanol is a particular promising liquid organic fuel (LOF) with a gravimetric hydrogen content of 12.6 wt %. The complete dehydrogenation of a methanol/water mixture (hydrogen content 12.4 wt %) gives rise to the release of CO$_2$ and three equivalents of H$_2$. Given an environmentally benign and cost-efficient source of hydrogen coupled with an efficient recycling process of CO$_2$, methanol might become a real alternative for sustainable hydrogen storage. In this scenario the system CH$_3$OH/CO$_2$ takes the role of recyclable hydrogen carrier. The overall process would in effect mimic the natural carbon cycle. However, the possibility of reversible hydrogen storage in methanol is still not broadly commercially developed. This is mainly due to the limitations related to the highly energetically costly full dehydrogenation and the reverse hydrogenation processes. Methanol reforming in the presence of heterogeneous catalysts typically requires high temperatures (>200 °C) and high pressures (25–50 bar). The main problems at these temperatures are the competing methanol decarboxylation and reverse water gas shift reaction (RWGS), leading to CO as an undesired product. Recent advances in the reforming of methanol under mild conditions based on homogeneous catalyzed processes have raised the expectations on this fuel. With the exception of the initial work of Cole–Hamilton on methanol reforming catalyzed by a rhodium bipyridine complex, only recently were two molecular Ru complexes found that act as homogeneous catalysts allowing the conversion of aqueous methanol below 100 °C and at atmospheric pressure. In both catalytic systems chemically “innocent” ligands bind to the metal center. The catalysis is efficient and highly selective, and no CO was detected. Several theoretical studies have been reported exploring possible mechanisms of these reactions. Subsequent experimental studies used other ruthenium, iridium, or rhodium complexes. More recently, the first catalytic reactions with first-row metal complexes (Fe and Mn) were reported. But the development of efficient...
catalysts for the additive-free dehydrogenation of methanol aqueous solutions under mild conditions remains a major challenge. To date, the only system that promotes the dehydrogenation of methanol in the absence of any additives (base, acid, or a secondary catalyst) is the diazadiene ruthenium complex \([\text{Ru(trop)}_{2}\text{dad}]\) (trop$_{2}$dad = 1,4-bis(5H-dibenzo[a,d]cyclohepten-5-yl)-1,4-diazabuta-1,3-diene) (see Figure 1d for a plot of the structure). If one does not want to rely on serendipity, a detailed understanding of the elementary steps involved in the catalytic reaction is necessary for the development of catalysts with improved efficiency. In this study, we present a plausible novel mechanism on the basis of DFT calculations, considering elementary steps based on metal–ligand cooperativity that were previously not considered. This allows us to propose a new mechanism for the dehydrogenation of a methanol–water mixture to CO$_2$ and hydrogen catalyzed by the [Ru(trop$_{2}$dad)] system, which is based on metal–ligand cooperativity. The dad ligand moiety plays a crucial and active role in catalysis, which first rearranges from a $\sigma$- to a $\pi$-bound coordination mode to allow essential catalytic elementary steps.

Dehydrogenation of methanol–water mixtures by the [Ru(trop)$_{2}$dad] catalyst proceeds via four separate steps (Scheme 1a): (a) methanol is dehydrogenated (oxidized) to formaldehyde under release of the first equivalent of H$_2$; (b) formaldehyde undergoes hydration to form methanediol; (c) methanediol is dehydrogenated to formic acid and a

Figure 1. (a) The electronic structure of complex 1 can be described with two contributing resonance structures: 1A, with ruthenium in oxidation state 0 coordinated to a neutral dad moiety, and 1B, with ruthenium in oxidation state +II coordinated to a dianionic dad moiety. (b) Potential energy surface (PES) scan to drive the H$_{\text{OH}}$ in methanol (H$_{\text{OH}}$(MeOH)) to one of the N$_{\text{dad}}$ atoms. In complex A, methanol prefers to bind via the OH group to the metal center, indicating a larger contribution of resonance structure 1A to the ground state of complex 1. The vertical axis represents the relative SCF energy ($\Delta E$) (BP86/def2-SVP), while the horizontal axis represents the distance between the N$_{\text{dad}}$ atom and MeOH hydroxyl proton. (c) Transformation of complex 1 to the $\pi$-coordinated complex 1’. (d) 3D structure plots of complexes 1 and 1’. (e) Highest occupied molecular orbital (HOMO) and lowest unoccupied molecular orbital (LUMO) of complex 1 (left) and complex 1’ along with their respective energy (right) (contour plots drawn at 0.08 au).
Scheme 1. (a) Four stages (1–4) of the Stepwise Dehydrogenation of a Methanol–Water Mixture to H2 and CO2; (b) Proposed Metal–Ligand Cooperative Pathway for Dehydrogenation of Methanol–Water Mixtures by Grützmacher and Co-workers; (c) Anionic Pathway Proposed by Li and Hall via Complex 1H’− Containing a Frustrated Lewis Acid Base Pair

