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Anirban Bhattacharjee, Alexander K. H. Weiss, Vincent Artero, Martin J Field, Thomas S. Hofer. Electronic structure and hydration of tetramine cobalt hydride complexes.. *Journal of Physical Chemistry B*, 2014, 118 (20), pp.5551-5561. 10.1021/jp502651s . hal-01069163

HAL Id: hal-01069163

<https://hal.science/hal-01069163v1>

Submitted on 29 May 2018

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Electronic Structure and Hydration of Tetramine Cobalt Hydride Complexes

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Abstract

In this work we have studied two hydrido-tetramine-cobalt(III) complexes using a mixture of computational techniques. These species were chosen as simple and computationally tractable models of the Co(III)-hydrido compounds that are known to be important intermediates in the catalytic cycles of hydrogen evolution mediated by the cobaloxime complexes. We have performed both static density functional theory (DFT) calculations of the complexes in implicit solvent and adaptive hybrid DFT/molecular mechanical (MM) molecular dynamics (MD) simulations in explicit solvent, and compared our results to the experimental structural and spectral data that is available for one of the compounds. A principal aim of the study has been to provide a benchmark for future work on cobaloxime and other hydrogen-evolving catalysts using adaptive DFT/MM MD methods.

Keywords

Cobalt hydrides, density functional theory, electronic structure, molecular dynamics simulations, qc/mm potentials

Introduction

Hydrogen evolution is a key process in the generation of fuel from solar energy, and much research has gone into the design of cheap and robust catalysts that can carry out this reaction. In this context, cobaloxime complexes containing the relatively abundant metal, cobalt, are among the most efficient molecular catalysts that are known.^{1,2} It is now generally accepted that the catalytic cycle for hydrogen evolution from these species proceeds via protonation of the Co(I) species,³⁻¹⁵ yielding a Co(III)-hydrido-cobaloxime (Co(III)-H) intermediate that, after further reduction to the Co(II)-H state,^{5,16-18} can evolve dihydrogen through either protonation of the hydride moiety or bimolecular reductive elimination.

In recent work, we have been interested in the mechanism of hydrogen evolution from a number of cobaloxime complexes and in the investigation of the intermediates that occur during their catalytic cycles. In one study, we characterized, using a mixture of experiment and theory, the crucial hydrido-cobaloxime compound that is central to hydrogen-evolution.^{19,20} The theoretical part of this work employed density functional theory (DFT) quantum chemical (QC) calculations with an implicit model of solvent. Such methods can give an accurate description of electronic structure, reactivity and other aspects of a complex, but they are essentially static and neglect potentially-important dynamical effects, either within the complex itself or between it and the surrounding solvent molecules.

DFT methods are probably the minimum level of QC theory that is necessary to obtain a reasonable description of transition-metal complexes. However, molecular dynamics (MD) simulations of sufficient length with pure DFT potentials are computationally very challenging and limited to quite small numbers of atoms. A viable alternative is to employ mixed DFT/molecular mechanical (MM) potentials that represent the complex and a small number of surrounding solvent molecules quantum chemically and treat the rest of the system with a simpler MM force-field approach.

In the current work, we employ a hybrid DFT/MM potential in an initial investigation of the dynamics and solvent structure of hydrido-Co(III) compounds. To start with, we have chosen to simulate two smaller hydrido-tetramine-cobalt(III) complexes, instead of the larger and more complicated cobaloxime compounds. The simulated complexes, which we denote by $[\text{HCo}(\text{en})_2]^+$ and $[\text{HCo}(\text{tmen})_2]^+$, are shown schematically in Figure 1. We note that the methylated tetramine complex, $[\text{HCo}(\text{tmen})_2]^+$, was synthesized by Rae *et al.*,²¹ is surprisingly stable, and is one of the few cobalt hydrides for which a crystal structure and full spectroscopic information in water exists. It therefore serves as a reference to which future studies of larger cobalt-complexes and other transition metal hydrides can be compared.

Figure 1

The outline of the paper is as follows. Section describes the methods that we have used to simulate the complexes and to analyse the results, section discusses the results of our

calculations, and section concludes.

Methods

We performed pure QC calculations on the complexes using DFT and more sophisticated *ab initio* methods, as well as hybrid potential QC/MM simulations within the quantum mechanical charge field molecular dynamics (QMCF-MD) approximation. We describe each in turn.

Electronic Structure Calculations

Standalone QC calculations were carried out with the ORCA quantum chemical program.²² For the DFT and time-dependent DFT (TDDFT) calculations, we employed both the BP86^{23,24} and B3LYP^{25,26} functionals, along with an Ahlrichs TZVP basis set.²⁷ The combination of the BP86 functional and triple- ζ quality basis sets has been shown to perform well for other cobalt complexes, such as the corrinoids in their ground and excited states.^{28–31} In our previous article, we also found that the B3LYP functional, with a TZVP basis, produces good transition energies for the hydrido tetramine compound using the TDDFT formalism. Solvent effects were included using the implicit COSMO model for water.³²

To complement the DFT calculations, we also employed an *ab initio* CASSCF method. Due to computational expense we employed a TZVP basis for the cobalt, hydride and nitrogen atoms and a smaller SVP basis for the remaining atoms. In itself, the CASSCF method is not adequate to describe dynamic correlation, so it is common to supplement it with a perturbative correction, such as CASPT2.²⁸ We chose the strongly contracted version of NEVPT2^{33,34} that has been implemented in the ORCA program and has an advantage of being free from intruder states. An alternative approach would be to employ a MRCI method, although these are computationally very demanding even for complexes of the size that we are studying. We have preferred, therefore, to leave these types of calculations for

the future.

The QMCF-MD Methodology

The QMCF-MD methodology is an adaptive QC/MM approach that has been used extensively for the study of solvated species. We describe it only briefly here as a number of detailed reports and reviews on the approach already exist. Readers are referred to references^{35–39} for further information.

In a QMCF-MD simulation, the system is partitioned into a QC and an MM region. The QC region is further subdivided into a central core, which contains the solute and some solvent molecules, and an external “intermediate” layer that contains only solvent species. The size of the intermediate layer is such that all non-Coulombic interactions between the solute species in the core and the solvent molecules in the MM region can be neglected. As a result, the particles within the core zone are affected only by forces from particles within the QC region, F^{QC} , and the Coulombic terms between QC and MM atoms:

$$F_q^{\text{Core}} = F_q^{\text{QC}} + \sum_{m=1}^M \frac{q_m q_q}{r_{mq}^2} \left[1 + 2 \left(\frac{\epsilon + 1}{2\epsilon - 1} \right) \left(\frac{r_{mq}}{r_c} \right)^3 \right] \frac{r_{mq}}{\|r_{mq}\|} \quad (1)$$

In this equation, the subscripts m and q refer to MM and QC particles, respectively, q is an atomic charge, r_{mq} is an internuclear distance, ϵ is the permittivity of water and M is the number of atoms in the MM region within the cutoff range, r_c . The charges on the MM atoms are fixed throughout the simulation and are determined by the force field that is being used. By contrast, the charges on the QC atoms are redetermined at every MD step using a population analysis, thereby accounting for the geometrical changes that occur during the dynamics. We have previously shown that the use of Mulliken populations⁴⁰ and a Dunning DZP basis set ensures compatibility between the charges on the QC atoms and those of the MM force field.^{41–44} The last part of the expression in square-brackets results from the application of a reaction-field correction⁴⁵ that adjusts for the error introduced by

the Coulombic cutoff.

The force expression with the QC intermediate layer is identical to that of the core region, except that there are in addition non-Coulombic coupling contributions, F_{mq}^{nC} :

$$F_q^{\text{Layer}} = F_q^{\text{QC}} + \sum_{m=1}^M \frac{q_m q_q}{r_{mq}^2} \left[1 + 2 \left(\frac{\epsilon + 1}{2\epsilon - 1} \right) \left(\frac{r_{mq}}{r_c} \right)^3 \right] \frac{r_{mq}}{\|r_{mq}\|} + F_{mq}^{nC} \quad (2)$$

Particles in the MM region of the simulation box experience forces according to:

$$F_m^{\text{MM}} = \sum_{m,n \text{ pairs}}^M F_{mn}^{\text{MM}} + \sum_{q=1}^{Q_1+Q_2} \frac{q_m q_q}{r_{mq}^2} \left[1 + 2 \left(\frac{\epsilon + 1}{2\epsilon - 1} \right) \left(\frac{r_{mq}}{r_c} \right)^3 \right] \frac{r_{mq}}{\|r_{mq}\|} + \sum_{q=1}^{Q_2} F_{mq}^{nC} \quad (3)$$

where Q_1 and Q_2 are the number of QC atoms in the the core and the intermediate layer regions of the QC region, respectively, and the term F_{mn}^{MM} indicates the force between the pair of MM atoms m and n .

