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A Mechanistic Rationale Approach Revealed the Unexpected Chemoselectivity of an Artificial Ru-dependent Oxidase - A Dual Experimental/Theoretical Approach

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Abstract

Artificial enzymes represent an attractive alternative to design abiotic biocatalysis. *Ec*NikA-**Ru1**, an artificial metalloenzyme developed by embedding a ruthenium-based catalyst into the cavity of the periplasmic nickel-binding protein NikA, was found to efficiently and selectively transform certain alkenes. The objective of this study was to provide a rationale on the enzymatic function and the unexpected substrate-dependent chemoselectivity of *Ec*NikA-**Ru1** thanks to a dual experimental/computational study.

We observed that the *de novo* active site allows the formation of the terminal oxidant *via* the formation of a ruthenium aquo species that subsequently reacts with the hypervalent iodine of phenyl iodide diacetic acid. The oxidation process relies on a Ru^{IV}=O pathway *via* a two-step reaction with a radical intermediate, resulting in the formation of either a chlorohydrin or an epoxide.

The results emphasize the impact of the protein scaffold on the kinetics of the reaction, through i) the promotion of the starting oxidizing species via the exchange of a CO ligand with a water molecule; and ii) the control of the substrate orientation on the intermediate structures, formed after the Ru^{IV}=O attack. When a $C\alpha$ attack is preferred, chlorohydrins are formed while an attack on $C\beta$ leads to an epoxide.

This work provides an evidence that artificial enzymes mimic the behavior of their natural counterparts.

Keywords

Artificial enzymes – catalysis – oxidation – hydroxychlorination – epoxidation - computational chemistry – reaction paths – spin flip

Introduction

In the search to improve the sustainability of organic chemistry, new green catalytic approaches are targeted.¹ When designing novel processes based around a metal-containing catalyst, the following features are desirable: abundant metal(s), easy-to-handle ligands, safe reactants/solvents, and minimal reaction steps (e.g. cascade reactions). Artificial enzymes (ArMs), based on the insertion of an inorganic catalyst as an active site into an unreactive protein scaffold, provide a promising alternative to purely biological catalysts by allowing the design of unnatural reactions.² While the catalytic efficiency of natural enzymes can be optimized by mutagenesis,³ ArMs offer an additional degree of freedom through the chemical modification of the inorganic catalyst. Because the active site of an ArM is formed by combining a protein environment with a stable inorganic complex, the catalytic properties of both the protein and the inorganic complex can be investigated separately, simplifying the process by which the resulting ArM can be optimized.

The flexibility in design afforded by ArMs allows reaction pathways to be controlled, and means that the repertoire of possible transformations can be expanded to unnatural reactions.

Our group has designed a Ru-based ArM by embedding a ruthenium complex (**Ru1**) (Figure 1) in the binding pocket of the nickel-binding protein NikA from *Escherichia coli* (*Ec*NikA) to form the *Ec*NikA-**Ru1** hybrid.⁴

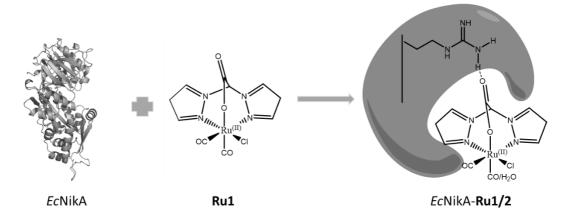


Figure 1. Drawings of *Ec*NikA, **Ru1** and *Ec*NikA-**Ru1/2**. Arg137 is included in *Ec*NikA-**Ru1/2** to show the salt bridge stabilizing the complex inside the protein.

*Ec*NikA-**Ru1** was shown to react with alkenes to give the corresponding β –chloro- α -hydroxy adducts. The understanding of the reaction mechanism at the molecular level is of great importance for the optimization of this kind of ArM. In particular, the nature of the oxidizing species and its impact on the reaction (enantio-, chemo-, and regioselectivity) must be determined to allow a better optimization strategy. Here, we propose a reaction mechanism that has been formulated using a combination of theoretical and experimental data. In particular, we focused on deciphering the overall mechanism

between the starting **Ru1** complex and the oxidant, revealing the transient formation of a genuine complex-oxidant adduct. Density functional theory (DFT) calculations confirmed the importance of the protein scaffold on the complex activation, giving a new example of the advantageous use of artificial enzymes. Moreover, the unexpected reaction chemoselectivity (epoxidation vs hydroxychlorination) is related to the competition between kinetic and thermodynamic control of the reaction. This reaction pathway is dependent on the substrate orientation in the computed radical intermediates, which is driven by the protein environment. Interestingly, the modeled formation of the β -chlorohydrins involves a chlorine transfer that is reminiscent to the one observed in iron chloroperoxidase catalyzing the formation of (di)chlorinated compounds.⁵

RESULTS

Catalytic studies

Scheme 1. Standard oxidation conditions for BMS.

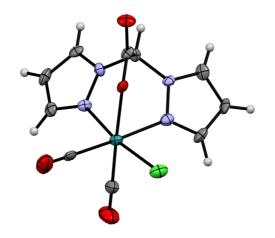
In a previous study, EcNikA-**Ru1** was shown to react with alkenes in the experimental conditions given by scheme 1.⁴ Typically, optimum catalytic efficiency was obtained using a 1/500/600/75 ratio for catalyst/substrate/oxidant/chloride, respectively, for a 37 μ M final concentration of either **Ru1** or EcNikA-**Ru1** for a 10 minutes reaction time at room temperature. Substrates were solubilized in acetonitrile prior mixing to the buffered medium but PIDA was added as a solid. The complex **Ru1** alone was inert, suggesting the activation of the catalyst via its binding to the protein scaffold.⁴ The hydroxychlorination of trans- β -methylstyrene (BMS) in the presence of PhI(OAc)₂ (PIDA) was performed with a complete regioselectivity to give the β -chloro- α -hydroxy adduct (Table S1) with a yield of 99% with respect to chloride that is the limiting reactant (15% with respect to substrate). The catalyzed hydroxychlorination had a turn over number (TON) of 74 and a turn over frequency (TOF) of 7 min⁻¹, which is highly efficient compared to similar systems,⁶ whereas apo-EcNikA is inactive. Likewise, an EcNikA-**Ru(DMSO)**₂Cl₂ analogue was found to be unreactive.

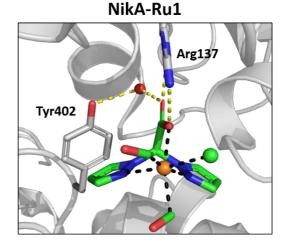
Then, the reactivity of *Ec*NikA-**Ru1** and **Ru1** was investigated for the oxidation of 4-methoxystyrene (MOS) under the conditions previously described for BMS (Scheme 2, Table S1). First, no product was detected by GC-MS either in the absence of the catalyst or in the presence of *Ec*NikA alone. Second,

Ru1 was found quasi inactive under these conditions. Product analysis revealed the exclusive formation of the unexpected 2-(4-methoxyphenyl)oxirane with a maximum yield of 35% after 10 minutes. *Ec*NikA-**Ru1** epoxidation was highly efficient, with a TON of 175 in 10 minutes, giving a TOF of 18 min⁻¹. No side product was observed. These values are unmatched when compared to those of the **Ru1** even in organic solvent (26 TON and TOF of 0.02 min⁻¹),⁷ or to the ones of the artificial streptavidin-OsO₄ (27 TON with an estimated TOF around 0.02 min⁻¹ but for diol formation),⁶ highlighting the ability of *Ec*NikA to speed-up reactions.

Scheme 2. Standard oxidation conditions for MOS.