"Only methanol is shown as substrate molecule but also methanediol (H2C(OH)2) or formic acid (HCOOH) can be converted."

second equivalent of H2; (d) formic acid is dehydrogenated to release CO2 and the third equivalent of H2. The active [Ru(trop2dad)] catalyst is produced by protonation of the anionic hydride precursor complex [RuH(trop2dad)]− (1H−) (Scheme 1b). Compared to most of the other first/second-row transition metal complexes that are known in the literature7,8,9c,11,12 for aqueous methanol reforming, the active catalyst complex [Ru(trop2dad)] (1) (Figure 1d) does not contain phosphorus donors in the ligand.13 Complex 1H− rather features relatively soft olefinic binding sites, which act as π-donor and acceptor groups coordinated to the metal from the rigid 1,4-bis(5H-dibenzo[a,d]cyclohepten-5-yl) (abbreviated as trop) ligand. Opposite the C=C trop units are hard N-donor atoms provided by the dad (1,4-diazabuta-1,3-diene) moiety.

In the original experimental paper reported by Grützmacher, Trincado, and co-workers, the dadi moiety of the ligand was already proposed to play an important role in the mechanism, but was considered to act as a unit that reversibly stores CO2. However, while plausible at the time, these experimental observations can also be interpreted differently. The mere fact that complexes 1 and 2 are both detectable does not mean that they are necessarily involved in the same catalytic cycle. They could act as independent catalysts, simply being in equilibrium.

In a recent computational study, Li and Hall primarily focused on a mechanism proposed in the original experimental paper, in which species 1 and 2 are intermediates in the same catalytic cycle. With density functional theory (DFT) methods, they computed a pathway for the formation of complex 2 from 1H− by hydrogen transfer from a substrate molecule (methanol/methanediol/formic acid), followed by a solvent- or substrate-assisted protonation step (see Scheme 1c).14 They found that dehydrogenation of complex 2 proceeds via a very high barrier at a transition state (TS) of >50 kcal mol−1. For that reason, they proposed an alternative pathway that involves anionic ruthenium complexes15 1H′− and 2− (Scheme 1c). In their proposal, the first step involves transfer of the hydride ligand of complex 1H− from Ru to the dadi moiety, generating complex 1H′−. The resulting amido and the imine (N=C) unit in the ligand of complex 1H′− form a Brønsted base and Lewis acid pair, respectively. This frustrated Lewis-acid–base pair acts as the catalytically active site and converts methanol or methanediol in a concerted protonation of the amido moiety and hydride transfer to the imine moiety. This reaction leads to complex 2−. In their mechanistic proposal the metal acts as a spectator, while all substrate activation steps occur at the ligand. On the contrary, our recent study of the dehydrogenation of formaldehyde–water mixtures by complex 1H− revealed an active involvement of the metal center.16 A similar metal-centered reactivity can therefore be expected in the steps leading to methanol dehydrogenation. Li and Hall considered the direct dehydrogenation of complex 2− as the hydrogen production step, which was proposed to be the turnover-limiting step of the catalytic cycle. A high barrier of about 35 kcal mol−1 for the transformation 2− → 1H− + H2 was computed, which is too high and not in agreement with a fast catalytic reaction performed at 363 K.