To ensure a continuous transition with respect to exchange of solvent molecules between the QC and MM regions, a smoothing function is applied to the forces of the atoms at the boundary of the intermediate layer. The forces on these atoms has the form:

$$F_q^{\text{Smooth}} = S(r) * (F_q^{\text{Layer}} - F_q^{\text{MM}}) + F_q^{\text{MM}} \quad (4)$$

where F_q^{Layer} is the force for the atom given in equation 2, F_q^{MM} is the force calculated as if the atom were in the MM region, and the smoothing function, applied between the distances r_{\min} and r_{\max} , is:

$$S(r) = \begin{cases} 1 & r \leq r_{\min} \\ \frac{(r_{\max}^2 - r^2)^2 (r_{\max}^2 + 2r^2 - 3r_{\min}^2)}{(r_{\max}^2 - r_{\min}^2)^3} & r_{\min} < r \leq r_{\max} \\ 0 & r > r_{\max} \end{cases} \quad (5)$$

Simulations

The two Co(III)-hydride complexes that we simulated were placed in cubic simulation boxes with 2000 water molecules, giving box lengths of 39.2 and 39.3 Å for the $[\text{HCo}(\text{en})_2]^+$ and $[\text{HCo}(\text{tmen})_2]^+$ complexes, respectively. Initial simulations of several hundred picoseconds and at constant temperature and pressure were done with an MM force-field using generalised Amber Force Field (GAFF)⁴⁶ force field parameters.

Following this initial equilibration phase, the QMCF-MD methodology was applied. Atoms in the QC region were treated at the DFT level of theory using the BP86 functional along with the 6-31G(d,p) basis for the remaining atoms. Diffuse basis functions were not included because they substantially increase the computation time without markedly affecting the results. The radii of the core, intermediate layer and smoothing zones were set to 0.0-2.0 Å, 2.0-7.0 Å and 6.8-7.0 Å, respectively. For the atoms in the MM region, the flexible SPC-mTR⁴⁷ water was employed. Charges on the atoms in the QC region, necessary for the QC/MM Coulombic interactions, were obtained via Mulliken analysis at each MD timestep and a long-range reaction-field correction⁴⁸ was applied to the same interactions using a water permittivity of 78.3. Simulations were performed at 298 K using Berendsen’s constant temperature algorithm⁴⁹ with a timestep of 0.2 fs for a duration of 25 ps, of which the first 5 ps were taken to be equilibration. To give an idea of the time required for these simulations, they each took approximately six months of computer time running in parallel on an eight-core machine using the Turbomole package as the engine for the QC calculations.⁵⁰

Structural Analyses

A range of standard structural analyses were performed, including the determination of radial (RDF) and angular (ADF) distribution functions. In addition, we employed the SLICE formalism^{39,51} to obtain three-dimensional distributions giving a detailed picture of the hydration of the complexes. For such analyses, the simulation trajectories were superimposed

by aligning the solute to a predefined reference geometry.

The SLICE method can be seen as the equivalent of computer tomography imaging applied to a simulation trajectory. Planes parallel to a predefined plane are first employed to segment the system and then the positions of selected target atoms are projected onto these planes. An atom’s contribution to the density distribution is then computed via a weighting function:

$$w(i, j) = e^{-\alpha_i*(r_i-r_j)^2} \tag{6}$$

where r_i denotes the coordinate of a target atom and r_j is the corresponding point on the plane. α_i is chosen empirically and determines how quickly the exponential function decays.

SLICE analysis yields much more information than spherical analysis schemes because of its use of a number of planes to collect projected data. A simplified version of SLICE is the two-dimensional particle mapping method⁵¹ in which the reference plane is defined with respect to two bond vectors issuing from a common atom. Taking an NH₂ group as an example, the plane would pass through the N atom and have as its normal the cross-product of the two N–H bond vectors.

Laguerre Tessellation and Dynamics

To analyze hydration layers around the solute, a space decomposition approach employing Laguerre tessellation^{52,53} was implemented using the Voro++⁵² library. In this method, the Voronoi decomposition⁵⁴ for indistinguishable particles is extended to describe distinguishable atoms using standard van der Waals radii as the distinguishing descriptors.⁵⁵ The analysis proceeds by centring the simulation boxes at the Co atoms and identifying individual layers of solvent hydration around the complexes — rather like the layers of an onion. Such layers are similar to hydration shells obtained from pair distribution functions, although small differences may occur due to the different definitions of nearest neighbours.

Once the hydration layers have been identified, further analysis was applied to obtain

layer-specific properties. In this work, we employed a mean residence time (MRT) analysis with direct counting⁵⁶ to describe the residence time of water molecules within distinct layers. We chose a window spacing of $t^*=0.5$ ps to reflect average hydrogen-bond lifetime, which has been reported to be of this magnitude.^{57,58} The number of attempts to achieve one lasting exchange event for a given time window (i.e. $t^*=0.5$ ps), with respect to a window size of $t^*=0.0$ ps, when all displacements are taken into account, is described by the ratio of exchange (R_{ex}) value:

$$R_{\text{ex}} = \frac{N_{\text{ex}}^t}{N_{\text{ex}}^{t^*}} \quad (7)$$

To analyze residence times within distinct hydration layers we modified an approach developed by Chanda *et al.* for analyzing hydrogen-bond properties.⁵⁹ Their method defines two time correlation functions (TCFs) as follows:

$$C(t) = \frac{\langle h(0)h(t) \rangle}{\langle h \rangle} \quad (8)$$

$$S(t) = \frac{\langle h(0)H(t) \rangle}{\langle h \rangle} \quad (9)$$

Here $C(t)$ is the intermittent hydrogen-bond time correlation function, which represents the structural relaxation of hydrogen-bonds (including breaking and re-forming events), whereas $S(t)$ is the continuous hydrogen bond time correlation, which provides a strict definition of their lifetime. The function $h(t)$ is unity when a particular pair of sites is hydrogen-bonded at time t and zero otherwise, whereas $H(t)$ is unity only if the pair of sites is continuously bonded between the times 0 and t .

In our version of the analysis, we redefine the functions $h(t)$ and $H(t)$ to reflect the presence of a molecule in a particular hydration layer, rather than the occurrence of a particular hydrogen-bond. With this modification, the $C(t)$ and $S(t)$ functions now measure intermittent and continuous hydration layer occupations, respectively. To aid in their interpretation

we also fit the $C(t)$ and $S(t)$ functions to double-exponential expressions of the form:

$$y = a \exp(-t/\tau_l) + (1 - a) \exp(-t/\tau_s) \quad (10)$$

where the τ values are the long (l) and short (s) time contributions to the correlation function, respectively.

Results and Discussion

Electronic Structure Calculations of $[\text{HCo}(\text{tmen})_2]^+$

For the pure QC calculations, we focused upon the methylated $[\text{HCo}(\text{tmen})_2]^+$ complex for which experimental data exists. The crystal structure of $[\text{HCo}(\text{tmen})_2]^+$ from the X-ray diffraction study²¹ was used as the starting point for our study. Hydrogens were added and the complex was geometry optimized at the DFT level of theory using an implicit solvent model of water. Note that the cobalt has, in addition to the hydride, a water molecule ligand that was treated explicitly. Agreement between the experimental and theoretical structures is good with the Co–H bond length after optimization being found to be 1.44 Å. The BP86 and B3LYP functionals produce very similar geometries with both the Ahlrichs TZVP basis and the larger, hybrid cc-pVTZ/TZVP set.

After these initial calculations we concentrated on an investigation of the spectra using both TDDFT and *ab initio* methods. A summary of these results is shown in Table 1. *Table 1* Overall, the performance of the TDDFT method is good. The higher energy band at 340 nm is well reproduced by all the methods. Agreement for the lower energy band at 444 nm is also satisfactory, with the B3LYP functional being slightly better than BP86 and, interestingly, the smaller 6-31G(d,p) basis better than TZVP.