The question of the starting active species then arose. Previously, we showed that the crystal structures of EcNikA-Ru1 and Ru1 displayed significant structural differences in ruthenium complex geometry (Figure 2). The major disparities are the elongation of the Ru-tCO length from 1.9 to 2.3 Å in EcNikA-Ru1 compared with Ru1, and the change of the Ru-tCO angle from 180° in Ru1 to 118° in EcNikA-Ru1.4 The metrics of the tCO are related to its loose binding to the ruthenium ion, as observed previously on iron systems. We also observed that, depending on the x-ray structures, the electron density for this region cannot be systematically modelled as a CO ligand. Here, the x-ray structure of EcNikA-Ru2 was solved at 1.9 Å by soaking EcNikA-FeEDTA crystals with Ru1 (PDB code: 6R4Q, Figure 2, Table S2). As observed previously, two molecules were present in the asymmetric unit, one of which contained the ruthenium complex at the NikA binding site with an occupancy of approximately 0.9. Only this molecule containing the ruthenium complex will be discussed hereafter. Compared to the crystal structure of EcNikA-Ru1, the major difference is the absence of the electron density corresponding to tCO. As expected, the loose binding of tCO to Ru1 led to its release over time. In its place, a weak electron density rather corresponding to a monoatomic species was observed, which could not be clearly modelled. However, based on the fact that Ru(II) prefers a six-coordinate geometry, this electron density was modelled as a water molecule with a Ru-H₂O distance of 1.9 Å, in agreement with literature and an occupancy of 0.9, forming EcNikA-Ru2. Embedding Ru1 into NikA led to the build-up of catalytic activity, pushing us to correlate the **Ru2** formation with the ArM reactivity.





Ru-Ru-Ru-Cl Ru-O tCO OH₂ (Å) (Å) (Ă) (Ă) Ru1 2.37 2.11 1.88 EcNikA-2.2 2.5 2.3 Ru1 EcNikA-2.2 2.5 1.9 Ru2

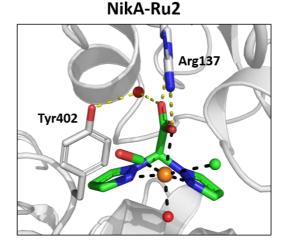


Figure 2. Structures of **Ru1** (CCDC: fsbu18) (top left), *Ec*NikA-**Ru1** (PDB code: 5L8D) (top right) and *Ec*NikA-**Ru2** (PDB code: 6R4Q) (bottom). In protein: Ru is depicted as an orange sphere, Cl as a green sphere and water molecules as red spheres. The ruthenium ligand and CO are in sticks. The figures were prepared with PyMOL (The PyMOL molecular graphic system, Version 2.0 Schrödinger, LCC).

The speciation in solution between **Ru1** and **Ru2** was also investigated using IR spectroscopy. As published previously, the IR spectrum of *Ec*NikA-**Ru1** displayed three resonances at 2082 cm⁻¹, 2017 cm⁻¹ and 2005 cm⁻¹.⁴ The 2082 and 2005 resonances were also observed in the spectrum of **Ru1** in solution, suggesting that free **Ru1** was present in addition to a new species. The vCO were calculated by DFT with B3LYP functional,⁹ a def2-SVP split valence polarization basis set¹⁰ and a Polarizable Continuum Model (PCM) for the solvent (Table S3) leading to two vibrations at higher energies (2150 and 2078 cm⁻¹); the differences between the two resonances were similar when comparing the observed and calculated data ($\Delta E = 72 \text{ vs } 77 \text{cm}^{-1}$). The impact of the protein on these vibrations was calculated, taking into account the already identified protein residues impacting the complex

stability.¹¹ Results showed that the ν CO were slightly affected (in a 5 cm⁻¹ range), suggesting that the third vibration observed at 2017 cm⁻¹ was issued from a distinct species, likely a mono CO species. Calculated **Ru2** vibration frequencies were found significantly lower than those of **Ru1** and similar to the observed one (2050 cm⁻¹ ν s 2017cm⁻¹).

Taken all together, we assume that the major species in solution is *Ec*NikA-**Ru2**. The mono-CO resonance at 2017 cm⁻¹ was not observed in the unbound **Ru1**, which is instead more likely to lose the chloride ligand (as shown by electrospray ionization/mass spectrometry studies), suggesting that the stabilization of this chloride group could also play a role in the activity of *Ec*NikA-**Ru1**.⁴

Reaction mechanism studies

*Ec*NikA-**Ru2** catalyzed the epoxidation of MOS in aqueous medium, whereas a chlorohydrin is formed with BMS. The hypervalent iodine compound PIDA was found to be the most effective stoichiometric oxidant to catalyze both the epoxidation of MOS and the hydroxychlorination of BMS (Table S4). This discrepancy in chemoselectivity led us to decipher the reaction mechanism using calculation based on experimental evidence.

Experimental studies

The activation of PIDA is essential for catalysis. The reaction steps for the alkene oxidation may follow several different pathways, distinguished by the involvement of metal-centered redox processes. Firstly, a transient amino acid-based oxidation may occur. Secondly, the protein scaffold or the metal complex may activate PIDA by its binding through a Lewis acid interaction with either an amino acid residue or the metal. Finally, the metal center may activate the oxidant *via* a redox process. In the following, we describe each possibility and confronted them to our experimental data.

Control reactions have been run with the open form of the protein apo *Ec*NikA (Table S1). As only a poor reactivity was observed, it seems that it is not an amino acid that is responsible of catalytic activity.

This was confirmed by a metal complex exchange strategy while keeping the closed form of the protein structure and the size of the cavity. Since *Ec*NikA is able to bind metal EDTA derivatives, ¹³ an *Ec*NikA-Mg(EDTA) hybrid was synthesized *in situ*. ¹⁴ The absence of reactivity for both chlorohydrin and epoxide formation (Table S1) confirmed the absence of amino acid based of Lewis acid pathway. Moreover, the possible aspecific binding of a free ruthenium atom issued from Ru1 decomposition to the protein was also ruled out by the absence of reactivity of the mixture *Ec*NikA and Ru(DMSO)₂Cl₂ (Table S1). While confirming that the Ru complex controlled the reaction, we could not discriminate between a

metal Lewis acid and a metal based redox mechanisms. Unfortunatly, any attemps to synthesize a Mg analog of **Ru1** was unsuccessful, because a Mg(bpza)₂ complex was the most stable species in solution.

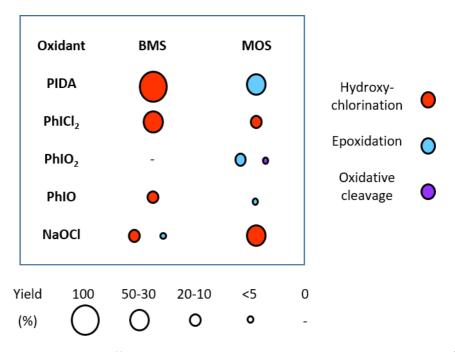


Figure 3. Oxidation reaction with different oxidants. Reaction conditions: Standard conditions: 1/500/600/75 *ratio* for catalyst/substrate/oxidant/chloride, respectively, for a final *Ec*NikA-**Ru1** concentration of 37 μ M in 40 mM Tris.HCl pH 7.5 and 10 mM HEPES pH 7.0; reaction at room temperature for 10 min.

The 73% diastereomeric excess, observed for the BMS β –chloro- α –hydroxychlorohydrin production (Table S1), is suggestive of a stepwise reaction involving the formation of an intermediate. In the case of MOS, the ring closing of the chlorohydrin issued from a putative hydroxychlorination was not considered since the chlorohydrin, made separately, was unreactive in our standard conditions of reaction. In the case of BMS, the reaction mechanism can correspond to either a typical electrophilic *anti* addition process, the intermediate being a chloronium bridge ion subsequently attacked in the α position of the phenyl ring by a water molecule, or to the formation of an epoxide, free or ligated to the metal, being attacked in the least-hindered β position by a chloride. As an epoxide was detected in the case of MOS oxidation, the behavior of the 2-methyl-3-phenyloxirane (BMS-epoxide) was investigated under oxidative conditions. When 500 eq. of epoxide was added to the standard reaction conditions in the presence of *Ec*NikA-**Ru1/2**, only 1-phenylpropane-1,2-diol was detected with a yield of 96%, whereas in the presence of *Ec*NikA or **Ru1**, the epoxide was recovered unchanged. Compared to the exclusive formation of chlorohydrin when BMS is the substrate, it is clear that free BMS-epoxide is not formed. Moreover, oxidant was required for the diol formation, indicating that the hybrid catalyst needs to be oxidized, either to bind the epoxide or to activate a bound water molecule.