More recently, Yang and co-workers calculated additional pathways for the dehydrogenation of methanol–water mixtures that likewise involved anionic pathways using 1H− as catalyst.17 In both of these studies the activation barriers leading to hydrogen production proved to be very high (>40 kcal mol−1).18 None of the previous computational studies took the possibility of metal–ligand cooperativity into account or considered pathways that involve neutral Ru complexes such as complex 1. Furthermore, these calculations used simplified models of the catalyst in which the dibenzo[a,d]cyclohepten-5-yl (trop) units were replaced by simple cycloheptatrienyl groups. In the present work we show that metal–ligand cooperative pathways calculated for neutral species using a full-atom treatment of the catalyst lead to reaction paths with much lower barriers. Additionally, we show that the use of simplified models of the catalyst leads to unsystematic errors and should be avoided.

**COMPUTATIONAL METHODS**

All DFT geometry optimizations were carried out with the Turbomole program19 coupled to the PQS Baker optimizer20 via the BOpt package.21 The B97D23 functional with a Turbomole def2-TZVP basis set24 for all atoms was used for optimization of geometries as minima or transition states. The resolution-of-identity (n) approximation25–27 was employed to speed up calculations. To include weak interactions, all calculations include Grimme’s...
Catalyst Activation and Electronic Structure of Complex 1. The anionic hydride complex 1H− was reported to react with a slight excess of water or a carboxylic acid to produce the active catalytic neutral complex 1 and H2.6,13 Complex 1 can be described as a resonance hybrid of two valence isomers, namely, 1A, with Ru0 coordinated to a neutral bis-imine form of the dad moiety, and 1B, which has Ru1I coordinated to a dianionic bis-amide (Figure 1a).6,13,16,30 The valence isomer 1B has larger negative charge densities on the Ndad atoms, which makes them favorable donor sites for hydrogen-bonding interactions. Resonance structure 1A has neutral Ndad atoms, which should be very weak hydrogen bond accepters. DFT calculations show that methanol prefers to interact with complex 1 via metal–proton-type interactions (Figure 1b) and does not tend to form hydrogen bonds with the Ndad atoms. This clearly points to a rather basic Ru center, as indicated in the resonance structure 1A with Ru0, which is likely the main contributor to the electronic structure of complex 1. This is further bolstered by the concave upward curvature of the potential energy surface (PES), which is obtained when the proton on the hydroxyl group of MeOH is transferred toward one of the Ndad atoms (Figure 1b). This observation clearly shows that the Ndad moieties in complex 1 are (if any) very poor proton acceptors.31 We therefore investigated the possibility that the dad moiety changes its planar coordination mode, in which only the N centers bind to the metal, to one in which the C=C double bond in the dad ligand undergoes π-coordination to the metal. This transition profoundly changes the electronic structure of the complex, leading to an increased Lewis acidity at the Ru center and simultaneously enhanced Bronsted basicity of the Ndad atoms.

None of the previous theoretical studies on this catalytic system considered the effect of π-coordination of the dad moiety to the metal center. π-Coordination of dad ligands has been experimentally observed by Vrieze and Van Koten and co-workers in low-valent ruthenium carbonyl complexes more than 30 years ago.32 Constrained geometry scan calculations were used to investigate the π-coordination of the C=Cdad bond of the dad backbone to the metal. This is a slightly endergonic process (+3.4 kcal mol−1), which proceeds over a relatively low barrier with TS1,π at +3.8 kcal mol−1 (Scheme 1c). Hence, complex [Ru(π-N,N-trop2dad)] 1 (containing a π-bound dad moiety) should be in rapid equilibrium with complex [Ru(π-trop2dad)] 1′, which contains a π-coordinated dad moiety with nonplanar and more basic Ndad atoms. Notably,...
the investigation of the interaction between \([\text{Ru(π-trop2dad)}]\) \(1'\) and methanol using similar constrained geometry scans resulted in a spontaneous rearrangement of the binding mode of the MeOH molecule (see Supporting Information for details). Starting from adduct \(A\), with the OH group of methanol bound to the Ru center, the PES scan showed that driving the C=Cdad backbone to interact with the metal center resulted in spontaneous formation of adduct \(A'\). In this complex the methanol is strongly hydrogen bonded to one of the "amido" Ndad atoms (Figure 2, bottom), while one C–H bond of the methyl group weakly interacts with Ru. This fact is highly relevant for catalysis, as it preorganizes the MeOH molecule for dehydrogenation via a cooperative mechanism, and is illustrative of the large change in electronic structure of both the metal and the ligand from the σ-bound (1) to π-bound (1’) coordination mode of the dad-ligand moiety (also see Figure 1e).

