

AEROBIC AQUEOUS REDOX CATALYSIS BY A RUTHENIUM-HYDRIDE
SPECIES AND EVIDENCE FOR THE REDUCING/HYDRIDE TRANSFER ABILITY
OF BIOLOGICAL OXIDANTS

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ABSTRACT

Reactive oxygen species (ROS), such as superoxide and hydrogen peroxide, have been classically viewed as being monolithically harmful to biological systems, only leading to disease and dysfunction. This understanding has evolved recently in light of new findings; 1) ROS are essential in fighting infections and 2) antioxidant enzyme, catalase that removes ROS (antioxidant activity) can also generate ROS (pro-oxidant activity) depending on the identity of the corresponding oxidized products.

We hypothesize that a complex that catalyzes oxidization of alcohols will be able to catalyze the reduction of radicals. We chose an organoruthenium catalyst (**Ru1**), air-stable and soluble in water-miscible solvents, as a catalyst for aerobic aqueous redox reaction and 2,2'-azino-bis(3-ethylbenzo-thiazoline-6-sulfonate) radical monoanion (ABTS^{•-}) as a model for radical species due to its comparable oxidizing potential to ROS. Because radicals can be reduced using alcohols as terminal reductants, we chose biologically-relevant alcohols that can be recycled to their reduced states by our body.

We have shown that **Ru1** catalyzed the reduction of ABTS^{•-} to its precursor ABTS²⁻ in phosphate buffered saline (pH 7.4). In doing so, **Ru1** used biologically-relevant alcohols containing CH-OH groups such as NAD⁺, amino acids, sugars, citric acid cycle metabolites as terminal reductants. Mechanistic evidence reveals that the catalytic radical reduction is achieved by a Ru-hydride intermediate formed by a β -hydride elimination from a ribose subunit in NAD⁺ and from a Ru-alkoxide species in citric acid cycles metabolites. These findings demonstrate the undiscovered reducing ability of biological oxidants. We have also revealed a new potential therapeutic strategy

by Ru catalysts that use the same type of Ru-hydride intermediate by demonstrating the catalytic antioxidant effects.

Collectively, these findings illustrate a central principle of redox therapeutics: it is not the identity of the redox catalyst itself that determines whether it produces pro- or antioxidant effects, but rather the identities and concentrations of the species being reduced (the terminal oxidant) and the species being oxidized (the terminal reductant).

DEDICATION

Never despair, keep smiling
Better than wealth with its carriage and pair
Better than rank or a face wondrous fair
Is a heart that life burdens can cheerfully bare
Just a brave loving heart that never despairs

B Wallis April 23rd, 1916

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CHAPTER ONE
CELLULAR REDOX SYSTEM: THE DUAL ROLES OF OXIDANTS AND
REDUCTANTS

“Life is nothing but an electron looking for a place to rest”.

“~Albert Szent-Gyrgyi”

1.1 HISTORICAL BACKGROUND AND SIGNIFICANCE

Life is a constant battle to overcome entropy. Living cells simultaneously synthesize thousands of molecules ranging from small inorganics to protein, carbohydrate, fat, and enzymes in the right proportions, only at the right time, in a very precise manner.¹ Energy is required to keep and maintain this well-regulated system in order. This required energy is retained from the flow of electrons during cellular redox reactions.¹ Redox reactions are generally linked to one another and altogether they form the basis of highly complex, interconnected and dynamic cellular network.¹

Cells proliferate in reducing environment² than in oxidizing state. In order to maintain this original state, cells use various redox couples.^{2,3} The most abundant cellular redox couple is glutathione (GSH) and its oxidized form, glutathione disulfide (GSSG). The endogenously synthesized tripeptide, GSH, is composed of glycine, cysteine and glutamine, and present at 1–11 mM concentrations in cells, far higher than other cellular redox active couples.^{2,4} Higher concentration of GSSG indicates oxidative shift at which biomolecules get oxidized and may be damaged if its original state is not re-established over time. Cells balance the redox state by producing more GSH. This is also true when cells are in more reducing state than its original balance; more GSSG is produced.

Enzymes catalyze the regeneration of GSH or GSSG: glutathione reductase, catalyzes the reduction of GSH from GSSG, and glutathione peroxidase catalyze the oxidation of GSSG from GSH. In cellular environment, NADPH is the major reducing agent and thus, provides necessary electrons for reduction of glutathione whereas NADP⁺ is the major oxidizing agent for regeneration of GSSG.⁴



Temporary perturbation of redox balance in cell, reductive shift or oxidative shift, is a necessary part of normal physiological processes.^{3,5} Regulated production of oxidants serves as a signal to produce more endogenous reductants and vice versa.^{3,6} Oxidants when present at high concentrations cause detrimental damages to important biomolecules by oxidizing them. This overproduction, also known as oxidative stress, is involved in over a hundred diseases.⁷ Yet, ironically, one of the main physiological roles of ROS is to act as a signal molecule to protect cells against oxidative stress.³ Therefore, the classical view of bad oxidants are destroyed by good antioxidants needs to be used with caution because redox active species can act as both oxidant and reductant (i.e. antioxidants) as will be illustrated in the following sections.

1.2. Endogenously produced redox active inorganic small molecules

1.2.1 Pathological and physiological roles of redox active oxygen species

Oxygen, being essential for our metabolism, is a radical itself and is the main producer of all reactive oxygen species.^{5,8} The dual nature of these metabolic species are apparent even in the case with oxygen. Oxygen when it is above its allowed concentration (>21%) becomes a poison to aerobic organisms.⁸

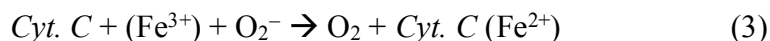
One electron reduction of oxygen produces superoxide (O_2^-). Superoxide itself in aqueous solution is not highly reactive and do not cause direct damage to cells. Most damage is usually caused by it reacting with other radicals (nitric oxide) or free metal ions, producing highly reactive hydroxyl radicals ($\text{HO}\bullet$). Hydroxyl radical is the most

reactive radical that can react readily with anything, change the structure and functions of any important biomolecules (DNA, protein, lipids).²



Exposure to reactive oxygen species (ROS) at low concentrations are normal and essential for our health as shown in **Table 1**. For example, cells use superoxide to kill environmental pathogens.³ When the level of superoxide exceeds the normal level, cells initiate the defense mechanism to reestablish the redox status.² The process signals is called “oxidative burst”, in which a large quantity of superoxide is produced. This superoxide production signals the defense system to produce more antioxidants.

Any oxidant given the appropriate conditions can also act as reductant. Standard reduction potential of O_2/O_2^- is -0.33 V ⁵ and thus, theoretically *in vitro*, it will reduce any redox couple whose potential is less negative, zero, or positive. Similarly, it can oxidize the redox couples whose potential is more negative. Superoxide in aqueous solution can act as a reducing agent⁵ when it reduces the iron complexes such as cytochrome c (Cyt. C) and ferri-EDTA.



Consequent reduction of O_2^- produces H_2O_2 that can exert intracellular damage. Being neutral in charge, H_2O_2 can diffuse into cells easily whereas negatively charged O_2^- cannot. Once inside the cell, hydrogen peroxide gets reduced to damaging hydroxyl radicals.⁵ All reactive oxygen species eventually form HO^\bullet and this chain reaction to form HO^\bullet is responsible for the detrimental effect of reactive oxygen species.





Similar to superoxide, when at physiological concentrations, H_2O_2 (0.001 to 0.7 μM) serves as cell signaling molecules for a number of important biochemical pathways as shown in **Table 1**.⁹ Hydrogen peroxide, when higher than 100 μM can be damaging to cells.

1.2.2 Pathological and physiological roles of redox active nitrogen species

Nitric oxide, another important endogenously produced radical, is known for its pro-oxidant role of generating other nitrogen-containing radicals as well as protecting oxidative damage of cell membranes¹⁰⁻¹² Nitric oxide (NO^\bullet), though can be toxic in excess, has many physiologically-important functions as shown in **Table 1**. Nitric oxide, though not very reactive, can produce a strong oxidant, peroxynitrite (ONOO^-), that can damage protein, DNA, lipids, and carbohydrate.¹² Nitric oxide inhibits lipid peroxidation as well as induces the peroxidation process. Lipid peroxidation is considered to be the major pathway leading toward free radical induced cellular damage. The facilitation or inhibition of the lipid peroxidation by NO^\bullet depends on the relative concentrations of nitric oxide and superoxide.¹¹

1.3 Enzymes as catalyst for redox active species

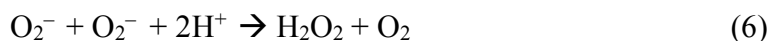
1.3.1 NADPH oxidase

Since radicals are part of our mechanism, they are produced by pro-oxidant enzymes in our body. Enzymes are designed to produce these ROS when needed to defend against environmental pathogens.⁷ This shows that pro-oxidant enzymes take the role of antioxidants. Oxidant enzymes such as NADPH oxidase and nitric oxide synthase,

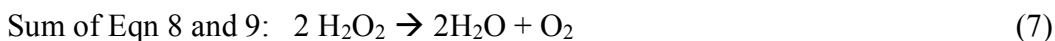
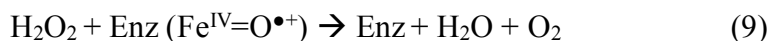
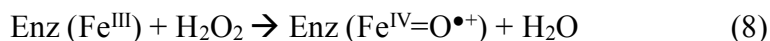
produces much needed superoxide and nitric oxide respectively, in a strictly controlled manner so that the concentration of ROS would not exceed the required amount.⁷ Mice with the knockout of NADH oxidase are found to be more susceptible to infection, implying the essential role of the oxidant enzyme in killing foreign organisms.²

1.3.2 Metalloenzymes

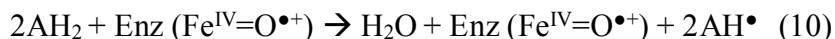
Superoxide dismutase (SOD) and catalase are antioxidant metalloenzymes that catalyze the disproportionation of superoxide (Eqn. 6) and hydrogen peroxide (Eqn. 7) respectively. SOD is the prime example of cellular antioxidants because SOD reduces radicals at the diffusion-controlled rate.²



However, these antioxidant enzymes can also exhibit pro-oxidant behavior.¹³ Catalase catalyzes the degradation of two molecules of hydrogen peroxides to water and oxygen. This reaction occurs by two steps in which H_2O_2 acts as oxidizing agent in the first step (8) and a reducing agent in the second step (9).^{14,15}



When the concentration of hydrogen peroxide is low, the enzyme deviates from the above two-step reactions (Eqn. 7, 8), and generate ROS in the presence of an electron donor.



1.4 Therapeutic candidates

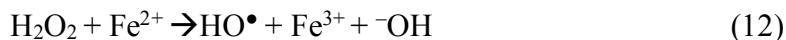
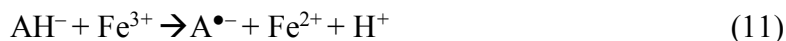
1.4.1 SOD Mimics

Superoxide dismutase can (SOD) scavenge superoxide radicals at the diffusion-controlled rate and thus, synthetic mimics of SOD that have comparable radical-scavenging activity are increasingly sought after.¹⁶ Even though the radical scavenging activity of most SOD mimics *in vitro* are as outstanding as the enzyme itself, some SOD mimics have shown pro-oxidant activity *in vivo*.¹⁷ This pro-oxidant behavior of SOD mimics is due to the formation of oxidizing intermediate similar to the dual action of catalase as discussed above.

1.4.2 Vitamin C

Chemically, powerful antioxidants are strong reducing agents and are marketed as “radical scavengers” in part due to their fast reaction rate to deplete oxidants. One such example is vitamin C or ascorbic acid. Vitamin C is the essential antioxidant in humans and deficiency of vitamin C causes scurvy.¹⁸ *In vitro*, at physiological pH, ascorbic acid (pK_{a1} 4.25) becomes ascorbate, its mono-anionic form, which is a very effective antioxidant that readily reduces all types of free radicals with the typical rate constants of $10^5 \text{ M}^{-1} \text{ s}^{-1}$ (pH 7.4).^{5,19} Despite the superiority of ascorbic acid as antioxidant *in vitro*, increasing number of human trials and some *in vivo* studies showed that vitamin C has no effect on oxidative stress.^{6,19,20} In fact, some *in vivo* studies suggest that administering ascorbic acid to smokers increases the cause of cancer: the evidence of vitamin C acting as a pro-oxidant.¹⁹⁻²¹

Radical scavengers can generate radicals. Ascorbate (AH^-) becomes a pro-oxidant in the presence of free metal ions, producing highly reactive hydroxyl radicals and hydrogen peroxide.^{18,20}



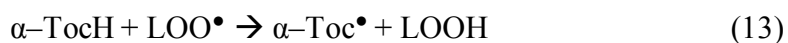
However, the possibility of the above reaction (i.e. Fenton reaction) occurring in physiological conditions has been questioned due to the lack of free metal ions *in vivo*. These metal ions are effectively sequestered by enzymes that are specifically designed to do so in extracellular fluids to prevent them catalyzing any free radical-producing reactions. Therefore, the concentration of free iron ions in blood plasma and body fluids is zero in normal healthy cells.^{5,22}

Even though vitamin C can act as a pro oxidant, it will more likely to act as an antioxidant rather than a prooxidant *in vivo*.^{6,19,20} Yet, *in vivo* studies have also shown that the deficiency of vitamin C, as well as the high dose (>500 mg per day) is associated with increased free radical induced damage to DNA, indicating how tightly the redox system is regulated. It is evident in the case of vitamin C that an antioxidant can also be a pro-oxidant and *vice versa*.

1.4.3 Vitamin E

Diet containing fruits and vegetables are known to have vitamins and minerals since Hippocrates (460–377 BC).²⁰ Since these diet-derived vitamins and minerals are essential for the normal growth of the body, their synthetic analogues are marketed as dietary supplements.²⁰ For example, Vitamin E is the major lipid-soluble antioxidant that

inhibits oxidation of lipoprotein, the process which leads to the hardening and blockage of blood vessels.²³ As a classical antioxidant, the biologically active form of Vitamin E, α -tocopherol, inhibits the lipid oxidation by scavenging lipid peroxy radicals (LOO^\bullet). The resulting radical can trap the second peroxy radical as in the reaction described below.



As in the case of catalase, when the concentration of LOO^\bullet is low, the resulting radical abstracts hydrogen from other lipids, deviating from its normal route.



In this case, vitamin E induces lipid oxidation, rather than inhibiting it. In addition, similar to ascorbate, tocopherol can reduce free metal ions (Fe^{3+} and Cu^{2+}) and exert pro-oxidant effect. However, the lack of free metal ions in human plasma and the very low amount in body fluids ($< \mu\text{M}$)⁵ disregards the possibility of this reaction happening *in vivo*. As expected, the pro-oxidant activity has been observed *in vitro* for other dietary supplements (vitamin A, vitamin C, and β -carotene).^{5,20}

CONCLUSIONS

Molecular oxygen that we breath in is the main source of reactive oxygen species and can be both beneficial when it is at a normal concentration (21%) and detrimental for us when it exceeds that level. Reactive species such as superoxide and nitric oxide are as essential to our health as therapeutic antioxidants when they are present at appropriate concentrations and when there are no other species (i.e. transition metal ions) that can perturb their normal route of reaction pathway. Superoxide and nitric oxide serve as cellular messengers in protecting cells against oxidative stress, participate in killing foreign organisms and fighting infections. Production of reactive species, is not only associated with the adverse health effects (cancer, neurodegenerative diseases) but this process is normal and beneficial when it generates the required reactive species (superoxide and nitric oxide) in a strictly-controlled manner. On the other hand, exogenously acquired therapeutics may act as antioxidants as well as pro-oxidants. Similarly, vitamins and metal-based enzyme mimics exhibit dual properties. The resulting biochemical process (i.e. either physiological or pathological) that these reactive species *in vivo* generate depends on the concentration of reactive species, and the nature (oxidizing vs. reducing) of intermediate involved.

OVERALL GOALS AND THESIS STRUCTURE

A well-functioned cell, thus a healthy life, inevitably requires both oxidants and reductants to maintain the dynamic cellular redox system. Cells use extremely complex, and interconnected meshwork of redox signaling system to control the biochemical processes involving redox active species. It is generally accepted that oxidants or

“reactive oxygen species (ROS)” are disease-causing and antioxidants or reductants are the answer to the diseased conditions. However, this oversimplified interpretation of the roles played by redox-active species, both ROS and antioxidant led to many pitfalls especially in the field of antioxidants.

Evidence that excessively high concentrations of ROS are found in diseased conditions is undeniable. However, scientific data has yet to confirm that ROS are the cause of diseases; ROS are implicated in pathogenesis but may or may not participate in pathological pathways. On the other hand, evidence that radical scavengers can also generate radicals have recently been discovered. Excessively increasing exogenous antioxidants will flood the cellular redox system with reductants and will lead to overexpression of antioxidative enzymes. The resulting reductive shift is as detrimental as oxidative shift caused by ROS. The complexity involved in performing thorough mechanistic studies of these redox-active and highly-reactive species in a dynamic system limit our understanding as evident in the clinical trials of Vitamin C. To this end, we study how an organoruthenium complex can act as a catalytic antioxidant in the radical reduction.

Chapter 2 reports a system in which an organoruthenium complex serves as a catalytic reductant, employing the chemically synthesized radical. The ruthenium complex catalyzes the radical degradation and inhibition under biologically-relevant condition via oxidation of biologically relevant alcohols. Amino acids, sugars and citric acid cycle metabolites are found to act as excellent terminal reductants that are essential for the radical reduction because replacing the O–H and C–H groups of alcohols

significantly slows down the reduction. **Chapter 3** describes the detailed mechanism of catalytic radical reduction by the N-heterocyclic carbene ruthenium complex in aqueous buffered solution. Here we describe the catalytic antioxidant mechanism that involves a reducing intermediate, rather than an oxidizing intermediate, in the catalytic reduction. The N-heterocyclic carbene ruthenium complex prevents the pro-oxidant effects that have been observed for current catalytic antioxidants. **Chapter 4** describes the use of nature's oxidizing agent, NAD⁺ (Nicotinamide Adenine Dinucleotide) as reductant, providing the evidence that biological oxidants have the ability to transfer hydride.

Table 1.1. Physiological and Pathological processes mediated by classical pro-oxidants

Type of reactive species	Source of Radical	Physiological process	Pathological process
<i>Reactive Oxygen Species (ROS)</i> 1. Superoxide (O_2^-) 2. H_2O_2 3. HO^\bullet	Oxygen NAD(P)H oxidase	Oxidative burst Superoxide: 10 pM ¹¹ H_2O_2 : 0.001-0.7 μM ¹¹	Diseases Superoxide: 0.01-0.1 μM ¹¹ H_2O_2 : 10-100 μM ¹¹
<i>Reactive Nitrogen Species (RNS)</i> 1. Nitric oxide (NO^\bullet) 2. $ONOO^-$	Nitric oxide synthase (NOS)	<i>Nervous system</i> Plays a role in long-term memory <i>Vascular system</i> Control blood pressure Signaling molecule in the cardiovascular system ¹⁰	<i>Nervous system</i> Stroke, epilepsy <i>Vascular system</i> Chronic inflammation, transplant rejection, septic shock
<i>Enzyme</i> 1. NAD(P)H oxidase 2. Nitric oxide synthase		Produce oxidants that act as cellular messengers	Excessive production of oxidants

Table 1.2. Physiological and Pathological processes mediated by classical reductants/antioxidants

Type of antioxidant	Physiological process	Pathological process
<i>Enzymes</i> Superoxide dismutase (SOD) Catalase Glutathione Peroxidase (GPx)	Catalyze in the reaction that depletes excess reactive species	Produce oxidants
<i>Dietary supplements</i> Vitamin C (Ascorbate) Vitamin E (α -tocopherol) <i>Inorganic/Organometallic</i> SOD mimics	Deplete reactive species	Produce oxidants

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CHAPTER TWO

CATALYTIC RADICAL REDUCTION IN AQUEOUS SOLUTION VIA OXIDATION OF BIOLOGICALLY-RELEVANT ALCOHOLS

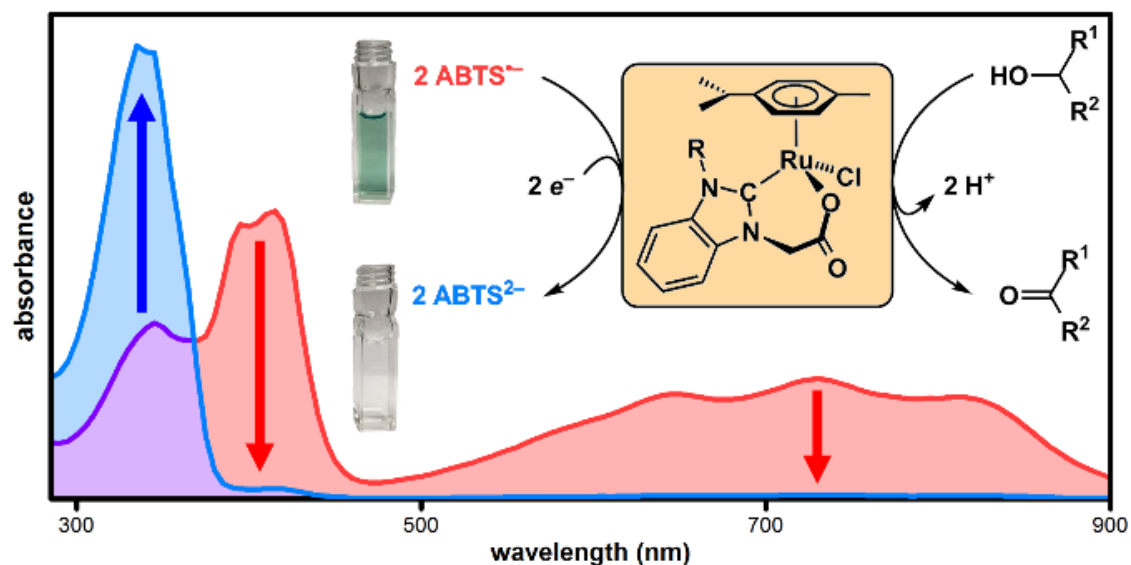


Figure 2. An organoruthenium complex catalyzed radical reduction in buffered aqueous solution using amino acids, sugars, or citric acid cycle metabolites as terminal reductants. These compounds are found abundant in our body and our cellular machinery exits to recycle them after being oxidized.

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ABSTRACT

Metalloenzymes that normally perform catalytic antioxidant or radical-degrading functions, as well as small-molecule complexes that mimic them, can also exert pro-oxidant or radical-forming effects depending on the identity of the terminal reductant. Because nitroxyl radicals function as redox active cocatalysts in the aerobic oxidation of alcohols, we hypothesized that catalytic radical reduction could be achieved via the oxidation of biologically-relevant alcohols. Herein we report an organoruthenium complex (**Ru1**) that catalyzed reduction of 2,2'-azino-bis(3-ethylbenzo-thiazoline-6-sulfonate) radical monoanion (ABTS^{•-}) to ABTS²⁻ in phosphate buffered saline (pH 7.4) using MeOH, EtOH, *i*-PrOH, serine, threonine, glucose, arabinose, methyl lactate or dimethyl malate as the terminal reductant. Replacing either the C–H and O–H groups of a –CHOH– moiety resulted in the loss of ABTS^{•-} reducing ability. Moreover, **Ru1** was able to inhibit the oxidation of ABTS²⁻ by H₂O₂ and horseradish peroxidase, even after multiple successive challenges with excess H₂O₂ or ABTS^{•-}. Collectively, these results demonstrate that **Ru1** inhibits the oxidative formation of and catalyzes the reduction of radicals in aqueous solution via oxidation of biologically-relevant alcohols.

INTRODUCTION

Reactive oxygen species (ROS) such as superoxide (O_2^-) and hydrogen peroxide (H_2O_2) can alter the structures and functions of biomolecules^{1,2} via DNA base lesion formation,³ transition metal ion release from proteins,⁴ and membrane phospholipid chain degradation.⁵ Antioxidants can prevent or attenuate this damage by reacting with ROS to yield less oxidizing, more stable radical or non-radical products. Superoxide dismutase (SOD) and catalase, for example, are metalloenzymes that protect cells by catalyzing the disproportionation of O_2^- and H_2O_2 , respectively.⁶ Synthetic Mn–salen and Mn–porphyrin complexes have demonstrated both SOD and catalase activity,⁷ and some have entered Phase 1 clinical trials as therapeutics for neurodegenerative and pulmonary diseases.⁸

Recent studies have shown that these classes of synthetic Mn complexes can also exert pro-oxidant effects at low ROS concentrations,^{9,10} dichotomous behavior that is consistent with the enzymes they mimic. Catalase converts $2 H_2O_2$ into $O_2 + 2 H_2O$ (Scheme 1.1, *i*),¹¹ wherein the first equivalent of H_2O_2 is reduced to H_2O and generates an $Fe(IV)=O^{\bullet+}$ intermediate, which then oxidizes the second equivalent of H_2O_2 to O_2 and releases H_2O .¹² At low H_2O_2 concentrations, this $Fe(IV)=O^{\bullet+}$ intermediate can abstract H^\bullet from other molecules (i.e., $A-H$) and thus exhibit peroxidase-like reactivity (Scheme 1.1, *ii*).¹³ Conversely, peroxidase can exhibit catalase-like reactivity at low $A-H$ / high H_2O_2 concentrations.¹⁴ Because the Mn–salen and Mn–porphyrin complexes proceed through high-valent Mn oxo/hydroxo intermediates¹⁵⁻¹⁸ with oxidizing power comparable to $Fe(IV)=O^{\bullet+}$, their reactivity will mimic catalase or peroxidase based on the availability

of H₂O₂. Therefore, the function of catalase and its biomimetic Mn complexes as either antioxidants or pro-oxidants will be determined by the nature of the terminal reductant.

A catalytic oxidation reaction important in synthetic chemistry is the aerobic oxidation of R¹-CHOH-R² to R¹-C(=O)-R² that does not cause any (1) over-oxidation to R-COOH or (2) undesired reactions at other functional groups.¹⁹⁻²³ One strategy employs a copper or ruthenium catalyst in conjunction with a nitroxyl radical, such as 2,2,6,6-tetramethylpiperidiny-*N*-oxyl (TEMPO), to oxidize R¹-CHOH-R² to R¹-C(=O)-R² with concomitant formation of TEMPO-H (Scheme 1.1, *iii*). Subsequent regeneration of the TEMPO radical occurs with the oxidation of TEMPO-H by O₂, which functions as the terminal oxidant for this reaction (Scheme 1.1, *iv*). Because the TEMPO radical is reduced by R¹-CHOH-R² in the aerobic alcohol oxidation catalytic cycle, we hypothesized that the catalytic reduction of other radicals could also be achieved via the oxidation (i.e., dehydrogenation) of non-tertiary alcohols (Scheme 1.1, *v*). Herein we report the validation of this hypothesis with an organoruthenium complex that catalyzes the reduction and inhibits the formation of radicals in buffered aqueous solutions via the oxidation of biologically-relevant non-tertiary alcohols. Given the successes of Group 8 transfer hydrogenation catalysts as chemotherapeutics^{24,25} in particular and the burgeoning scope of medicinal applications for transition metal-based catalytic systems²⁶⁻²⁸ in general, we anticipate that the reduction of radicals via the Ru-catalyzed oxidation of alcohols will lead to new strategies for protecting against ROS *in vivo*.

RESULTS AND DISCUSSION

Approach and rationale

We recently reported a Ru complex supported by a chelating N-heterocyclic carbene–carboxylate ligand (**Ru1**, Scheme 1.2A) that catalyzed the hydrogenation of C=O, C=N and C=C bonds, in which the H₂ transferred was obtained from the oxidation of *i*-PrOH to acetone.³² Based on this catalytic chemical reactivity observed with **Ru1** and previous reports of other Ru complexes with chelating carboxylate ligands exerting antioxidant effects in cells, which derived from irreversible stoichiometric reactions with nitric oxide,^{33,34} we hypothesized that **Ru1** would function as a catalytic antioxidant, reducing radicals using non-tertiary alcohols as terminal reductants. Using methods previously reported by our group,^{32,35} the N-heterocyclic carbene (NHC) ligand precursor [1H₂][Br] was treated with Ag₂O to afford the Ag–NHC complex [Ag(**1**)]_n (**2**). The NHC ligand was subsequently transferred to Ru via the reaction of **2** with [{RuCl(η⁶-cymene)}₂(μ-Cl)₂], which yielded the Ru–NHC complex [RuCl(**1**)(η⁶-cymene)] (**Ru1**).

As a radical substrate to probe antioxidant activity, 2,2'-azino-bis(3-ethylbenzothiazoline-6-sulfonate) radical monoanion (ABTS^{•-}, Scheme 1.2B) was selected on the basis of its large extinction coefficient (ε) at long wavelengths (ε₇₅₀ = 1.6 × 10⁴ M⁻¹ cm⁻¹ in EtOH and ε₇₃₄ = 1.5 × 10⁴ M⁻¹ cm⁻¹ in aqueous buffer),³⁰ its reversible one-electron reduction to ABTS²⁻ (*E*_{1/2} = 0.68 V vs. NHE) falling within the range of potentials accessible in biological systems (−0.4 to +0.8 V vs. NHE),³⁶ and its stability in aerobic, protic media. The longer absorption wavelength and greater extinction coefficient for ABTS^{•-} enables spectroscopic analysis of radical-degrading and -forming

reactions (1) with less interference from other optically absorbing species present in biological systems and (2) at lower concentrations[‡] more relevant to oxidative stress than would be possible with direct measurements of $\text{O}_2^{\bullet-}$ ($\epsilon_{250} = 1.9 \times 10^3 \text{ M}^{-1} \text{ cm}^{-1}$) or H_2O_2 ($\epsilon_{240} = 43.6 \text{ M}^{-1} \text{ cm}^{-1}$).^{37,38} Furthermore, the $1e^-$ reduction potential for $\text{ABTS}^{\bullet-}$ is lower than or comparable to the standard reduction potentials for several of the oxidizing species associated with oxidative stress in biological systems (e.g., $E^\circ = 1.78 \text{ V}$ for H_2O_2 , 1.6 V for RO^\bullet , 1.0 V for ROO^\bullet , 0.92 V for Cys-S^\bullet , 0.70 for O_2 , etc.),³⁶ therefore $\text{ABTS}^{\bullet-}$ is a reasonable substrate for approximating biologically-relevant oxidants.