We then investigated the possible formation of a chloronium bridge ion as an intermediate of the reaction. PhICl₂ has been described to generate such an intermediate when reacted with carbon-carbon double bonds. Moreover, the presence of small amounts of chlorohydrin in the uncatalyzed reaction could be related to the presence of additional hypervalent iodine species in aqueous solution when PIDA is mixed with Tris. HCl buffer. Indeed, H NMR study of a scaled-up, uncatalyzed reaction mixture revealed that a new species slowly appeared as a function of time with chemical shifts of 7.20 (t), 7.43 (t) and 7.80 (d) ppm for aromatic protons. These shifts values are identical those of PhICl₂, suggesting the presence of such a species in solution in the absence of the hybrid catalyst. The formation of PhICl₂ in buffered solution results from a nucleophilic attack of chlorides on the hypervalent iodine and the removal of the acetate anions. This hypervalent iodine transformation could also be catalyzed by the hybrid, as has been observed in CH₃CN:H₂O mixtures with RuCl₃. Therefore, we replaced PIDA with PhICl₂ under catalytic conditions (Figure 3). When MOS was the substrate, PhICl₂ was able to catalyze the formation of the previously unobserved corresponding chlorohydrin (35% yield) without forming any epoxide, attesting of the absence of PhICl₂ as a free oxidative species in our standard catalytic conditions.

Finally, it is well documented that the reaction of a metal with a hypervalent iodine leads mostly to the formation of a high-valent metal-oxo species, with the oxygen atom coming from a water molecule (except for PhIO) or an oxygenated substituent such as acetate in PIDA. 15b,18 In fact, it has been demonstrated that Ru-OH₂ systems could be oxidized with a two-electron oxidation and the release of protons. Two competitive reactions between free Cl and a high-valent ruthenium-oxo (Rul (Cl)=O) can then be considered i) the production of free Cl+ by direct chloride oxidation, ii) the formation of hypochlorite, the latter being described in several examples of bioinspired chloroperoxidase systems. Nevertheless, the Ru^{IV}(CI)=O could lead to the chlorination of alkenes in a similar fashion to iron α -ketoglurate dependent enzymes.¹⁹ In these cases, the metal-oxo species was only an oxidant species without any oxygen atom transfer (HAT mechanism). This putative metal-oxo species is also analogous to iron-based oxidizing species involved in cis-dihydroxylation catalyzed by the macrocyclic tetraaza ligand. 5c,20 In our case, the formation of a chlorohydrin as the sole product during oxidation of MOS in the presence of sodium hypochlorite (Figure 3) led us to finally neglect the presence of a chloronium ion bridge as an intermediate. Conversely, when PhIO, a reactant known to generate only metal-oxo species, 18c,21 replaced PIDA (Figure 3), the reaction specificity was conserved while the yield dropped to 12% (probably due to the polymeric form of PhIO). In the case of MOS, the formation of an epoxide is also consistent with a classic oxidation pathway in the presence of Ru^{IV}(CI)=O.²²

The hydroxychlorination mechanism may also be related to the existence of other oxidizing species. First, the literature describes a metal-based mechanism that allows the disproportionation of PIDA into

iodobenzene (PhI) and iodylbenzene (PhIO₂).²³ PhIO₂ is known to oxidize alcohols and alkanes and could then be present in our reaction medium. The involvement of PhIO₂, generated by RuCl₃, was ruled out by the formation of only traces of chlorohydrin in the case of BMS oxidation (Figure 3).¹⁷ Second, the formation of a dioxo Ru(VI) species is rejected on the basis of its propensity to cleave alkenes into aldehydes or to generate diols.²⁴

The hypothesis of a unique oxidizing species for both formation of chlorohydrin and epoxide was challenged by UV-visible kinetic experiments (Figure S1). Treating EcNikA-Ru1 with 50 eq. of PIDA led to the formation of a new species characterized by an absorption at 560 nm, while no change was observed in the presence of Ru1 alone. Addition of 500 eq. of either BMS or MOS to this solution led to the rapid loss of this new species upon time, supporting that it is an oxidizing intermediate in both reaction pathways (Figure S1A). The same species could be observed when NikA-Ru1 was treated with a same amount of PhIO (Figure S1B). The absorbance was then lower in accordance with the lower reactivity observed when PhIO was used as the oxidant (Figure 3). Consequently, this common behavior suggest the hypothesis of the formation of a Ru=O oxidizing species.^{21d} Following our modelisation approach for the IR spectrum (see above), we calculated the UV-visible spectrum of the suggested L(CO)(CI)Ru=O species using the Gaussian 09 program package and the hybrid exchangecorrelation B3LYP functional in conjunction with the 6–311++G(d,p) basis set and long range dispersion correction. Optimization was carried in the presence of chlorobenzene solvent by applying the integral equation formalism for the polarizable continuum model (IEF-PCM). Then, spectra were obtained using time-dependent density functional theory (TD-DFT) both in absence and in presence of five protein residues (Arg137, Tyr22, Tyr402, Trp100 and Trp398) maintaining their α and β carbons fixed to mimic the presence of the protein. In vacuum, two transitions were visible in the 500-700 nm range (λ = 563 and 605 nm) resulting in an average large band centered around 585 nm when a dispersion corresponding to $\sigma = 0.2$ eV was applied to the spectrum (Figure S1C). The band at 563 nm corresponds to a transition from the HOMO β anti-bonding orbital between Ru and Cl atoms to the second β antibonding virtual orbital located between Ru and O (π -> π *). A second transition was also calculated around 450 nm, that was hidden by the absorption of excess PIDA in the experimental spectrum and the Ru1 and NikA as well (Figure S1). In presence of the protein residues, similar observations were made but resulting in an even broader band. The concordance of energy bands between theory and experiments supports the hypothesis of a the formation of an oxidizing L(CO)(CI)Ru=O species.

The absence of more direct positive experimental proofs urged us to explore both the epoxidation and chlorohydrination reaction mechanisms at a theoretical level as a function of substrates to define both the nature of the active species and the reaction specificity. All the above pathways including PIDA

adducts, were then confronted to extensive calculations in order to discriminate their occurrence. When done, their compatibility with experimental data should help to elucidate the reaction pathway.

Computational studies

The possible reaction pathways leading from MOS or BMS to epoxide and chlorohydrin were evaluated using quantum chemistry computations. Since all-electron calculations are not accessible for *Ec*NikA-Ru1/2, our approach consisted of modelling the Ru1 or Ru2 complex either alone or in the presence of the close amino acids residues that are involved in the ruthenium complex stabilization *via* supramolecular interactions.

Calculations were run using the Gaussian g09 program.²⁵ Both singlet and triplet states were calculated for all the species presented here.