Comparison of charges obtained by natural population analysis (NPA) reveals that the transformation of complex 1 to complex 1’ results in a net increase of negative charge density on the Ndad moieties. The metal center becomes more positive during this transformation. These effects are more pronounced in the MeOH adducts \(A\) and \(A'\). For example, in complex \(A'\), the Ndad atom involved in hydrogen bonding to the MeOH molecule has a 0.10 unit higher net negative natural charge than in complex \(A\). As such, the transformation of a σ-coordinated dad moiety to a π-coordinated one significantly changes the electronic structure of the complex. The Brønsted basicity of the nitrogen atoms and the Lewis acidity of the metal ion are
enhanced. This allows binding of the methanol molecule in a way that dehydrogenation via a cooperative mechanism becomes easily feasible (vide infra).

**Simplified Atom Model vs Full-Atom Model.** The differences of the relative free energies between the κ-N-sigma bound and π-bound forms were studied using a simplified model with cycloheptenyl units instead of the complete trop-dad ligand using various functionals (Figure 3). The results clearly show that the simplified atom model systematically overestimates the relative free energy of π-complex 1' relative to σ-complex 1 (Figure 3). This may explain why previous MERPs explored with the simplified atom model did not identify complex 1' as a likely intermediate. Further analysis (see Supporting Information) shows that the complexation of methanol over complex 1 and the TS for dehydrogenation of MeOH (TS-1-MeOH) are not described satisfactorily by the simplified atom model either. Li and Hall showed examples where the simplified atom model underestimates the TS barriers.15 Our own calculations show that there are also cases where the simplified model overestimates the TS barriers (see Supporting Information). This comparative study therefore reveals that the use of simplified models of the catalyst leads to unsystematic errors and shows that a full-atom treatment is essential. We therefore performed all further mechanistic studies using only the full-atom models of the catalyst species.

**Acceptorless Dehydrogenation of MeOH to Formaldehyde.** The computed pathway for dehydrogenation of methanol to formaldehyde promoted by complex 1 is shown in

---

Figure 4. Computed free energy profile for dehydrogenation of methanol to formaldehyde over complex 1 (BP86//B3LYP (Water)). Transient bonds in transition states are drawn as dashed brown lines.

| Relative Gibbs free energy (ΔG°298 K) values (BP86//B3LYP (Water)) are reported in kcal mol⁻¹ with respect to B-HCOOH-H₂O. The enthalpy (ΔH°298 K) for the reaction B + (HCCOH-H₂O) is given in parentheses.33 Transient bonds in transition states are drawn as dashed brown lines. |

**Scheme 3. Direct (Unassisted) and Formic Acid Mediated Pathways for Dehydrogenation of Complex B**

(Original content continues with figures and equations related to the pathways and calculations.)
Scheme 4. Dehydrogenation of Methanediol to Formic Acid

Scheme 2 and Figure 4. In the first exothermic step, methanol binds with its Lewis-basic oxygen center to the Lewis-acidic Ru center of 1 to give 1-CH$_3$OH. There is another isomer A, only slightly higher in energy ($\Delta G^\circ_{298 K} = +1.2$ kcal mol$^{-1}$), in which the OH group is hydrogen bonded to the metal center. This is a reflection of the remarkable electronic flexibility of the [Ru(trop$_3$dad)] complex. There is a third possible binding mode of the CH$_3$OH ligand that leads to A’. In this isomer, the...
producing complex 1-H$_2$ generated complex 1 substrate-mediated transition states leading to the dehydro-
unassisted reaction, $\Delta$G° = +15.5 kcal mol$^{-1}$. Under the experimental conditions the process is driven by constant removal of H$_2$ (and CH$_3$O; vide infra).