As a non-catalytic antioxidant control, Trolox (Scheme 1.2C) was employed given its use as a benchmark in a variety of radical degradation and antioxidant studies.³⁹ Trolox can serve as a $1e^-$ or $2e^-$ reductant, whereby the first $1e^-$ oxidation is accompanied by rapid H^+ loss to form a phenoxyl radical (TrO^\bullet), which can then undergo a second $1e^-$ oxidation to form a phenoxonium cation (TrO^+). However, in methanol (MeOH) or ethanol (EtOH) solutions, these processes converge into a single $2e^-$ oxidation.⁴⁰ Subsequent hydrolysis of TrO^+ can then cleave the tertiary carbon–oxonium bond, which renders the $2e^-$ oxidation of Trolox irreversible.

Because ROS and other oxidants can cause damage to critical biomolecules, inhibiting the formation of these oxidizing species in biological systems can potentially prevent the onset of harmful pathologies.² As a model system, we selected the oxidation of ABTS^{2-} to $\text{ABTS}^{\bullet-}$ by horseradish peroxidase (HRP) and H_2O_2 (i.e., Scheme 1.1, *ii*).³¹ However, before studying the ability of **Ru1** to inhibit $\text{ABTS}^{\bullet-}$ formation, it was first necessary to explore the fundamental reactivity of **Ru1** with $\text{ABTS}^{\bullet-}$ under controlled

conditions and achieve the catalytic reduction of $\text{ABTS}^{\bullet-}$ to ABTS^{2-} in buffered aqueous solution.

Catalytic radical reduction in EtOH

Addition of 5 μM **Ru1** (CH_3CN stock) to 50 μM $\text{ABTS}^{\bullet-}$ in EtOH caused a 100% decay in $\text{ABTS}^{\bullet-}$ concentration within 15 min (Fig. 2.1, red line), accompanied by the formation of 50 μM ABTS^{2-} (Fig. 2.2), indicating the 1:1 conversion of $\text{ABTS}^{\bullet-}$ to ABTS^{2-} by **Ru1**. No $\text{ABTS}^{\bullet-}$ degradation occurred when CH_3CN alone was added (Fig. 2.1, dotted green line), indicating that **Ru1** was essential for the reduction of $\text{ABTS}^{\bullet-}$ to ABTS^{2-} and that CH_3CN did not impact $\text{ABTS}^{\bullet-}$ stability. No $\text{ABTS}^{\bullet-}$ formation was observed in an aerobic solution containing 5 μM **Ru1** and 50 μM ABTS^{2-} , indicating that **Ru1** does not oxidize ABTS^{2-} to $\text{ABTS}^{\bullet-}$ under these conditions. In contrast to **Ru1**, the addition of 5 μM Trolox caused a rapid (within mixing time) 22% decrease in $\text{ABTS}^{\bullet-}$ concentration (Fig. 2.1, blue line), which corresponded to the reduction of 10 μM $\text{ABTS}^{\bullet-}$ (2 equiv. vs Trolox) and was consistent with the ability of Trolox to serve as a non-catalytic $2e^-$ reductant in EtOH solution.⁴⁰

To assess its catalytic potential and corresponding regeneration, the reactivity of **Ru1** with multiple sequential aliquots of excess $\text{ABTS}^{\bullet-}$ was examined. After the reduction of 50 μM $\text{ABTS}^{\bullet-}$ by 5 μM **Ru1** was complete (Fig. 2.3, red line), 2 additional 50 μM $\text{ABTS}^{\bullet-}$ aliquots were introduced[§] and complete $\text{ABTS}^{\bullet-}$ reduction was observed in each instance, indicating that **Ru1** remained catalytically competent for the complete reduction of 150 μM $\text{ABTS}^{\bullet-}$ (30 turnovers). Unlike **Ru1**, the addition of successive 10 μM aliquots[§] of $\text{ABTS}^{\bullet-}$ following the initial reduction by 5 μM Trolox (Fig. 2.3, blue

line) only caused the radical absorbance to increase by amounts equivalent to the concentration of ABTS^{•-} added, indicating that the antioxidant capacity of Trolox had been exhausted after the reduction of only 2 equiv. of ABTS^{•-}. However, subsequent addition of 5 μM **Ru1** to the exhausted Trolox solution containing 70 μM ABTS^{•-} resulted in 100% radical reduction within 30 min.

Addition of 5 μM ascorbate (AscH⁻) or 5 μM glutathione (GSH) to a solution of 50 μM ABTS^{•-} in EtOH caused the ABTS^{•-} concentration to decrease by 5% or 60% (Figs. 2.4–2.5), values that were consistent with their ability to function as 1e⁻ or 6e⁻ reductants, respectively, under these experimental conditions.⁴¹⁻⁴³ Similar to the experiments with Trolox, subsequent addition of ABTS^{•-} only afforded increases in radical absorbance equal to the concentration of ABTS^{•-} added, which likewise indicated that the antioxidant capacity of AscH⁻ and GSH had been exhausted. Quantitative radical reduction could nonetheless be achieved via addition of 5 μM **Ru1** to the exhausted AscH⁻ or GSH solutions that contained excess ABTS^{•-}. Collectively, these data demonstrate that **Ru1** can catalytically reduce ABTS^{•-} and, because it is regenerated after each reaction cycle, that a catalytic antioxidant can offer significantly greater protection than traditional stoichiometric antioxidants, such as Trolox, AscH⁻ and GSH.

Although ABTS^{•-} degradation has been previously observed with other Ru complexes,⁴⁴⁻⁴⁶ **Ru1** is the first to demonstrate catalytic ABTS^{•-} reduction. It is important to note that the previous studies measured percentage of ABTS^{•-} degradation with respect to Ru complex concentration, for the purpose of determining dose dependence. However, the reported absorbance vs. time plots displayed significantly slower ABTS^{•-}

degradation for the Ru complexes compared to Trolox (much like **Ru1** vs. Trolox, Fig. 2.1) and 100% ABTS^{•-} degradation at multiple Ru concentrations (demonstrating dose *independence*). Thus, it is possible that a previously reported Ru complex may have degraded ABTS^{•-} catalytically, but its significantly slower reactivity compared to Trolox created the appearance that the percent of ABTS^{•-} degraded was dependent on Ru complex concentration and led to the conclusion that the observed radical degradation was non-catalytic.

Catalytic radical reduction in aqueous buffer

As a catalyst for the $1e^-$ reduction of ABTS^{•-} to ABTS²⁻, **Ru1** itself cannot serve as the terminal reductant for this reaction, a role most likely played by the EtOH solvent for the experiments displayed in Fig. 2.1. To test this hypothesis and identify the chemical features required for terminal reductant function, it was first necessary to determine experimental conditions under which no ABTS^{•-} degradation occurred in the presence of **Ru1** alone. Phosphate-buffered saline (PBS, pH 7.4) was selected as a suitable reaction medium because neither the solvent (H₂O) nor the buffer components (Na₂HPO₄, KH₂PO₄, NaCl, KCl) would undergo oxidation to supply the electrons necessary for the reduction of ABTS^{•-} (i.e., Fig. 2.6, *i*). No degradation of ABTS^{•-} in PBS was observed after treatment with 5 μ M **Ru1** (Fig. 2.6, *ii*), indicating this solution lacked a suitable terminal reductant. Subsequent addition of 50 mM EtOH to this PBS solution containing 5 μ M **Ru1** and 50 μ M ABTS^{•-} caused the ABTS^{•-} concentration to decrease (Fig. 2.6, *iii*), evidence that the electrons needed for the ABTS^{•-} reduction observed in Fig. 1 were ultimately supplied by the EtOH solvent.

If the hypothesis that the oxidation of $R_1\text{-CHOH-R}_2$ to $R_1\text{-C(=O)-R}_2$ provides the electrons necessary for the reduction of $\text{ABTS}^{\bullet-}$ to ABTS^{2-} is correct (i.e., Scheme 2.1, v), then both the O–H and C–H groups in a –CHOH– moiety will be essential for terminal reductant function. To test this hypothesis, PBS solutions containing 5 μM **Ru1** and 50 μM $\text{ABTS}^{\bullet-}$ were treated with 50 mM MeOH, *i*-PrOH, *t*-BuOH, ethylene glycol (EG) or 1,2-dimethoxyethane (DME) and the $\text{ABTS}^{\bullet-}$ concentration was measured. With MeOH, *i*-PrOH and EG, the $\text{ABTS}^{\bullet-}$ concentration began to decrease immediately following addition of the alcohol, indicating that these alcohols could serve as terminal reductants (Figs. 2.7–2.9). In contrast, no $\text{ABTS}^{\bullet-}$ reduction was observed with DME or *t*-BuOH (e.g., Fig. 2.10, *ii*). However, when the addition of 50 mM *t*-BuOH was followed by 50 mM EtOH, $\text{ABTS}^{\bullet-}$ reduction did occur (Fig. 2.10, *iii*), revealing that the lack of reactivity with *t*-BuOH was not due to catalyst deactivation but rather the inability of *t*-BuOH to serve as a terminal reductant. Collectively, these results show that the C–H and O–H groups of a –CHOH– moiety are both essential for terminal reductant function, which is consistent with the hypothesis that oxidation of $R_1\text{-CHOH-R}_2$ to $R_1\text{-C(=O)-R}_2$ provides the reducing equivalents necessary to convert $\text{ABTS}^{\bullet-}$ to ABTS^{2-} . Replacing either the C–H group with C–Me or the O–H group with O–Me precludes this oxidation and thus the ability to function as a terminal reductant, which is supported by the lack of $\text{ABTS}^{\bullet-}$ reduction observed with *t*-BuOH and DME.

Biologically–relevant terminal reductants

A wide variety of biomolecules comprise –CHOH– groups, therefore we hypothesized that sugars, amino acids and citric acid cycle metabolites could serve as

suitable terminal reductants in **Ru1**-catalyzed ABTS^{•-} reduction. To test this hypothesis, PBS solutions containing 5 μM **Ru1** and 50 μM ABTS^{•-} were treated with 50 mM serine, threonine, glucose, arabinose, malic acid or lactic acid. Reduction of ABTS^{•-} was markedly slower following the addition of the amino acids compared to the sugars, consistent with the greater number of CH–OH groups per molecule in the latter (Fig. 2.11). No significant differences in reactivity were observed between serine and threonine or between glucose and arabinose.

Malic acid and lactic acid are metabolites of the citric acid cycle and each comprises a –CHOH– group. However, no ABTS^{•-} reduction occurred following the addition of 50 mM malic acid or lactic acid to 5 μM **Ru1** and 50 μM ABTS^{•-} in PBS (Fig. 2.12). Malate and lactate will be the predominant species in pH 7.4 PBS, and these carboxylate anions can function as chelating bidentate ligands, which could potentially disrupt the catalytic activity of **Ru1**. To prevent catalyst deactivation, the methyl esters of these substrates were used. Addition of 50 mM methyl lactate (Me-lactate) or dimethyl malate (Me₂-malate) to PBS solutions containing 5 μM **Ru1** and 50 μM ABTS^{•-} caused the ABTS^{•-} concentration to decrease rapidly. None of the biologically-relevant alcohols afforded ABTS^{•-} reduction in the absence of **Ru1**, and no oxidation of ABTS²⁻ to ABTS^{•-} by **Ru1** was observed under any experimental conditions in PBS.

Inhibition of oxidative radical formation

Having obtained unambiguous evidence that **Ru1** catalyzes ABTS^{•-} reduction using biologically-relevant alcohols as terminal reductants, we next sought to demonstrate that **Ru1** could also inhibit radical formation. As the model system, we

utilized the *in situ* oxidation of ABTS^{2-} to $\text{ABTS}^{\bullet-}$ by horseradish peroxidase (HRP) and H_2O_2 (i.e., Scheme 2.1, *ii*).³¹ Addition of 10 μM H_2O_2 to a PBS solution of 10 nM HRP, 20 μM ABTS^{2-} , 5 μM **Ru1**, and 50 mM EtOH caused the $\text{ABTS}^{\bullet-}$ concentration to increase immediately. The $\text{ABTS}^{\bullet-}$ concentration peaked at 1.0 μM after 2.3 min, which then gradually declined and complete $\text{ABTS}^{\bullet-}$ reduction was observed 13 min after the peak (Fig. 2.13A, red line). No $\text{ABTS}^{\bullet-}$ formation occurred when 10 μM H_2O_2 was added to 5 μM **Ru1** and 20 μM ABTS^{2-} in PBS, indicating that **Ru1** did not cause any pro-oxidant reactions. Because **Ru1** and Trolox were added as 30 μL aliquots from 5.0 mM stock solutions in CH_3CN , 10 μM H_2O_2 was added to a PBS solution containing 10 nM HRP, 20 μM ABTS^{2-} and 0.19 M CH_3CN to control for the influence of this solvent. The radical absorbance immediately began to increase and reached a maximum of 17.9 μM after 50 min (Fig. 2.13A, dotted green line and Fig. 2.14). In contrast, the addition of 10 μM H_2O_2 to a PBS solution of 5 μM Trolox, 10 nM HRP and 20 μM ABTS^{2-} resulted in no change in $\text{ABTS}^{\bullet-}$ concentration for the first 7.6 min, demonstrating that Trolox completely halted $\text{ABTS}^{\bullet-}$ formation (Fig. 2.13A, blue line). After 7.6 min, however, the $\text{ABTS}^{\bullet-}$ concentration began to increase and reached a maximum of 9.3 μM at 12 min after the onset of $\text{ABTS}^{\bullet-}$ formation. Notably, the maximum $[\text{ABTS}^{\bullet-}]$ observed in the presence of Trolox was 8.6 μM lower than the control, which was consistent with the ability of Trolox to function as a $2e^-$ reductant (cf., Fig. 2 or Scheme 2.2). For the control and Trolox experiments, the decline in $\text{ABTS}^{\bullet-}$ concentration after peaking was consistent with normal $\text{ABTS}^{\bullet-}$ thermal decay.

To determine if **Ru1** remained catalytically competent after the complete reduction of $\text{ABTS}^{\bullet-}$ formed by 10 nM HRP and 10 μM H_2O_2 , two additional 10 μM aliquots of H_2O_2 were introduced (Fig. 2.13B, red line). Impressively, the concentration of $\text{ABTS}^{\bullet-}$ never exceeded 0.4 μM and decreased back to zero in less than 16 min after the introduction of the second and third H_2O_2 aliquots. In contrast, addition of the second 10 μM H_2O_2 aliquot to the Trolox experiment caused the $\text{ABTS}^{\bullet-}$ concentration to gradually increase by 7.9 μM over the course of 38 min (Fig. 2.13B, blue line), effectively resulting in complete oxidation of ABTS^{2-} to $\text{ABTS}^{\bullet-}$ (96% of the 17.9 μM observed for control). It was therefore unsurprising that the third 10 μM H_2O_2 aliquot produced no change in radical absorbance. Subsequent addition of 50 mM EtOH and 5 μM **Ru1** caused the $\text{ABTS}^{\bullet-}$ concentration to decrease rapidly, affording quantitative $\text{ABTS}^{\bullet-}$ reduction in less than 30 min.

To confirm that the results obtained with **Ru1** arose from its ability to catalyze $\text{ABTS}^{\bullet-}$ reduction, rather than inhibiting HRP or another chemical reaction, two 10 μM aliquots of chemically synthesized $\text{ABTS}^{\bullet-}$ were introduced after the $\text{ABTS}^{\bullet-}$ concentration had decreased to zero following the initial addition of 10 μM H_2O_2 (Fig. 2.13C, red line). The first and second $\text{ABTS}^{\bullet-}$ aliquots produced immediate increases in $\text{ABTS}^{\bullet-}$ concentration corresponding to 6.3 and 6.7 μM , respectively, followed by rapid decreases in radical absorbance that resulted in quantitative radical reduction within 75 s of aliquot addition. With the Trolox experiment, however, the first and second $\text{ABTS}^{\bullet-}$ aliquots caused $[\text{ABTS}^{\bullet-}]$ to increase by 9.9 and 9.7 μM , respectively, with no observable changes in radical absorbance other than thermal decay (Fig. 2.13C, blue line). Addition

of 50 mM EtOH and 5 μ M **Ru1** after the second ABTS \bullet^- aliquot afforded complete ABTS \bullet^- reduction within 30 min, similar to the experiment with multiple H₂O₂ aliquots. Both AscH $^-$ and GSH inhibited the HRP-induced oxidation of ABTS $^{2-}$, albeit for shorter periods of time and less effectively than Trolox. The introduction of additional H₂O₂ (Figs. 2.14–2.15) or ABTS \bullet^- aliquots (Figs. 2.16–2.17) only afforded increases in ABTS \bullet^- concentration that mirrored the behaviors observed for Trolox (i.e., Figs. 2.13B–C). Subsequent addition of 50 mM EtOH and 5 μ M **Ru1** to the AscH $^-$ and GSH solutions containing excess ABTS \bullet^- produced quantitative radical reduction, consistent with the Trolox experiments.

Synthetic Procedures

General synthetic considerations. N-(*p*-tolyl)benzimidazole and ABTS \bullet^- were prepared as previously described.^{29,30} All other materials were of reagent quality and used as received. All solvents used were HPLC grade. ^1H and $^{13}\text{C}\{^1\text{H}\}$ NMR spectra were recorded using a Bruker 500 MHz spectrometer. Chemical shifts δ (in ppm) for ^1H and ^{13}C NMR are referenced to SiMe₄ using the residual protio-solvent as an internal standard. For ^1H NMR: CDCl₃, 7.26 ppm; DMSO-*d*₆, 2.50 ppm. For ^{13}C NMR: CDCl₃, 77.16 ppm; DMSO-*d*₆, 39.52 ppm. Coupling constants (*J*) are expressed in hertz (Hz). Infrared spectra were recorded with 4 cm⁻¹ resolution on a Shimadzu IRAffinity-1S spectrometer equipped with a Pike Technologies MIRacle ATR (diamond) sampling accessory. Elemental analyses were performed at Atlantic Microlab, Inc. (Norcross, GA). All reactions were performed under an inert atmosphere under an N₂ atmosphere using standard Schlenk or glovebox techniques with the exclusion of light. All subsequent

manipulations were performed under ambient conditions using standard benchtop techniques without the exclusion of light. When required, solvents were dried and deoxygenated using an Innovative Technologies solvent purification system, and then stored over molecular sieves (3 Å) in a drybox.

Synthesis of 2-(3-*p*-tolyl)-benzimidazol-1-ium-1-yl)-acetate hydrobromide [1H₂][Br]. Bromoacetic acid (503 mg, 3.62 mmol) and N-(*p*-tolyl)benzimidazole (500 mg, 2.41 mmol) were dissolved with 20 mL of toluene in a vial equipped with a stir bar. The vial was then sealed with a Teflon-lined cap and the clear, dark orange-brown solution was heated to 120 °C. After 16 h, a white precipitate had formed, at which point the reaction mixture was allowed to cool to room temperature and the precipitate was collected by filtration. The resulting solid was washed with minimum toluene followed by CH₂Cl₂ and was then dried *in vacuo* to afford 514 mg (1.48 mmol, 80% yield) of the desired product as a white powder. ¹H NMR (500 MHz, DMSO-*d*₆): δ = 13.95 (s, 1H), 10.18 (s, 1H), 8.18 (d, *J* = 8.0, 1H), 7.85 (d, *J* = 8.0, 1H), 7.78–7.73 (m, 4H), 7.57 (d, *J* = 8.0, 2H), 5.58 (s, 2H), 2.47 (s, 3H). ¹³C NMR (125 MHz, DMSO-*d*₆): δ 167.5, 143.3, 140.5, 131.5, 130.8, 130.6, 130.4, 127.4, 127.1, 124.8, 114.3, 113.4, 47.8, 20.7. FT-IR (KBr): 3421 (vw), 2953 (m), 2372 (m), 1718 (m), 1560, 1508 (m), 1381 (m), 1300 (m), 1225 (m), 822, 754, 742, 482 (vw) cm⁻¹. Anal. Calcd. for C₁₆H₁₆BrN₂O_{2.5} ([1H₂][Br]•0.5H₂O): C, 53.95; H, 4.53; N, 7.86. Found: C, 54.01; H, 4.35; N, 7.79.

Synthesis of [Ag(1)]_n (2). To a solution of [1H₂][Br] (500 mg, 144 μmol) in anhydrous CH₂Cl₂ (50 mL) under nitrogen was added Ag₂O (502 mg, 217 μmol), and the resulting white suspension was allowed to stir at room temperature in the absence of

light. After 24 h, the reaction mixture was filtered and the resulting grey solid was washed with CHCl_3 (10 mL). The filtrate solvent was removed under reduced pressure and the resulting solids were then dried *in vacuo* to afford 452 mg (121 μmol , 84% yield) of the desired product as an off-white powder. ^1H NMR (500 MHz, $\text{DMSO-}d_6$): δ 7.71 (d, $J = 8.5$, 1H), 7.58 (d, $J = 8.0$, 2H), 7.44–7.38 (m, 5H), 5.09 (s, 2H), 2.39 (s, 3H). ^{13}C NMR (125 MHz, $\text{DMSO-}d_6$): δ 138.6, 135.2, 134.1, 133.4, 130.3, 125.6, 124.1, 112.7, 111.6, 99.5, 79.1, 52.1, 20.6. FT-IR (KBr): 3416 (w), 1608, 1516, 1475, 1439, 1385, 1308, 1250, 1207 (m), 1109 (m), 1094 (m), 1018 (m), 835, 814, 746, 715, 482 (w) cm^{-1} . Anal. Calcd. for $\text{C}_{16}\text{H}_{14}\text{N}_2\text{O}_{2.5}\text{Ag}$ ($2 \cdot 0.5\text{H}_2\text{O}$): C, 50.36; H, 3.77; N, 7.49. Found: C, 50.23; H, 4.09; N, 7.05.

Synthesis of $[\text{RuCl}(\mathbf{1})(\eta^6\text{-cymene})]$ (Ru1**).** To a stirred solution of **2** (100 mg, 134 μmol) in anhydrous CH_2Cl_2 (10 mL) under nitrogen was added $[\text{RuCl}(\eta^6\text{-cymene})(\mu\text{-Cl})_2]$ (82 mg, 134 μmol) dissolved in 10 mL of CH_2Cl_2 . During the course of the addition, the mixture changed first to a clear red and then to a clear orange solution with concomitant formation of a white precipitate, and the reaction was allowed to stir at room temperature. After 24 h, the reaction mixture was filtered through a 0.2 μm PTFE filter (to remove AgCl) with the aid of CH_2Cl_2 (4 mL). The filtrate solvent was removed under reduced pressure and the resulting solids were then dried *in vacuo* to afford 71.7 mg (134 μmol , 98%) of the desired product as a light red-orange powder. ^1H NMR (500 MHz, CDCl_3): δ 7.95 (d, $J = 8.0$, 1H), 7.49–7.44 (m, 2H), 7.41 (d, $J = 10.0$, 1H), 7.37–7.32 (m, 2H), 7.25–7.23 (m, 1H), 7.04 (d, $J = 8.5$, 1H), 5.43 (d, $J = 6.0$, 1H), 5.18 (d, $J = 6.0$, 1H), 5.16 (d, $J = 6.5$, 1H), 5.04 (d, $J = 5.5$, 1H), 4.91 (d, $J = 17.0$, 1H), 4.71 (d, $J = 18.5$, 1H),

3.95 (d, $J = 5.5$, 1H), 2.60 (sep, 1H), 2.52 (s, 3H), 2.03 (s, 3H), 1.08 (d, $J = 7.5$, 3H). ^{13}C NMR (125 MHz, CDCl_3): δ 186.6, 169.2, 140.3, 136.8, 136.0, 134.0, 131.1, 131.0, 129.7, 127.4, 124.2, 124.1, 111.3, 110.2, 104.7, 103.2, 90.0, 89.4, 83.6, 80.3, 49.9, 30.9, 23.3, 22.0, 21.5, 19.0. FT-IR (KBr): 3429 (vw), 3055 (w), 2959 (m), 2872 (w), 2230 (w), 2100 (w), 1628 (vs), 1514, 1475, 1443, 1373 (w), 1356, 1300, 1244, 1207 (m), 1190 (m), 1014, 930, 862, 843 (m), 827 (m), 804, 748, 663 (m), 615, 555, 476 (m), 470, 424, 405 cm^{-1} . Anal. Calcd. for $\text{C}_{26}\text{H}_{23}\text{ClN}_2\text{O}_2\text{Ru}$: C, 58.26; H, 5.08; N, 5.23. Found: C, 57.98; H, 4.95; N, 5.17.

Spectroscopic and Kinetic Analysis Procedures

General spectroscopic considerations. UV–visible absorption spectra were acquired on a Varian Cary 50 Bio spectrometer equipped with a water-cooled Quantum Northwest TC-125 peltier temperature controller. All solution measurements were performed at 25.0 ± 0.1 °C in matched gas-tight quartz cuvettes (Precision Scientific) with 1 cm path lengths and 3.0 mL analyte solution volumes. Absorption spectra were acquired from 900 to 200 nm with a scanning speed of 300 nm min^{-1} and a resolution of 0.5 nm. Each kinetic analysis experiment (5 second intervals) was performed in quadruplicate on four different days. Stock solutions were prepared fresh daily and filtered (0.2 μm PTFE) immediately prior to use.

Kinetic analysis of $\text{ABTS}^{\bullet-}$ degradation in EtOH. An aliquot of $\text{ABTS}^{\bullet-}$ (40 μL from a 3.75 mM stock solution in H_2O) was added to EtOH to afford the $\text{ABTS}^{\bullet-}$ working solution in EtOH. The absorbance spectrum of this working solution was acquired to confirm $\text{ABTS}^{\bullet-}$ concentration and the single wavelength kinetics program

was initiated. After 20 s, the cuvette was removed from the spectrometer without stopping the kinetics program, an aliquot of **Ru1**, Trolox, ascorbate or glutathione was added, the cuvette was covered and mixed via repeated inversion for 3 s, placed back in the spectrometer, and the kinetics program was allowed to continue. After the kinetics program had completed, the absorbance spectrum was acquired to confirm formation of ABTS^{2-} from the peak at 340 nm.

Representative procedure for kinetic analysis of $\text{ABTS}^{\bullet-}$ degradation by Ru1 in PBS. An aliquot of $\text{ABTS}^{\bullet-}$ (40 μL from a 3.75 mM stock solution in H_2O) was added to PBS to afford the $\text{ABTS}^{\bullet-}$ working solution in PBS. The absorbance spectrum of this working solution was acquired to confirm $\text{ABTS}^{\bullet-}$ concentration and the single wavelength kinetics program was initiated. After 20 s, the cuvette was removed from the spectrometer without stopping the kinetics program, an aliquot of **Ru1** (30 μL from 0.5 mM stock solution in CH_3CN) was added, the cuvette was covered and mixed via repeated inversion for 3 s, placed back in the spectrometer, and the kinetics program was allowed to continue. After 5 min, an aliquot of EtOH (30 μL from 5.0 M stock solution in H_2O) was added, the cuvette was covered and mixed via repeated inversion for 3 s, placed back in the spectrometer, and the kinetics program was allowed to continue. After the kinetics program had completed, the absorbance spectrum was acquired to confirm formation of ABTS^{2-} from the peak at 340 nm.

Kinetic analysis of $\text{ABTS}^{\bullet-}$ formation by HRP in PBS. Derived from a literature procedure.³¹ Aliquots of ABTS^{2-} (30 μL from a 2.0 mM stock solution in H_2O), HRP (30 μL from a 1.0 μM stock solution in H_2O), and **Ru1**, Trolox, ascorbate or

glutathione were added to PBS (pH 7.4) to afford the ABTS²⁻/**Ru1** (or Trolox or ascorbate or glutathione)/HRP working solution in PBS. The absorbance spectrum of this working solution was acquired to confirm no ABTS^{•-} - had formed. An aliquot of H₂O₂ (30 μL from 1.0 mM stock solution in H₂O) was added, the cuvette was covered and mixed via repeated inversion for 3 s, placed back in the spectrometer, and the single wavelength kinetics program was initiated. For the control enzyme experiments, the aliquot of H₂O₂ was added directly after HRP, and the analysis was performed with no other additives.

CONCLUSIONS

An organoruthenium complex (**Ru1**) that was previously reported to catalyze the hydrogenation of unsaturated organic substrates, using the oxidation of *i*-PrOH to acetone to supply the needed H₂, can also catalyze the reduction of ABTS^{•-} to ABTS²⁻ in buffered aqueous solution. By itself, **Ru1** was unreactive towards ABTS^{•-} and the presence of a non-tertiary alcohol was essential for ABTS^{•-} reduction to occur. Replacing either the C–H or O–H group in a –CHOH– moiety with C–Me or O–Me, respectively, resulted in the loss of ABTS^{•-} reducing ability. Given the previously reported transfer hydrogenation activity of **Ru1** and the fact that **Ru1**-catalyzed ABTS^{•-} reduction required a reactant comprising a –CHOH– moiety, we deduced that the reducing equivalents necessary for the 1e⁻ reduction of ABTS^{•-} to ABTS²⁻ were supplied by the oxidation of R₁–CHOH–R₂ to R₁–C(=O)–R₂. Consistent with this deduction, a diverse array of biologically-relevant non-tertiary alcohols, including amino acids, sugars and citric acid cycle metabolites, were found to be suitable terminal reductants for the reduction of ABTS^{•-} catalyzed by **Ru1**. Furthermore, **Ru1** was able to inhibit the oxidation of ABTS²⁻ to ABTS^{•-} by H₂O₂ and horseradish peroxidase with the help of a non-tertiary alcohol. In contrast, non-catalytic antioxidants (Trolox, ascorbate, glutathione), only offered limited protection that, once exhausted, had no impact on ABTS^{•-} formation or stability. Impressively, no ABTS^{•-} formation or other pro-oxidant effects by **Ru1** were observed under any experimental conditions. The mechanism for the catalytic reduction of ABTS^{•-} by **Ru1** and non-tertiary alcohols, along with the biological applications, will be detailed in subsequent reports.