If we consider the B3LYP/TZVP spectrum, the experimental band at 444 nm corresponds to a superposition of two approximately equally intense transitions at 415 and 418 nm,

whereas the band at 340 nm is comprised of an intense band at 326 nm and a less intense one at 322 nm. All these bands involve transitions between the HOMO and HOMO-1, which are of metal d-type, and the LUMO and LUMO+1. The LUMO is mainly a σ anti-bonding type orbital between metal d- and donor nitrogen p-orbitals in the equatorial plane, whereas the LUMO+1 is the σ anti-bonding orbital between the axial hydrido ligand and metal d-orbitals. These orbitals are illustrated in Figure 2. The lower energy transitions at 415 and 418 nm are dominated by transitions of type Co(d) to Co(d)-N(p) σ anti-bonding, whereas the higher energy bands at 322 and 326 nm are mainly of type Co(d) to Co(d)-H(s) σ anti-bonding. *Figure 2*

For the *ab initio* calculations, the choice of an active space in the reference CASSCF is crucial to the success of the method. In this paper, we have decided to focus on smaller active spaces and leave discussion of larger spaces to related work that we have been doing on the spectra of cobalt-hydride complexes.⁶⁰ These larger spaces include those that incorporate an additional set of correlating cobalt “4d” orbitals to account for the so-called “double-shell effect”.

As an initial guess in this work, we assumed a d_6 configuration for the cobalt, and started with the minimum active space of 6 electrons in 5 orbitals, which we denote CAS(6,5). The resulting transition energies were then further refined with the NEVPT2 correction to correct for the lack of dynamic correlation. We found that state averaging among at least the lowest 10 singlet and 3 triplet roots was essential for obtaining good transition energies. In this case, there is a single lower energy band at 415 nm and two higher energy bands of comparable intensity at 339 and 341 nm. Although the energies are reasonable, the intensity of the lower energy band is severely underestimated.

Increasing the size of the active space by the inclusion of two occupied and two virtual orbitals gives a CAS(10,9) model. This greatly improves the intensities compared to the CAS(6,5) model, while maintaining the quality of the transition energies. The important orbitals involved in these transitions are shown in Figure 3. For convenience we label them as *Figure 3*

HOMO and LUMO even though all the orbitals in the active space can have partial occupancies. In this case, the HOMO has an occupancy of 1.53, with all the lower orbitals having an occupancy greater than this. By contrast the LUMO and LUMO+1 have 0.89 and 0.41 electrons, respectively, whereas the occupancies of the higher orbitals are much smaller (~ 0.01). The orbitals HOMO-2 to HOMO are predominantly of metal d-type, whereas the HOMO-3 and HOMO-4 are Co-H and Co-N σ bonding orbitals, and the LUMO and LUMO+1 are Co(d)-N(p) and Co-H σ anti-bonding orbitals, respectively. The CASSCF ground state is dominated ($> 90\%$) by the reference, doubly-occupied configuration, [222220000]. The transitions at 415 and 421 nm are principally of metal d-orbital to LUMO (Co(d)-N(p) σ anti-bonding) character, whereas the higher energy bands at 313 and 317 nm are mostly of Co(d) to Co-H σ anti-bonding type.

A comparison of the TDDFT and CASSCF/NEVPT2 results shows that they are of comparable quality and give very similar interpretations of the transitions. In this case, therefore, TDDFT is much more efficient than the *ab initio* method (even with relatively small active spaces), as it is much less computationally expensive, as well as being much more straightforward to apply.

QMCF-MD Simulations

This section discusses the simulation results. Equilibration in terms of system energy and temperature were achieved during the first 5 ps of the 25 ps simulations and so we have only used the last 20 ps of data in our analysis.

Properties of the Complex

A summary of the properties of some pertinent internal coordinates of the complexes during the simulation are provided in Table 2. The two molecules display similar behaviours with significant variations of the internal coordinates around their average values. Perhaps the most marked difference between the complexes concerns the maximum and minimum values *Table 2*

of the Co–N bond lengths that are observed. Closer inspection indicates that these extremes are attained only once in the $[\text{HCo}(\text{tmen})_2]^+$ trajectories in events that last 100 fs or less.

The dihedrals in the tables show that the central CoN_4 cores of the complexes maintain planarity throughout the simulations although they have substantial flexibility. Plots of the C–N–N–C dihedrals in the two complexes are illustrated in Figure 4. These indicate that there are two peaks for each complex that are centred around $\pm 45^\circ$, respectively. In the case of the methylated complex these peaks are separate with no exchange between them on the timescale of the simulation. However, the unmethylated complex flips conformations several times with quite significant periods spent with both dihedrals within the same peak. *Figure 4*

The remaining heavy atom C–C–C–C and C–C–C–N dihedral degrees of freedom for the methylated complex have similar dynamics to their C–N–N–C counterparts with oscillations within well-defined peaks but no exchange between different conformations. By contrast, the C–C–C–H dihedral angles show very broad distributions within the range $[-180^\circ, 180^\circ]$, indicating that the methyl groups can readily rotate during the simulation.

Statistics for the partial charges of the central atoms in the complexes from the simulations are presented in Table 3. Charges are calculated with both the B3LYP and BP86 functionals and using three different charge approximations, namely electrostatic potential fitting (ESP), Löwdin and Mulliken analyses. As might be expected the charges values vary quite widely, depending upon the analysis. The hydride has a small magnitude charge in all cases, although only the ESP method makes it negative. The cobalt atom is always positive and the nitrogen atoms are strongly negative except with the Löwdin method for which they are approximately neutral. Overall, the cobalt is less positive in the $[\text{HCo}(\text{tmen})_2]^+$ complex and the nitrogens are more negative, whereas the B3LYP functional favors larger magnitudes for the cobalt and nitrogen charges than BP86. The decrease in partial charge of the cobalt upon methylation of the complex, irrespective of the employed functional and population analysis scheme, is consistent with the weak electron donor property of the methyl groups. *Table 3*

Power spectra of the complexes’ vibrational modes were computed via Fourier transform

of the atoms' velocity auto-correlation functions. The spectra are presented in Figure 5 and *Figure 5* depict both common features and clear differences in the fine structure. The methylated species has more pronounced peaks overall, suggesting a more rigid structure and better defined modes. The Co-Hydride spectra of the methylated complex depicts a slight blue-shift compared to the unmethylated species (bottom of the Figure). Experimentally, the Co-H stretching frequency has been reported to be in the range 1930-1970 cm^{-1} by Saillant *et al.*⁶¹ although for complexes containing phosphine ligands as stabilizers. The relatively high Co-H stretching frequency in the present case could be attributable to the absence of such axial ligands.

To investigate how the electronic transitions of the complexes change along the QMCF-MD simulations, we performed TDDFT calculations on 500 equally-spaced snapshots taken from the $[\text{HCo}(\text{tmen})_2]^+$ simulation, using the B3LYP functional and both the 6-31G(d,p) and TZVP basis sets. The results are plotted as histograms in Figure 6 and are to be *Figure 6* compared with the static B3LYP spectra calculated at the optimized geometries in Table 1. In both histograms, two peaks can be discerned which correspond to those of the two principal bands in the static spectra. These occur at slightly longer wavelengths in the dynamic spectra than in their static counterparts with values at approximately 360 and 340 nm and at 470 and 450 nm with the 6-31G(d,p) and TZVP basis sets, respectively. The major difference, however, in the dynamic spectra is that the shorter wavelength bands are now dominant, whereas the longer wavelength bands are almost washed out with amplitudes that are only slightly above the background.

Solvation Structure

As initial analyses we examined the spherical radial distribution functions (RDFs) of the cobalt and hydride with the water molecules. The cobalt-water oxygen RDFs are shown in Figure 7, from which it can be seen that the solvation around $[\text{HCo}(\text{tmen})_2]^+$ has a more *Figure 7* pronounced shell structure than $[\text{HCo}(\text{en})_2]^+$. The figure also shows the distribution of

the smallest Co–O (water) distances that are observed in the MD simulations. The most likely distance for both complexes is approximately 2.2 Å, which is what is observed in a structure that is geometry optimized with water as a coordinating ligand. For $[\text{HCo}(\text{en})_2]^+$ the distribution drops off sharply after this although there is still a substantial fraction of the population ($\sim 8\%$) that has longer Co–O coordination distances than 3.0 Å. By contrast, the $[\text{HCo}(\text{tmen})_2]^+$ distribution is clearly bimodal with a first peak containing about 65% of the population up to about 2.9 Å, and a second peak with the remainder of the structures with a maximum at 3.3 Å. The different distributions of the shortest Co–O distance show that the water-coordination site of the methylated complex is not populated in a rather larger number of configurations. This is not the case in the unmethylated system, where the coordination site is predominantly occupied by a water molecule. This clearly shows the important steric effects that the methyl groups have on water coordination.