Using the DFT method with B3LYP functional, 9 a def2-SVP split valence polarization basis set 10 and a Polarizable Continuum Model (IEFPCM) 26 for the solvent, we first examined the geometry of the starting **Ru1** complex (Figure 4, top). We compared the metrics obtained from the x-ray structure of *Ec*NikA-**Ru1** 4 (PDB code: 5L8D, Figure 2 top right) with the computed ones of the complex in the singlet (S=0) and triplet (S=1) state. The involvement of the protein scaffold has been partially modelled by adding five residues (Arg137, Tyr22, Tyr402, Trp100 and Trp398) to the calculations. Only the modelled triplet state showed similar CO bond lengths to the crystal structure. Most importantly, the modelled triplet state predicted the curvature of the *t*CO at the cost of the Ru-O_{carb} length. However, the triplet state **Ru1** is nearly 50 kcal mol $^{-1}$ higher in energy than the singlet state, prohibiting its existence in solution. Subsequently, **Ru2**, observed by x-ray crystallography, was modelled (Figure 4, bottom). The modelled structure more closely matches the bond lengths and angles observed in the crystal structure of *Ec*NikA-**Ru2** (PDB code: 6R4Q, Figure 2, bottom right) when the singlet state is considered. This result was replicated when the complex was modelled in the absence of the nearby residues (Figure S2). Altogether, these observations validate the calculation methodology employed and lead us to describe **Ru1** and **Ru2** as S=0 species.

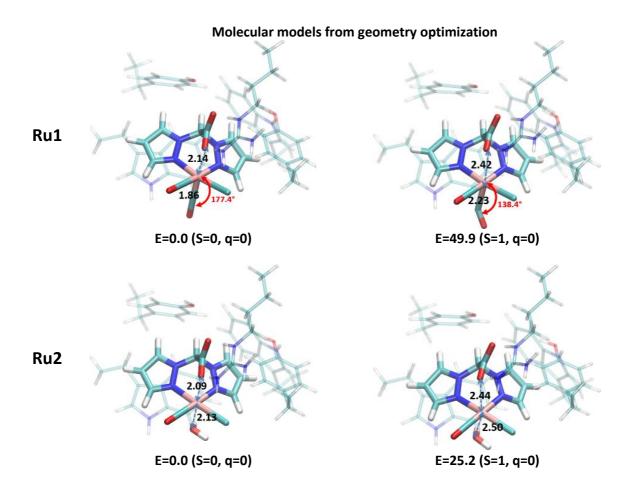


Figure 4. Comparison of bond lengths (black) and angles (red) of modelled **Ru1** in the presence of five amino acids in the singlet (S=0) and triplet (S=1) states (first line) and modelled **Ru2** in the presence of five amino acids in the singlet and triplet states (second line). All energies in kcal mol⁻¹. The Ru-OH₂ and Ru-O_{carb} bonds are present even if not drawn by the program.

In order to discriminate between active species, two relevant forms of the ruthenium complex (activated by PIDA in the presence of NaCl) were considered: Ru^{IV}(Cl)=O and Ru^{II}(Cl)(OCl) (Figure 5). Epoxidation is well described in the literature as being related to the presence of hypervalent oxo species, while Ru^{II}OCl can be a source of chloronium for chlorination. ^{18d,24,27}

Ru^{II}(CI)(OCI) is more stable in the singlet state (28.0 kcal mol⁻¹ lower in energy than the triplet state) but the charge on the chloronium atom is only 0.07, making it a weak electrophile. Similarly, calculations attempted to make Ru^{II}(CI)(OCI) react with alkenes led to a non-conclusive pathway. Considering these results, the hypothesis of Ru^{II}(CI)(OCI) being the active oxidation species was discarded. The Ru^{IV}(CI)=O complex is more stable in the triplet spin state than in the singlet state by 15.7 kcal mol⁻¹ but in both cases, atomic partial charges do not account for an electrophilic character of the double-bonded oxygen atom or the chlorine atom.

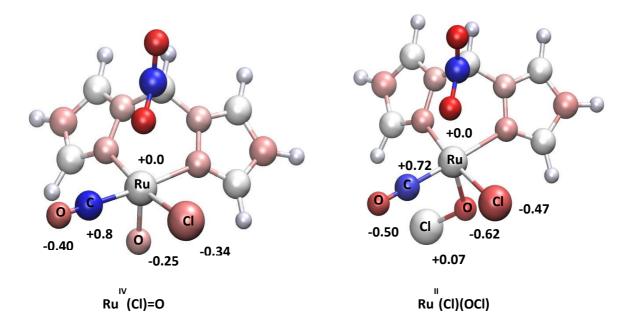


Figure 5. Atomic partial charges and geometry of the Ru^{IV}(CI)=O (charge 0, spin 1) and Ru^{II}(CI)(OCI) (charge -1, spin 0) activated complexes. Sizes and colors of balls are representative of relative atomic partial charges. The Ru-O_{carb} bond is present even if not drawn by the program.

The reaction mechanisms of hypervalent iodine compounds with metal complexes are poorly described in the literature, with only few examples of iron complexes and even fewer with ruthenium. 27-28 Although no active ruthenium oxidant has been directly observed in EcNikA-Ru1/2, all empirical evidence points towards the existence of a metal-centered species responsible for transferring an oxygen atom during catalyzed epoxidation. To assess the process by which the oxidant acts on the ruthenium center, it was necessary to probe the reaction between Ru1/2 and PIDA using small QM models. These calculations did not evidence any reactivity between Ru1 and PIDA. Conversely, after geometry optimization of Ru2, a symmetrical hydrogen bonding structure was obtained prior to any reactivity (species (a), Figure 6). The C=O groups of each acetate moiety became angled towards the protons of the bound water. The reduction in energy caused by such an interaction is magnified due to the absence of explicit water molecules, which would provide alternative hydrogen bonding opportunities when the oxidant and Ru2 are well separated. Explorative calculations were trialed using explicit solvent (1–5 molecules of H₂O); however, the additional degrees of freedom associated with explicit solvent molecules led to sampling issues with false energy minima, along with a significant increase in computational cost. The formation of a bond between the bound water and iodine of the oxidant led to the release of a single acetate group (species (b), Figure 6), maintaining the triply bonded hypervalent iodine motif and liberating a proton from the water. Similar energetic barriers were found for the initial attack of the iodine on the water in the S=0 and S=1 states (21.0 and 19.3 kcal mol⁻¹, respectively). However, the intermediate Ru^{II} species was found to be relatively unstable in the triplet state. When the iodine was pulled away from the ruthenium complex using a potential energy surface (PES) scan, species (**b**) underwent simultaneous breakage of both I-O bonds, releasing free AcOH and PhI. The iodine gained two electrons from the oxygen of the Ru-bound water, at which point the formal oxidation state of the ruthenium center changed from +2 to +4, and the Ru-O bond shrank from 2.01 Å to 1.81 Å, indicating that a Ru=O had formed. Unlike the singlet-state pathway, no intermediate species were observed between (**b**) and the Ru^{IV}(CI)=O (species (**d**), Figure 6). In the singlet state, species (**b**) was found to be stable, but significantly higher in energy than its separate components (species (**a**), Figure 6). Upon removal of the second acetate group and proton (species (**c**), S=O pathway), the energy increased by a further 15.3 kcal mol⁻¹, yet the O-I bond remained intact. The stability of such a relatively high-energy species is testament to the resistance of the singlet state complex to form an oxo species. When the remaining PhI group was removed by PES scan, followed by a reoptimization, the Ru^{IV}(CI)=O was observed (species (**d**)).

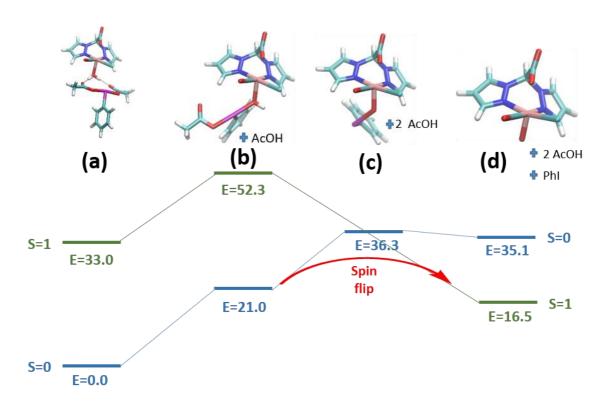


Figure 6. Thermodynamics of conversion of **Ru2** to Ru^{IV}(CI)=O using PIDA. The red arrow indicates the path of least resistance to a stable oxo, assuming a change in spin state is possible. All energies in kcal mol⁻¹. The Ru-O_{carb} bond is present even if not drawn by the program.