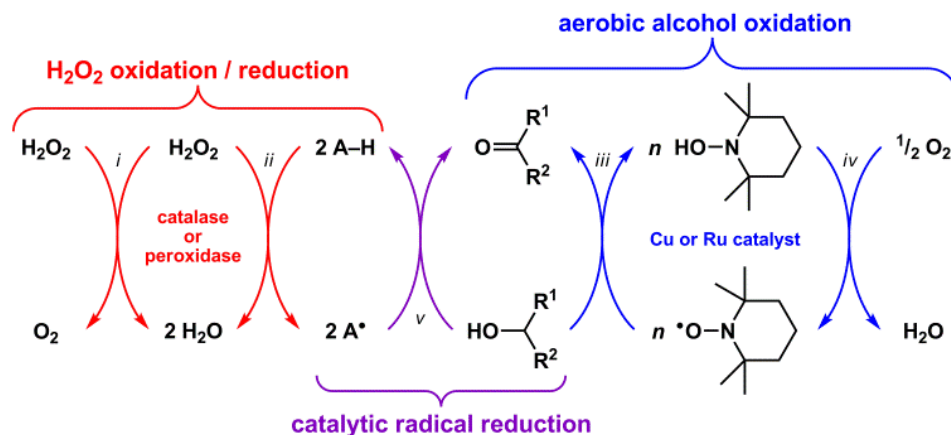
Acknowledgements

We are grateful to A. Mangalum for preceding work and helpful insight into the synthesis of **2** and **Ru1**.

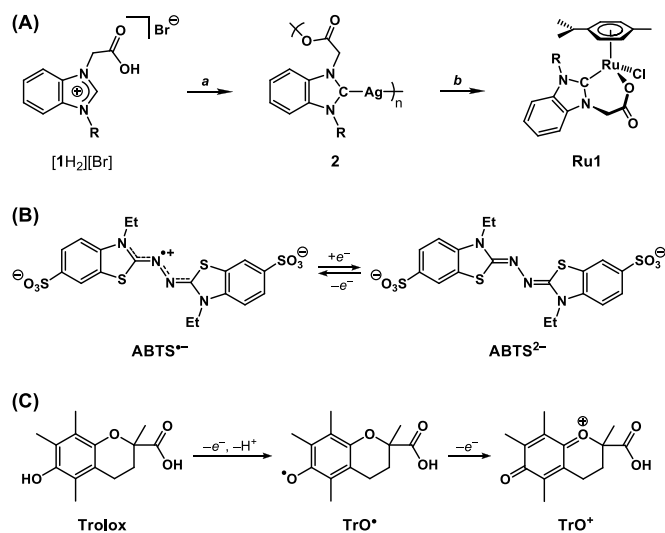
Notes

‡ In PBS, an ABTS^{•-} concentration of 6.7 μM would give rise to an absorbance value of 0.1. Achieving a similar value with O₂^{•-} or H₂O₂ would require a concentration of 53 μM or 2.29 mM, respectively.

§ The concentration of the successive ABTS^{•-} aliquots was selected based on the concentration of the initial 50 μM ABTS^{•-} that was degraded by the antioxidant. Because **Ru1** degraded all of the initial 50 μM ABTS^{•-} and Trolox only degraded 10 μM , the concentration of the multiple sequential aliquots added was 50 μM with **Ru1** and 10 μM with Trolox.



Scheme 2.1. Catalase normally catalyzes the disproportionation of $2 \text{H}_2\text{O}_2 \rightarrow \text{O}_2 + 2 \text{H}_2\text{O}$ (*i*), but at low H_2O_2 concentrations it can abstract H^\bullet from other molecules (A-H) and produce radicals (A^\bullet) in a manner similar to peroxidase (*ii*). Copper or ruthenium complexes can catalyze the oxidation of $\text{R}^1\text{-CHOH-R}^2$ to $\text{R}^1\text{-C(=O)-R}^2$ using a nitroxyl radical (e.g., TEMPO, $n = 1$ for Cu, $n = 2$ for Ru) as the H^\bullet abstracting reagent (*iii*) and O_2 as the terminal oxidant (*iv*). By a similar mechanism, a complex could catalyze radical reduction using a biologically relevant non-tertiary alcohol as the terminal reductant (*v*).



Scheme 2.2. (A) Synthesis of $[1H_2][Br]$, **2** and **Ru1**. Reagents and conditions: (a) Ag_2O (1.5 equiv.) or (b) $\{[RuCl(\eta^6\text{-cymene})]_2(\mu\text{-Cl})_2\}$ (1:1 Ag/Ru), CH_2Cl_2 , room temperature, 24 h. R = *p*-tolyl. (B) One-electron redox interconversion between $ABTS^{\bullet-}$ and $ABTS^{2-}$ (NH_4^+ counterions omitted for clarity). (C) First and second one-electron oxidations of Trolox.

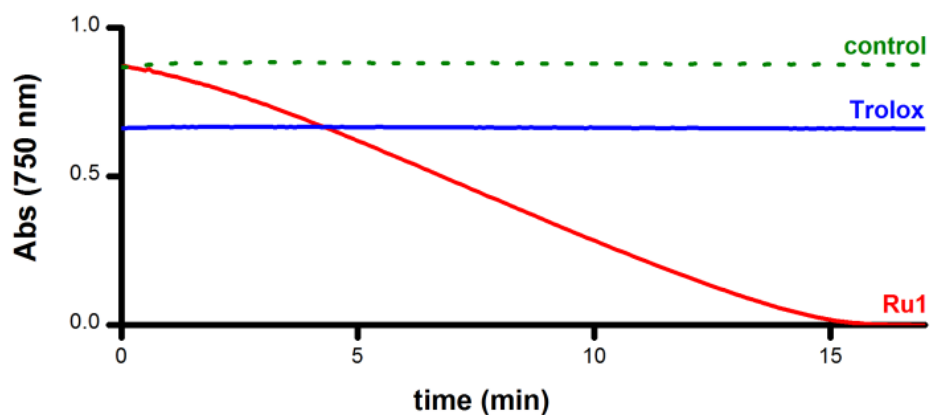


Fig. 2.1. Plot of relative $[\text{ABTS}^{\bullet-}]$ vs. time in EtOH following the addition of **Ru1** (CH_3CN stock, red line), Trolox (CH_3CN stock, blue line), or CH_3CN alone (dotted green line). Conditions: $[\text{Ru1}]_0$ or $[\text{Trolox}] = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, EtOH, $25 \text{ }^\circ\text{C}$; $[\text{ABTS}^{\bullet-}]$ determined using absorbance measured at 750 nm and $\epsilon_{750} = 1.6 \times 10^4 \text{ M}^{-1} \text{ cm}^{-1}$ (ref 33)

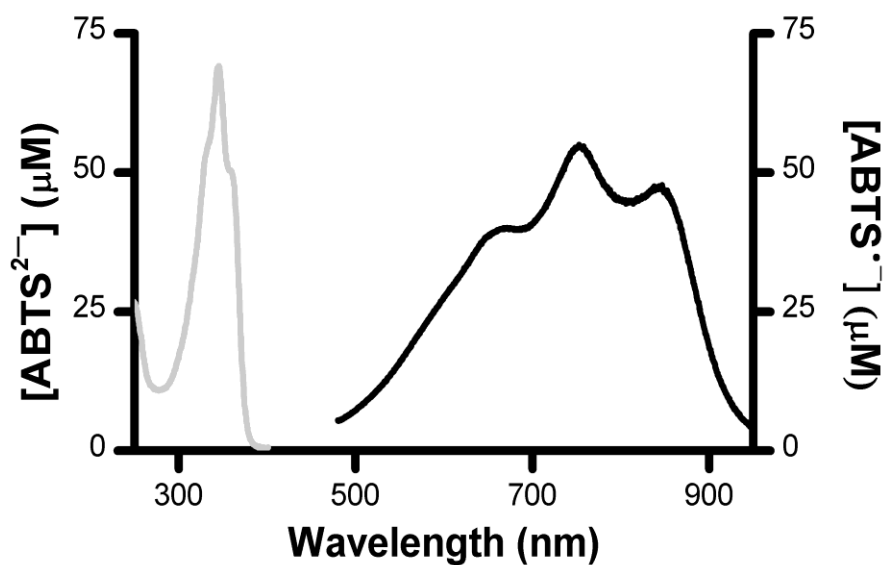


Fig. 2.2. UV/vis of ABTS²⁻ formed from ABTS^{•-} degradation by **Ru1** in EtOH. The absorbance at 340 nm (grey line) represents the total amount of ABTS²⁻ which includes the additional ABTS²⁻ formed and ABTS²⁻ from the original stock solution. Conditions: [Ru1]₀ = 5 µM, [ABTS^{•-}]₀ = 50 µM, EtOH, 25 °C, absorbance measured from 200–950 nm.

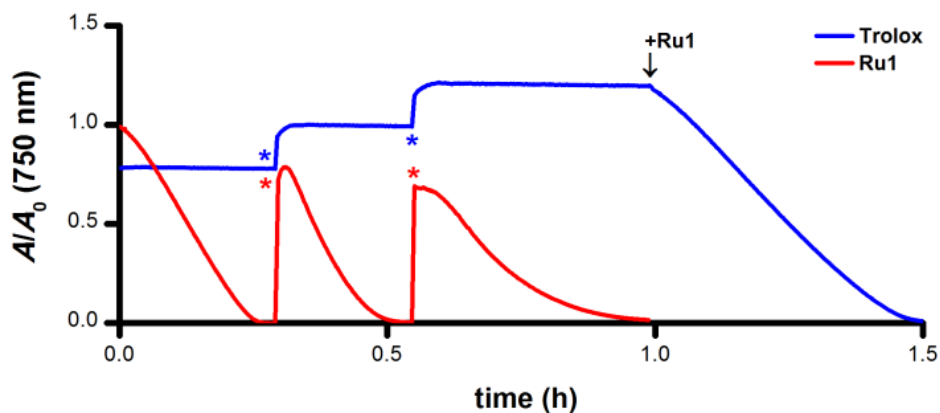


Fig. 2.3. Plot of relative $[ABTS^{\bullet-}]$ vs. time which shows the reduction of $ABTS^{\bullet-}$ after adding **Ru1** (red line) followed by 2 additional $50 \mu\text{M}$ $ABTS^{\bullet-}$ aliquots (*) and Trolox (blue line) followed by 2 additional $10 \mu\text{M}$ $ABTS^{\bullet-}$ aliquots (*), then $5 \mu\text{M}$ **Ru1**. Conditions: $[Ru1]_0$ or $[Trolox]_0 = 5 \mu\text{M}$, $[ABTS^{\bullet-}]_0 = 50 \mu\text{M}$, EtOH, $25 \text{ }^\circ\text{C}$; $[ABTS^{\bullet-}]$ determined using absorbance measured at 750 nm and $\epsilon_{750} = 1.6 \times 10^4 \text{ M}^{-1} \text{ cm}^{-1}$ (ref 33)

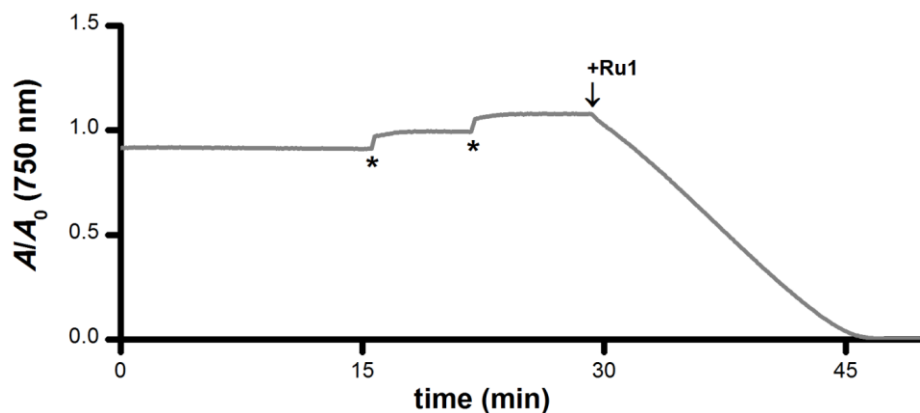


Fig. 2.4. Addition of 5 μM ascorbate (AscH^-) to an EtOH solution containing 50 μM $\text{ABTS}^{\bullet-}$ caused a rapid (within mixing time) 10% decrease in the absorbance at 750 nm, consistent with ascorbate functioning as a $1e^-$ reductant (i.e., $\text{AscH}^- \rightarrow \text{Asc}^\bullet$). Two additional 5 μM $\text{ABTS}^{\bullet-}$ aliquots were then introduced (*), which only caused the absorbance at 750 nm to increase by amounts equivalent to the concentration of $\text{ABTS}^{\bullet-}$ added, indicating that the antioxidant capacity of ascorbate had been exhausted after the reduction of only 1 equiv. of $\text{ABTS}^{\bullet-}$. Subsequent addition of 5 μM **Ru1** produced a 100% decrease in radical absorbance, indicating that **Ru1** could achieve quantitative $\text{ABTS}^{\bullet-}$ reduction under experimental conditions in which ascorbate (a traditional antioxidant) had been exhausted. Conditions: $[\text{AscH}^-]_0 = 5 \mu\text{M}$, $[\text{Ru1}] = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, EtOH, 25 $^\circ\text{C}$; absorbance measured at 750 nm.

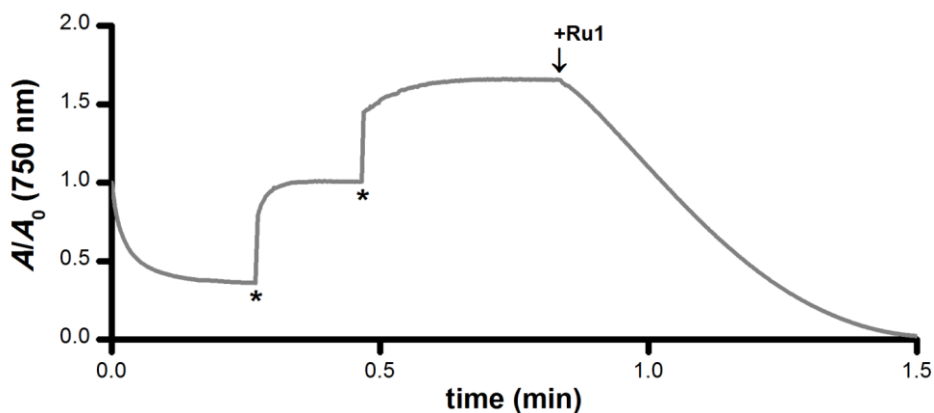


Fig. 2.5. Addition of 5 μM glutathione (G-SH) to an EtOH solution containing 50 μM $\text{ABTS}^{\bullet-}$ caused a gradual 60% decrease in the absorbance at 750 nm, consistent with glutathione functioning as a $6e^-$ reductant (i.e., $\text{G-SH} \rightarrow \text{G-SO}_3\text{H}$). Two additional 30 μM $\text{ABTS}^{\bullet-}$ aliquots were then introduced (*), which only caused the absorbance at 750 nm to increase by amounts equivalent to the concentration of $\text{ABTS}^{\bullet-}$ added, indicating that the antioxidant capacity of glutathione had been exhausted after the reduction of only 6 equiv. of $\text{ABTS}^{\bullet-}$. Subsequent addition of 5 μM **Ru1** produced a 100% decrease in radical absorbance, indicating that **Ru1** could achieve quantitative $\text{ABTS}^{\bullet-}$ reduction under experimental conditions in which glutathione (a traditional antioxidant) had been exhausted. Conditions: $[\text{G-SH}]_0 = 5 \mu\text{M}$, $[\text{Ru1}] = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, EtOH, 25 $^\circ\text{C}$; absorbance measured at 750 nm.

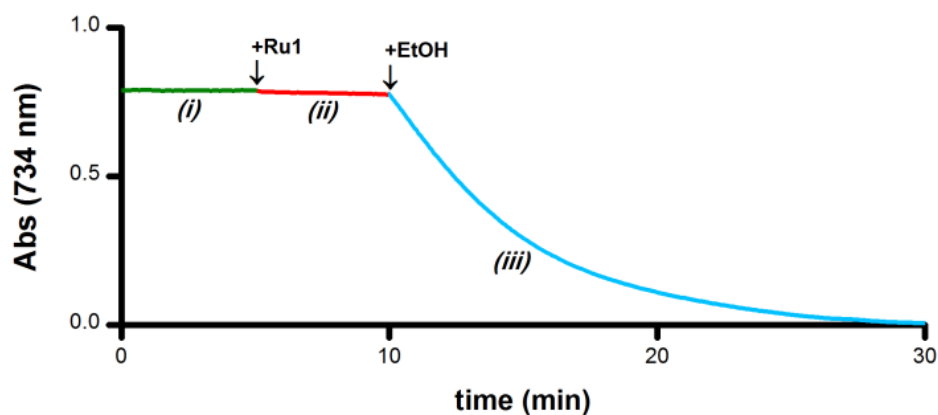


Fig. 2.6. Plot of relative $[\text{ABTS}^{\bullet-}]$ vs. time which shows the catalytic reduction of $\text{ABTS}^{\bullet-}$ by **Ru1** in PBS. By itself, $\text{ABTS}^{\bullet-}$ was stable in PBS (*i*, green line). Addition of **Ru1** did not cause $\text{ABTS}^{\bullet-}$ reduction (*ii*, red line). Subsequent addition of EtOH caused the absorbance to decrease (*iii*, blue line), which indicated that EtOH functioned as a terminal reductant. Conditions: $[\text{Ru1}]_0 = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[\text{EtOH}]_0 = 50 \text{mM}$, PBS (pH 7.4), 25 °C; $[\text{ABTS}^{\bullet-}]$ determined using absorbance measured at 734 nm and $\epsilon_{734} = 1.5 \times 10^4 \text{M}^{-1} \text{cm}^{-1}$ (ref 33)

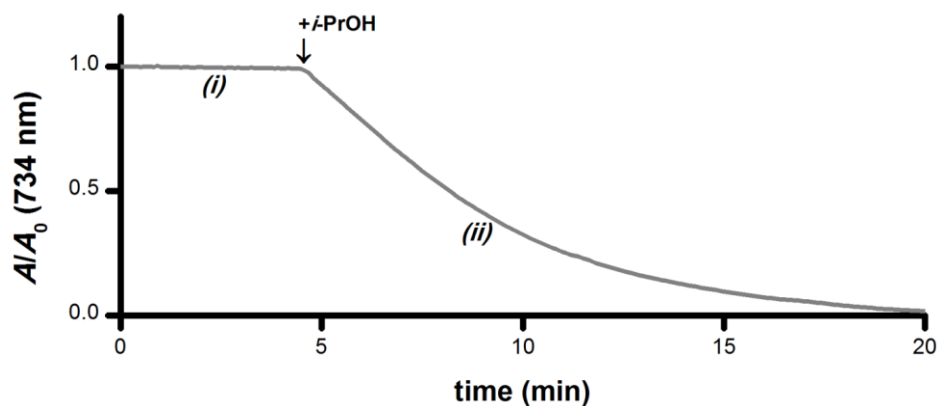


Fig. 2.7. Catalytic ABTS^{•-} reduction in PBS by **Ru1** with *i*-PrOH as the terminal reductant. No ABTS^{•-} degradation was observed in a PBS solution containing only **Ru1** (i), but subsequent addition of *i*-PrOH caused the absorbance at 734 nm to decrease (ii). Conditions: [**Ru1**]₀ = 5 μM, [ABTS^{•-}]₀ = 50 μM, [*i*-PrOH]₀ = 50 mM, PBS (pH 7.4), 25 °C, absorbance measured at 734 nm.

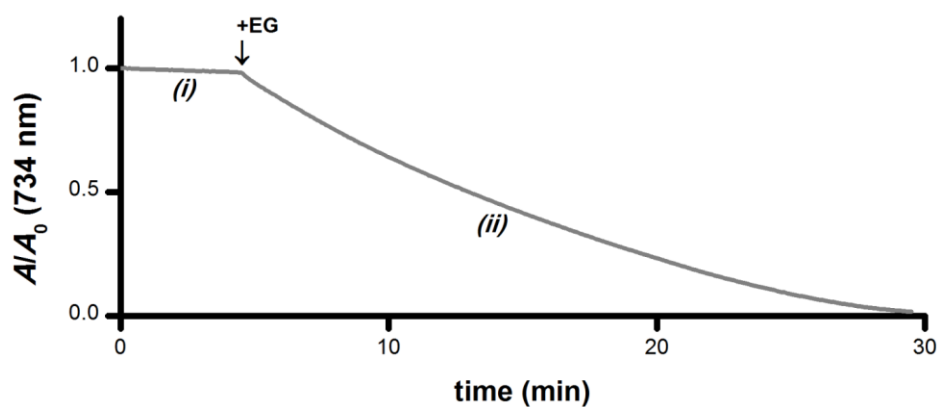


Fig. 2.8. Catalytic $\text{ABTS}^{\bullet-}$ reduction in PBS by **Ru1** with ethylene glycol (EG) as the terminal reductant. No $\text{ABTS}^{\bullet-}$ degradation was observed in a PBS solution containing only **Ru1** (i), but subsequent addition of EG caused the absorbance at 734 nm to decrease (ii). Conditions: $[\text{Ru1}]_0 = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[\text{EG}]_0 = 50 \text{mM}$, PBS (pH 7.4), 25 °C, absorbance measured at 734 nm.

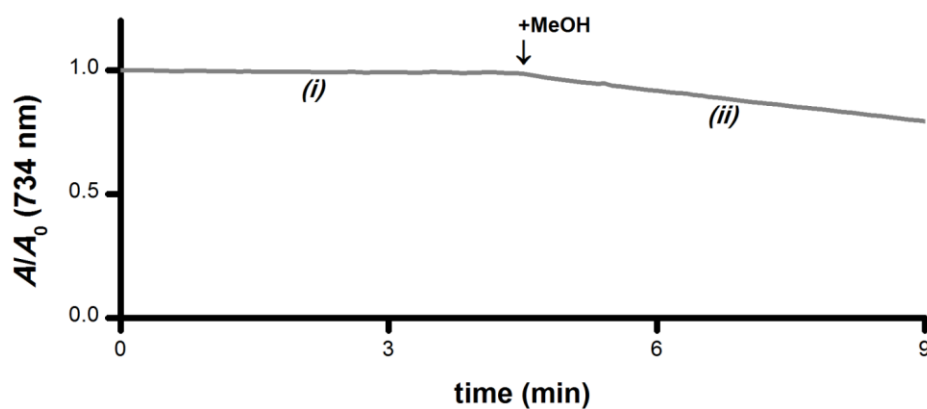


Fig. 2.9. Catalytic $\text{ABTS}^{\bullet-}$ reduction in PBS by **Ru1** with MeOH as the terminal reductant. No $\text{ABTS}^{\bullet-}$ degradation was observed in a PBS solution containing only **Ru1** (i), but subsequent addition of MeOH caused the absorbance at 734 nm to decrease (ii). Conditions: $[\text{Ru1}]_0 = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[\text{MeOH}]_0 = 50 \text{mM}$, PBS (pH 7.4), 25 °C, absorbance measured at 734 nm.

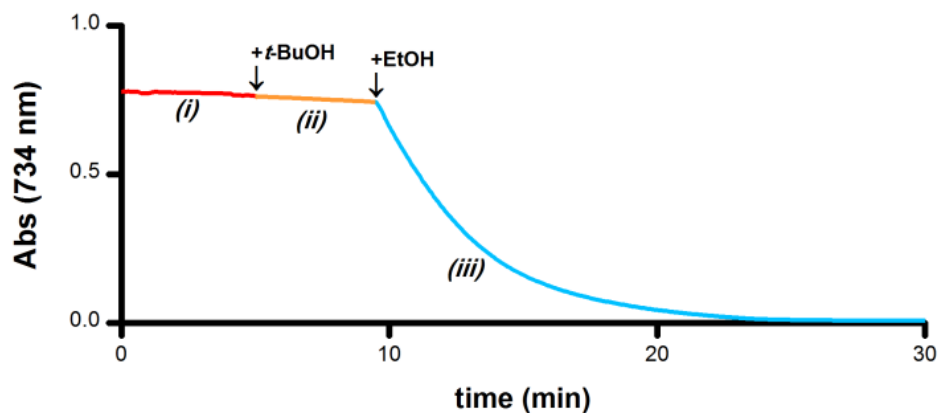


Fig. 2.10. Plot of relative $[\text{ABTS}^{\bullet-}]$ vs. time which shows the catalyst is not deactivated by *t*-BuOH. In a PBS solution containing only **Ru1**, $\text{ABTS}^{\bullet-}$ was stable (*i*, red line). Addition of *t*-BuOH did not cause any $\text{ABTS}^{\bullet-}$ reduction (*ii*, orange line). Subsequent addition of EtOH resulted in $\text{ABTS}^{\bullet-}$ degradation (*iii*, blue line), indicating that the lack of reactivity with *t*-BuOH was not due to catalyst deactivation. Conditions: $[\text{Ru1}]_0 = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[t\text{-BuOH}]_0 = 50 \text{mM}$, $[\text{EtOH}]_0 = 50 \text{mM}$, PBS (pH 7.4), 25 °C; $[\text{ABTS}^{\bullet-}]$ determined using absorbance measured at 734 nm and $\epsilon_{734} = 1.5 \times 10^4 \text{M}^{-1} \text{cm}^{-1}$ (ref 33).

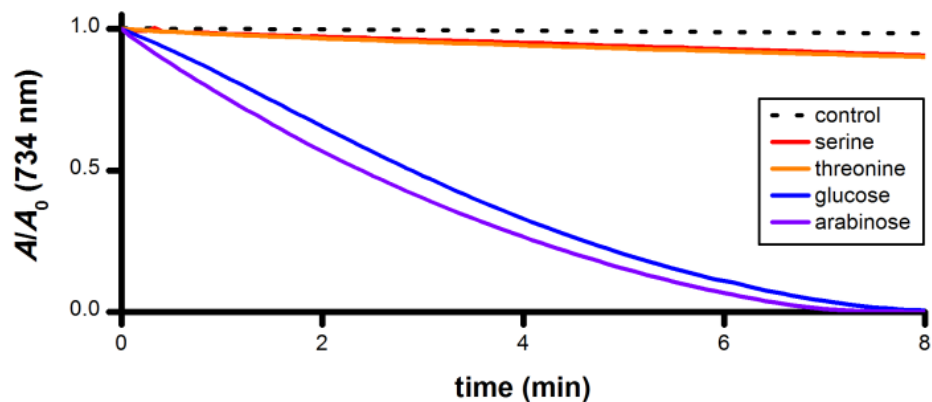


Fig. 2.11. Plot of relative $[\text{ABTS}^{\bullet-}]$ vs. time following the addition of amino acid or sugar ($t = 0$) to PBS solutions containing $\text{ABTS}^{\bullet-}$ and **Ru1**. Conditions: $[\text{Ru1}]_0 = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[\text{ROH}]_0 = 50 \text{ mM}$, PBS (pH 7.4), $25 \text{ }^\circ\text{C}$; $[\text{ABTS}^{\bullet-}]$ determined using absorbance measured at 734 nm and $\epsilon_{734} = 1.5 \times 10^4 \text{ M}^{-1} \text{ cm}^{-1}$ (ref 33).