The RDFs for the hydride–water oxygen distances are shown in Figure 8. These depict a decreased intensity of hydration in the vicinity of the Co center, within ~ 6 Å. The relative peak intensities indicate that the hydride is more exposed to the solvent in the unmethylated complex, whereas the methylated complex shows a clear peak at about 8 Å, which is associated with the oxygens that interact with the methyl groups.

Figure 8

SLICE projections of the complexes are given in Figure 9. Each shows three distributions: the water-oxygen distribution around the complexes in green and blue, the complexes in gray, and the hydride explicitly in red. The molecular plane employed for the projection was determined from the four nitrogen atoms and the spacing of the projection planes was 1.5 Å. Starting at the central panels that represent the molecular plane, panels '+1' to '+4' show the hydration above the plane, and panels '-1' to '-4' the hydration below the plane.

Figure 9

The most striking difference between the two complexes is the much more highly organized solvation structure for $[\text{HCo}(\text{tmen})_2]^+$ with well-defined solvation hot-spots occurring in all planes, except '+4'. By contrast, the only major solvation peaks for the unmethylated $[\text{HCo}(\text{en})_2]^+$ occur in the '+1' and, to a lesser extent, the '-1' planes in positions that bridge

the two terminal amine groups of the $[\text{HCo}(\text{en})_2]^+$ ligands. These results emphasize the significant ordering effects that methyl groups can have on solvent structure. Concerning the complexes themselves, the major difference is that the hydride distribution is much broader in the unmethylated case, indicating a higher degree of flexibility when the hydride is not engaged by the complex. This agrees with the results of the spherical radial distribution analysis.

Hydration layer statistics from the Laguerre tessellation analysis is presented in Table 4 *Table 4* and the corresponding TCFs in Figure 10. As might be expected the hydration layers are *Figure 10* larger for the methylated complex. The number of exchanges and residence times indicate that the inner solvation layers of $[\text{HCo}(\text{tmen})_2]^+$ are more structured than those of $[\text{HCo}(\text{en})_2]^+$ although this difference is less notable in the outer layer. Differences between the complexes are clearer in a comparison of the TCFs, $C(s)$ and $S(s)$, which decay much more quickly for $[\text{HCo}(\text{en})_2]^+$ than $[\text{HCo}(\text{tmen})_2]^+$. The short-time correlation times, τ_s , from the double exponential fit are only slightly greater for $[\text{HCo}(\text{tmen})_2]^+$ in each of the layers, whereas the long-time contributions, τ_l , are between 50% and 100% greater. This indicates that the minimum residence times are similar for both simulations, whereas the maximum residence times clearly differ with the methylated species featuring a higher degree of structure in the bulk region, and $[\text{HCo}(\text{en})_2]^+$ having a structure that is closer to that of bulk water.^{57,58}

As a final illustration of the data that is provided by Laguerre tessellation, some molecular and atomic volumes and surfaces for the complexes are presented in Table 5. The effective *Table 5* surface is the surface that an atom or a molecule shares with the surrounding water, and provides an estimate of the solvent accessible surface. Not surprisingly the volumes and areas obtained by this analysis are larger for the methylated species, although this is not reflected by the atomic volume and surface of the hydride as only minor decreases in these two properties are observed.

Conclusions

In this study we have simulated two tetramine Co(III)-hydrido compounds that were chosen as models of the intermediates that are known to be important in the hydrogen-evolving catalytic cycles of the cobaloxime complexes. We used both static DFT and hybrid potential molecular dynamical QMCF-MD techniques and compared our results to the available experimental structural and spectral data. A major goal of this work has been to provide a foundation for future simulations of hydrogen-evolving catalysts that we intend to carry out using adaptive QC/MM molecular dynamics methods of the QMCF-MD type. Such dynamical techniques appear to us to be essential for obtaining a full and accurate understanding of the behaviours and properties of these complexes in their functional environments.

Acknowledgement

Financial support for AB, MJF and VA from the French National Research Agency (NiFe-Cat, ANR-10-BLAN-7-11), the FP7 CEA-Eurotalent COFUND Programme, and the CEA's DSV-ENERGIE 2011 Programme is gratefully acknowledged. AKHW thanks the Leopold-Franzens-University of Innsbruck (Rector Prof. Dr. Dr.hc. Tilmann Märk) for a PhD scholarship. AKHW and THH acknowledge the support by the Austrian Ministry of Science BMWF UniInfrastrukturprogramm as part of the Research Platform Scientific Computing at the University of Innsbruck. The COST Action CM1202 PERSPECT-H2O is also acknowledged.

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Table 1: Comparison of experimental and theoretical UV-VIS absorption spectra for $[\text{HCo}(\text{tmen})_2]^+$. Relative intensities with respect to the strongest transition are in parentheses after the wavelength values.

Method	λ_1/nm	λ_2/nm
Experiment	444 (1.00)	340 (0.53)
<i>DFT</i>		
BP86/6-31G(d,p)	430 (1.00), 436 (0.10)	354 (0.35)
BP86/TZVP	401 (0.65), 398 (1.00)	334 (0.10), 337 (0.10)
B3LYP/6-31G(d,p)	439 (1.00), 442 (0.25)	336 (0.75)
B3LYP/TZVP	418 (0.25), 415 (1.00)	326 (0.50), 322 (0.03)
<i>CAS/NEVPT2</i>		
CAS(6,5)/NEVPT2	415 (< 0.01)	341 (1.00), 339 (1.00)
CAS(10,9)/NEVPT2	422 (0.40), 415 (0.40)	317 (1.00), 313 (0.70)

Table 2: Table of various internal coordinates calculated from 20 ps of simulation. Distances are in Å, and angles and dihedrals in degrees. The values in the table are computed for all possible internal coordinates of the same type.

Internal Coordinate	[HCo(en) ₂] ⁺				[HCo(tmen) ₂] ⁺			
	Avrg. & Std. Dev.	Max.	Min.		Avrg. & Std. Dev.	Max.	Min.	
Co-H (hyd.)	1.42 ± 0.09	1.84	1.15		1.41 ± 0.09	1.79	1.15	
Co-N	2.04 ± 0.12	2.62	1.71		2.04 ± 0.11	2.87	1.62	
N-Co-H (hyd.)	89.8 ± 9.2	140.3	54.5		90.7 ± 10.3	141.6	54.7	
Co-N-H (amine)	110.0 ± 10.3	155.6	60.6		109.4 ± 11.1	160.9	61.0	
Co-N-N-N	0.4 ± 6.3	22.6	-33.0		-1.0 ± 6.1	24.1	-23.6	
N-N-N-N	-0.1 ± 8.6	28.8	-32.7		0.1 ± 7.8	30.6	-24.4	

Table 3: Table of the means and standard deviations of partial charges for atoms in the complexes, computed from 20 ps of simulation. Charges are calculated with the TZVP basis set, with both the B3LYP and BP86 functionals, and using the ESP, Löwdin and Mulliken approximations.