The formation of an oxo species in the triplet state, although involving a net decrease in energy of 16.5 kcal mol⁻¹, is clearly thermodynamically disfavored. Therefore, three possibilities for reactivity

were pursued. Firstly, it is possible that singlet multiplicity is maintained throughout, and that the active oxidant in the *Ec*NikA-**Ru1/2** catalytic cycle is the species (**b**), S=0 (blue pathway).

Secondly, in the triplet state (green pathway), both the initial energy and the energy barriers seem to preclude a pathway between reactants and products. However, the calculations present several shortcomings. The implicit solvent conditions may not account for the dielectric constant inside the protein cavity, and the calculations do not take into account the gain in entropy resulting from the liberation of the AcOH and PhI molecules. The size of the considered molecules excluded rigorous QM/MM calculations with the protein present. It has been shown in the literature that similar transition metal species can change their spin state during redox reactions.^{27,29} It is therefore possible that the ruthenium center undergoes a change in spin state upon oxidation, with the reactants beginning in the singlet state, and ending in the triplet state oxo, as shown by the red arrow and a third path on Figure 6.

Therefore, the active oxidizing species could be the Ru^{IV}(CI)=O species (**d**) (singlet or triplet state), or the ruthenium-iodine adduct (**c**). In order to make a decision, the MOS carbon-carbon double bond was moved towards the oxygen atom of the singlet state species (**c**), minimizing the energy at each step along a PES scan. No thermodynamically viable reaction pathway was obtained for this system. Indeed, a single well-defined minimum energy path for the approach of the olefin was difficult to obtain and resulted in very high-energy barriers. This calculation was repeated, scanning the alkene towards the iodine atom in an attempt to form the iodinium cation species proposed by Yang *et al.*,³⁰ but again, no low energy reaction path was found. These results led us to consider Ru^{IV}(CI)=O (**d**) as the only possible active oxidizing species for the oxidation of MOS and BMS. To cope with the problem of weak, non-bonded interactions between Ru^{IV}(CI)=O and the substrates before the reaction (absent with the B3LYP density functional), an empirical dispersion correction (*GD3BJ*)³¹ term was added in the calculations.

We then investigated substrate oxidation reactions. An energetic pathway was mapped out for the complex in both S=0 and S=1 states. In the case of MOS, a summary of the computations is presented in Figure 7 (see Figure S3 for all the calculated structures and Table S5). The starting point is species (e), the Ru^{IV}(CI)=O complex (species (d), Figure 6) in the presence of MOS. In the case of S=0, close-shell calculations led to only one transition state (TS1) slightly uphill (6.4 kcal mol⁻¹), which gave the epoxide product (g) downhill by 37.1 kcal mol⁻¹. In the case of S=1, two competing pathways were observed, following an attack of the oxygen atom on either the α or the β carbon atom of the double bond. In both cases, we observed a two-steps mechanism. Initial attack on the C β led to an overall lower energy pathway. The first transition state (TS2) was significantly lower than when C α attack was

considered (7.5 vs 17.1 kcal mol⁻¹) and the second energy barrier was computed to be 11.7 kcal mol⁻¹ lower than for the attack on the C α . As expected, spin and NPA charge calculations demonstrated that a radical species is formed on the carbon atoms and a Ru(III) in (**TS2**), (species (**f**) and (**h**), Figure 7, Table S6). The intermediate is better stabilized (-13.3 vs 3.5 kcal mol⁻¹) when the radical is located on the C α atom (**f**). Nevertheless, the two pathways give the same epoxide product (species (**g**), Figure 7). Comparing all pathways, the only observed product is the unique epoxide (**g**).

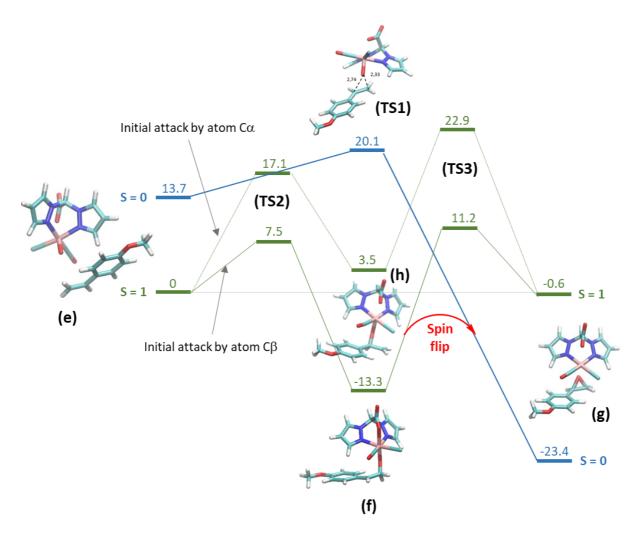


Figure 7. Reaction diagram showing the B3LYP calculated energies for epoxidation of MOS, catalyzed by Ru^{IV}(CI)=O. S=1 state calculations are shown in green, and the close shell S=0 process in blue. Energies given in kcal mol⁻¹ relative to the energy of the starting species in S=1 state. The Ru-O_{carb} bond is present even if not drawn by the program.

Next, we examined the reaction mechanism on BMS. Again, an energetic pathway was mapped out for the complex in both S=0 and S=1 states (Figure 8; see Figure S4 and Table S5 for all calculated structures). The starting point is the species (i): the Ru^{IV}(CI)=O complex (species (d), Figure 6) in the presence of BMS. In the case of S=0, close shell calculations led to a nearby transition state (**TS4**) (only

2.9 kcal mol⁻¹ uphill), followed directly by product (j) downhill by 33.1 kcal mol⁻¹. Surprisingly, this product is not the predicted epoxide, but rather a chlorohydrin. In the case of S=1, we again compared the two competing pathways following an attack of the oxygen on either the α or β carbon. As expected, an initial attack on the C β led to lower energy barriers, with (TS8) lying 7.4 kcal mol⁻¹ lower than (TS5), and an energy lowered by 12.0 kcal mol⁻¹ for the radical intermediate (k) (Table S6).

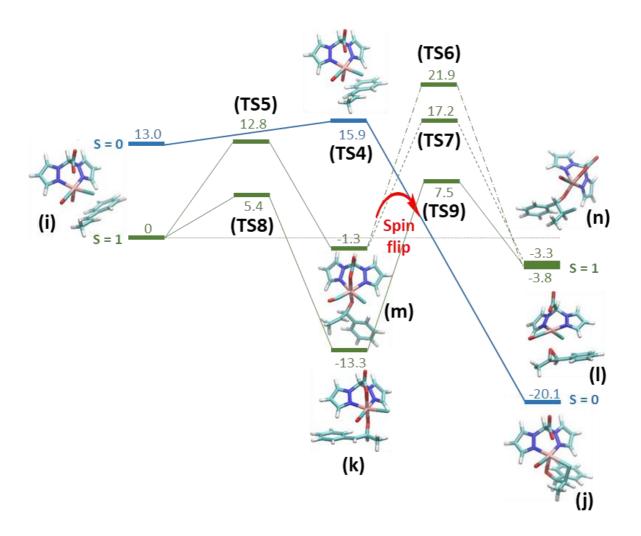


Figure 8. Reaction diagram showing the B3LYP calculated energies for epoxidation and hydroxychlorination of BMS, catalyzed by Ru^{IV}(CI)=O. S=1 state calculations are shown in green, and the close shell S=0 process in blue. Energies given in kcal mol⁻¹ relative to the energy of the starting species in S=1 state. The Ru-O_{carb} bond is present even if not drawn by the program.