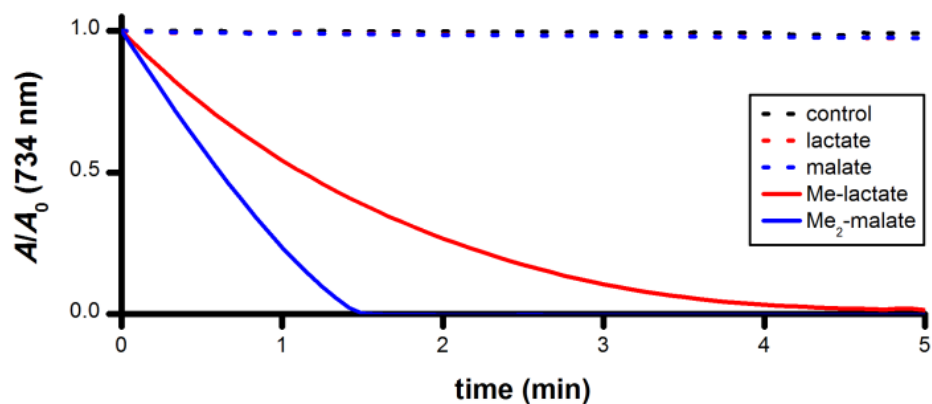


Fig. 2.12. Plot of relative $[\text{ABTS}^{\bullet-}]$ vs. time following the addition of lactate, malate or their methyl esters ($t = 0$) to PBS solutions containing $\text{ABTS}^{\bullet-}$ and **Ru1**. Conditions: $[\text{Ru1}]_0 = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[\text{ROH}]_0 = 50 \text{mM}$, PBS (pH 7.4), $25 \text{ }^\circ\text{C}$; $[\text{ABTS}^{\bullet-}]$ determined using absorbance measured at 734 nm and $\epsilon_{734} = 1.5 \times 10^4 \text{ M}^{-1} \text{ cm}^{-1}$ (ref 33)

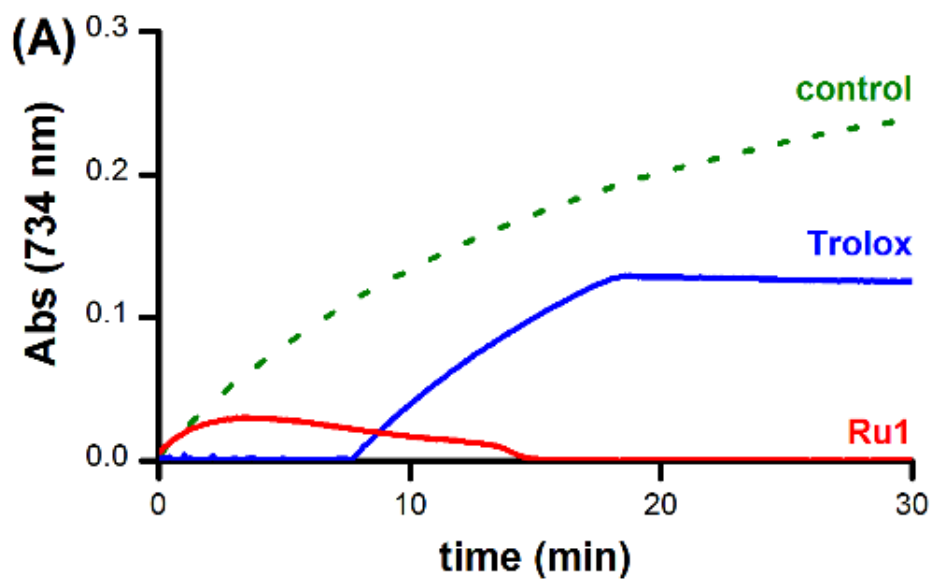


Fig. 2.13 (A) Plot of relative $[\text{ABTS}^{\bullet-}]$ vs. time which shows the *in situ* oxidation of ABTS^{2-} to $\text{ABTS}^{\bullet-}$ via HRP and H_2O_2 in the presence of **Ru1** and EtOH (red line), Trolox (blue line) or 0.19 M CH_3CN as a control (dotted green line). Conditions: $[\text{HRP}]_0 = 10 \text{ nM}$, $[\text{Ru1}]_0$ or $[\text{Trolox}]_0 = 5 \text{ }\mu\text{M}$, $[\text{H}_2\text{O}_2]_0 = 10 \text{ }\mu\text{M}$, $[\text{ABTS}^{2-}]_0 = 20 \text{ }\mu\text{M}$, $[\text{EtOH}]_0 = 50 \text{ mM}$, PBS (pH 7.4) at $25 \text{ }^\circ\text{C}$; $[\text{ABTS}^{\bullet-}]$ determined using absorbance measured at 734 nm and $\epsilon_{734} = 1.5 \times 10^4 \text{ M}^{-1} \text{ cm}^{-1}$ (ref 33)

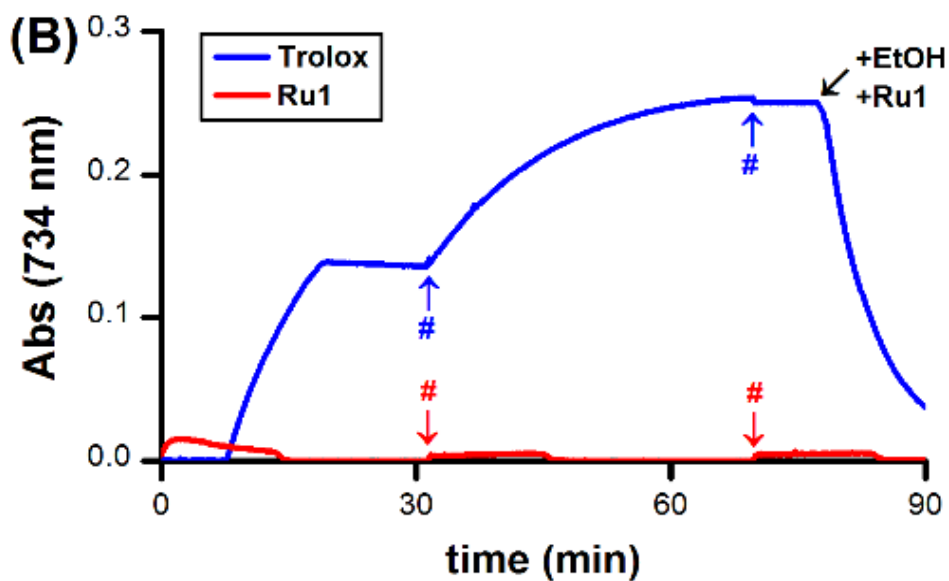


Fig. 2.13.B Plot of relative $[\text{ABTS}^{\bullet-}]$ vs. time which shows the $\text{ABTS}^{\bullet-}$ formation by HRP in the presence of **Ru1** and EtOH (red line) or Trolox (blue line) followed by two additional aliquots of $10 \mu\text{M}$ H_2O_2 (#). For the Trolox experiments shown, 50 mM EtOH and $5 \mu\text{M}$ **Ru1** were added after the final aliquot of H_2O_2 or $\text{ABTS}^{\bullet-}$, respectively. Conditions: $[\text{HRP}]_0 = 10 \text{ nM}$, $[\text{Ru1}]_0$ or $[\text{Trolox}]_0 = 5 \mu\text{M}$, $[\text{H}_2\text{O}_2]_0 = 10 \mu\text{M}$, $[\text{ABTS}^{2-}]_0 = 20 \mu\text{M}$, $[\text{EtOH}]_0 = 50 \text{ mM}$, PBS (pH 7.4) at $25 \text{ }^\circ\text{C}$; $[\text{ABTS}^{\bullet-}]$ determined using absorbance measured at 734 nm and $\epsilon_{734} = 1.5 \times 10^4 \text{ M}^{-1} \text{ cm}^{-1}$ (ref 33)

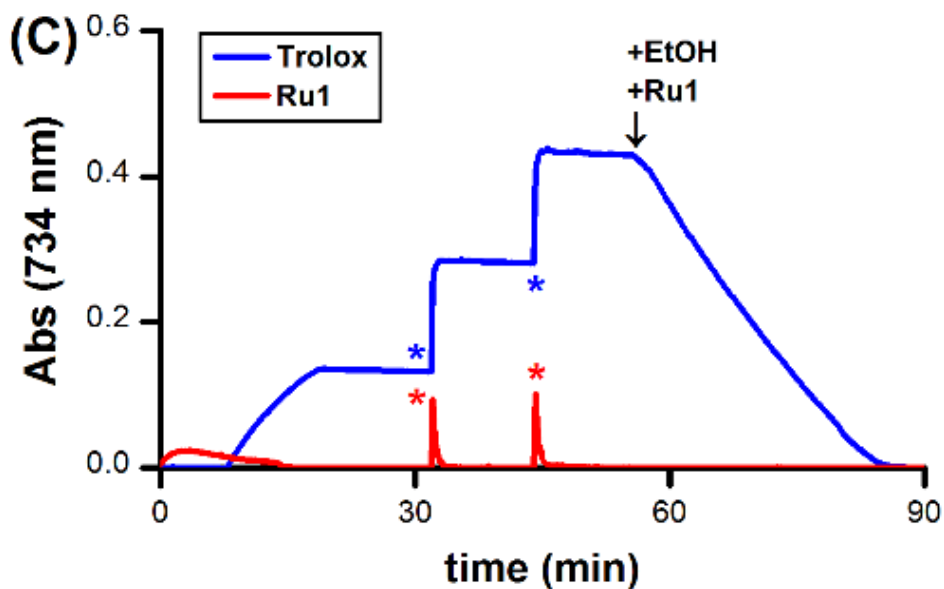


Fig. 2.13 C. Plot of relative $[\text{ABTS}^{\bullet-}]$ vs. time which shows the two $10 \mu\text{M}$ aliquots of chemically synthesized $\text{ABTS}^{\bullet-}$ (*). For the Trolox experiments shown, 50 mM EtOH and $5 \mu\text{M Ru1}$ were added after the final aliquot of H_2O_2 or $\text{ABTS}^{\bullet-}$, respectively. Conditions: $[\text{HRP}]_0 = 10 \text{ nM}$, $[\text{Ru1}]_0$ or $[\text{Trolox}]_0 = 5 \mu\text{M}$, $[\text{H}_2\text{O}_2]_0 = 10 \mu\text{M}$, $[\text{ABTS}^{2-}]_0 = 20 \mu\text{M}$, $[\text{EtOH}]_0 = 50 \text{ mM}$, PBS (pH 7.4) at $25 \text{ }^\circ\text{C}$; $[\text{ABTS}^{\bullet-}]$ determined using absorbance measured at 734 nm and $\epsilon_{734} = 1.5 \times 10^4 \text{ M}^{-1} \text{ cm}^{-1}$ (ref 33)

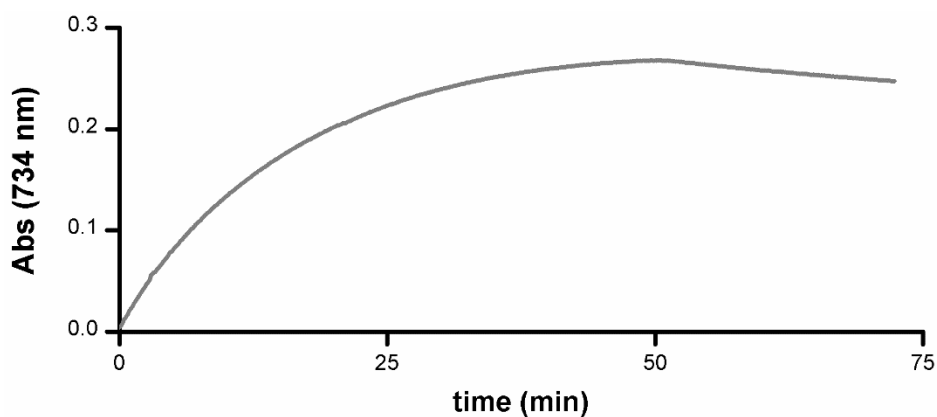


Fig. 2.14. *In situ* oxidation of ABTS^{2-} to $\text{ABTS}^{\bullet-}$ via horseradish peroxidase (HRP) and H_2O_2 in the presence of 0.19 M CH_3CN . Formation of $\text{ABTS}^{\bullet-}$ began immediately following addition of H_2O_2 ($t = 0$). After 50 min, $[\text{ABTS}^{\bullet-}]$ reached a maximum of 17.9 μM . The subsequent gradual decline in A_{734} was due to normal $\text{ABTS}^{\bullet-}$ thermal decay. Conditions: $[\text{CH}_3\text{CN}]_0 = 0.19$ M, $[\text{HRP}]_0 = 10$ nM, $[\text{H}_2\text{O}_2]_0 = 10$ μM , $[\text{ABTS}^{2-}]_0 = 20$ μM ; PBS (pH 7.4), 25 °C; absorbance measured at 734 nm.

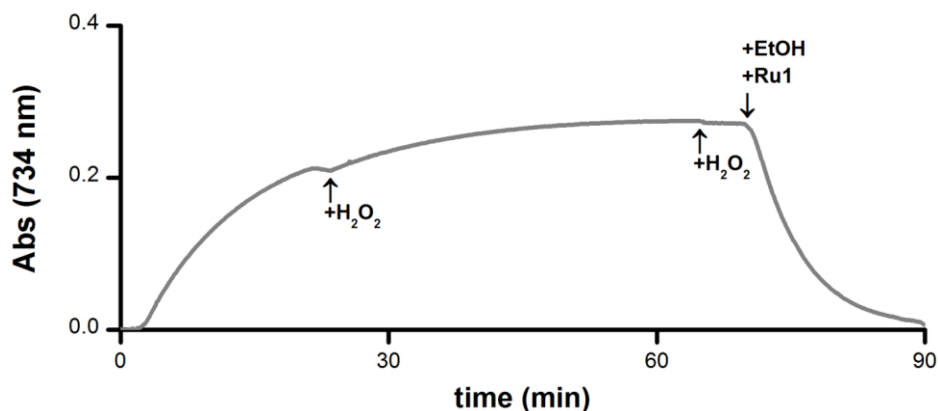


Fig. 2.15. *In situ* oxidation of ABTS^{2-} to $\text{ABTS}^{\bullet-}$ via horseradish peroxidase (HRP) and H_2O_2 in the presence of $5 \mu\text{M}$ glutathione (G-SH) followed by two additional aliquots of $10 \mu\text{M}$ H_2O_2 . No $\text{ABTS}^{\bullet-}$ formation was observed for the first 1.4 min after the addition of the first H_2O_2 aliquot. After 1.4 min, $\text{ABTS}^{\bullet-}$ began to form and, 20 min after onset, $[\text{ABTS}^{\bullet-}]$ reached a maximum of $14.1 \mu\text{M}$. The subsequent gradual decline in A_{734} was due to normal $\text{ABTS}^{\bullet-}$ thermal decay. Addition of the second $10 \mu\text{M}$ H_2O_2 aliquot resulted in complete oxidation of the remaining ABTS^{2-} over the course of 41 min, and addition of the third $10 \mu\text{M}$ H_2O_2 aliquot afforded no change in A_{734} . Subsequent addition of 50 mM EtOH and $5 \mu\text{M}$ **Ru1** caused $\text{ABTS}^{\bullet-}$ reduction to occur that was complete within 20 min of the addition of **Ru1**. Conditions: $[\text{G-SH}]_0 = 5 \mu\text{M}$, $[\text{HRP}]_0 = 10 \text{ nM}$, $[\text{H}_2\text{O}_2]_0 = 10 \mu\text{M}$, $[\text{ABTS}^{2-}]_0 = 20 \mu\text{M}$; $[\text{Ru1}] = 5 \mu\text{M}$, $[\text{EtOH}] = 50 \text{ mM}$; PBS (pH 7.4), $25 \text{ }^\circ\text{C}$; absorbance measured at 734 nm.

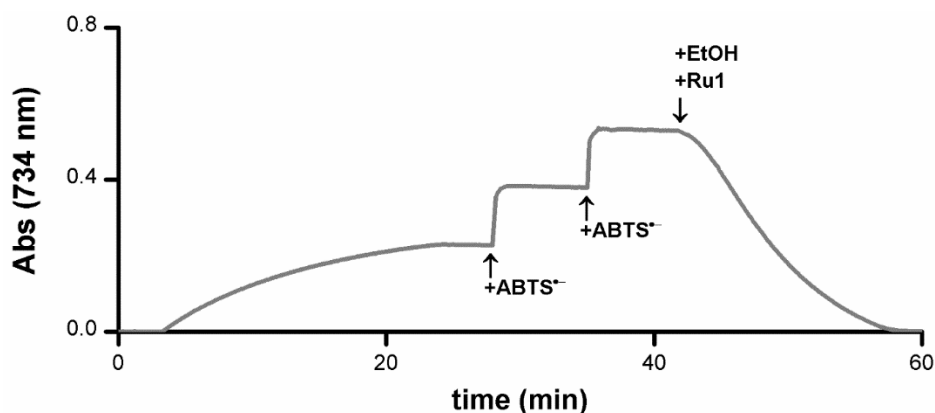


Fig. 2.16. *In situ* oxidation of ABTS^{2-} to $\text{ABTS}^{\bullet-}$ via horseradish peroxidase (HRP) and H_2O_2 in the presence of $5 \mu\text{M}$ ascorbate (AscH^-) followed by two aliquots of $10 \mu\text{M}$ $\text{ABTS}^{\bullet-}$. No $\text{ABTS}^{\bullet-}$ formation was observed for the first 3.3 min after the addition of the first H_2O_2 aliquot. After 3.3 min, $\text{ABTS}^{\bullet-}$ began to form and, 21 min after onset, $[\text{ABTS}^{\bullet-}]$ reached a maximum of $15.3 \mu\text{M}$. The subsequent gradual decline in A_{734} was due to normal $\text{ABTS}^{\bullet-}$ thermal decay. Addition of the first and second $10 \mu\text{M}$ $\text{ABTS}^{\bullet-}$ aliquots caused A_{734} to increase by amounts proportional to the $\text{ABTS}^{\bullet-}$ added. The gradual decline in A_{734} following each $\text{ABTS}^{\bullet-}$ aliquot was due to normal $\text{ABTS}^{\bullet-}$ thermal decay. Subsequent addition of 50 mM EtOH and $5 \mu\text{M}$ **Ru1** caused $\text{ABTS}^{\bullet-}$ reduction to occur that was complete within 18 min of the addition of **Ru1**. Conditions: $[\text{AscH}^-]_0 = 5 \mu\text{M}$, $[\text{HRP}]_0 = 10 \text{ nM}$, $[\text{H}_2\text{O}_2]_0 = 10 \mu\text{M}$, $[\text{ABTS}^{2-}]_0 = 20 \mu\text{M}$; $[\text{Ru1}] = 5 \mu\text{M}$, $[\text{EtOH}] = 50 \text{ mM}$; PBS (pH 7.4), $25 \text{ }^\circ\text{C}$; absorbance measured at 734 nm.

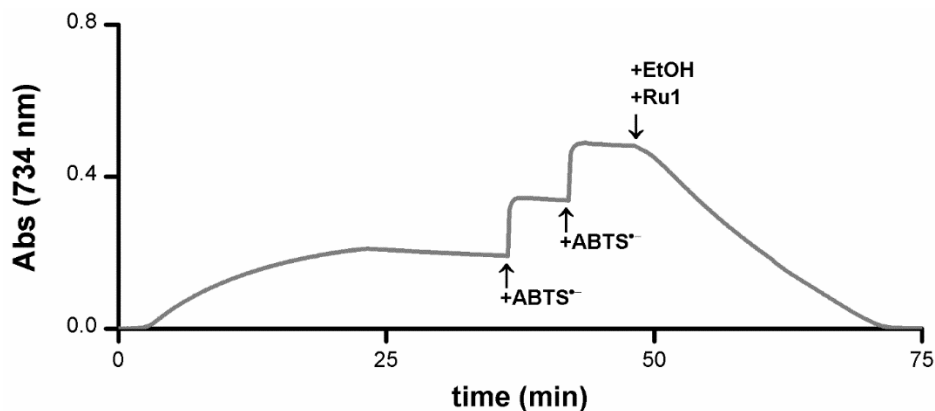
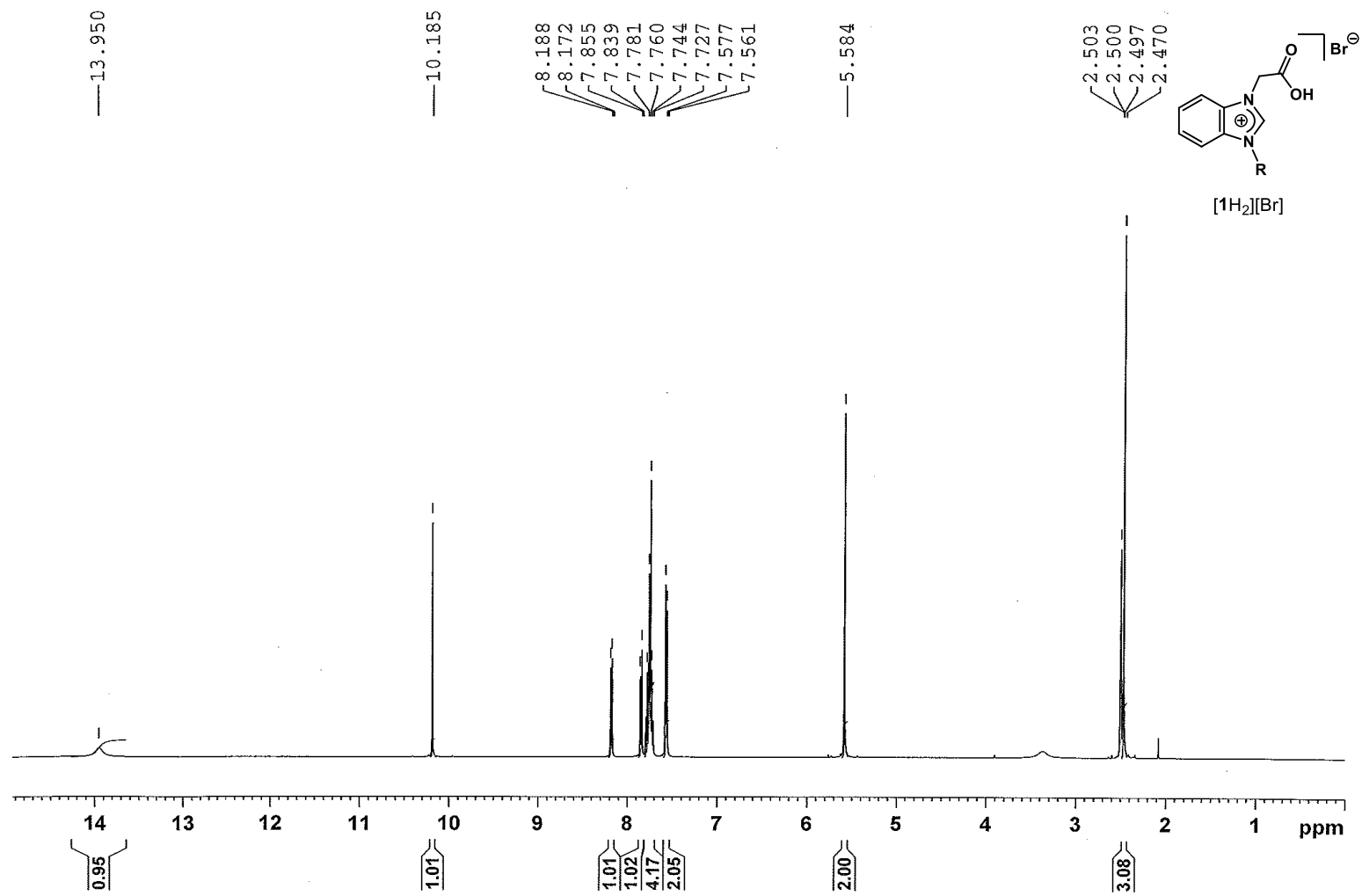
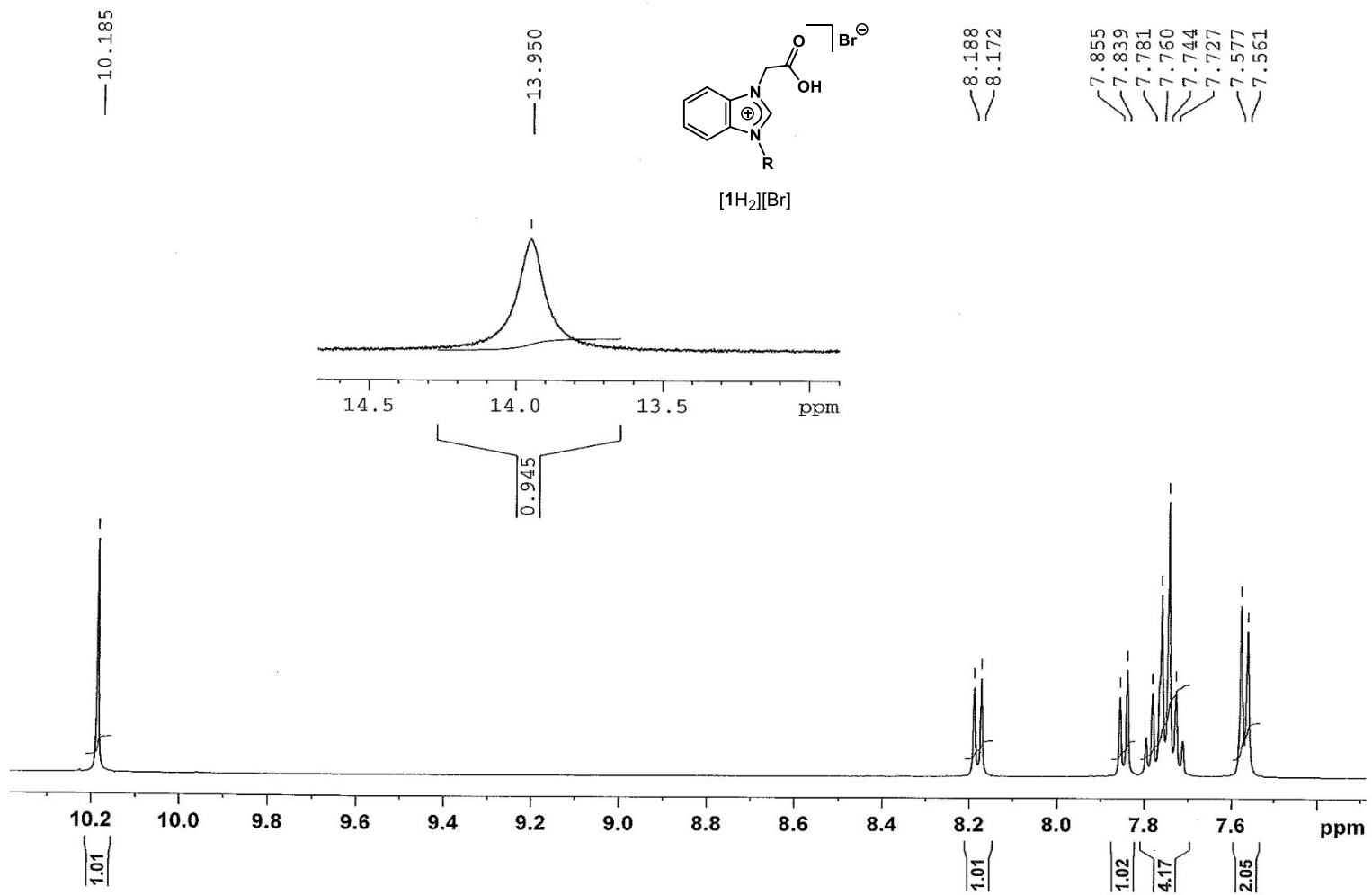
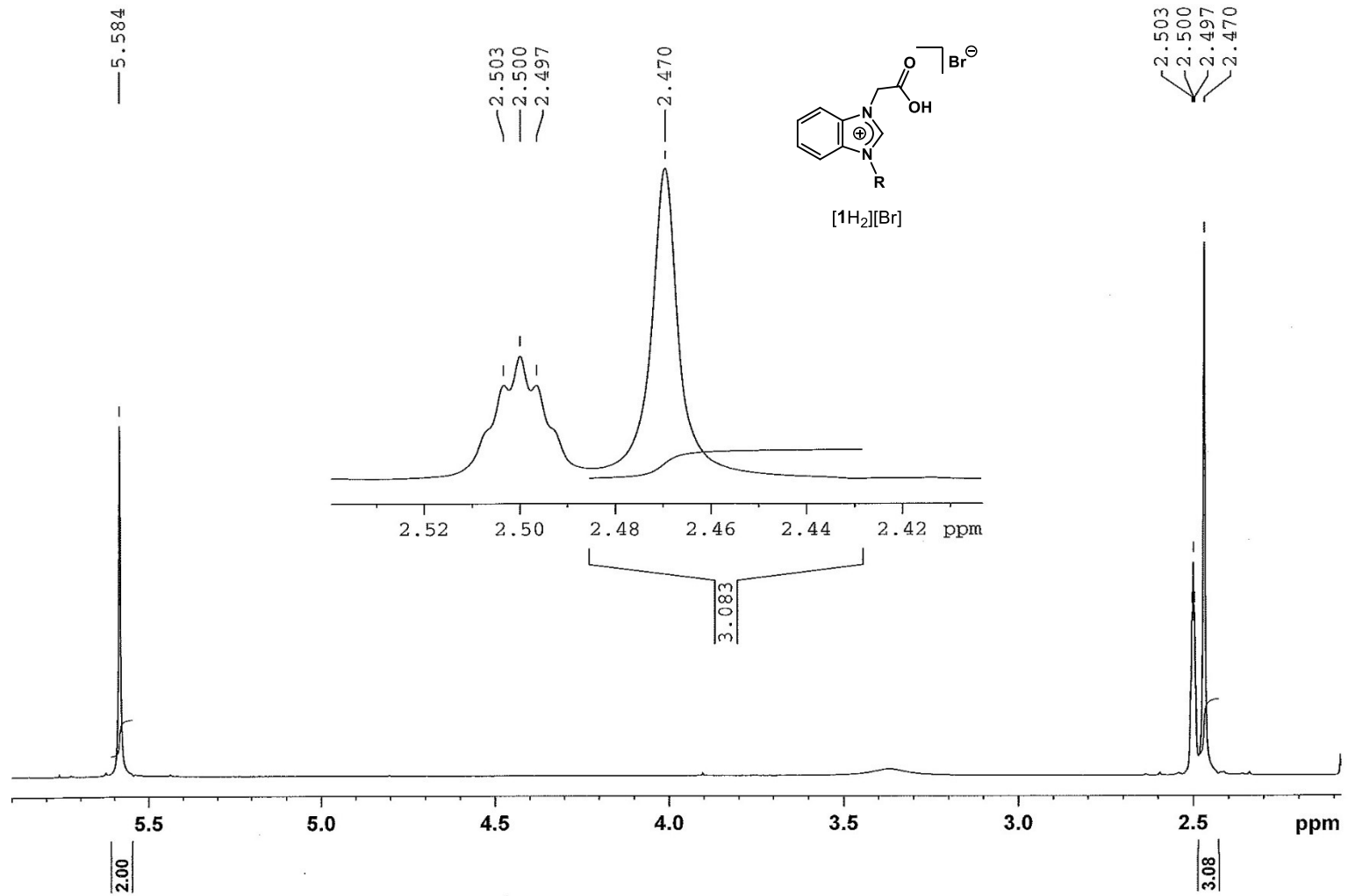
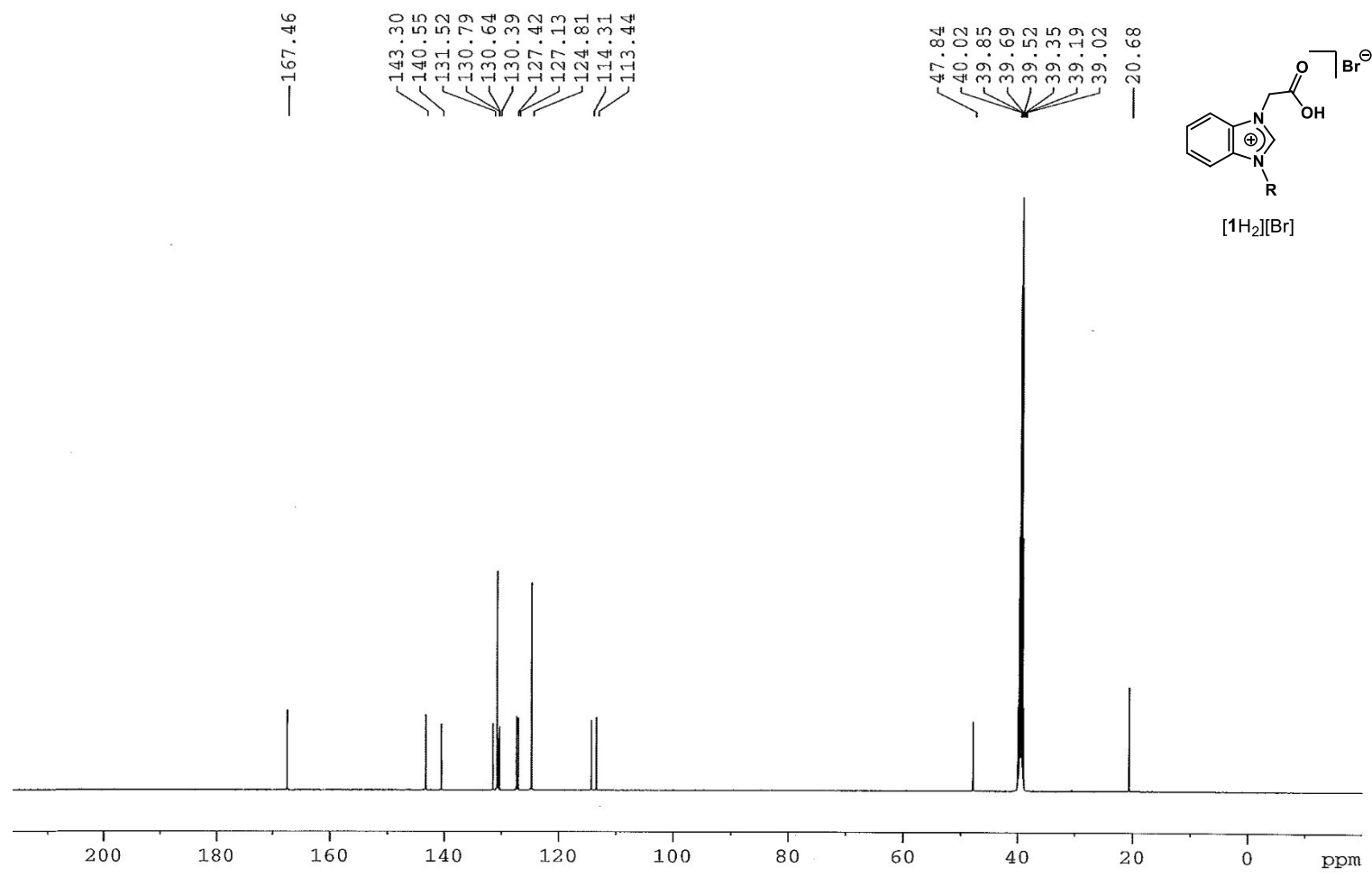