Atom	ESP	Löwdin	Mulliken
[HCo(en) ₂] ⁺ — B3LYP			
Co	0.74 ± 0.12	0.29 ± 0.03	0.34 ± 0.05
H (hyd.)	-0.10 ± 0.03	0.08 ± 0.01	0.06 ± 0.01
N (avrg.)	-0.52 ± 0.21	-0.03 ± 0.03	-0.37 ± 0.05
[HCo(en) ₂] ⁺ — BP86			
Co	0.65 ± 0.12	0.21 ± 0.03	0.23 ± 0.05
H (hyd.)	-0.09 ± 0.03	0.09 ± 0.01	0.07 ± 0.01
N (avrg.)	-0.50 ± 0.21	-0.02 ± 0.03	-0.35 ± 0.04
[HCo(tmen) ₂] ⁺ — B3LYP			
Co	0.68 ± 0.20	0.24 ± 0.03	0.33 ± 0.07
H (hyd.)	-0.09 ± 0.04	0.08 ± 0.01	0.06 ± 0.02
N (avrg.)	-0.74 ± 0.14	-0.00 ± 0.04	-0.41 ± 0.06
[HCo(tmen) ₂] ⁺ — BP86			
Co	0.58 ± 0.21	0.16 ± 0.03	0.22 ± 0.08
H (hyd.)	-0.08 ± 0.04	0.09 ± 0.01	0.06 ± 0.02
N (avrg.)	-0.70 ± 0.14	0.01 ± 0.03	-0.37 ± 0.06

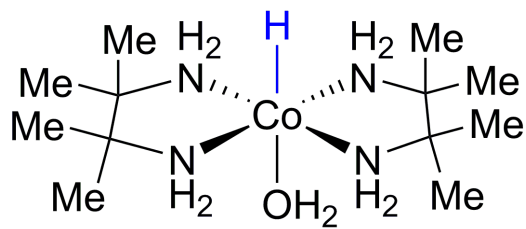
Table 4: Top: table of coordination numbers (CND), exchanges, mean residence times (MRT) and the ration of exchanges (R_{ex}) of water molecules within distinct hydration layers, obtained via Laguerre tessellation. The exchange data of the i th layer refers to events with the $i + 1$ th layer. This data is based on 20 ps of sampling. Bottom: table of computed $C(t)$ and $S(t)$ correlation times for the residence time of water molecules within distinct hydration layers, obtained via Laguerre tessellation. This data is based on 20 ps of sampling. All times are in ps.

Layer	Avrg. CND	Exchanges		MRT		R_{ex}
		$t^*=0.5$	$t^*=0.0$	$t^*=0.5$	$t^*=0.0$	
$[\text{HCo}(\text{en})_2]^+ - 1$	43.6	390	5996	2.5	0.160	15.4
$[\text{HCo}(\text{en})_2]^+ - 2$	137.1	1523	33671	2.0	0.092	22.1
$[\text{HCo}(\text{en})_2]^+ - 3$	252.4	1836	103657	3.1	0.055	56.5
$[\text{HCo}(\text{tmen})_2]^+ - 1$	63.3	359	6030	3.4	0.200	16.8
$[\text{HCo}(\text{tmen})_2]^+ - 2$	167.1	1304	30598	2.5	0.110	23.5
$[\text{HCo}(\text{tmen})_2]^+ - 3$	290.2	1912	90475	2.9	0.062	47.3

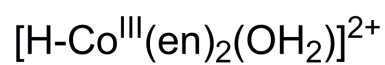
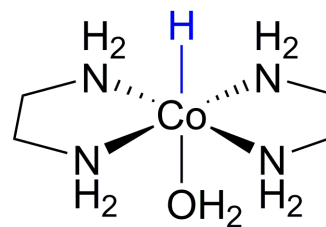
	Layer	$[\text{HCo}(\text{en})_2]^+$		$[\text{HCo}(\text{tmen})_2]^+$	
		τ_l	τ_s	τ_l	τ_s
$C(t)$	1	2.9	0.046	5.5	0.055
	2	2.8	0.032	4.8	0.037
	3	2.8	0.025	4.3	0.029
$S(t)$	1	0.60	0.045	1.2	0.060
	2	0.43	0.037	0.74	0.049
	3	0.34	0.031	0.50	0.038

Table 5: Table of atomic and molecular volumes (V) and surfaces (A) obtained via Laguerre tessellation for the hydride atom and the complexes. Data is based on 20 ps of sampling and the averages and standard deviations for each property are shown.

Complex	$V_{Hydride} \text{ \AA}^3$	$A_{Hydride} \text{ \AA}^2$	$V_{complex} \text{ \AA}^3$	$A_{complex} \text{ \AA}^2$
$[\text{HCo(en)}_2]^+$	13.6 \pm 3.4	31.9 \pm 5.0	318.3 \pm 24.8	320.5 \pm 18.2
$[\text{HCo(tmen)}_2]^+$	12.0 \pm 3.1	29.8 \pm 4.8	559.1 \pm 26.8	461.7 \pm 15.9



1



2

Figure 1: The hydrido-tetramine-cobalt(III) complexes studied in this work. Left: $[\text{HCo}(\text{tmen})_2]^+$; right: $[\text{HCo}(\text{en})_2]^+$.

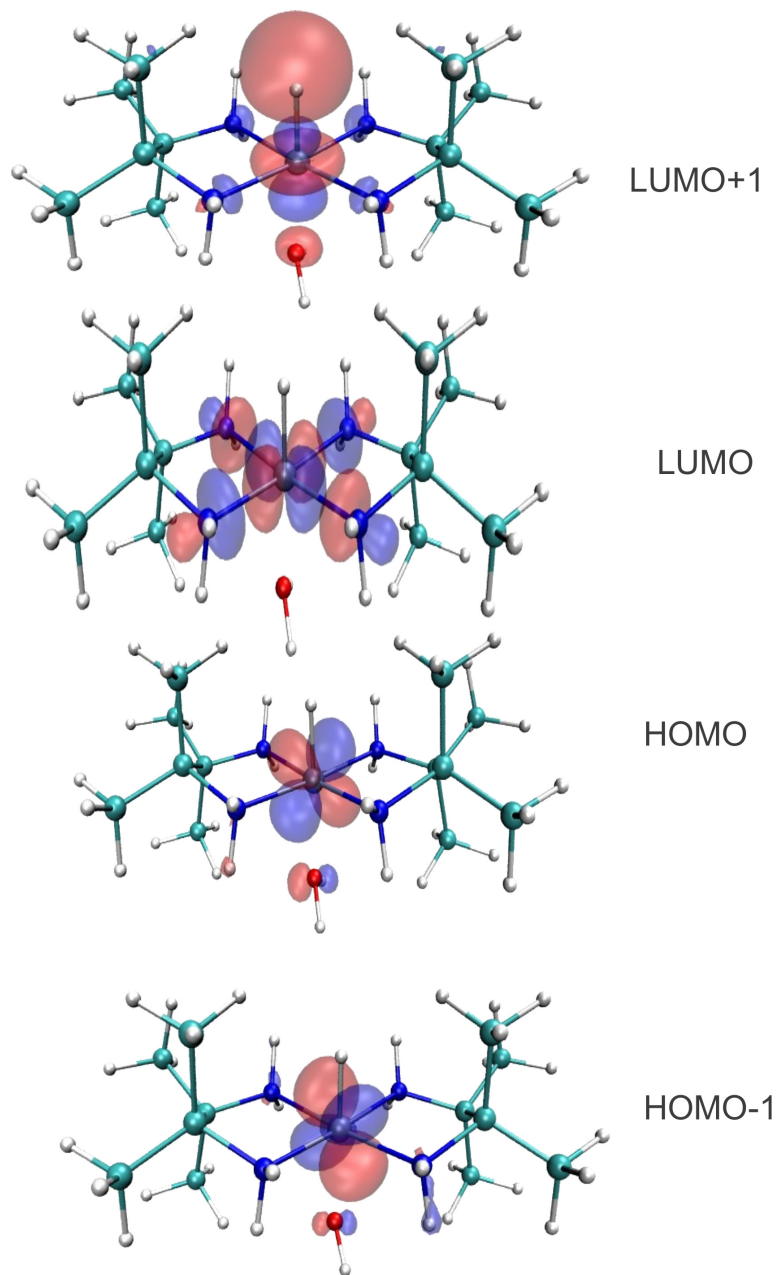


Figure 2: Isosurface plots of the Kohn-Sham orbitals from the B3LYP/TZVP DFT calculations that are important in the UV-VIS absorption spectrum transitions.