**Dehydrogenation of Formaldehyde–Water Mixtures to Formic Acid.** The computed pathway for dehydrogenation of methanediol (formed by hydration of the formaldehyde released in a previous step$^{6,16}$) is shown in Scheme 4. As computed for the methanol dehydrogenation steps, methanediol can bind to the metal center in three different modes: (1) via its Lewis basic oxygen center (complex 1-CH$_2$(OH)$_2$); (2) via a metal–proton interaction involving one of the OH groups (complex A-CH$_3$(OH)$_2$), and (3) as in the methanediol adduct A$^\prime$-CH$_2$(OH)$_2$, wherein the dad moiety adopts a $\pi$-coordination mode, rendering the N$_{dad}$ atoms sufficiently basic to interact with both protic OH groups via H-bond interactions and the Ru center sufficiently Lewis acidic to interact with one of the C–H units of the substrate (Ru–HC = 2.16 Å). The latter isomer (A$^\prime$-CH$_2$(OH)$_2$) is the entry to product formation. Concerted metal–ligand cooperation allows dehydrogenation of methanediol over the TS-1-CH$_2$(OH)$_2$ at +17.6 kcal mol$^{-1}$ to give the hydrogen-bonded formate complex B$^\prime$-HCOO$^-$ in an exergonic reaction (−12.1 kcal mol$^{-1}$). The whole reaction sequence is thus very similar to the one described above for methanol dehydrogenation over TS-1-CH$_3$OH.

**Dissociation of formic acid (HCOOH) from B$^\prime$-HCOO$^-$ to form the hydrogenated complex B most likely proceeds via complex B$^\prime$-HCOOH (endergonic; +7.3 kcal mol$^{-1}$), which rearranges to B-HCOOH (exergonic; −1.3 kcal mol$^{-1}$) upon decoordination of the Ru center from the C=C$_{dad}$ backbone (Scheme 4). From there, the reaction proceeds by dissociation of formic acid (−4.1 kcal mol$^{-1}$) to form B and release of H$_2$ (+2.3 kcal mol$^{-1}$) to regenerate complex 1 (vide supra). The reaction from B$^\prime$-HCOOH to 1 is overall slightly endergonic (+4.2 kcal mol$^{-1}$), but the process is driven by continuous...
removal of gaseous H2 under the experimental conditions. The overall dehydrogenation of methanediol to formic acid is found to be exergonic by $-7.9 \text{ kcal mol}^{-1}$ with respect to complex 1.

**Dehydrogenation of Formic Acid to CO2 over Complex 1.** The computed mechanism for the dehydrogenation of formic acid to produce CO2 and H2 is shown in Scheme 5. Dehydrogenation of formic acid over complex 1 proceeds in a similar manner to that of methanol (vide supra). In contrast to methanediol and methanol, protonation of one of the N_{dad} moieties by formic acid and subsequent coordination of the formate to the metal center via one of the oxygen atoms to produce complex 1-HCOOH is exergonic. This species is the resting state of the catalytic cycle for dehydrogenation of formic acid. Complex 1-HCOOH undergoes an endergonic rearrangement to form A-HCOOH, where formic acid interacts with the metal center via a metal−proton interaction, analogous to complex A. Coordination of the C═C_{dad} backbone to the metal center generates the π-coordinated complex A’-HCOOH. Complexes A-HCOOH and A’-HCOOH should exist in equilibrium with each other. In a similar manner to that computed for the dehydrogenation of methanol via TS-1-CH$_3$OH and dehydrogenation of methanediol via TS-1-CH$_2$(OH)$_2$, a concerted Noyori−Morris-type mechanism via transition state TS-1-HCOOH at +15.8 kcal mol$^{-1}$ leads to the

**Figure 7.** Comparison of transition state for hydride transfer (TS-1-HCOOH) with previous work$^{16}$ and involving an anionic system (BP86//B3LYP (Water)). Transient bonds in the transition states are drawn as dashed brown lines.

$^{33}$All relative Gibbs free ($\Delta G_{298 K}^{\circ}$) energy values (BP86//B3LYP (Water)) are reported in kcal mol$^{-1}$. The number in parentheses is the enthalpy change ($\Delta H_{298 K}^{\circ}$) of this step$^{33}$. Transient bonds in transition states are drawn as dashed brown lines.
formation of the CO$_2$ adduct B$'$. Exergonic loss of CO$_2$ from B$'$-CO$_2$ generates the complex B$, which rearranges in an exergonic process to complex B. As discussed above, this complex finally loses H$_2$ to regenerate complex 1 (vide supra), thus completing the catalytic cycle. The overall dehydrogenation of formic acid to CO$_2$ and H$_2$ is found to be exergonic by $-7.5$ kcal mol$^{-1}$.