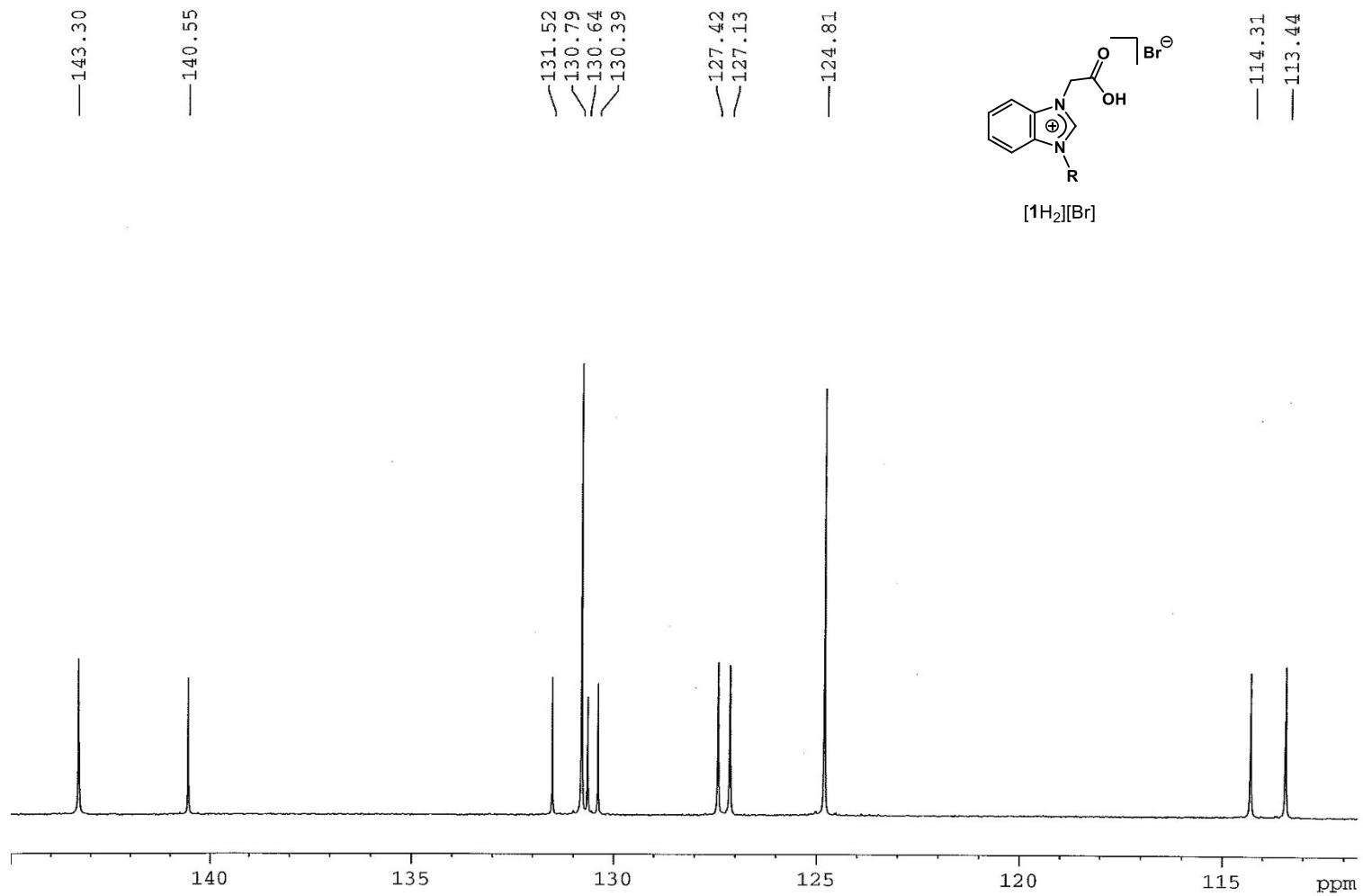
Fig. 2.17. *In situ* oxidation of ABTS^{2-} to $\text{ABTS}^{\bullet-}$ via horseradish peroxidase (HRP) and H_2O_2 in the presence of $5 \mu\text{M}$ glutathione (G-SH) followed by two aliquots of $10 \mu\text{M}$ $\text{ABTS}^{\bullet-}$. No $\text{ABTS}^{\bullet-}$ formation was observed for the first 1.4 min after the addition of the first H_2O_2 aliquot. After 1.4 min, $\text{ABTS}^{\bullet-}$ began to form and, 22 min after onset, $[\text{ABTS}^{\bullet-}]$ reached a maximum of $14.0 \mu\text{M}$. The subsequent gradual decline in A_{734} was due to normal $\text{ABTS}^{\bullet-}$ thermal decay. Addition of the first and second $10 \mu\text{M}$ $\text{ABTS}^{\bullet-}$ aliquots caused A_{734} to increase by amounts proportional to the $\text{ABTS}^{\bullet-}$ added. The gradual decline in A_{734} following each $\text{ABTS}^{\bullet-}$ aliquot was due to normal $\text{ABTS}^{\bullet-}$ thermal decay. Subsequent addition of 50 mM EtOH and $5 \mu\text{M}$ **Ru1** caused $\text{ABTS}^{\bullet-}$ reduction to occur that was complete within 24 min of the addition of **Ru1**. Conditions: $[\text{G-SH}]_0 = 5 \mu\text{M}$, $[\text{HRP}]_0 = 10 \text{ nM}$, $[\text{H}_2\text{O}_2]_0 = 10 \mu\text{M}$, $[\text{ABTS}^{2-}]_0 = 20 \mu\text{M}$; $[\text{Ru1}] = 5 \mu\text{M}$, $[\text{EtOH}] = 50 \text{ mM}$; PBS (pH 7.4), $25 \text{ }^\circ\text{C}$; absorbance measured at 734 nm.

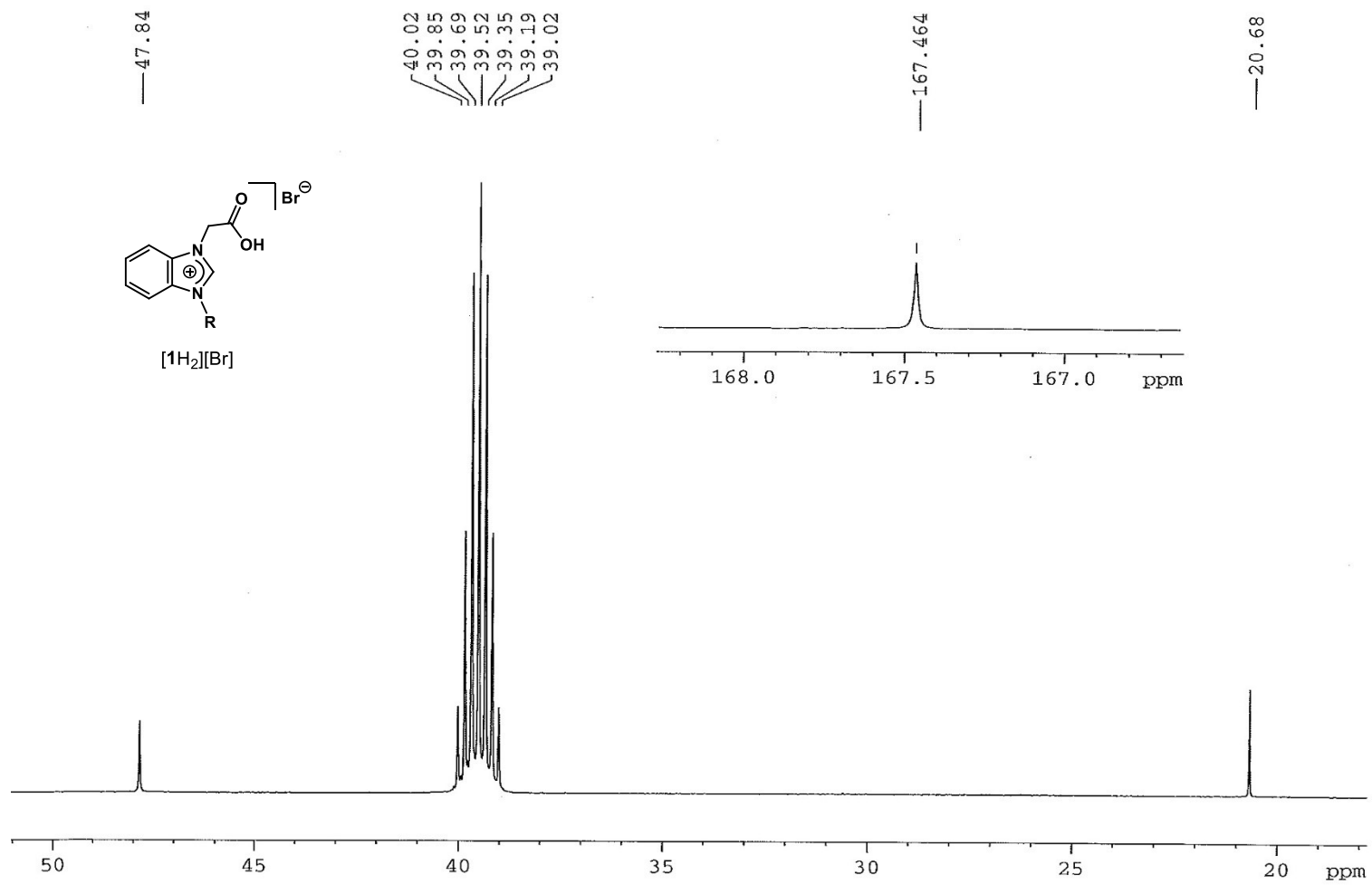


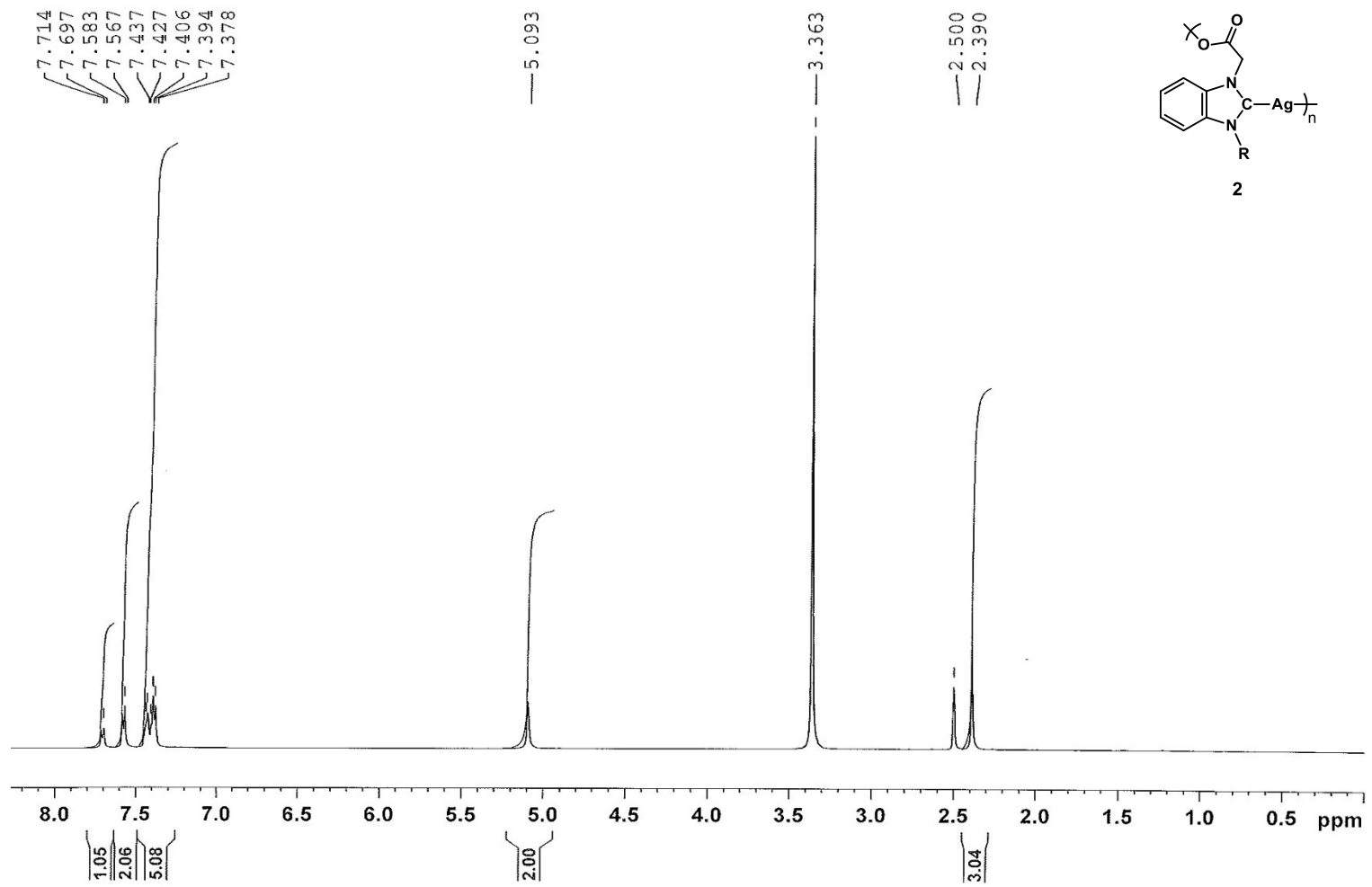


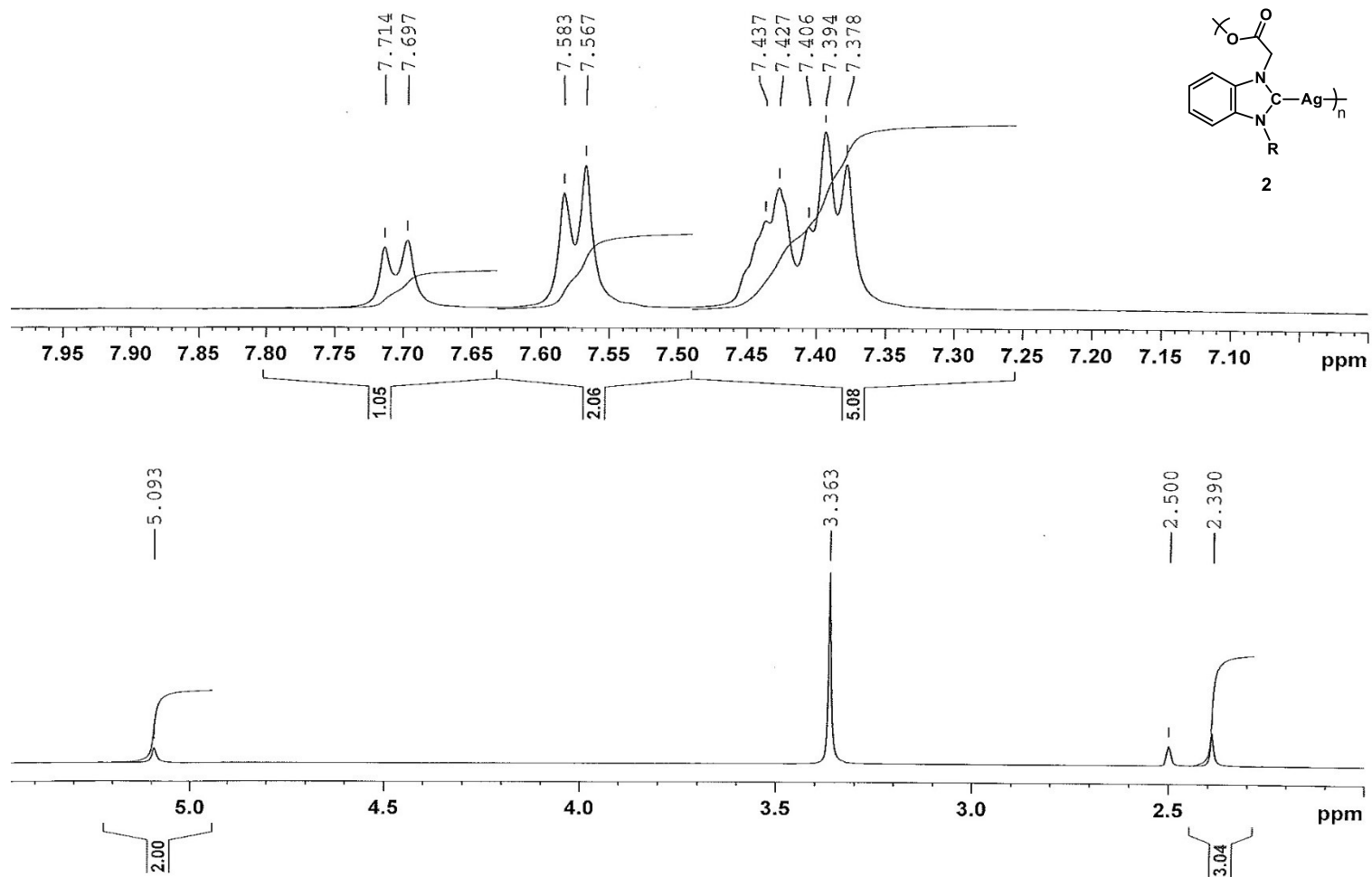


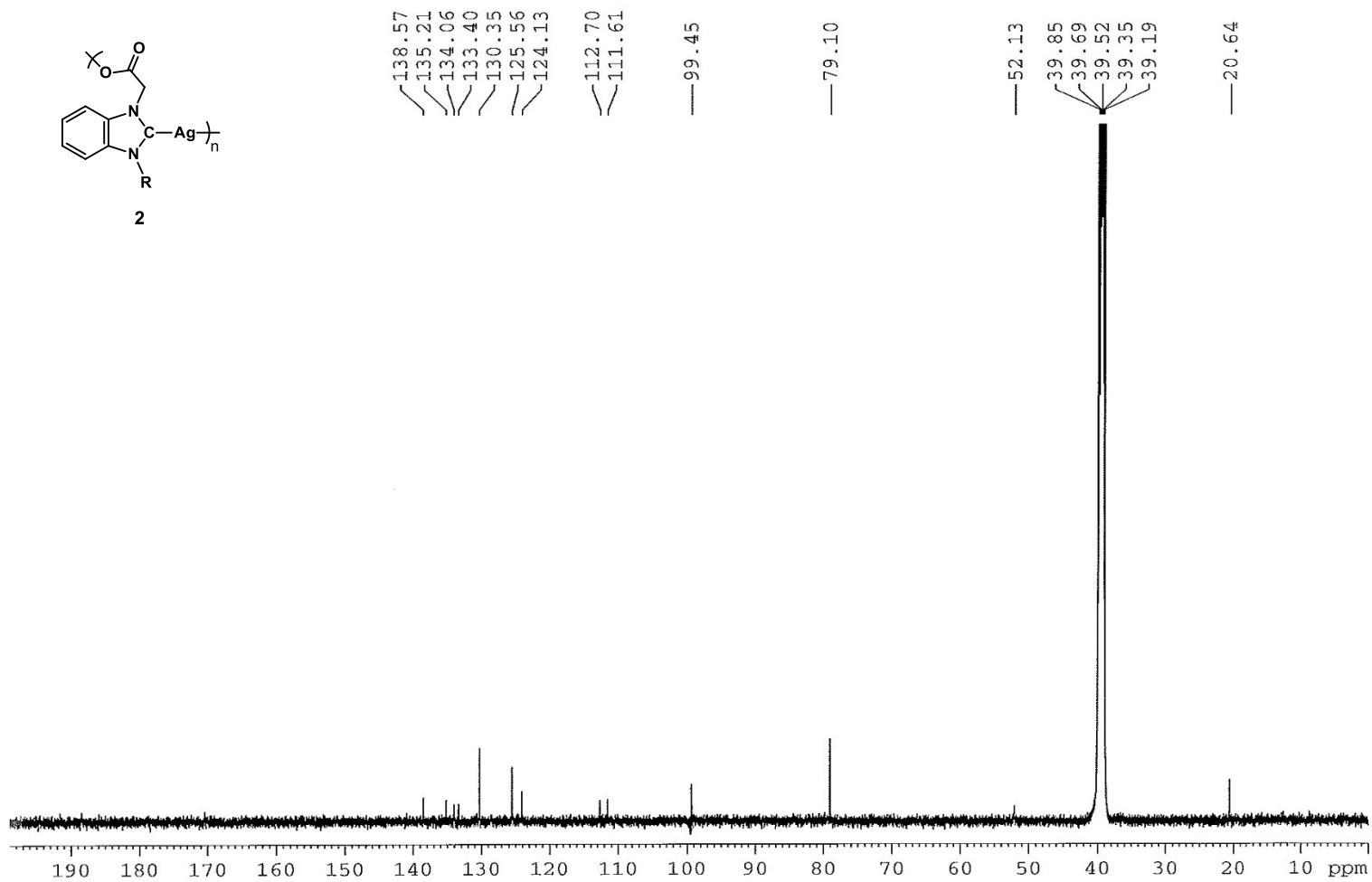
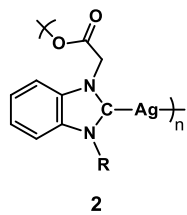


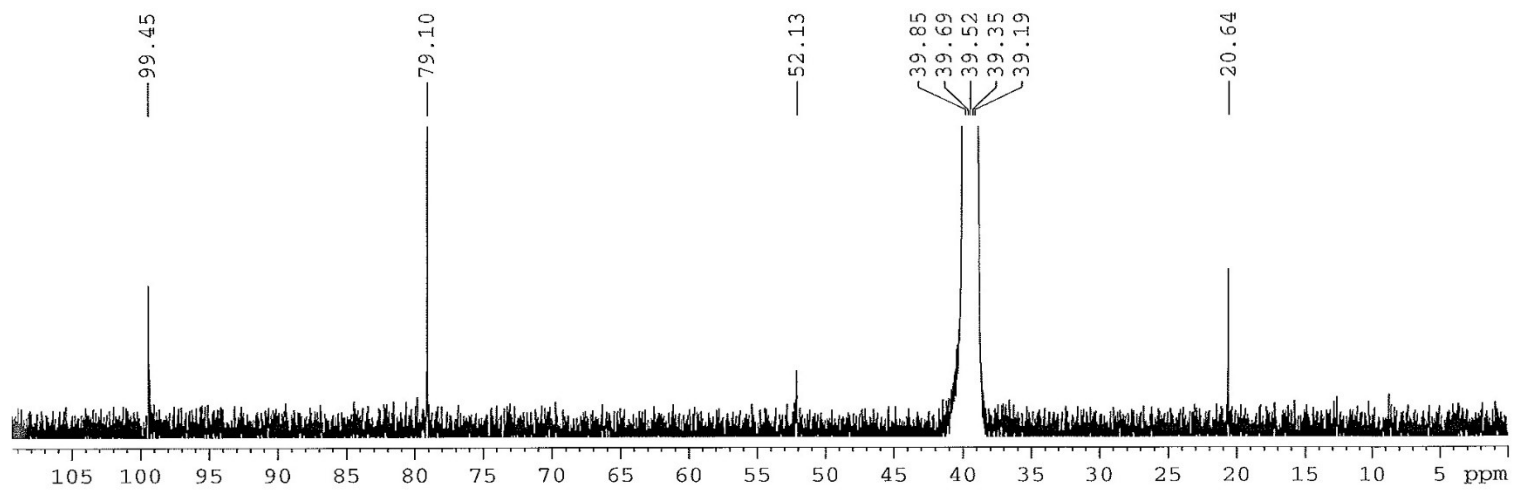
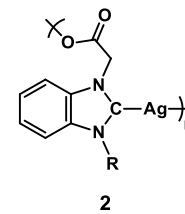
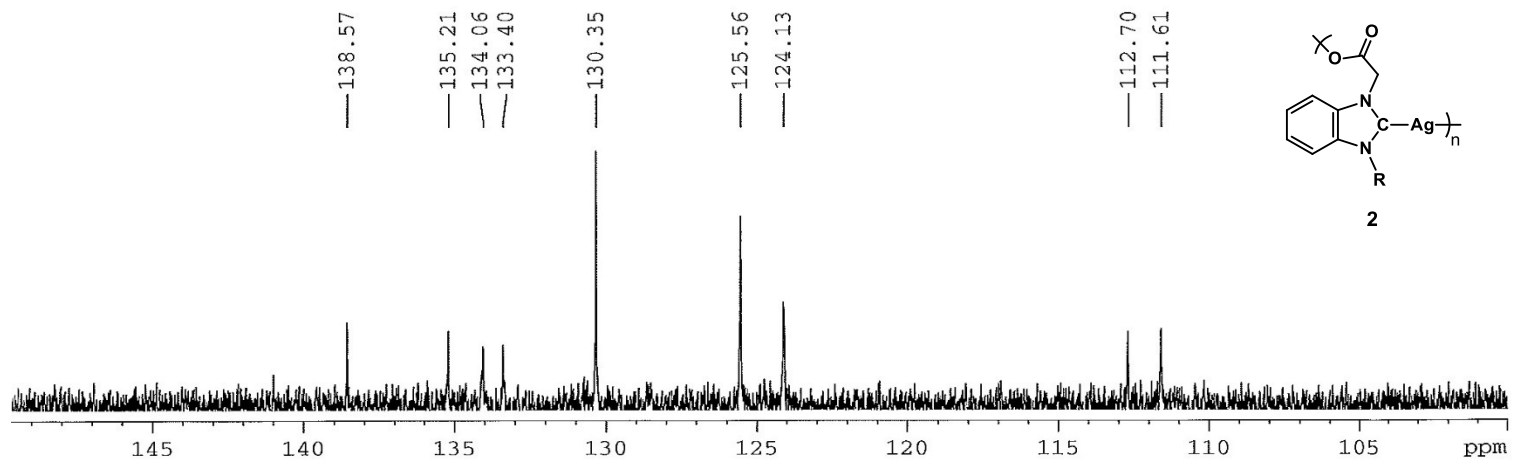