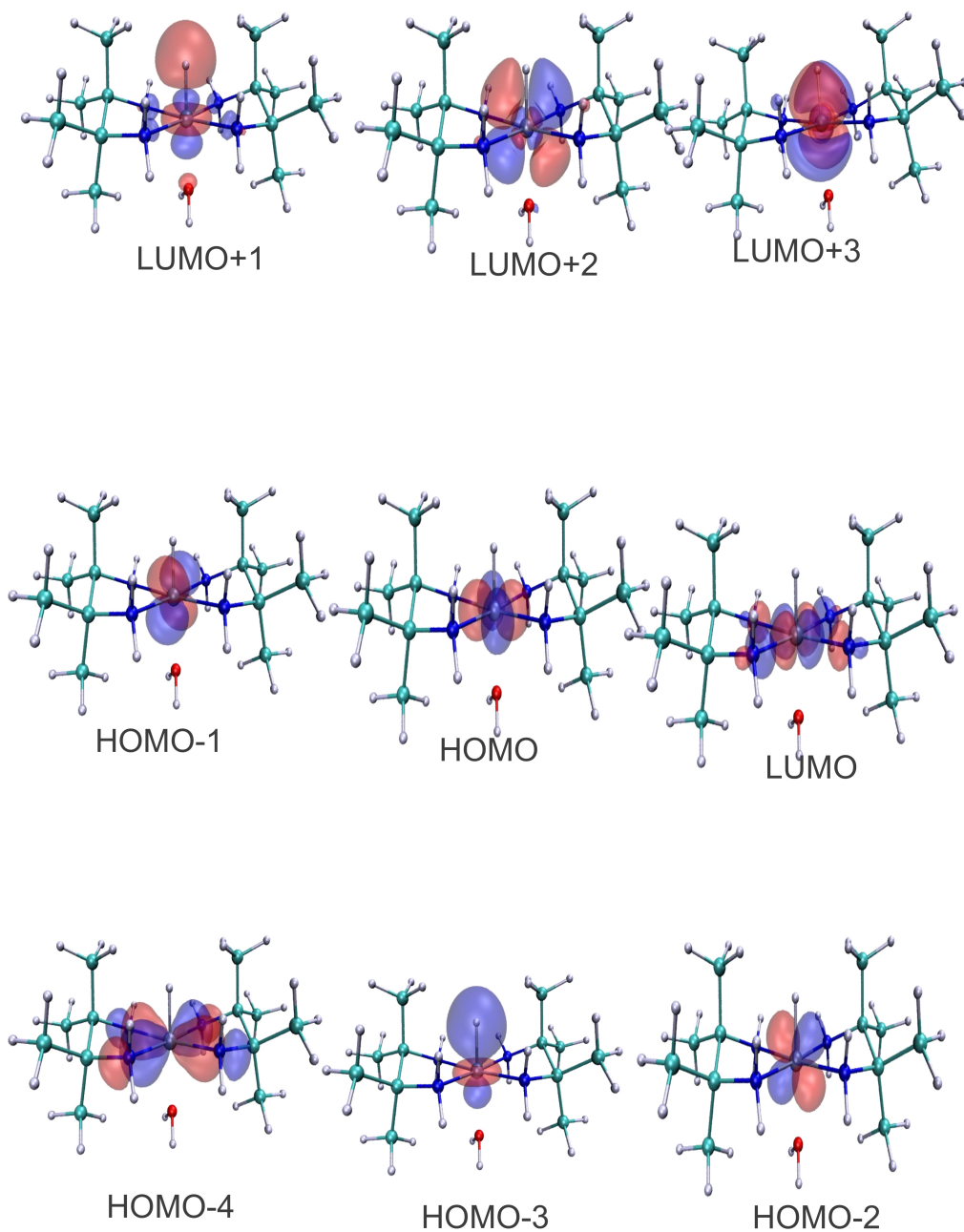


Figure 3: Isosurface plots of the orbitals from the CAS(10,9) CASSCF calculations that are important in the UV-VIS absorption spectrum transitions.

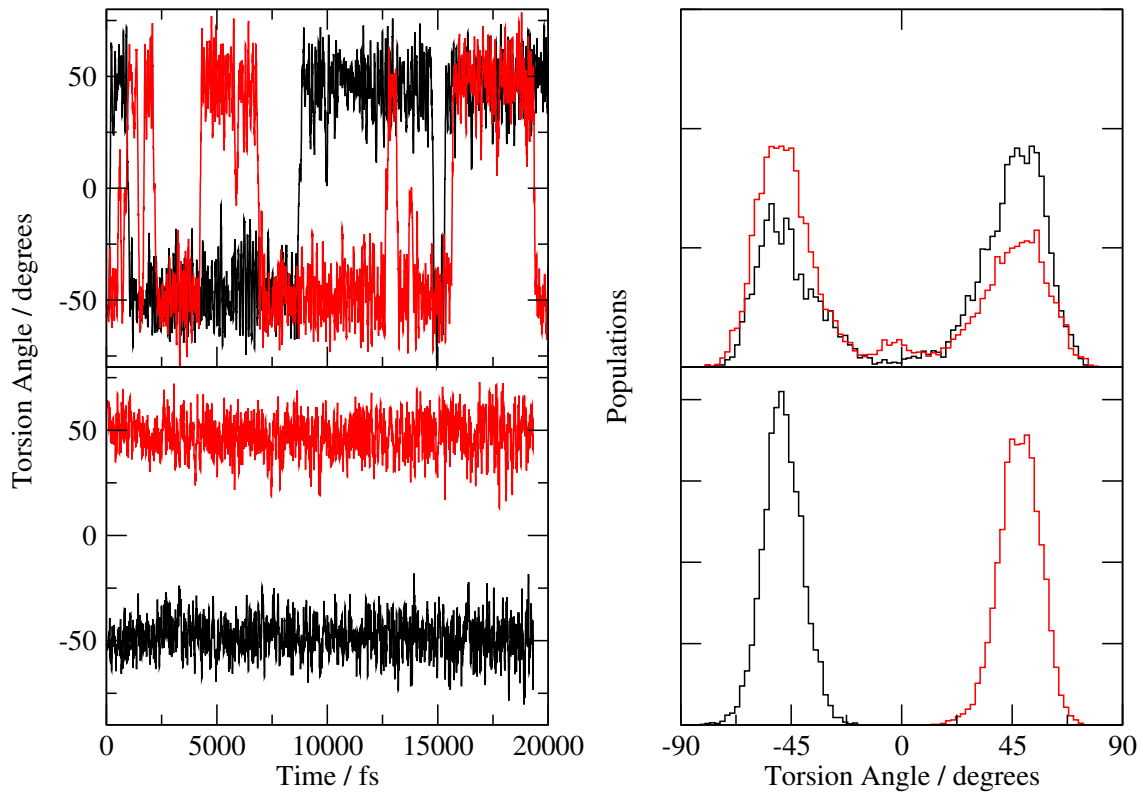


Figure 4: Trajectories (left) and histogram representations (right) of the N-C-C-N dihedral angles. Top: $[\text{HCo}(\text{en})_2]^+$; bottom: $[\text{HCo}(\text{tmen})_2]^+$.

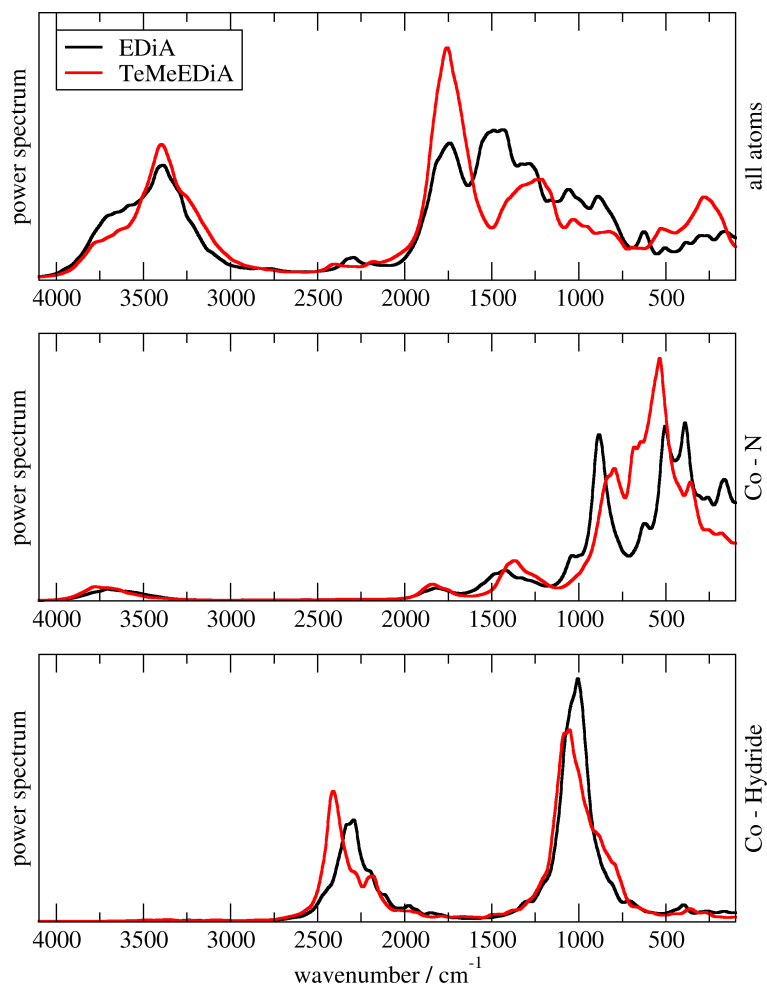


Figure 5: Power spectra of the complexes' vibrational modes, obtained via Fourier transform of the relevant velocity auto-correlation functions. Top: all atoms of the complexes; middle: Co-N bonds; bottom: Co-H (hyd.) bond.