**Quantifying the Effect of Metal–Ligand Cooperativity.** Dehydrogenation of formic acid by complex 1 provides an exemplary case to quantify the effect of metal–ligand cooperativity arising from the transformation of a σ-bound (complex 1) to a π-bound (complex 1') dad moiety. The experimental turnover frequency (TOF) for formic acid dehydrogenation over complex 1 is one of the highest reported (24 000 h$^{-1}$), indicating a relatively low barrier for oxidation of formic acid to CO$_2$ promoted by complexes 1/1'. In our earlier work, we calculated the dehydrogenation of formic acid by complex 1 in the absence of π-coordination of the Ru center with the C=C$_{dad}$ backbone.$^{16}$ In addition, Yang and co-workers have reported an anionic TS for hydride transfer from formate to complex 1 to produce CO$_2$ and H$_2$ using a simplified atom model.$^{17}$ We recomputed this cycle using a full-atom model of complex 1. Figure 7 summarizes these findings. Within the manifold of calculated reactions, formic acid dehydrogenation via TS-1'-HCOOH proceeds with the lowest barrier of +15.8 kcal mol$^{-1}$. All of the transition states considered in Figure 7 involve a hydride transfer from an anionic formate moiety to the metal center. To reach an energetically favorable TS for the dehydrogenation of formic acid/formate, the Ru center should be highly Lewis acidic in order to be a strong hydride acceptor. Furthermore, little steric hindrance in the TS geometry and a favorable stabilization of the formate moiety by hydrogen bond interactions will lower the activation barrier. TS-1'-HCOOH involves a classical beta-hydride transfer from an O-coordinated formate to Ru. Not only does the molecular geometry of TS-1'-HCOOH suffer from steric hindrance, it also lacks hydrogen bond stabilization of the formate moiety in the TS. Moreover, there is no particular enhancement in the Lewis acidity of the metal center. It is therefore a little surprising that TS-1'-HCOOH has the highest energy (+20.1 kcal mol$^{-1}$) among the activated complexes shown in Figure 5. The anionic TS-1''-HCOO$^-$ is sterically less hindered compared to TS-1'-HCOOH, thus leading to a somewhat lower barrier of +18.7 kcal mol$^{-1}$. However, it should be noted that in the absence of any H-bond stabilization of the formate anion by explicit solvent molecules, the computed energy of the activated complex TS-1''-HCOO$^-$ is actually unrealistic and most likely considerably underestimated.$^{39}$ The geometry of the neutral transition state TS-1-HCOOH features a sterically favored arrangement of catalyst and substrate in a six-membered ring, a stabilization of the formate moiety by hydrogen bonds, and a π-bound dad ligand, which increases the Lewis acidity of the Ru$^{11}$ center. The combination of these factors leads to a relatively low barrier for CO$_2$ production.$^{34}$ The calculated barrier of +15.8 kcal mol$^{-1}$ is consistent with the experimentally observed high TOF values for formic acid dehydrogenation promoted by complex 1. Similar arguments can be used to rationalize the relatively low barriers for dehydrogenation of methanol and methanediol (see the Supporting Information).

**SUMMARY AND CONCLUSIONS**

In summary, the mechanism for complete dehydrogenation of aqueous methanol to CO$_2$ and three equivalents of H$_2$ catalyzed by [Ru(trop,dad)] was investigated with DFT employing the full-atom model of the catalyst. Previous computational mechanistic studies on this system using a simplified model of the catalyst proposed a fully ligand centered anionic pathway, with the metal acting as merely a spectator. This pathway requires crossing of high TS barriers for dehydrogenation of aqueous methanol, inconsistent with the experimental results. In this study we show that the dad ligand moiety plays an active role as a cooperative ligand with the metal in the dehydrogenation mechanism, leading to much lower TS barriers. We find that the dad moiety rearranges from a σ-bound to a π-bound coordination mode, resulting in an enhanced Brunsveld basicity of the amido nitrogen atoms and a stronger Lewis acidity of the metal center. This crucial rearrangement allows a favorable coordination of the substrate to the Ru center to open effective catalytic reaction channels for substrate dehydrogenation. The geometric and electronic fluxionality of the dad ligand moiety gives access to metal–ligand cooperativity, providing pathways with low activation barriers for alcohol activation and H$_2$ production in an aqueous medium. Overall, dehydrogenation of the substrate (methanol/methanol/methanediol/formic acid) occurs via a Noyori–Morris-type mechanism to form complex [Ru(H)(tropNH-CH==CH-Ntrop)] B, which is a hydrogenated form of complex 1. Subsequent dehydrogenation of complex B can occur directly or via substrate/solvent-assisted pathways, both having relatively low activation barriers. The cooperative function of π-bound dad ligand moieties might well be a more general feature of catalysts with diazadiene ligands, and it could therefore be key for the design of improved catalysts for hydrogen production from methanol–water mixtures.