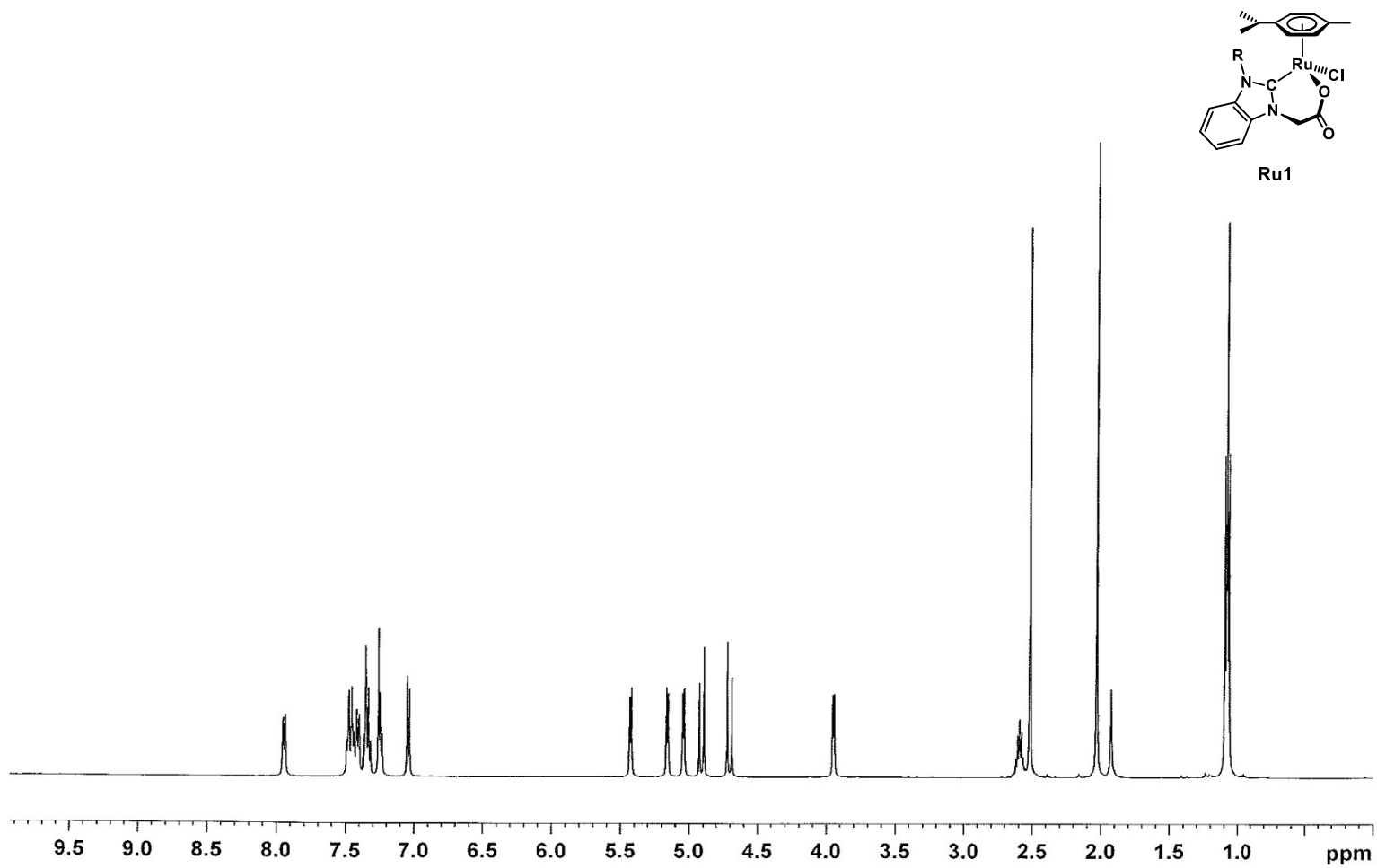


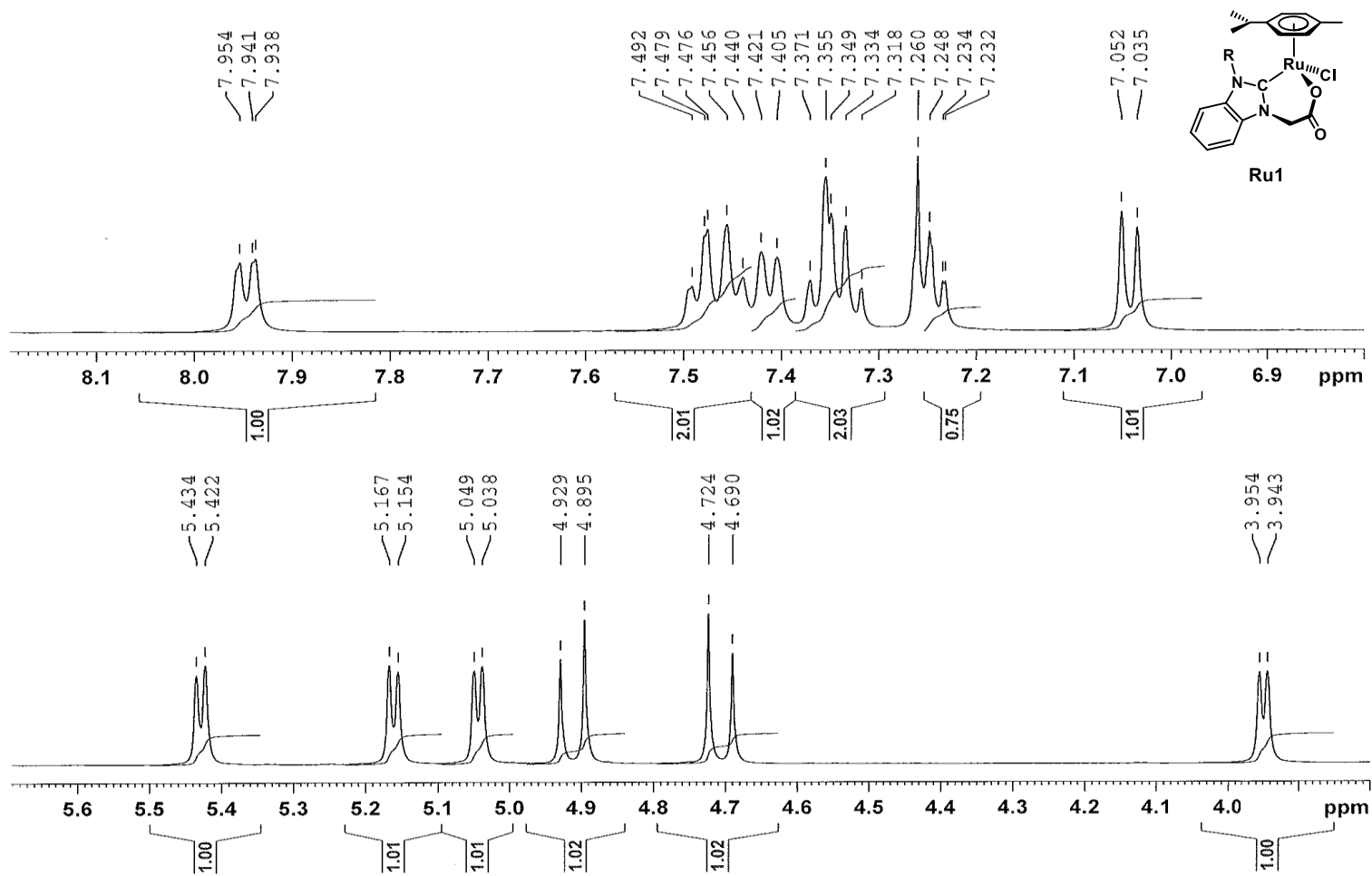


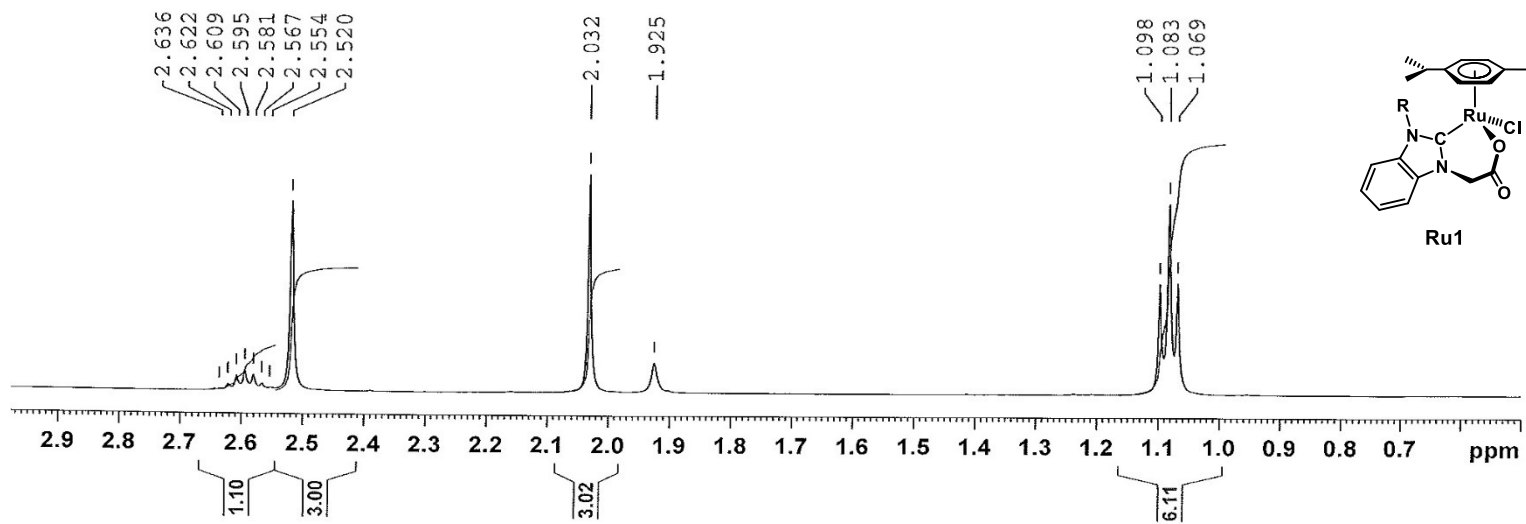


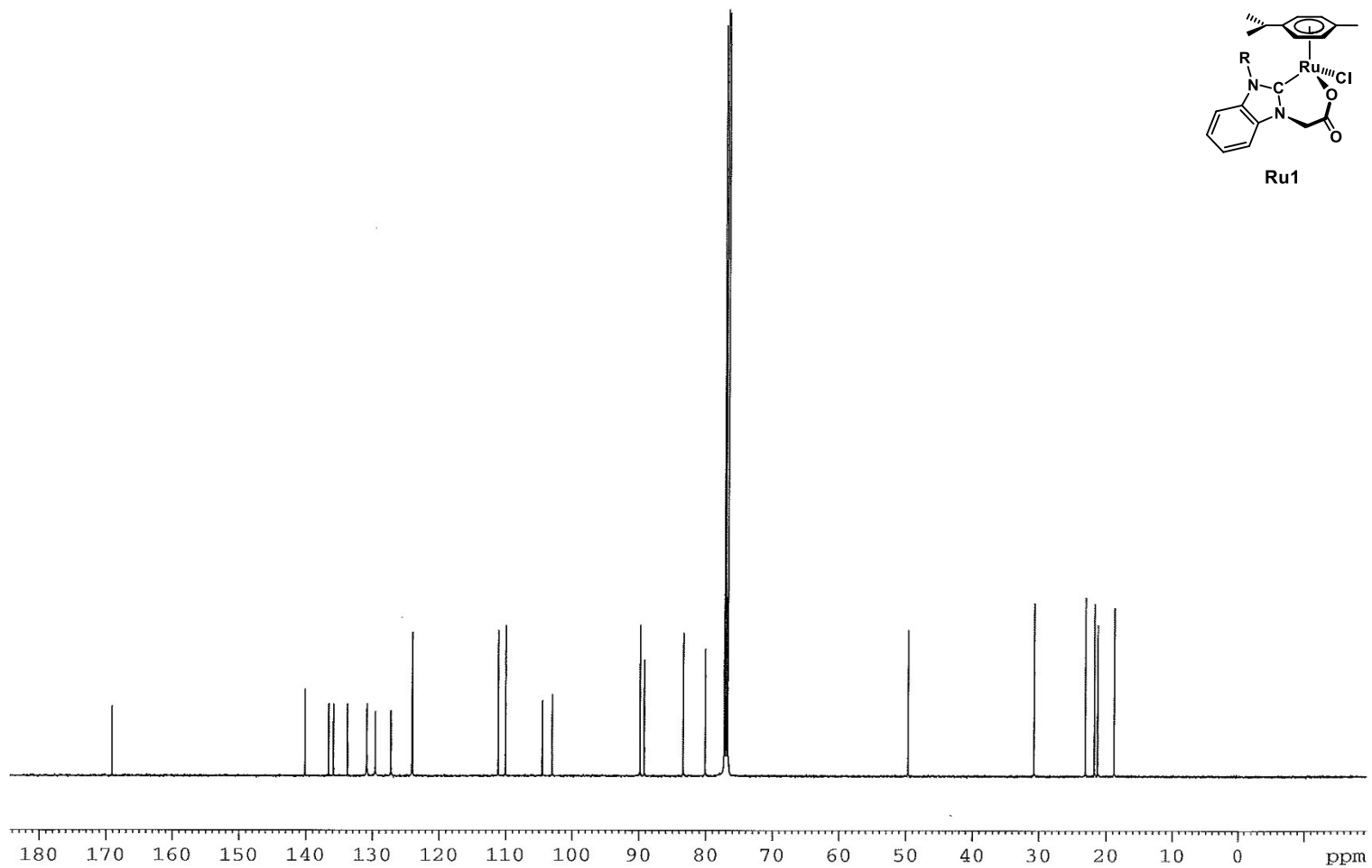


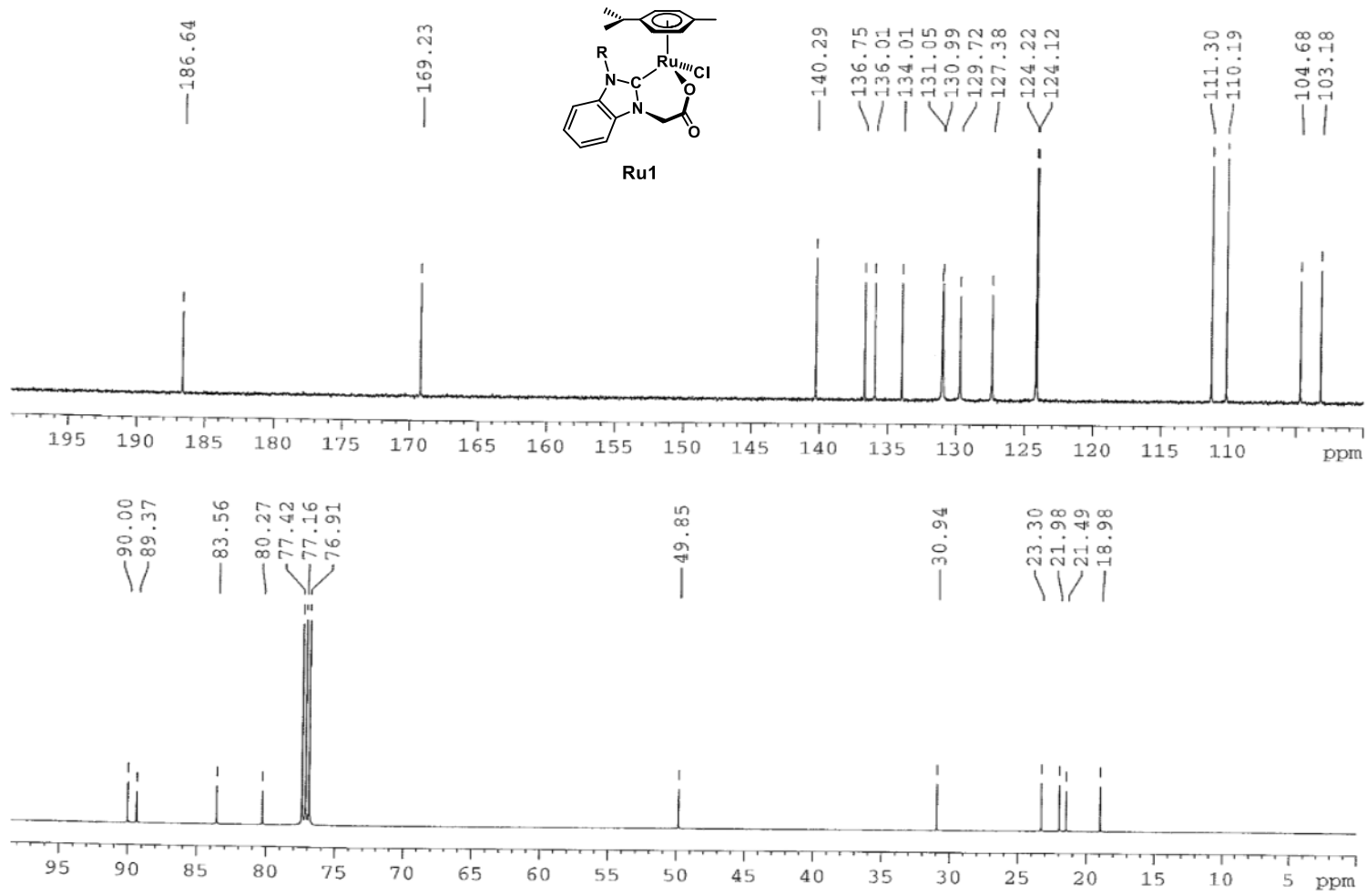












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CHAPTER THREE

CATALYTIC RADICAL REDUCTION IN AQUEOUS SOLUTION BY A RUTHENIUM HYDRIDE INTERMEDIATE

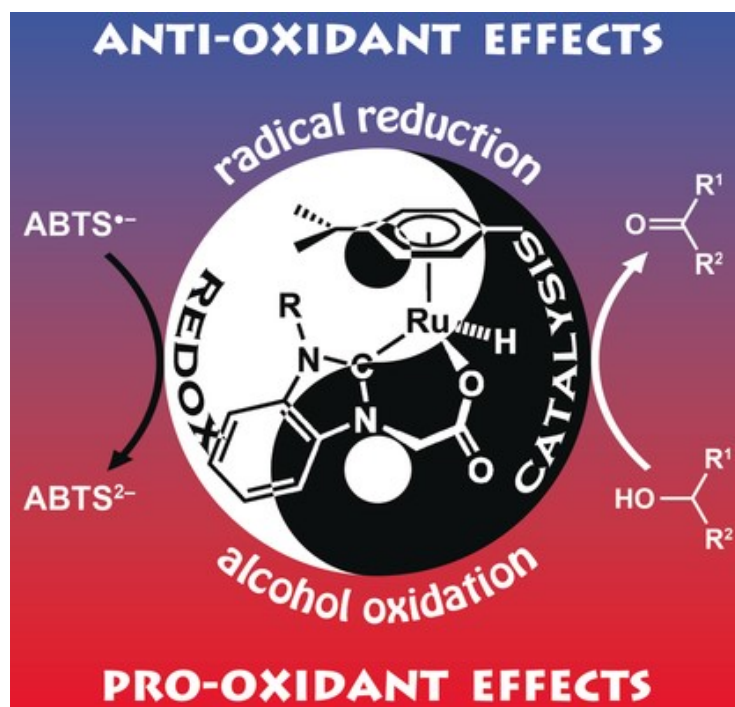


Figure 3. Catalytic radical reduction in buffered aqueous solutions was achieved by a Ru complex with non-tertiary alcohol terminal reductants. Primary kinetic isotope effects were observed with both C–H and O–H bonds. Radical reduction was faster at higher pH, and no reaction occurred in pure H₂O. Mechanistic studies suggest that the catalytically active species is a Ru-hydride intermediate formed by β-hydride elimination from a Ru-alkoxide.

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ABSTRACT

Some manganese complexes can catalyze both antioxidant and pro-oxidant reactions whereby the disparate reactivity modes are determined by the catalyst environment and afford distinct therapeutic effects. We recently reported the reduction of radicals in buffered aqueous solution catalyzed by a ruthenium complex with biologically relevant non-tertiary alcohols as terminal reductants. Mechanistic evidence is presented, indicating that this catalytic radical reduction is achieved by a Ru-hydride intermediate formed by a β -hydride elimination from a Ru-alkoxide species. A similar mechanism and Ru-hydride intermediate was previously reported to kill cancer cells with catalytic pro-oxidant effects. Therefore, our demonstration of catalytic pro-oxidant effects by the same type of intermediate reveals new potential therapeutic strategies and applications for catalytic systems that form Ru-hydride intermediates.

INTRODUCTION

Reactive oxygen species (ROS) and other free radicals can damage critical biomolecules in living systems and are implicated in a wide variety of pathologies.^{1,2} Antioxidants can reduce ROS or oxidizing radicals and thus protect against damage, but they typically proceed through stoichiometric reductions that deplete their protective capacity.³ However, a catalytic reduction could regenerate an antioxidant after each turnover and thus enable it to provide significantly greater protection. Manganese porphyrin-based complexes can catalyze $O_2^{\bullet-}$ and H_2O_2 disproportionation reactions and have shown beneficial antioxidant effects in stroke and diabetes therapy by reducing ROS levels.⁴⁻⁹ Interestingly, some of these Mn complexes (two of which are in Phase 1 clinical trials) can also produce pro-oxidant effects in cancer cells, which endows them with potent chemotherapeutic activity.⁴⁻¹¹ The ability of a Mn complex to produce antioxidant or pro-oxidant effects in different environments is derived from the same catalytic cycle and intermediates, whereby the catalyst environment determines the antioxidant/pro-oxidant outcome.⁴⁻¹⁵ Knowledge of the mechanism and intermediates by which a complex catalyzes a redox reaction in a biological system is thus essential for developing its therapeutic applications.

We recently reported the **Ru1**-catalyzed reduction of $ABTS^{\bullet-}$ to $ABTS^{2-}$ in buffered aqueous solution with biologically relevant non-tertiary alcohols as terminal reductants (Scheme 3.1 A–B).¹⁶ Although $ABTS^{\bullet-}$ is not found in biological systems, it is an ideal radical substrate because its oxidizing power ($ABTS^{\bullet-}/ABTS^{2-} E_{1/2} = +0.68$ V vs. NHE)¹⁷ is comparable to radicals that are generated during biological oxidative stress

(e.g., $E^\circ = 1.6$ V for RO^\bullet , 1.0 V for ROO^\bullet , 0.92 V for Cys-S^\bullet , and so forth)^{*,18} and its high absorptivity ($1.5 \times 10^4 \text{ M}^{-1} \text{ cm}^{-1}$)¹⁹ permits quantification at lower concentrations than would be achievable with $\text{O}_2^{\bullet-}$ or H_2O_2 .^{†,20,21} No $\text{ABTS}^{\bullet-}$ reduction occurred with **Ru1** alone if either the O–H or C–H group of a CH–OH moiety was absent, or in pure H_2O solutions, which is consistent with alcohol oxidation by β -hydride elimination from a Ru–alkoxide. A similar process has been proposed for the aerobic oxidation of alcohols catalyzed by Ru-oxide/hydroxide materials, in which the resulting Ru–H intermediate reacts with O_2 , the terminal oxidant, by 1,2–insertion (Scheme 3.1 C, top).²²

If a Ru complex that can oxidize a biomolecule to yield a Ru–H species, and if that Ru–H species can then be oxidized by O_2 to yield H_2O_2 , then that Ru complex can exert catalytic pro-oxidant effects in biological systems, which has been shown to result in potent anticancer properties by Sadler et al.²³⁻²⁵ Compounds such as *trans*- $[\text{RuCl}_4(\text{imidazole})(S\text{-dms})]$ (NAMI-A) and related coordination complexes that also possess anticancer properties, have similarly been shown to function by exertion of pro-oxidant effects.²⁶⁻²⁸ Antioxidant effects have been observed with some Ru coordination complexes, but none exhibited catalytic antioxidant activity.^{29,30} Inspired by the fact that 1) a single Mn porphyrin–based complex can produce either antioxidant or pro-oxidant effects in different environments through a conserved catalytic cycle and set of intermediates and, 2) a catalytic cycle involving a Ru–H intermediate can produce pro-oxidant effects, we hypothesized that a comparable catalytic cycle and Ru–H

* All redox potentials are reported relative to NHE unless specified otherwise.

† Achieving the same absorbance value as $\text{ABTS}^{\bullet-}$ requires a 7.9-fold higher concentration of $\text{O}_2^{\bullet-}$ ($\epsilon = 1.9 \times 10^3 \text{ M}^{-1} \text{ cm}^{-1}$ at 250 nm) or a 344-fold higher concentration of H_2O_2 ($\epsilon = 43.6 \text{ M}^{-1} \text{ cm}^{-1}$ at 240 nm).

intermediate could also give rise to antioxidant effects. Herein, we present mechanistic evidence indicating that catalytic antioxidant effects by **Ru1** in the form of catalytic radical reduction (Scheme 3.1 C, bottom) are derived from a similar catalytic cycle and Ru–H intermediate previously shown to produce catalytic pro–oxidant effects.

RESULTS AND DISCUSSION

An organoruthenium–chloride complex and the corresponding aquo complex exist in H₂O solution that is determined by the relative ligand affinities and total Cl[–] ion concentration, but ligand exchange is rapid for both the aquation and anation reactions.^{31,32} Therefore, the Cl[–] ligand in **Ru1** will be similarly labile toward substitution with other ligands in phosphate–buffered saline (PBS) solutions, such as ABTS^{•–}, H₂O and EtOH. The chemical synthesis of ABTS^{•–} from ABTS^{2–} does not reach 100% completion,¹⁹ therefore an additional 100 μM of ABTS^{2–} was included to control the variation in rates caused by unreacted ABTS^{2–}. Addition of 5 μM **Ru1** to an aqueous solution containing 100 μM of ABTS^{2–} will result in rapid ligand exchange (relative to turnover) to afford [L_nRu–A_{red}]^{1–} and begin the catalytic cycle (Scheme 3.2). Substitution of ABTS^{2–} with H₂O will yield [L_nRu–OH₂]¹⁺ (step 1), which will undergo ligand exchange with EtOH to produce [L_nRu–(EtOH)]¹⁺ (step 2). Complexes similar to [L_nRu–OH₂]¹⁺ exhibit pK_a values > 9,^{33,34} therefore [L_nRu–OH₂]¹⁺ is expected to be the predominant form at pH 7.4 rather than [L_nRu–OH]. Proton transfer from the EtOH ligand to the buffered solution will form [L_nRu–OEt] (step 3). Subsequent β–hydride elimination from the ethoxide ligand (via **TS4**) and acetaldehyde dissociation will afford the catalytically-active [L_nRu–H] intermediate (step 4). This Ru^{II}–hydride species will

then undergo $1e^-$ oxidation by $\text{ABTS}^{\bullet-}$ to yield a Ru^{III} -hydride cation (step 5). Deprotonation by buffer, oxidation by a second equivalent of $\text{ABTS}^{\bullet-}$, and subsequent coordination of ABTS^{2-} to Ru will regenerate $[\text{L}_n\text{Ru}-\text{A}_{\text{red}}]^{1-}$ and restart the cycle (dashed arrow, Scheme 3.2).

To elucidate the steps in the catalytic cycle, we conducted rate law studies by varying the initial concentration of $\text{ABTS}^{\bullet-}$, ABTS^{2-} , EtOH and **Ru1**, as well as solution pH and temperature.[‡] The initial rate (v_0) increased linearly with $[\text{ABTS}^{\bullet-}]_0$ (Fig. 3.1A), consistent with a reaction that is first-order in $[\text{ABTS}^{\bullet-}]$ [Eq. (1)]. This result also suggests that the 2 equiv. of $\text{ABTS}^{\bullet-}$ per turnover were reduced in separate steps (that is, step 5 followed by the dashed arrow in Scheme 3.2). A linear relationship was apparent in the plot of k_{obs} versus $1/[\text{ABTS}^{2-}]_0$ (Fig. 3.1B), which suggested that ABTS^{2-} dissociation from Ru is necessary for the reaction to proceed (consistent with step 1). An inverse relationship is observed between k_{obs} and $[\text{H}^+]$ (Fig. 3.1C), which is expected for H^+ transfer from the EtOH ligand to the (buffered) solution before or during the rate-determining step (consistent with step 3). In contrast, the values of k_{obs} increased as $[\text{EtOH}]_0$ increased (Fig. 3.1 D), which implied that ligand substitution of the Ru-aquo intermediate by EtOH is a necessary step in the catalytic cycle (consistent with step 2). The plot of k_{obs} vs. **[Ru1]** is also linear (Fig. 3.1E), which suggests that the predominant Ru-containing species in solution leading up to the rate-determining step is mononuclear. However, the possibility that multinuclear species were present in solution

[‡] For experimental details and data plots, see synthetic procedures and figures. For equation derivations, see “Derivation of General Rate Law”

in minor amounts cannot be definitively excluded. Overall, the rate of ABTS^{•-} reduction is proportionally dependent on [ABTS^{•-}]₀, [EtOH]₀, and [Ru1]₀, but inversely dependent on [ABTS²⁻]₀ and [H⁺]₀ [Eq. (2)], which is similar to the rate laws derived by Sigman et al.^{35,36} and Stahl et al.^{37,38} for the Pd- and Cu-catalyzed aerobic oxidation of alcohols.

$$v_0 = -\frac{d[\text{ABTS}^{\bullet-}]}{dt} = k_{\text{obs}}[\text{ABTS}^{\bullet-}] \quad (1)$$

$$-\frac{d[\text{ABTS}^{\bullet-}]}{dt} \propto \frac{[\text{EtOH}]_0[\text{Ru1}]_0}{[\text{ABTS}^{2-}]_0[\text{H}^+]_0} [\text{ABTS}^{\bullet-}] \quad (2)$$

Plotting $\ln(k_{\text{obs}}/T)$ versus $1/T$ (Fig. 3.1F) yields a positive ΔS^\ddagger value (28.9 ± 1.7 cal mol⁻¹ K⁻¹), which reveals that disorder increases during the rate-determining step and could arise from ligand dissociation or fragmentation (consistent with step 4). Positive ΔS^\ddagger values were also observed for Ru1-catalyzed ABTS^{•-} reduction with *i*-PrOH (32.2 ± 2.3 cal mol⁻¹ K⁻¹; Fig. 3.2), MeOH (11.4 ± 2.9 cal mol⁻¹ K⁻¹; Fig. 3.3), and ethylene glycol (32.8 ± 2.1 cal mol⁻¹ K⁻¹; Fig. 3.4),[§] which suggests that these alcohols undergo β-hydride elimination (via a TS4-like transition state). The presence of acetone did not inhibit the rate of ABTS^{•-} reduction with *i*-PrOH, therefore we conclude that β-hydride elimination with concomitant dissociation of acetone to generate [L_{*n*}Ru-H] (step 4) is irreversible. A negative ΔS^\ddagger value would be expected if step 5 (a bimolecular reaction) was rate-determining but v_0 would not vary with [ABTS^{•-}]₀ if step 4 was rate-determining.[§] Therefore, neither step 4 nor 5 is the unique rate-determining step for the Ru1-catalyzed reduction of ABTS^{•-} into ABTS²⁻. Instead, both steps are likely to exert a comparable influence on catalyst turnover. No ABTS^{•-} reduction occurred with *t*-BuOH,

[§] For experimental details and data plots, see synthetic procedures and figures. For equation derivations, see “Derivation of General Rate Law”

consistent with the inability of $[L_nRu-O-t-Bu]$ to undergo β -hydride elimination (step 4 is blocked). This lack of reactivity with t -BuOH also suggests that $ABTS^{\bullet-}$ reduction does not occur by oxidation of $[L_nRu-OR]$, because a t -BuO⁻ ligand would render the Ru center more electron-rich, and thus easier to oxidize, than either EtO⁻ or i -PrO⁻. A similar lack of reactivity with 1,2-dimethoxyethane was unsurprising, given that no $[L_nRu-OR]$ intermediate can be formed (step 3 is blocked).