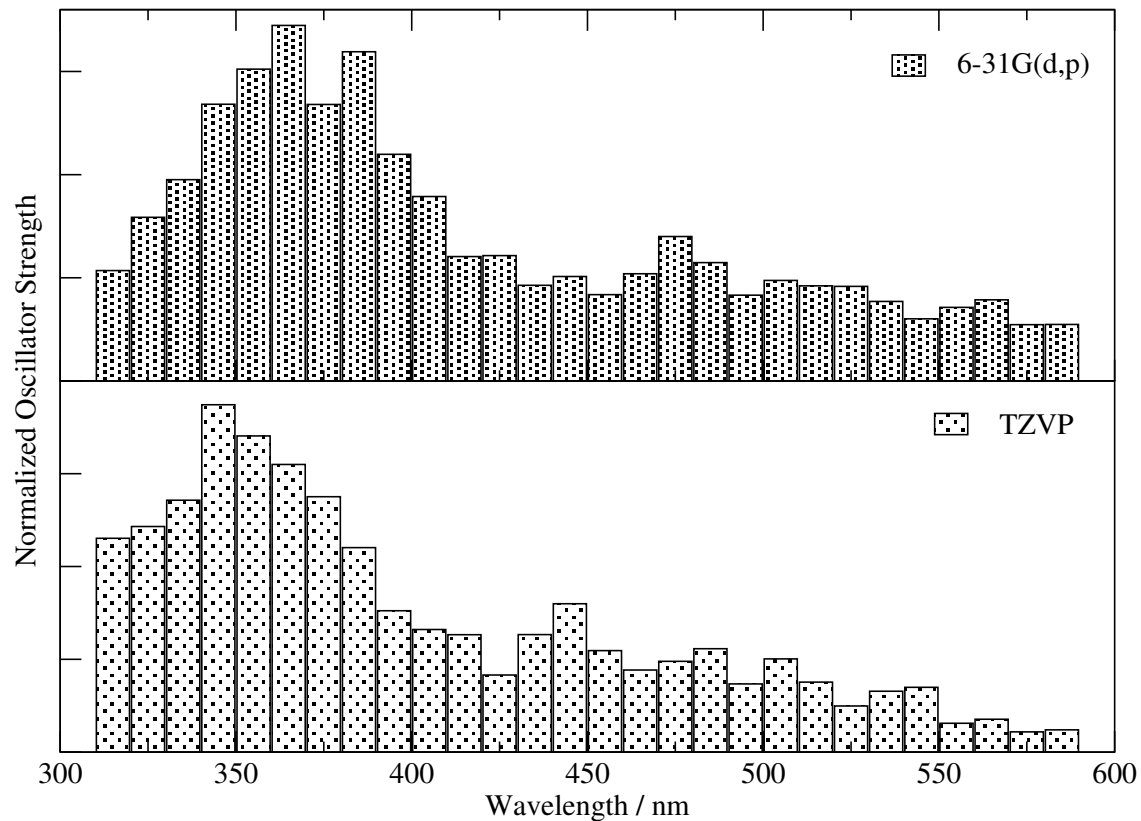


Figure 6: A histogram of the time-averaged wavelengths of the electronic transitions for the $[\text{HCo}(\text{tmen})_2]^+$ complex calculated at the TDDFT level of theory using the B3LYP functional and the 6-31G(d,p) (top) and TZVP (bottom) basis sets. The intensities are the normalized oscillator strengths and the histogram bin width is 10 nm.

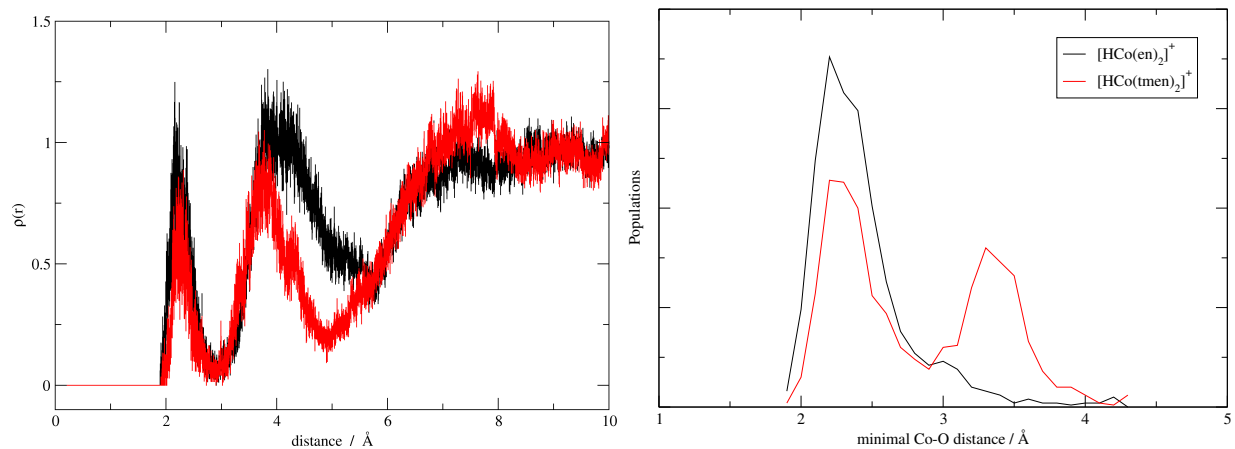


Figure 7: Left: Co–O (water) radial distribution functions; right: distributions of the shortest Co–O (water) distances. All data calculated from the 20 ps MD trajectories. Black: $[\text{HCo}(\text{en})_2]^+$; red: $[\text{HCo}(\text{tmen})_2]^+$.

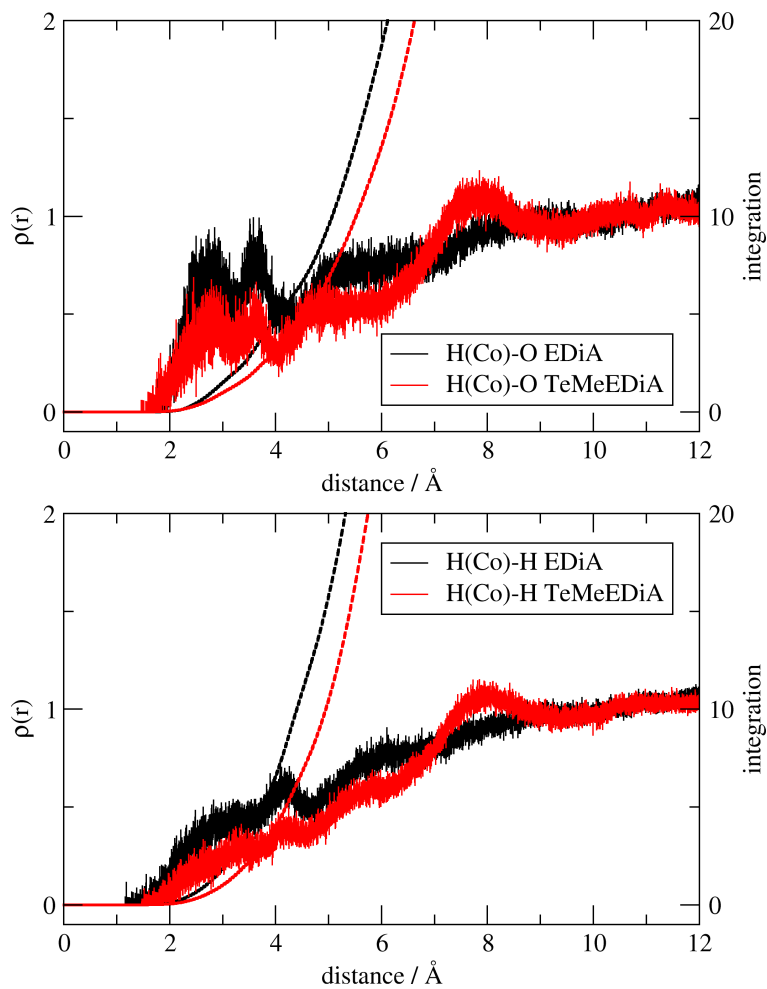


Figure 8: Spherical pairwise radial distribution functions. Top: H (hyd.)–O (water); bottom: H (hyd.)–H (water).