Nevertheless, following the reaction from (**k**) *via* **TS9** was found to lead only to the epoxide (**I**). Looking carefully at the structure of (**k**) (Figure S4C, spacefill representation), it was possible to identify an important interaction between the chlorine atom held by the ruthenium and the methyl group of BMS, avoiding the rotation necessary to a subsequent chlorohydrination. The other possible pathway

followed an attack on the $C\alpha$ through (**TS5**). The radical intermediate (**m**) could evolve to both epoxide (**I**) and chlorohydrin (**n**) *via* activation barriers of 18.5 kcal mol⁻¹ (**TS7**) and 23.2 kcal mol⁻¹ (**TS6**), respectively. Comparing all the pathways, the thermodynamic product is the unique chlorohydrin (**j**) in the singlet state.

Looking for differences between the reactivity of radical intermediates (\mathbf{m}), (\mathbf{k}) and (\mathbf{f}) coming from BMS and MOS, respectively, we explored the PES depending on the orientation of the substrate facing the chlorine atom (Table S7). In the case of MOS, the shortest distance between the chlorine and the radical carbon is 4.37 Å, while in the case of BMS, the distance ranges between 3.38 and 5.46 Å. One of the shortest (3.64 Å) is found for intermediate (\mathbf{m}) (Figure 8) at -1.3 kcal mol⁻¹ which may lead to the formation of a Cl-C β bond and hence the chlorohydrin.

Finally, we recalculated the majority of the reaction paths in the presence of the protein. Quantum calculations in the presence of protein residues surrounding the active site demand high computational resources. They were thus restricted to the closest environment in the reaction pathways. Residues in immediate proximity to the complex are arginine 137 (interacting with the carboxylate of the Ru complex), tyrosines 22 and 402, tryptophans 100 and 398. Ru IV (CI)=O was used as a template to correctly position BMS and MOS (in complex with Ru IV (CI)=O) in the different conformations presented above but inside the protein. The coordinates of BMS and MOS were obtained by least square superimposition of the template in vacuum versus in the protein (reduced to 5 amino acids). The C α and C β atoms of the five protein residues (with coordinates taken from the x-ray structure) were fixed during the calculations. All of the results are presented in Table S8.

For BMS, the energies along the reaction paths were comparable with or without the protein residues. For example, the transition state corresponding to O-C α bond formation ((**TS4**) in Figure 8) was retrieved (Figure 9, top left). This transition state led to the 5-atom ring with $\Delta E = -36.7$ kcal mol⁻¹ (comparable to -36 kcal mol⁻¹ from (**TS4**) to (**j**) in the absence of the protein). In the case of MOS oxidation, the singlet state could not be calculated, as we were unable to characterize a transition state where MOS would not clash with Trp398 (Figure 9, top right). Another limitation imposed by the protein is illustrated in the case of the formation of an epoxide from BMS: no analog of product (**g**') could be computed when replacing MOS with BMS, since the methyl group clashes with the same Trp398 (Figure 9, Bottom right).

Exploring the PES for the most stable species, we found another impact of the protein on the reaction. There are relative conformations of the substrate and Ru complex, equivalent to states (e) or (i), with lower energies but non-reactive. For example, relaxation of (TS2) for MOS led to an equivalent of (e) but lying 2.6 kcal mol⁻¹ downhill (Table S9). The effect is even clearer for BMS and MOS in open-shell

singlet state configurations, where energy reductions of 9.9 and 12.2 kcal mol⁻¹ were obtained, respectively, when allowing structures incompatible with the presence of the protein. Note also from Table S9 that the lowest reactive MOS orientation in the singlet state, compatible with the protein, has an energy of 16.4 kcal mol⁻¹, 2.7 kcal mol⁻¹ higher than structure (e) presented in Figure 7. When these lowest energy species are superimposed on the 5 amino acids representing the protein, the stabilized substrate always clashes with Trp398 (Figure S5). We therefore concluded that the main role of the protein might be to preclude the formation of these stable unreactive species, rather than lower the barrier between reactants (reactive species) and products.

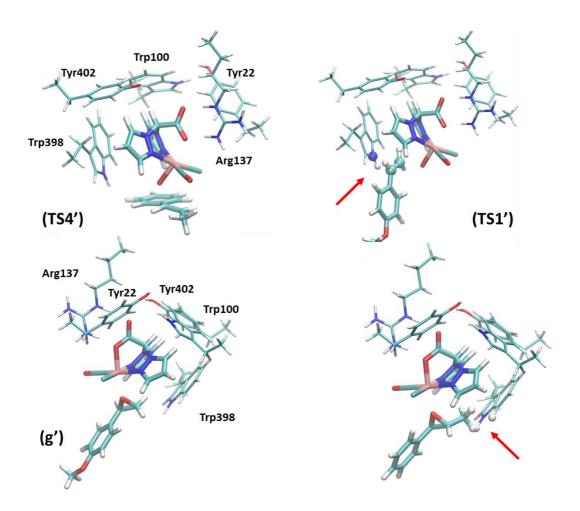


Figure 9. <u>Top</u>: Transition states for product formation between Ru^{IV}(CI)=O and BMS ((**TS4'**), left) or MOS ((**TS1'**), right), open shell singlet states, in the presence of surrounding protein residues. Note the steric clash between MOS and Trp398 highlighted with CPK spheres for atoms closer than 2 Å from each other. <u>Bottom</u>: Epoxide complexes between Ru^{IV}(CI)=O and MOS ((**g'**), left) or BMS (putative, right), singlet state, in the presence of surrounding protein residues. Note again the steric clash between BMS and Trp398. The Ru-O_{carb} bond is present even if not drawn by the program.

Discussion

All experimental observations led us to rule out a mechanism with a chloronium bridge ion as an intermediate, and instead suggested the formation of Ru^{IV}(CI)=O as the sole (terminal) oxidizing species (common behavior between PhIO and PIDA, observation of an electronic transition for an oxidizing intermediate indicative of high valent ruthenium and finally the formation of epoxide). The catalytic behavior of EcNikA-Ru1 was then evaluated computationally in order to decipher the reaction pathway, taking into account the chemoselectivity, regioselectivity and stereoselectivity observed. First of all, it has to be noted that, due to computational costs, it would have been impossible to examine all models encountered in the complete pathways using the full hybrid structure by QM/MM. Accordingly, Ru1 and the reactivity of Ru2 with PIDA were investigated. Computations showed that Ru1 is not an active species. Subsequently, it was proposed that the combined electron withdrawing effects of the ruthenium center of Ru2 and the bound proton could all contribute to a heightened electrophilicity of an 'activated water' species. This facilitates a nucleophilic attack on the hypervalent iodine atom of PIDA to form the Ru^{IV}(CI)=O species, through a Ru-O-IPh intermediate, shown in a genuine calculated energy profile. All other possible reaction pathways were found too costly energetically. Hence, the Ru^{IV}(CI)=O was determined as the active species for the epoxidation of MOS and hydroxychloration of BMS.

We first examined the case of MOS. In the S=0 state, the formation of the epoxide proceeds in a single step with a low barrier of 6.4 kcal mol⁻¹, and is highly exothermic ($\Delta E = -37.1$ kcal mol⁻¹). The spin-state and NPA charge calculations evidenced the oxidation state conservation on the ruthenium and the absence of radical or carbocation character on the MOS molecule (Table S6). Moreover, the final product (**g**) shows a spin increase on the ruthenium, attesting of the reduction of the ruthenium ion during the process. This was expected for two reasons: firstly, the Ru^{IV}(CI)=O complex is far less stable in the singlet state than in the triplet ground state ($\Delta E = 13.7$ kcal mol⁻¹), leading to a higher initial energy of the reactants. Secondly, following the reduction of Ru(IV) to Ru(II), the singlet state becomes notably more stable than the triplet ($\Delta E = -22.8$ kcal mol⁻¹). Interestingly, open shell energies for the singlet states (**e**) and (**TS1**) are significantly reduced compared to close shell values reported above from 13.7 to 4.2 and from 20.1 to 15.6 kcal mol⁻¹, respectively (Table S5). The initial barrier thus increases from 6.4 to 11.4 kcal mol⁻¹ and the triplet pathway remains more favorable.