To confirm that C-H and O-H bond breakage occurred before or during the rate-determining step, **Ru1**-catalyzed $ABTS^{\bullet-}$ reduction was performed with EtOH, EtOH- d_6 , i -PrOH, and i -PrOH- d_8 in protio and deutero PBS.** Significantly greater kinetic isotope effects (KIEs) were observed for the C-H/D and O-H/D isotopic substitutions in protio versus deutero PBS, consistent with the role of H₂O in multiple steps of the proposed mechanism: displacement of $ABTS^{2-}$ by H₂O (step 1), ligand exchange with EtOH (step 2), and H⁺ transfer from the EtOH ligand to the buffered solution (step 3). Substituting H₂O with D₂O can affect the equilibrium for each step, therefore it is unsurprising to observe a solvent KIE value for **Ru1**-catalyzed $ABTS^{\bullet-}$ reduction. The solvent KIE values calculated using EtOH and i -PrOH as terminal reductants (1.76 ± 0.39 and 1.72 ± 0.37 , respectively) were consistent with base-catalyzed H⁺ transfer (that is not rate-determining) and agreed with previously-reported values for catalytic redox reactions involving $ABTS^{\bullet-}$,³⁹ Ru-mediated alcohol oxidation,⁴⁰ and enzymes⁴¹⁻⁴³ in H₂O and D₂O solutions.

** For experimental details and data plots, see synthetic procedures and figures. For equation derivations, see "Derivation of General Rate Law"

The KIEs for the C–H/D and O–H/D isotopic substitutions in EtOH and *i*–PrOH were unambiguous for primary KIEs,⁴⁴ which demonstrated that O–H and C–H bond breakage both occurred before or during the rate–determining steps (consistent with step 3 and **TS4**). Notably, the C–H/D KIE values measured for **Ru1**–catalyzed ABTS^{•–} reduction with EtOH (2.88 ± 0.27) and *i*–PrOH (2.86 ± 0.31) were nearly identical to the corresponding C–H/D values reported by Bäckvall et al. (2.57 ± 0.26) and Casey et al. (2.86 ± 0.20) for respective H₂ transfer from 1-(4-fluorophenyl)ethanol and *i*–PrOH to other Ru complexes similar to **Ru1**.^{45,46} In contrast, the O–H/D KIE values observed for **Ru1**–catalyzed ABTS^{•–} reduction with EtOH (2.92 ± 0.51) and *i*–PrOH (4.18 ± 0.60) were significantly greater than the corresponding values reported by Bäckvall et al. (1.87 ± 0.17) and Casey et al. (1.79 ± 0.07). One possible cause for this difference is that [L_{*n*}Ru–H] is formed by H⁺ transfer from the O–H group in a separate step before β–hydride elimination (that is, step 3 followed by **TS4**), whereas the systems studied by Bäckvall et al. and Casey et al. involved H₂ transfer from the CH–OH moiety in a single, concerted step. The slightly smaller O–H/D KIE value measured for EtOH versus *i*–PrOH with **Ru1** could be due to the greater acidity of EtOH, which lowers the activation barrier to H⁺ dissociation and renders ionic O–H bond breakage less sensitive to H/D isotopic substitution.

We propose that [L_{*n*}Ru–H] undergoes 1e[–] oxidation by ABTS^{•–} to produce a cationic 17 e[–] Ru^{III} complex, [L_{*n*}Ru–H]¹⁺ (Scheme 3.3A). Deprotonation of [L_{*n*}Ru–H]¹⁺ followed by coordination of an additional ligand L' from solution (for example, H₂O) generates [L_{*n*}Ru–L']. This 19 e[–] Ru^I complex is oxidized by a second equivalent of

ABTS^{•-} and subsequently combines with the resultant ABTS²⁻ to produce [L_nRu-A_{red}]¹⁻ and complete the catalytic cycle. Direct observation of these subsequent steps was not possible because these reactions occurred after the rate-determining step. However, literature precedent (see below) suggests that the proposed 1 e⁻ oxidation of [L_nRu-H] and [L_nRu-L'] by ABTS^{•-}, as well as deprotonation of [L_nRu-H]¹⁺ at pH 7.4, will be thermodynamically favorable.

Tilset and Norton et al. have shown that **M1_{red}** undergoes 1 e⁻ oxidation at +0.060 V to form **M1_{ox}**, which has a pK_a 20–25 log units lower than **M1_{red}** (Scheme 3.3B).^{47,48} Oxidation of **M1_{red}** to **M1_{ox}** by ABTS^{•-} is expected to be thermodynamically feasible because the ABTS^{•-}/ABTS²⁻ redox couple occurs at +0.68 V.¹⁷ Tilset and Norton et al. also noted that the deprotonation of **M1_{ox}** will occur even if it is thermodynamically unfavorable by 4 pK_a units, provided that the 19 e⁻ Ru^I conjugate base (**M2_{red}**) is irreversibly oxidized. Even a weak oxidant, such as [Co(η⁵-C₅Me₅)₂]¹⁺ (E_{1/2} = -1.51 V),⁴⁹ could oxidize **M2_{red}** because the 1 e⁻ reduction of **M2_{ox}** to **M2_{red}** occurs at -2.02 V (Scheme 3.3C),⁴⁷ Oxidation of the 19 e⁻ Ru^I intermediate [L_nRu-L'] (analogous to **M2_{red}**) by ABTS^{•-} should be similarly favorable, and would likewise facilitate deprotonation of [L_nRu-H]¹⁺.

An anionic η⁵-C₅H₅ ligand is more electron-donating than a neutral η⁶-cymene, thus it is more relevant to compare the oxidation of [L_nRu-H] to other Ru complexes with η⁶-cymene ligands rather than **M1_{red}**. Unsurprisingly, the oxidation of **M3_{red}** (+0.16 V; Scheme 3.3D) occurred at a higher potential than **M1_{red}**,⁵⁰ but still well below the ABTS^{•-}/ABTS²⁻ redox couple. The difference in oxidation potentials between [L_nRu-H]

and **M3_{red}** is expected to be smaller than that between **M3_{red}** and **M1_{red}**, therefore ABTS^{•-} should be a sufficient strong oxidant to oxidize [L_nRu-H] and [L_nRu-L].

The complex most similar to [L_nRu-H]¹⁺ for which a pK_a value has been reported is **M4_{red}** (pK_a = 22.5 ± 0.1 in CH₃CN; Scheme 3.3 E)⁵¹ The pK_a for **M4_{red}** in H₂O can be estimated as 15.9 using Morris' method to estimate acidity in different solvents.⁵² However, **M4_{red}** is an 18 e⁻ Ru^{II} species and [L_nRu-H]¹⁺ is a 17 e⁻ Ru^{III} species, therefore a more relevant comparison would involve **M4_{ox}**, which has not been reported. Using the observations of Tilset and Norton et al. that 1e⁻ oxidation of an 18 e⁻ Ru^{II} complex typically lowers the pK_a by 20–25 log units,^{47,48} the pK_a for **M4_{ox}** in H₂O is estimated to be in the range of -4 to -9. Similarly dramatic decreases in pK_a have been observed following 1e⁻ oxidation of [RuH(η⁵-C₅H₅)(CO)(PPh₃)] (from 27–28 to 4–5)⁵³ and [WH(η⁵-C₅H₅)(CO)₂] (from 16 to -3).⁵⁴ Therefore, [L_nRu-H]¹⁺ should be sufficiently acidic to undergo complete deprotonation in pH 7.4 aqueous solution.

Synthetic Procedures

General synthetic considerations. N-(*p*-tolyl)benzimidazole and ABTS^{•-} were prepared as previously described.^{19,55} All other materials were of reagent quality and used as received. All solvents used were HPLC grade. ¹H and ¹³C {¹H} NMR spectra were recorded using a Bruker 500 MHz spectrometer. Chemical shifts δ (in ppm) for ¹H and ¹³C NMR are referenced to SiMe₄ using the residual protio-solvent as an internal standard. For ¹H NMR: CDCl₃, 7.26 ppm; DMSO-*d*₆, 2.50 ppm. For ¹³C NMR: CDCl₃, 77.16 ppm; DMSO-*d*₆, 39.52 ppm.⁵⁶ Coupling constants (*J*) are expressed in hertz (Hz). Infrared spectra were recorded with 4 cm⁻¹ resolution on a Shimadzu IRAffinity-1S

spectrometer equipped with a Pike Technologies MIRacle ATR (diamond) sampling accessory. Elemental analyses were performed at Atlantic Microlab, Inc. (Norcross, GA). All reactions were performed under an inert atmosphere under an N₂ atmosphere using standard Schlenk or glovebox techniques with the exclusion of light. All subsequent manipulations were performed under ambient conditions using standard benchtop techniques without the exclusion of light. When required, solvents were dried and deoxygenated using an Innovative Technologies solvent purification system, and then stored over molecular sieves (3 Å) in a drybox.

Synthesis of 2-(3-*p*-tolyl)-benzimidazol-1-ium-1-yl)-acetate hydrobromide [1H₂][Br]. Bromoacetic acid (503 mg, 3.62 mmol) and N-(*p*-tolyl)benzimidazole (500 mg, 2.41 mmol) were dissolved with 20 mL of toluene in a vial equipped with a stir bar. The vial was then sealed with a Teflon-lined cap and the clear, dark orange-brown solution was heated to 120 °C. After 16 h, a white precipitate had formed, at which point the reaction mixture was allowed to cool to room temperature and the precipitate was collected by filtration. The resulting solid was washed with minimum toluene followed by CH₂Cl₂ and was then dried *in vacuo* to afford 514 mg (1.48 mmol, 80% yield) of the desired product as a white powder. ¹H NMR (500 MHz, DMSO-*d*₆): δ = 13.95 (s, 1H), 10.18 (s, 1H), 8.18 (d, *J* = 8.0, 1H), 7.85 (d, *J* = 8.0, 1H), 7.78–7.73 (m, 4H), 7.57 (d, *J* = 8.0, 2H), 5.58 (s, 2H), 2.47 (s, 3H). ¹³C NMR (125 MHz, DMSO-*d*₆): δ 167.5, 143.3, 140.5, 131.5, 130.8, 130.6, 130.4, 127.4, 127.1, 124.8, 114.3, 113.4, 47.8, 20.7. FT-IR (KBr): 3421 (vw), 2953 (m), 2372 (m), 1718 (m), 1560, 1508 (m), 1381 (m), 1300 (m),

1225 (m), 822, 754, 742, 482 (vw) cm^{-1} . Anal. Calcd. for $\text{C}_{16}\text{H}_{16}\text{BrN}_2\text{O}_{2.5}$ ($[\text{1H}_2][\text{Br}] \cdot 0.5\text{H}_2\text{O}$): C, 53.95; H, 4.53; N, 7.86. Found: C, 54.01; H, 4.35; N, 7.79.

Synthesis of $[\text{Ag}(\mathbf{1})]_n$ (2**).** To a solution of $[\text{1H}_2][\text{Br}]$ (500 mg, 144 μmol) in anhydrous CH_2Cl_2 (50 mL) under nitrogen was added Ag_2O (502 mg, 217 μmol), and the resulting white suspension was allowed to stir at room temperature in the absence of light. After 24 h, the reaction mixture was filtered and the resulting grey solid was washed with CHCl_3 (10 mL). The filtrate solvent was removed under reduced pressure and the resulting solids were then dried *in vacuo* to afford 452 mg (121 μmol , 84% yield) of the desired product as an off-white powder. ^1H NMR (500 MHz, $\text{DMSO}-d_6$): δ 7.71 (d, $J = 8.5$, 1H), 7.58 (d, $J = 8.0$, 2H), 7.44–7.38 (m, 5H), 5.09 (s, 2H), 2.39 (s, 3H). ^{13}C NMR (125 MHz, $\text{DMSO}-d_6$): δ 138.6, 135.2, 134.1, 133.4, 130.3, 125.6, 124.1, 112.7, 111.6, 99.5, 79.1, 52.1, 20.6. FT-IR (KBr): 3416 (w), 1608, 1516, 1475, 1439, 1385, 1308, 1250, 1207 (m), 1109 (m), 1094 (m), 1018 (m), 835, 814, 746, 715, 482 (w) cm^{-1} . Anal. Calcd. for $\text{C}_{16}\text{H}_{14}\text{N}_2\text{O}_{2.5}\text{Ag}$ ($\mathbf{2} \cdot 0.5\text{H}_2\text{O}$): C, 50.36; H, 3.77; N, 7.49. Found: C, 50.23; H, 4.09; N, 7.05.

Synthesis of $[\text{RuCl}(\mathbf{1})(\eta^6\text{-cymene})]$ (Ru1**).** To a stirred solution of **2** (100 mg, 134 μmol) in anhydrous CH_2Cl_2 (10 mL) under nitrogen was added $[\text{RuCl}(\eta^6\text{-cymene})(\mu\text{-Cl})_2]$ (82 mg, 134 μmol) dissolved in 10 mL of CH_2Cl_2 . During the course of the addition, the mixture changed first to a clear red and then to a clear orange solution with concomitant formation of a white precipitate, and the reaction was allowed to stir at room temperature. After 24 h, the reaction mixture was filtered through a 0.2 μm PTFE filter (to remove AgCl) with the aid of CH_2Cl_2 (4 mL). The filtrate solvent was removed under

reduced pressure and the resulting solids were then dried *in vacuo* to afford 71.7 mg (134 μmol , 98%) of the desired product as a light red-orange powder. ^1H NMR (500 MHz, CDCl_3): δ 7.95 (d, $J = 8.0$, 1H), 7.49–7.44 (m, 2H), 7.41 (d, $J = 10.0$, 1H), 7.37–7.32 (m, 2H), 7.25–7.23 (m, 1H), 7.04 (d, $J = 8.5$, 1H), 5.43 (d, $J = 6.0$, 1H), 5.16 (d, $J = 6.5$, 1H), 5.04 (d, $J = 5.5$, 1H), 4.91 (d, $J = 17.0$, 1H), 4.71 (d, $J = 17$, 1H), 3.95 (d, $J = 5.5$, 1H), 2.60 (sep, 1H), 2.52 (s, 3H), 2.03 (s, 3H), 1.08 (d, $J = 7.5$, 6H). ^{13}C NMR (125 MHz, CDCl_3): δ 186.6, 169.2, 140.3, 136.8, 136.0, 134.0, 131.1, 131.0, 129.7, 127.4, 124.2, 124.1, 111.3, 110.2, 104.7, 103.2, 90.0, 89.4, 83.6, 80.3, 49.9, 30.9, 23.3, 22.0, 21.5, 19.0. FT-IR (KBr): 3429 (vw), 3055 (w), 2959 (m), 2872 (w), 2230 (w), 2100 (w), 1628 (vs), 1514, 1475, 1443, 1373 (w), 1356, 1300, 1244, 1207 (m), 1190 (m), 1014, 930, 862, 843 (m), 827 (m), 804, 748, 663 (m), 615, 555, 476 (m), 470, 424, 405 cm^{-1} . Anal. Calcd. for $\text{C}_{26}\text{H}_{27}\text{ClN}_2\text{O}_2\text{Ru}$: C, 58.26; H, 5.08; N, 5.23. Found: C, 57.98; H, 4.95; N, 5.17.

Spectroscopic and Kinetic Analysis Procedures

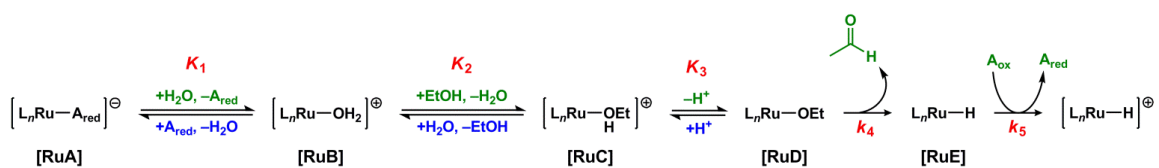
General spectroscopic considerations. UV–visible absorption spectra were acquired on a Varian Cary 50 Bio spectrometer equipped with a water-cooled Quantum Northwest TC-125 peltier temperature controller. All solution measurements were performed at 25.0 ± 0.1 $^\circ\text{C}$ in matched gas-tight quartz cuvettes (Precision Scientific) with 1 cm path lengths and 3.0 mL analyte solution volumes. Absorption spectra were acquired from 900 to 200 nm with a scanning speed of 300 nm min^{-1} and a resolution of 0.5 nm. Each kinetic analysis experiment (5 second intervals) was performed in quadruplicate on four different days. Stock solutions were prepared fresh daily and filtered (0.2 μm PTFE) immediately prior to use.

General procedure for kinetic analysis of ABTS^{•-} degradation by Ru1 in PBS. An aliquot of ABTS²⁻ (30 μ L from stock solution in H₂O) and an aliquot of ABTS^{•-} (35 μ L from stock solution in H₂O) were added to PBS (pH 7.4) and this working solution was allowed to equilibrate at 25 °C for 10 min. The absorbance spectrum of this working solution was acquired to confirm ABTS^{•-} concentration and the single wavelength kinetics program was initiated. After 20 s, the cuvette was removed from the spectrometer without stopping the kinetics program, an aliquot of **Ru1** (30 μ L from stock solution in CH₃CN) was added, the cuvette was covered and mixed via repeated inversion for 3 s, placed back in the spectrometer, and the kinetics program was allowed to continue. After 5 min, an aliquot of EtOH (30 μ L from stock solution in H₂O) was added, the cuvette was covered and mixed via repeated inversion for 3 s, placed back in the spectrometer, and the kinetics program was allowed to continue. After the kinetics program had completed, the absorbance spectrum was acquired to confirm the formation of ABTS²⁻ from the peak at 340 nm. Standard conditions (unless specified otherwise): 100 μ M ABTS²⁻, 50 μ M ABTS^{•-}, 5 μ M **Ru1**, 50 mM EtOH, PBS (pH 7.4), 25 °C; absorbance measured at 734 nm.

Rate law experiments. The concentrations of ABTS²⁻, ABTS^{•-}, **Ru1** and EtOH employed in the rate law experiments (cf. Figure 1) were obtained using this general kinetic procedure by varying stock solution concentration and holding aliquot volume constant. The H⁺ concentrations were obtained using PBS adjusted to different pH values before the addition of any aliquots. The temperatures were obtained by allowing the working solution to equilibrate at different temperatures before the addition of **Ru1**.

Kinetic isotope effect experiments. The general kinetic procedure and standard conditions were employed, with minor modifications. Experiments were performed in either protio PBS (pH 7.4) or deutero PBS (pD 7.4). Protio and deutero alcohol stock solutions were prepared in H₂O and D₂O (5.0 M) to obtain all 4 permutations of C–H and O–H isotopic substitution. For example, EtOH-*d*₆ stock solutions in H₂O and D₂O enabled comparison of *k*_{obs} with CD₃CD₂OH and CD₃CD₂OD.

Derivation of General Rate Law



Proposed Mechanism: Adapted from Scheme 2 in manuscript. The intermediates [L_{*n*}Ru–A_{red}]^{1–}, [L_{*n*}Ru–OH₂]¹⁺, [L_{*n*}Ru–(EtOH)]¹⁺, [L_{*n*}Ru–OEt] and [L_{*n*}Ru–A_{red}]^{1–} have been abbreviated in the following equations as [RuA], [RuB], [RuC], [RuD] and [RuE], respectively, to enhance clarity.

- (1) Based on the observation that the rate is directly proportional to the concentration of ABTS^{•–} (cf. Fig.3.1 A) the underlying reaction is the oxidation of [L_{*n*}Ru–H] by ABTS^{•–}:

$$\text{rate} = -\frac{d[\mathbf{A}_{\text{ox}}]}{dt} = k_5[\mathbf{A}_{\text{ox}}][\mathbf{RuE}]$$

- (2) The sum of the concentrations of the Ru-containing species leading up to the rate-determining step is equal to the total concentration of Ru1 added at the beginning of the experiment:

$$[\mathbf{Ru1}]_0 = [\mathbf{RuA}] + [\mathbf{RuB}] + [\mathbf{RuC}] + [\mathbf{RuD}] + [\mathbf{RuE}]$$

(3) Assuming that steps 1-3 achieve equilibrium rapidly (with respect to turnover):

$$K_1 = \frac{[\text{RuB}][\text{A}_{\text{red}}]}{[\text{RuA}]} \quad K_2 = \frac{[\text{RuC}]}{[\text{RuB}][\text{EtOH}]} \quad K_3 = \frac{[\text{RuD}][\text{H}^+]}{[\text{RuC}]}$$

(4) Assuming the system achieves steady-state rapidly (with respect to turnover):

$$\frac{d[\text{RuE}]}{dt} = k_4[\text{RuD}] - k_5[\text{A}_{\text{ox}}][\text{RuE}]$$

$$[\text{RuD}] = \frac{k_5[\text{A}_{\text{ox}}]}{k_4} [\text{RuE}]$$

(5) Solving (2) for $[\text{RuE}]$ using the relationships established in (3) and (4):

$$[\text{Ru1}]_0 = \left(\frac{[\text{A}_{\text{red}}]}{K_1} + 1 \right) [\text{RuB}] + [\text{RuC}] + [\text{RuD}] + [\text{RuE}]$$

$$[\text{Ru1}]_0 = \left(\frac{[\text{A}_{\text{red}}]}{K_1 K_2 [\text{EtOH}]} + \frac{1}{K_2 [\text{EtOH}]} + 1 \right) [\text{RuC}] + [\text{RuD}] + [\text{RuE}]$$

$$[\text{Ru1}]_0 = \left(\frac{[\text{A}_{\text{red}}][\text{H}^+]}{K_1 K_2 K_3 [\text{EtOH}]} + \frac{[\text{H}^+]}{K_2 K_3 [\text{EtOH}]} + \frac{[\text{H}^+]}{K_3} + 1 \right) [\text{RuD}] + [\text{RuE}]$$

(6) For convenience, quotients are substituted with terms Q_1 , Q_2 and Q_3 :

$$Q_1 = \frac{K_1}{[\text{A}_{\text{red}}]} \quad Q_2 = K_2 [\text{EtOH}] \quad Q_3 = \frac{K_3}{[\text{H}^+]}$$

$$\frac{1}{Q_1} = \frac{[\text{A}_{\text{red}}]}{K_1} \quad \frac{1}{Q_2} = \frac{1}{K_2 [\text{EtOH}]} \quad \frac{1}{Q_3} = \frac{[\text{H}^+]}{K_3}$$

(7) Continuing from (5):

$$[\text{Ru1}]_0 = \left(\frac{1}{Q_1 Q_2 Q_3} + \frac{1}{Q_2 Q_3} + \frac{1}{Q_3} + 1 \right) [\text{RuD}] + [\text{RuE}]$$

$$[\text{Ru1}]_0 = \left(\frac{1 + Q_1 + Q_1 Q_2 + Q_1 Q_2 Q_3}{Q_1 Q_2 Q_3} \right) \left(\frac{k_5 [\text{A}_{\text{ox}}]}{k_4} \right) [\text{RuE}] + [\text{RuE}]$$

$$[\text{Ru1}]_0 = \left\{ \frac{(1 + Q_1 + Q_1 Q_2)(k_5 [\text{A}_{\text{ox}}]) + (k_5 [\text{A}_{\text{ox}}] + k_4)(Q_1 Q_2 Q_3)}{k_4 Q_1 Q_2 Q_3} \right\} [\text{RuE}]$$

$$[\mathbf{RuE}] = \left\{ \frac{k_4 Q_1 Q_2 Q_3}{(1 + Q_1\{1 + Q_2\})(k_5[\mathbf{A}_{\text{ox}}]) + (k_5[\mathbf{A}_{\text{ox}}] + k_4)(Q_1 Q_2 Q_3)} \right\} [\mathbf{Ru1}]_0$$

(8) Plugging this result into (1) gives the general rate law:

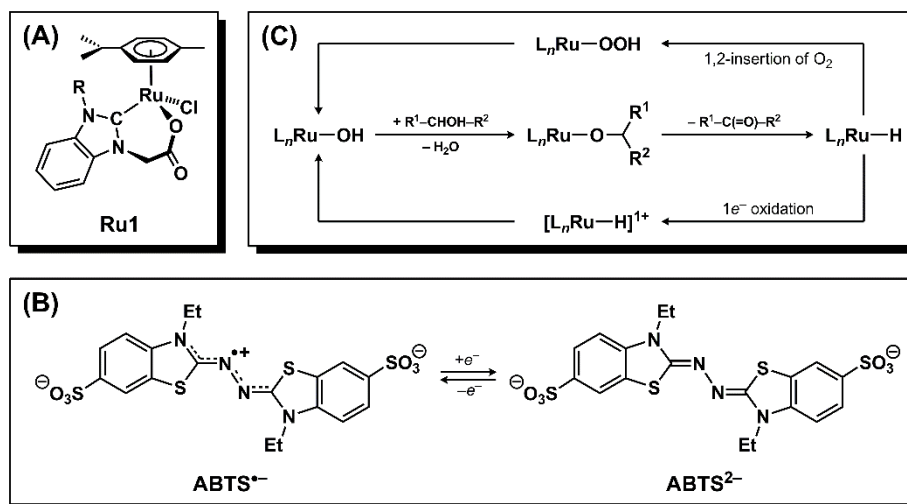
$$\text{rate} = \frac{k_4 k_5 Q_1 Q_2 Q_3 [\mathbf{Ru1}]_0 [\mathbf{A}_{\text{ox}}]}{(1 + Q_1\{1 + Q_2\})(k_5[\mathbf{A}_{\text{ox}}]) + (k_5[\mathbf{A}_{\text{ox}}] + k_4)(Q_1 Q_2 Q_3)}$$

CONCLUSIONS

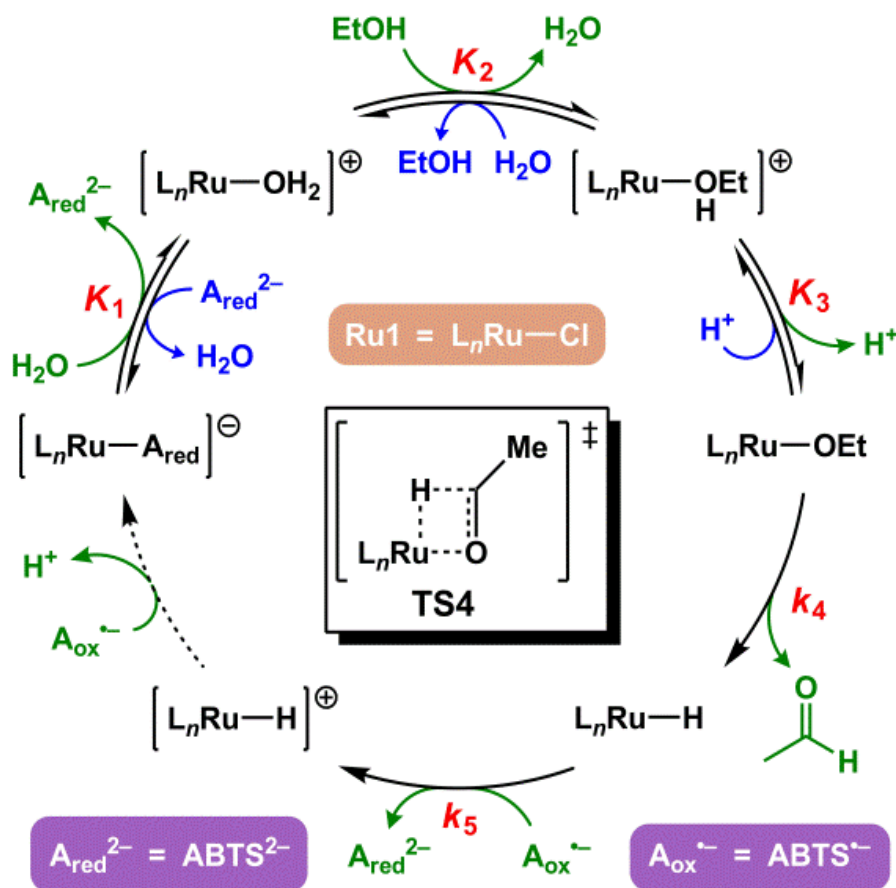
In summary, mechanistic data for the catalytic reduction of $\text{ABTS}^{\bullet-}$ into ABTS^{2-} by **Ru1**, with EtOH as the terminal reductant, suggests that β -hydride elimination from a Ru-alkoxide species generates the catalytically active Ru-H intermediate. In light of previous reports indicating that, 1) Ru-H intermediates can achieve anticancer properties by exerting catalytic pro-oxidant effects,²³⁻²⁵ and 2) some Mn porphyrin-based complexes can produce catalytic antioxidant or pro-oxidant effects in different environments with distinct therapeutic uses,⁴⁻¹⁵ we believe that new therapeutic strategies and applications can potentially be developed that utilize the catalytic radical reducing ability we have demonstrated for complexes that form Ru-hydride intermediates.