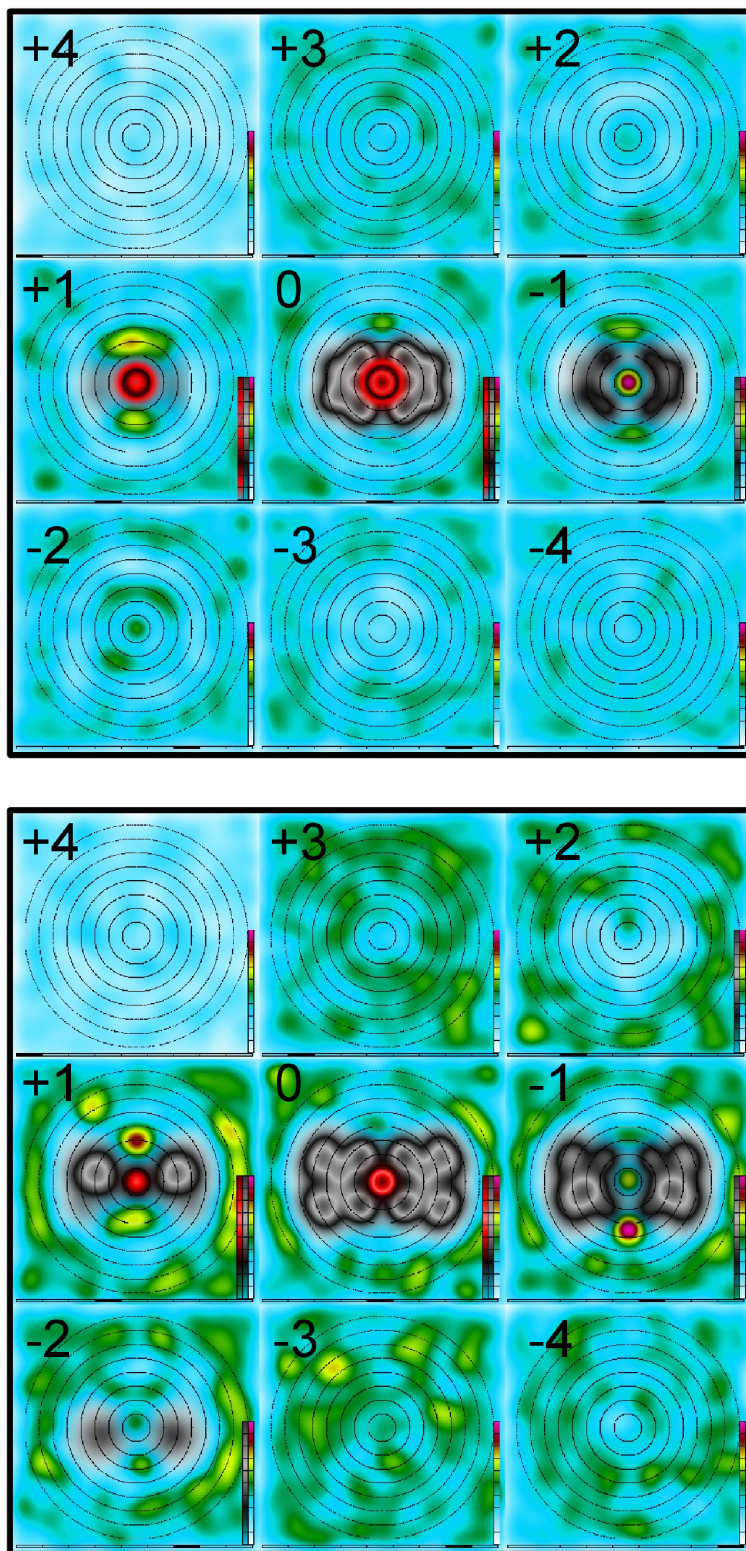


Figure 9: SLICE projection analysis based on superimposed trajectories of the complexes. Top: $[\text{HCo}(\text{en})_2]^+$; bottom: $[\text{HCo}(\text{tmen})_2]^+$. The three colorbars are as follows: blue/green — water oxygens; gray — complex; red — hydride.

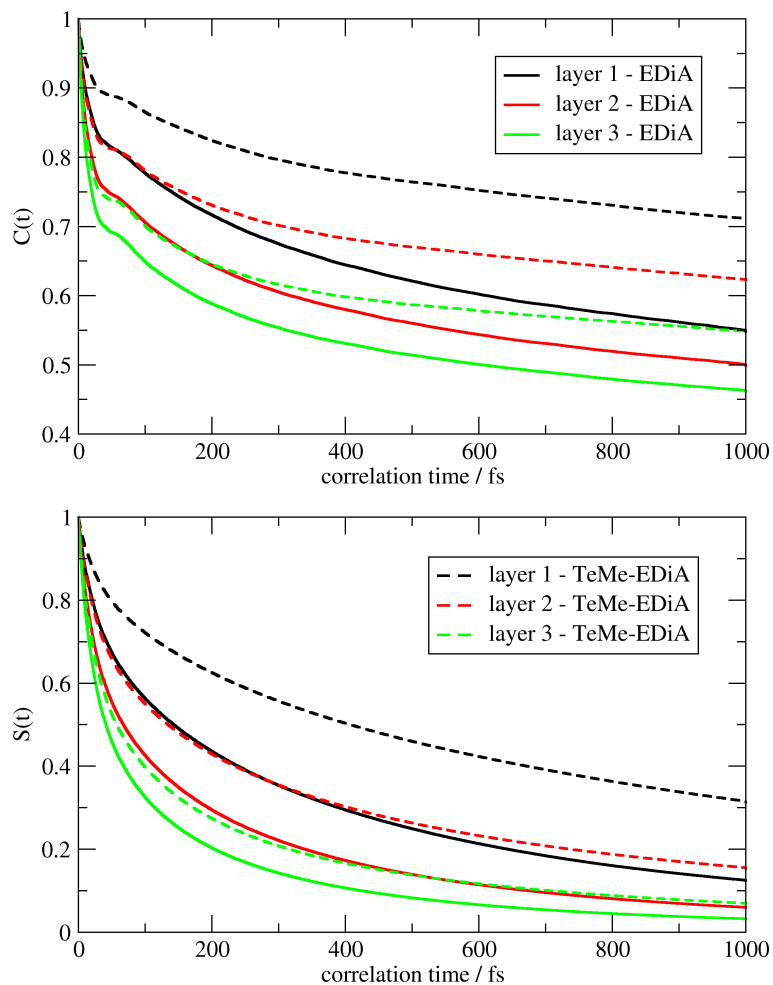
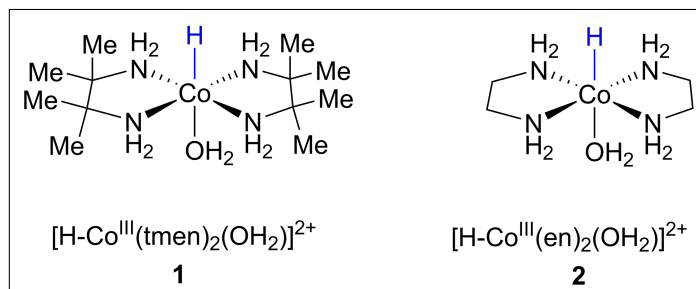


Figure 10: $C(t)$ and $S(t)$ residence time correlation functions of water molecules within individual hydration layers based on a Laguerre tessellation. Full lines: $[\text{HCo}(\text{en})_2]^+$; dotted lines: $[\text{HCo}(\text{tmen})_2]^+$. Black, red and green refer to hydration layers 1, 2 and 3, respectively.

Graphical TOC Entry



Adaptive QC/MM molecular dynamics simulations have been performed of two models of the cobalt(III)-hydride intermediate that is important in the catalytic cycles of hydrogen evolution mediated by cobaloximes.