In the triplet state, the path of α -carbon attacking first is clearly disfavored when compared with the β -carbon attack. Not only does the stability of the triplet-state intermediate increase ($\Delta E = 16.8$ kcal mol⁻¹), but the initial transition state (**TS2**) is also lowered in energy ($\Delta E = 9.6$ kcal mol⁻¹). The second transition state (**TS3**) then requires 19.4-24.5 kcal mol⁻¹ relative to the stable intermediates. The height of this barrier is problematic for a catalytic process, and begs the question of why these stable

intermediates have never been observed. A possible solution rests on the ability of the ruthenium center to change its spin state. Once the first low barrier (TS2) has been overcome, the oxo ligand is inserted into the double bond and the ruthenium center is no longer Ru(IV). Analysis of the (TS3) and the final product indicates a Ru(II) state, as the spin on the ruthenium atom increases to close to 2 and no residual spin is present on the Cβ. Therefore, the enhanced stability afforded by the S=1 state disappears, and S=0 becomes strongly favored. In summary, if the ruthenium center was able to switch from S=1 to S=0 following the formation of the first C-O bond, the processes afterwards would be barrierless, avoiding (**TS3**) (red arrow). Taken as a whole, this would result in an overall $\Delta E = -23.4$ kcal mol⁻¹ for MOS epoxidation, much more favorable than the -0.6 kcal mol⁻¹ found when the reaction is constrained to the triplet state. Iron systems have previously been described which exhibit a remarkably similar reaction profile. Iron porphyrin systems have been shown to perform barrierless, 1-step oxidations in the low spin state and stepwise radical rebound oxidations in the high spin state. ^{27,29a} Spin inversions have also been proposed to be crucial in epoxidation reactions catalyzed by manganese, chromium and ruthenium. 27,29b,29c For example, to understand the mechanism of epoxidation by the complex Cp*Cr(O)Cl2, Hess et al. turned to first-principles theory to calculate the minimum energy crossing point between the two PES of the chromium III and V potential oxidant in doublet and quartet spin multiplicity, respectively. They determined that the surface crossing is fast and opens a lower barrier pathway. ^{29c} However, taking into account the whole reaction path starting from Ru2, envisaging a reaction from species (a) to (g) in the S=0 state through species (d) and (e) in the S=1 state represents two successive spin flips. A whole pathway in the S=0 state could also be envisaged, with the terminal oxidant being the Ru^{IV}(CI)=O complex (\mathbf{d}) in the S=0 state and the driving force the strong stabilization of the product.

Finally, no intermediates leading to a chlorohydrin type product or the chlorohydrin itself was observed along all the calculated pathways. A chlorohydrin product can be modelled *via* a chloride attack on the epoxide, but occurs *via* a supplementary energy barrier of 27.4 kcal mol⁻¹ (Figure S3, A), excluding its formation. When the Ru^{IV}(CI)=O complex reacts with MOS, the overall effect of introducing the olefin is a rapid, concerted epoxidation, driven by a strongly favorable two-electron reduction of Ru(IV) to Ru(II).

In the case of BMS, the most favorable path appears to be the direct formation of a chlorohydrin in the singlet spin state. In the S=0 state, the formation of the chlorohydrin proceeds in a single step with a low barrier of 2.9 kcal mol⁻¹ and is highly exothermic ($\Delta E = -33.1 \text{ kcal mol}^{-1}$). Open-shell energy for state (i) is reduced from 13.0 to 6.4 kcal mol⁻¹, while the energy of the transition state (TS4) is unchanged. The initial barrier for chlorohydrin formation would thus increase from 2.9 to 9.5 kcal mol⁻¹. In the triplet state, an attack on the β -carbon leads to a lower energy pathway but with an

epoxide as the sole final product, and a higher energy level for the second transition state. The latter offers the possibility for ($\bf k$) to come back to state ($\bf i$) and follow the pathway of α -carbon attacking first, affording the regioselectivity experimentally observed. Reaching (**TS5**) requires 12.8 kcal mol⁻¹, roughly the same amount of energy as a spin flip from S=1 to S=0 for the ($\bf i$) species. Then, even if (**TS6**) is disfavored when compared with (**TS7**), ($\Delta E = 4.7$ kcal mol⁻¹), starting from species ($\bf m$), two pathways coexist leading to the epoxide ($\bf l$) and the chlorohydrin ($\bf n$), respectively. Again, spin and charge calculations attest of the radical character of intermediate ($\bf m$), similar to intermediate ($\bf f$) in the case of MOS oxidation. According to these calculations, reaction of species ($\bf i$) in the triplet state should result in a mixture of products with an overall ΔE around -4 kcal mol⁻¹, and not the 100% chemoselectivity observed in favor of the BMS-chlorohydrin. In summary, as observed for MOS, the best overall pathway for BMS oxidation would follow a pathway from ($\bf a$) to ($\bf j$), both in the singlet state after two spin flips from S=0 to 1 and back to 0, but with an initial reaction between the Ru^{IV}(CI)=O complex and the α -carbon of the substrate. It is noteworthy that ($\bf j$), in the presence of chloride and water, can dissociate barrierless into ($\bf a$) and the β -chlorohydrin (Figure S6).

The influence of the protein on this reactivity was then evaluated through the introduction of five amino acids close to the ruthenium complex. No significative changes were observed in terms of energy barriers. Conversely, some intermediate species were found to be conflicting with the presence of Trp398, which helps to discriminate between reaction pathways by controlling the orientation of the substrate and, therefore, the kinetics of the reaction. Thermodynamic energies favor an epoxide as the major species in most of the computed cases, whereas the kinetic β –chlorohydrin results from an attack on the substrate $C\alpha$, whereby the chlorine is the closest to the $C\beta$ radical carbon. Moreover, the presence of Trp398 helps to destabilize the enzyme-substrate complex, leading to catalysis. This aspect reveals that our system possesses allmost of the characteristics of an enzyme, *i.e.* a control of the substrate binding and orientation in an active site (see Figures S7 and S8 for a comprehensive point of view). This role adds to the promotion of the starting **Ru2** species, as the hydrophobicity and the neutral charge of the protein environment affect the lability of the tCO.

The calculations have shed light on the important energy barriers for most of the reaction pathways based on Ru1/2 studied here. For example, the activation of PIDA seems unrealistic unless we consider a spin crossover during the process. This outcome is related to the stability difference between the Ru(II) starting complex and the Ru(IV)=O oxidative species, the former being S=O and the latter S=1. This difference is also present in the second set of calculations for the reactivity of Ru^{IV}(CI)=O with the alkene substrate. There are then two ways to depict the dynamics of the reactions: one considering the whole process from PIDA activation to product release. In this case the reaction is driven by the large exothermic final step, and can be considered to occur at constant spin S=O. An alternative consists

of two spin crossovers (S=0 to S=1 and then S=1 to S=0) during the two main reactions. In this case, the energy barriers for all transformations will stay below 21 kcal mol⁻¹ (21.0 for the activation of PIDA, 12.8 and 7.5 kcal mol⁻¹ for the oxygen transfer reactions). The second spin flip has also been proposed for iron oxene for OAT³² and recently for a Ru(IV)=O species.²⁷ The latter example displayed the same free energy profiles as a function of spin states as the ones in our study but for the epoxidation of styrene. Here, in the case of MOS, the two-step process in the S=1 state consists of formation of a Ru-O-C β bonded intermediate with a radical on C α , followed by the closing of the epoxide ring. The second step was calculated with a 24.5 kcal mol⁻¹ barrier, too high to be realistic. The proposed spin crossover at this stage makes this step more feasible. As we cannot discriminate between the two processes, we can concentrate our effort, rather, on conciliating calculated reactivity and the different selectivities observed experimentally.