**ASSOCIATED CONTENT**

Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/jacs.8b09011.

Additional computational data, other possible reaction pathways, xyz coordinates, and benchmark calculations (PDF)

**AUTHOR INFORMATION**

Corresponding Authors

*hgruetzmacher@ethz.ch
*b.debruin@uva.nl

ORCID

Vivek Sinha: 0000-0002-6856-9469
Bas de Bruin: 0000-0002-3482-7669

Notes

The authors declare no competing financial interest.

**ACKNOWLEDGMENTS**

This work was supported by the Schweizer Nationalfonds (SNF 200021_162437/1), Eidgenössische Technische Hochschule (ETH) Zürich, The Netherlands Organization for Scientific Research NWO-CW (VICI project 016.122.613), the NWO-Shell Computational Sciences for Energy Research Initiative (project 13CSER003), and the RPA Sustainable Chemistry of the University of Amsterdam (UvA). The calculations were carried out on the Dutch national e-infrastructure with the support of the SURF cooperative.
Anionic complexes are rarely used for (or even observed to
be active in) catalytic hydrogen transfer or dehydrogenation processes. See also ref 9a.


(18) Li and Hall calculated proton abstraction from complex 2 to generate 2+ to be uphill by 8.6 kcal mol⁻¹ in aqueous media, meaning that 2 must be considered as the resting state for dehydrogenation to regenerate complex 1H⁺. Yang computed the deprotonation of complex 2 to generate 2− to be uphill by 3.4 kcal mol⁻¹.

(19) University of Karlsruhe and Forschungszentrum Karlsruhe GmbH. TURBOMOLE V7.1; TURBOMOLE GmbH, 2016.


(30) In our previous work (ref 13) we found that one of the possible valence isomers of complex 1 can be described as Ru⁺ coordinated to a monoanionic d-d radical, with spin densities on the metal and ligand center being antiparallel to each other. Detailed calculations showed that this singlet-biradical character does not influence the catalytic reaction pathway for dehydrogenation of a methanol–water mixture. Detailed analysis is presented in the Supporting Information (this paper). For the sake of simplicity, we have not involved this singlet-biradical character in the discussion here.

(31) We also considered C₄ as internal base, but the resulting TS barriers for the formation of formaldehyde were found to be relatively high (>30 kcal mol⁻¹). Further details are presented in the Supporting Information.

(33) The substrate methanol is also the solvent in the system. Therefore, the translational entropy contribution to the methanol binding step should be virtually zero, making enthalpy a better estimate for the free energy of this step in solution than the gas phase free energy based on vibrational analyses only.

(34) Note that introducing explicitly solvent interactions has only a minor influence on the MERP pathways and relative free energies of the minima and transition states (see Supporting Information). The trop ligands in complex 1 form a relatively hydrophobic pocket around the metal center, which explains the minimal effects of introducing explicit solvent molecules on the computed MERP of complex 1.

(35) We also computed the pathways for methanol dehydrogenation to formaldehyde by [Ru(trop-dae)] complex 2 (derived from complex 1 by hydrogenation of the dad backbone). See Supporting Information for details.


(37) The direct hydrogen production from complex B’ was found to occur via a high TS barrier process (\(\sim\)40 kcal mol\(^{-1}\), BP86/def2-TZVP) and was not investigated further.

(38) The mechanistic step forming a nonclassical dihydrogen complex which leads to the regeneration of the catalyst is commonly proposed in dehydrogenation (or the microscopically reverse hydrogenation) reactions. See, for example: Jessop, P. J.; Morris, R. H. Reactions of transition metal dihydrogen complexes. *Coord. Chem. Rev.* 1992, 121, 155–284.