In summary, we have deciphered the reaction mechanism keeping in mind the selectivities observed. All the calculations are in line with the experimental observations. First, the thermodynamics may explain the difference in chemoselectivity between the two substrates. In the case of MOS, the formation of the chlorohydrin is too costly in energy compared to the epoxide formation. With BMS, access to both products is possible but the chlorohydrin is preferred (ΔE ± 17 kcal mol⁻¹ considering the spin state). Second, the regioselectivity for the chlorohydrin formation is explained by the radical character of the reaction. The slightly negative partial charge calculated for the double bonded oxygen atom in Ru^{IV}(CI)=O precludes any electrophilic addition on alkenes. The radical attack of the Ru^{IV}(CI)=O on the $C\alpha$ of the aryl alkene double bond forms a Ru-O- $C\alpha$ bonded intermediate, (m) or (f) for BMS and MOS, respectively. This Ru(III) intermediate has a free radical character on Cβ (Table S5) that may potentially rotate along the $C\alpha$ - $C\beta$ bond before the chlorine atom insertion or the epoxide ring formation. In the case of BMS, two structures of the chlorinated product have been modelled depending upon the spin state considered ((j) and (n) for S=0 and S=1, respectively), revealing the direct insertion of the chlorine ligand from the ruthenium into the radical substrate. They differ only by the breaking of the Ru-Cl bond. Their existence, in addition to the radical character of the reaction, should exclude the pathway implicating the insertion of a chloride anion from the buffer. Finally, in terms of diastereoselectivity, the S=0 pathway does not fit the 73% diastereomeric excess obtained for the BMS oxidation that should be 100% if the oxygen and chlorine atoms were both inserted in a concerted way. Then, it is assumed that the S=1 pathway should be privileged in this case.

A plausible reaction mechanism is displayed in Figure 10 based on the common intermediate Ru^{IV}(CI)=O, as determined by the QM calculations. The selectivity between hydroxychlorination and epoxidation is driven by a competition between a thermodynamic and a kinetic pathway. In the case of MOS, the reaction follows a thermodynamic pathway, passing by the lowest energy barriers and the

more stable intermediate, a $C\alpha$ radical stabilized by the methoxyphenyl substituent. This stabilization allows the reorganization necessary to the formation of the epoxide from the intermediate. In the case of BMS, the reaction follows the kinetic pathway passing by an attack on the less-hindered $C\alpha$ atom leading to an unstable $C\beta$ radical that reacts rapidly with the near chlorine atom. In both cases, the presence of a radical intermediate allows rotation along the $C\alpha$ - $C\beta$ bond (black arrows in Figure 10) precluding the obtaining of any ee. At the end, the starting Ru(II) is formed again, confirmed by spin and NPA charge, after the coordination of a chloride, if necessary, and a water molecule to start a new cycle.

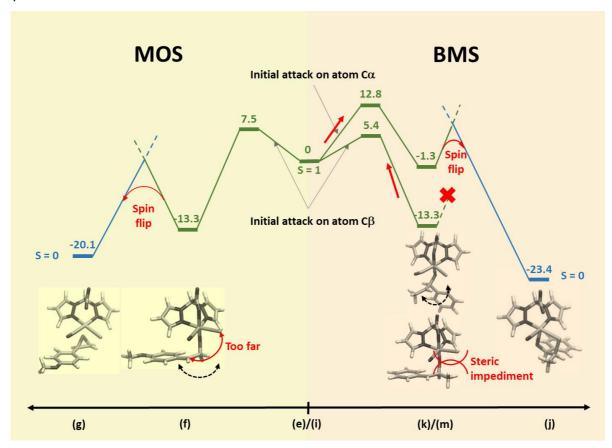


Figure 10. Comparison between MOS and BMS reactivity.

Conclusion

The design of new artificial enzymes is of great interest for abiotic chemical reactions. In this study, thanks to x-ray crystallographic data and combined experimental and computational studies, we have tried to clarify the impact of the insertion of a Ru(II) complex into a protein cavity in remarkable abiotic oxidation catalysis conditions. While a full QM/MM study was unreachable, the QM evaluation of the complex alone and the experimental results helped us to define reasonable reaction pathways. We were able to propose a mechanism, revealing the different reaction pathways between two close substrates. The full reaction pathways were examined, revealing for the first time

a reasonable route for $Ru^{IV}=O$ formation from a hypervalent-iodine-based oxidant. During the calculations, the question of the spin transitions of the ruthenium at the two modelled stages of the whole process was not settled. The role of the protein environment was highlighted, and it was demonstrated that the selectivity of the artificial enzyme is due to i) the promotion of the starting oxidizing species via the exchange of a CO ligand, and ii) the control of the substrate orientation on the intermediate structures, formed after the $Ru^{IV}=O$ attack: when $C\alpha$ attack is preferred, chlorohydrin is formed while attack on $C\beta$ leads to an epoxide. This work contributes to the promotion of artificial metalloenzymes as alternative to natural enzymes for the purpose of non-natural reactions.

Experimental part

Hybrid preparation

*Ec*NikA was purified and *Ec*NikA-**Ru1** was prepared as previously described.⁴ The synthesis of *Ec*NikA-**Mg(EDTA)** was performed by incubating 500 μL of a 500 μM protein solution with a 2-fold excess of Mg(EDTA) in 40 mM Tris-HCl pH 7.5 at 4°C overnight. Complex excess was removed using a desalting column (PD10), followed by concentration on a centricon filter with a 30 kDa cut-off (4 cycles) which led to the separation of the protein-complex hybrid from the complex-free solution.

Catalytic standard procedure

10 μ L of a 496 μ M *Ec*NikA solution (5 nmol) in a Tris.HCl buffer 40 mM pH 7.0 were mixed with 30 μ L of a 8.10⁻⁴ M complex solution (25 nmol) for 30 minutes. Then, 88 μ L of HEPES buffer 10 mM pH 7.0, 6 μ L of a 0.41 M solution of 4-methoxystyrene (2.46 μ mol) and 1 mg of iodobenzene diacetate (3 μ mol) were added. The reaction was performed at room temperature for 10 minutes, and the reaction mixture was extracted with AcOEt (200 μ L). The organic phase was analyzed by GC/MS.

Theoretical calculations methods

Oxidation of *trans*-β-methyl- and 4-methoxy-styrene were studied using computational quantum chemistry with the Gaussian g09 program.²⁵ The DFT method and B3LYP functional⁹ with the def2-SVP split valence polarization basis set¹⁰ and empirical dispersion correction (*GD3BJ*)³¹ were used by default for most calculations. The Polarizable Continuum Model (PCM) was used to describe the solvating water. All the calculations were done in open shell mode (UB3LYP) but the Gaussian singlet state wave functions and / or optimization algorithms seem to always result in strict 0.0 spin contributions for all atoms. To enforce singlet state open shell calculations with non-zero spins on all atoms, we optimized

singlet state geometries with the corresponding triplet state wave function as initial guess (Guess=Read). Stability of the wave functions has been carefully checked in each case.

Most calculations have been reproduced in the absence of the dispersion correction term and relative energies are similar to those presented in the text.

Calculations were also checked by changing basis set and functional and compare the energy differences between singlet and triplet states for most optimized geometries and transitions states reported with the def2-SVP basis set. The meta-GGA M11-L dual-range exchange functional³³ was used with the triple zeta Def2TZVP basis set.^{10,34} Some variations are observed between the two methods but the conclusions are identical.

Supporting information. Catalysis tables, x-rays structure refinement and statistics, computational details. This information is available free of charge on the ACS Publication website.

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