Acknowledgements

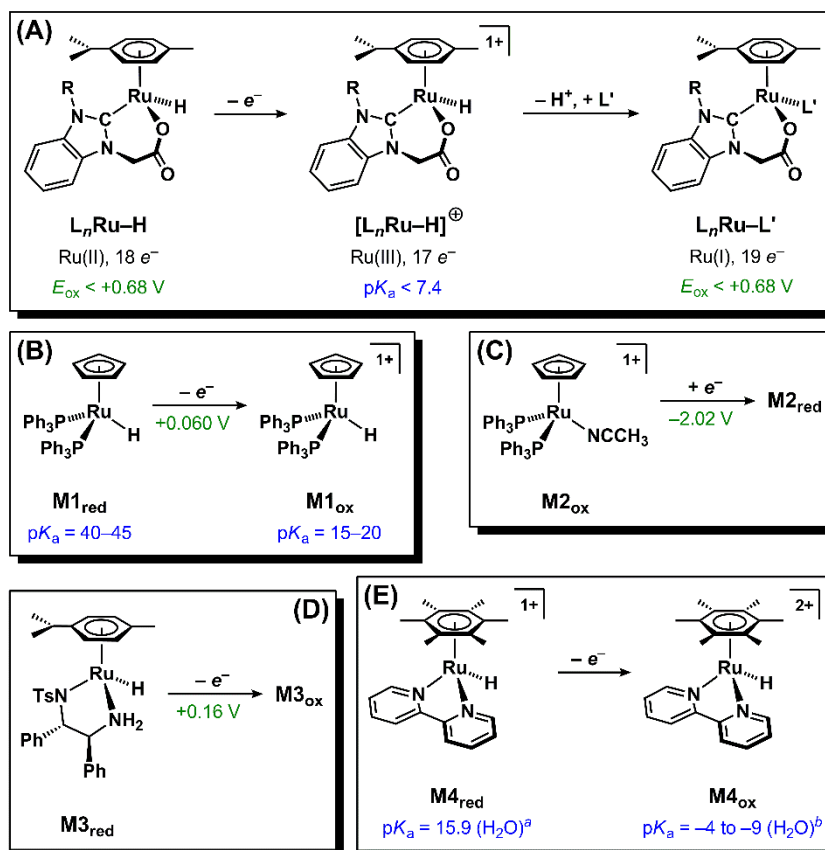
This work was supported by the National Science Foundation (DMR-1555224). We thank Dr. A. Mangalum for prior work with **Ru1** and helpful discussions.



Scheme 3.1. (A) Structure of **Ru1**. (B) 1 e⁻ interconversion between ABTS^{•+} and ABTS²⁺. (C) Formation of Ru–hydride in catalytic aerobic alcohol oxidation (top) and ABTS^{•+} reduction (bottom).



Scheme 3.2. Proposed mechanism for **Ru1**-catalyzed reduction of $\text{ABTS}^{\bullet-}$ to ABTS^{2-} , with EtOH as the terminal reductant. Forward reactions (clockwise green), reverse reactions (counter-clockwise, blue). Each K_n or k_n corresponds to the equilibrium or rate constant, respectively, for the forward reaction in step “n” (step 1 has equilibrium constant K_1 , and so forth). The transition-state structure for step 4 is shown as **TS4** in the center. The dashed arrow includes multiple transformations that occur after the rate-determining steps. The spectator ligand set “ L_n ” comprises the η^6 -cymene and κ^2 -(C,O)-benzimidazolylidene-carboxylate ligands, but their hapticity/denticity may decrease to accommodate additional ligands.



Scheme 3.3. A) Proposed completion of the catalytic cycle starting from $[L_nRu-H]$, and B–E) literature precedent. Unless specified otherwise, pK_a values and redox potentials were measured in CH_3CN solutions. ^a pK_a value for $M4_{red}$ in H_2O estimated using ref 52. ^b pK_a value for $M4_{ox}$ estimated using ref. 47, 48.

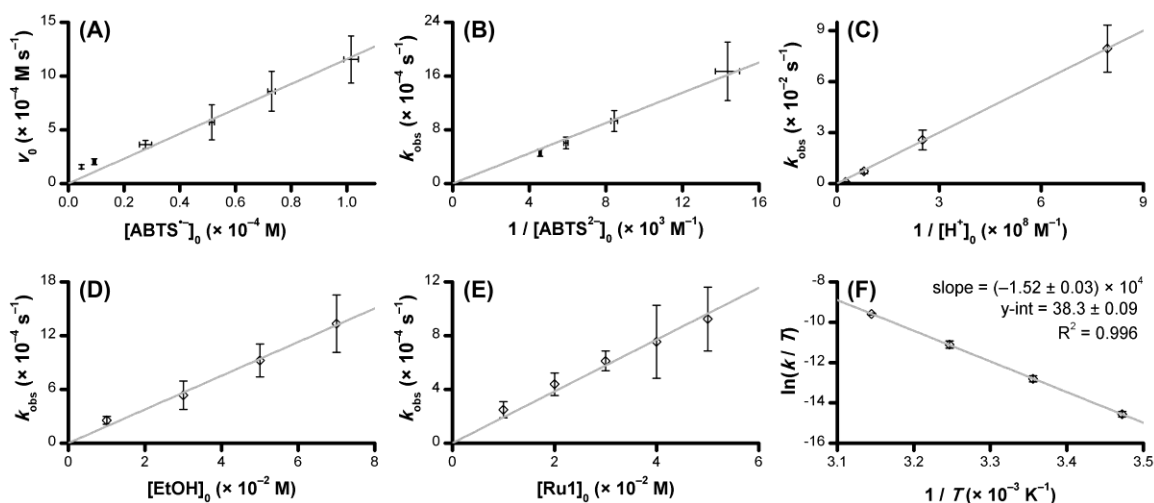


Fig. 3.1. (A) Dependence of initial rate (v_0) of **Ru1**-catalyzed $\text{ABTS}^{\bullet-}$ reduction on $[\text{ABTS}^{\bullet-}]_0 = 5, 10, 25, 50, 75$ or $100 \mu\text{M}$. Dependence of observed rate constant (k_{obs}) for **Ru1**-catalyzed $\text{ABTS}^{\bullet-}$ reduction on: (B) $[\text{ABTS}^{2-}]_0 = 50, 100, 150$ or $200 \mu\text{M}$; (C) $\text{pH} = 7.4, 7.9, 8.4$ or 8.9 ; (D) $[\text{EtOH}]_0 = 10, 30, 50$ or 70 mM ; (E) $[\text{Ru1}]_0 = 1, 2, 3, 4$ or $5 \mu\text{M}$; (F) $T = 15, 25, 35$ or $45 \text{ }^\circ\text{C}$. Each data point (\diamond) is the average of 4 experiments. Conditions: $[\text{Ru1}]_0 = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[\text{ABTS}^{2-}]_0 = 100 \mu\text{M}$, $[\text{EtOH}]_0 = 50 \text{ mM}$, PBS ($\text{pH} 7.4$), $25 \text{ }^\circ\text{C}$; absorbance measured at 734 nm .

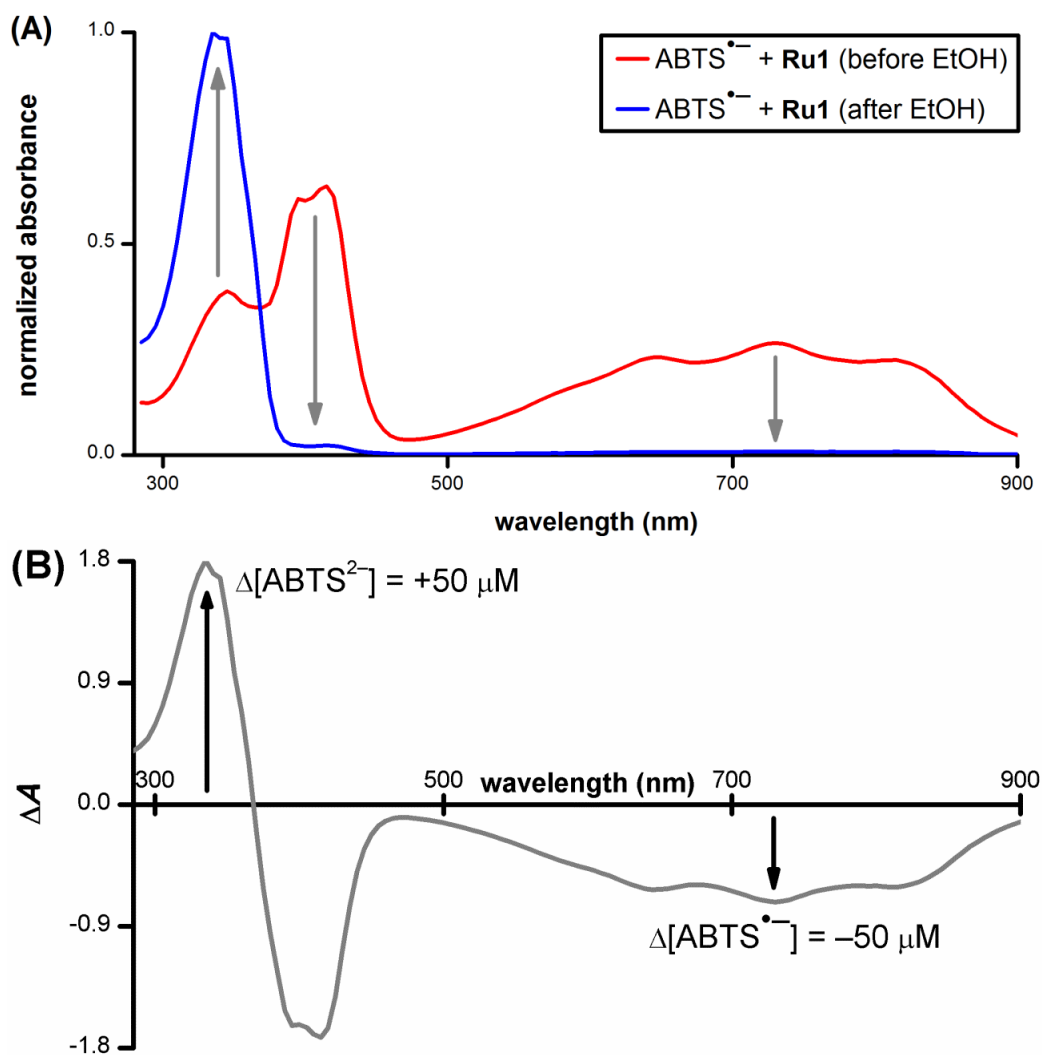


Fig. 3.2. (A) Normalized absorption spectrum of 50 μM $\text{ABTS}^{\bullet-}$ and 5 μM **Ru1** in PBS (red) and after 50 mM EtOH was added and 30 minutes had elapsed (blue). (B) Difference spectrum obtained by subtracting red trace from blue trace, which showed that the 50 μM decrease in $\text{ABTS}^{\bullet-}$ concentration was accompanied by a 50 μM increase in ABTS^{2-} concentration. Conditions: $[\text{Ru1}]_0 = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[\text{ABTS}^{2-}]_0 = 100 \mu\text{M}$, $[\text{EtOH}]_0 = 50 \text{ mM}$, PBS (pH 7.4), 25 $^\circ\text{C}$; absorbance measured from 285–900 nm

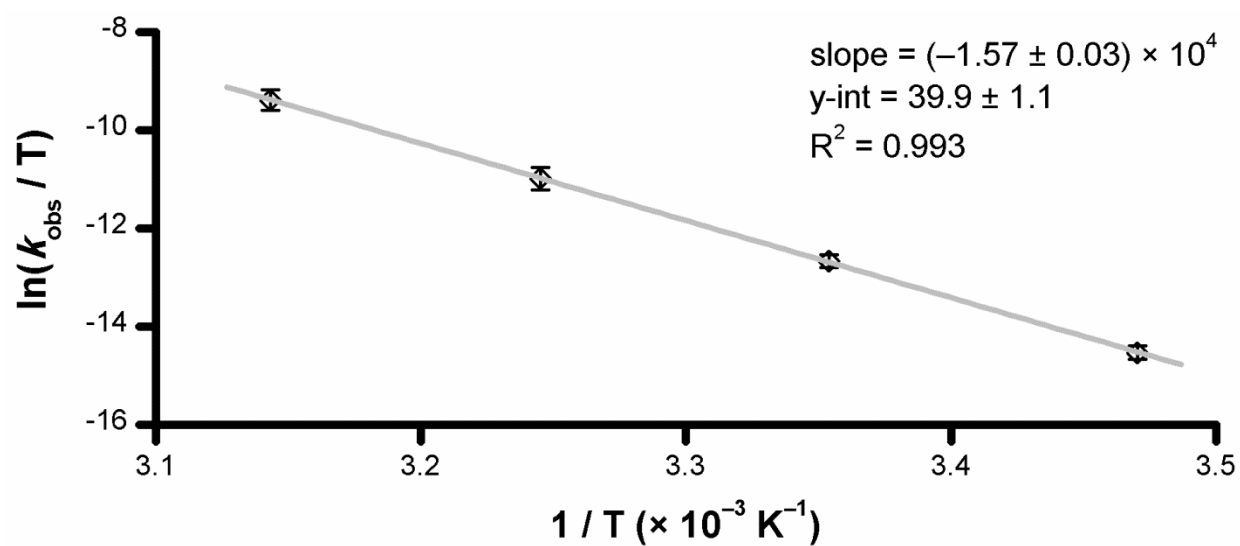


Fig. 3.3. Eyring plot for **Ru1**-catalyzed $\text{ABTS}^{\bullet-}$ reduction with *i*-PrOH ($T = 15, 25, 35$ or 45 °C). Each data point (\diamond) is the average of 4 experiments. Conditions: $[\text{Ru1}]_0 = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[\text{ABTS}^{2-}]_0 = 100 \mu\text{M}$, $[i\text{-PrOH}]_0 = 50 \text{ mM}$, PBS (pH 7.4); absorbance measured at 734 nm.

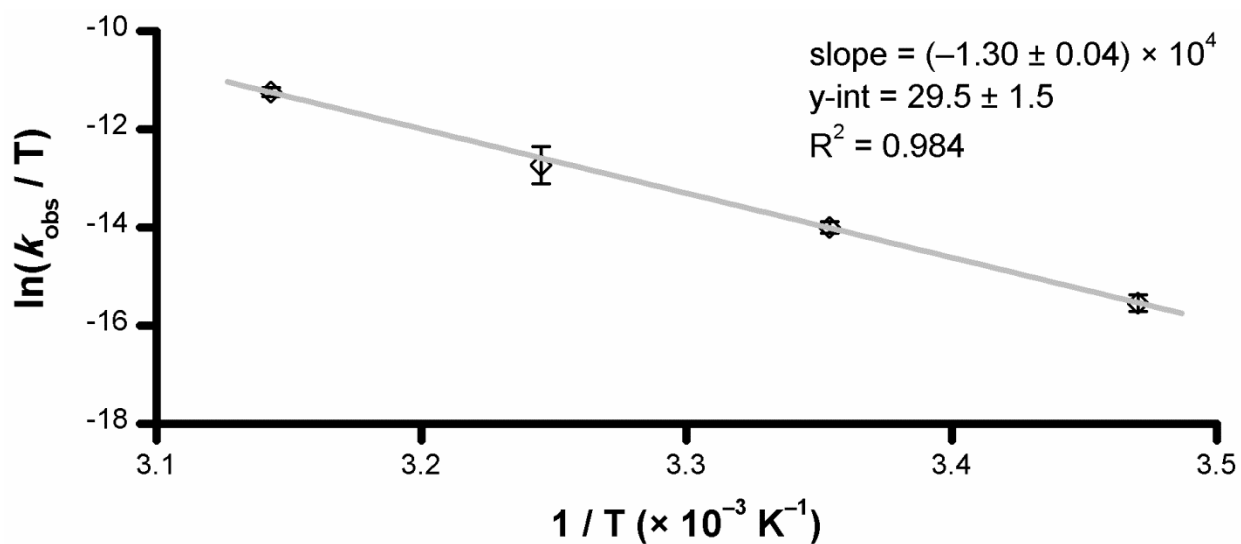


Fig. 3.4. Eyring plot for **Ru1**-catalyzed $\text{ABTS}^{\bullet-}$ reduction with MeOH ($T = 15, 25, 35$ or 45 °C). Each data point (\diamond) is the average of 4 experiments. Conditions: $[\text{Ru1}]_0 = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[\text{ABTS}^{2-}]_0 = 100 \mu\text{M}$, $[\text{MeOH}]_0 = 50 \text{ mM}$, PBS (pH 7.4); absorbance measured at 734 nm.

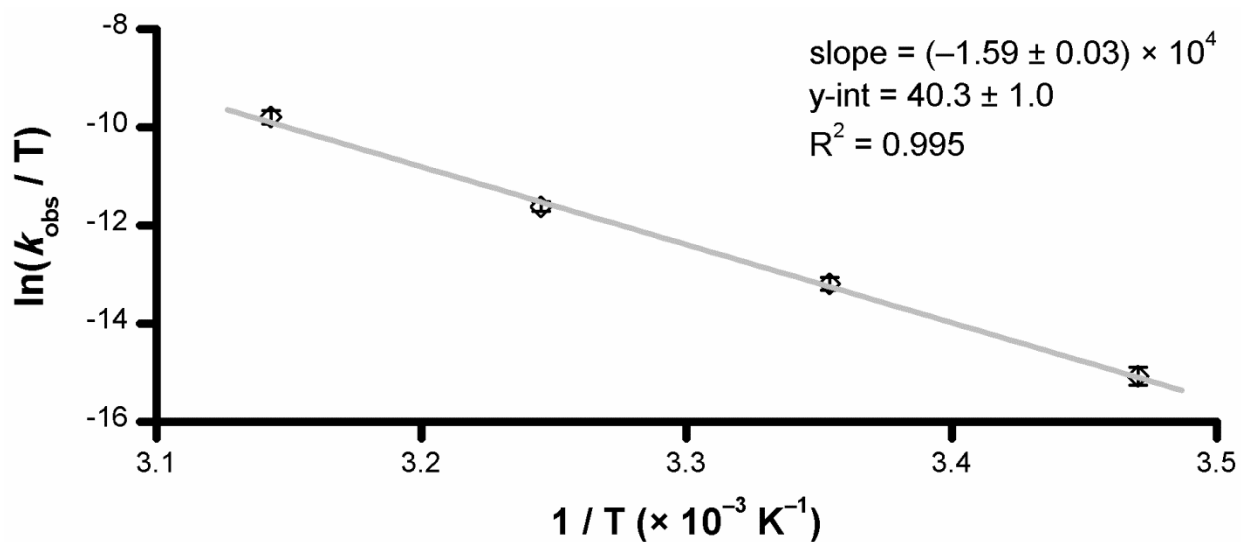


Fig. 3.5. Eyring plot for **Ru1**-catalyzed $\text{ABTS}^{\bullet-}$ reduction with ethylene glycol (EG) ($T = 15, 25, 35$ or 45 °C). Each data point (\diamond) is the average of 4 experiments. Conditions: $[\text{Ru1}]_0 = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[\text{ABTS}^{2-}]_0 = 100 \mu\text{M}$, $[\text{EG}]_0 = 50 \text{ mM}$, PBS (pH 7.4); absorbance measured at 734 nm.

Table 3.1. Ratio of observed rate constants (k_{obs}) for **Ru1**-catalyzed ABTS^{•-} reduction with protio and deuterio EtOH and *i*-PrOH measured in PBS-buffered H₂O and D₂O solutions

	X = H	X = D
CH ₃ CH ₂ OX / CD ₃ CD ₂ OX (C–H/C–D for EtOH)	5.05 ± 1.02 ^a	2.88 ± 0.27 ^b
(CH ₃) ₂ CHOX / (CD ₃) ₂ CDOX (C–H/C–D for <i>i</i> -PrOH)	4.93 ± 0.93 ^a	2.86 ± 0.31 ^b
CX ₃ CX ₂ OH / CX ₃ CX ₂ OD (O–H/O–D for EtOH)	5.13 ± 0.71 ^c	2.92 ± 0.51 ^d
(CX ₃) ₂ CXOH / (CX ₃) ₂ CXOD (O–H/O–D for <i>i</i> -PrOH)	7.21 ± 1.18 ^c	4.18 ± 0.60 ^d

^a Ratio of k_{obs} measured for protio alcohol ÷ deuterio alcohol in protio PBS (O–H groups conserved). ^b Ratio of k_{obs} measured for protio alcohol ÷ deuterio alcohol in deuterio PBS (O–D groups conserved). ^c Ratio of k_{obs} measured for protio alcohol in protio PBS ÷ protio alcohol in deuterio PBS. ^d Ratio of k_{obs} measured for deuterio alcohol in protio PBS ÷ deuterio alcohol in deuterio PBS. Values reported as the average ± standard deviation of 4 independent experiments performed on 4 different days. Alcohols were added from 5.0 M stock solutions in H₂O or D₂O (e.g., CH₃CH₂OD was added from 5.0 M EtOH stock solution in D₂O). Conditions: [**Ru1**]₀ = 5 μM, [ABTS^{•-}]₀ = 50 μM, [ABTS²⁻]₀ = 100 μM, [alcohol]₀ = 50 mM, PBS in H₂O (pH 7.4) or PBS in D₂O (pD 7.4), *T* = 25 °C.

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CHAPTER FOUR

NAD⁺ AS A HYDRIDE DONOR AND REDUCTANT

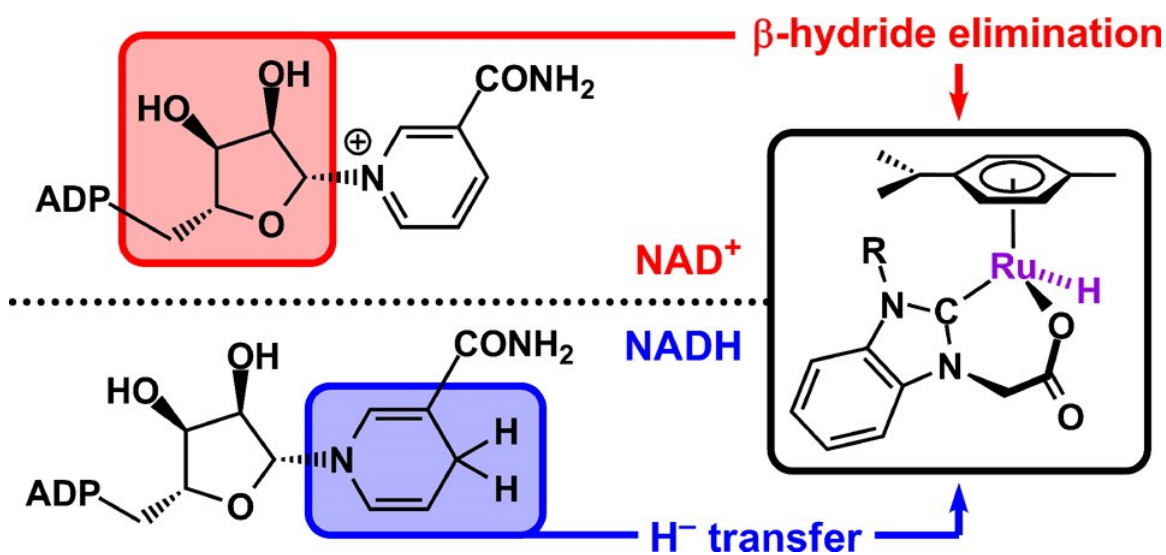


Figure 4. An organoruthenium complex reduced radicals catalytically using NAD⁺, a nature's oxidizing agent, as a hydride donor.

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ABSTRACT

Reduced nicotinamide adenine dinucleotide (NADH) can generate a ruthenium–hydride intermediate that catalyzes the reduction of O₂ to H₂O₂, which endows it with potent anticancer properties. A catalyst that could access a Ru–H intermediate using oxidized nicotinamide adenine dinucleotide (NAD⁺) as the H[−] source, however, could draw upon a supply of reducing equivalents 1000-fold more abundant than NADH, which would enable significantly greater H₂O₂ production. Herein, it is demonstrated, using the reduction of ABTS^{•−} to ABTS^{2−}, that NAD⁺ can function as a reductant. Mechanistic evidence is presented that suggests a Ru–H intermediate is formed via β-hydride elimination from a ribose subunit in NAD⁺. The insight gained from the heretofore unknown ability of NAD⁺ to function as a reductant and H[−] donor may lead to undiscovered biological carbohydrate oxidation pathways and new chemotherapeutic strategies.

INTRODUCTION

Redox reactions provide the chemical motive force essential for all forms of life.^{1,2} Reduced nicotinamide adenine dinucleotide (NADH) supplies two e^- to mitochondrial electron transport by donating H^- from its hydroxyridine moiety (Scheme 4.1, blue box) to flavin mononucleotide.^{3,4} Conversely, oxidized nicotinamide adenine dinucleotide (NAD^+) accepts two e^- from glyceraldehyde-3-phosphate or lactate dehydrogenase by accepting H^- into its pyridinium moiety (Scheme 4.1, purple box).^{5,6} The H^- donating ability of NADH has been harnessed for catalytic applications ranging from the reduction of O_2 to cytotoxic H_2O_2 in cancer cells⁷⁻⁹ to petroleum-free H_2 production¹⁰ to alcohol deracemization.¹¹

Free NAD^+ is 640–1100 times more abundant in cells than free NADH;¹²⁻¹⁵ therefore, a catalyst that could utilize NAD^+ would have access to a significantly greater H^-/e^- supply than NADH. Because catalytic carbohydrate oxidation can be performed by enzymatic¹⁶ and transition metal-based systems,^{17,18} we reasoned that oxidation of a ribose subunit (Scheme 4.1, red boxes) could enable NAD^+ to function as a reductant. Catalytic oxidation of ribose has been achieved by a Ru complex to afford ribonolactone with concomitant H_2 transfer to an alkene.¹⁹ We therefore hypothesized that (1) a ribose subunit in NAD^+ could similarly undergo oxidation by a Ru complex via H^- transfer to the metal center and (2) the resulting Ru–H species would exhibit catalytic reduction activity. Herein, we report the first instance of H^- donation via β -hydride elimination from a ribose subunit of NAD^+ , which enables NAD^+ to function as a reductant.

RESULTS AND DISCUSSION

To probe for Ru–H formation, the conversion of 2,2'-azino-bis(3-ethylbenzothiazoline-6-sulfonate) radical monoanion ($\text{ABTS}^{\bullet-}$, Scheme 4.2.A) to ABTS^{2-} was selected as a spectroscopically more convenient reduction reaction than the conversion of O_2 to H_2O_2 , given that $\text{ABTS}^{\bullet-}$ consumption can be quantified at significantly lower concentrations and longer wavelengths than H_2O_2 production.^{††,‡‡ 20,21} Furthermore, the reduction of $\text{ABTS}^{\bullet-}$ to ABTS^{2-} (0.68 V vs NHE)²² occurs at nearly the same potential as the reduction of O_2 to H_2O_2 (0.70 V vs NHE).²³ Therefore, from a thermodynamic perspective, the reactivity of Ru–H with $\text{ABTS}^{\bullet-}$ can provide insight into its reactivity with O_2 .

We recently reported the catalytic reduction of $\text{ABTS}^{\bullet-}$ to ABTS^{2-} in aqueous solution using biologically relevant alcohols as terminal reductants, including arabinose, a diastereomer of ribose (Scheme 4.2.B).²⁴ Subsequent kinetic studies elucidated the mechanism: in aqueous solution, **Ru1** converts to Ru–aquocomplex **I_A**, followed by ligand exchange with a nontertiary alcohol ($\text{R}^1\text{–CHOH–R}^2$) and deprotonation to afford Ru–alkoxide species **I_B**, which then undergoes β -hydride elimination (**TS_{B/C}**) to generate the catalytically active Ru–H intermediate **I_C** that reduces $\text{ABTS}^{\bullet-}$.²⁵ We reasoned that a ribose subunit of NAD^+ could likewise undergo β -hydride elimination to produce **I_C**, whose presence could be inferred from the reduction of $\text{ABTS}^{\bullet-}$ to ABTS^{2-} .

^{††} A 3.3 μM decrease in $[\text{ABTS}^{\bullet-}]$ would produce a measurable change in absorbance (0.050), but a comparable absorbance change would require a 1.1 mM increase in $[\text{H}_2\text{O}_2]$. If a catalyst concentration of 5 μM is used, it would be possible to observe the reduction of less than 1 equiv of $\text{ABTS}^{\bullet-}$, but O_2 reduction would not be observable until more than 200 equiv of H_2O_2 had been produced.

^{‡‡} In aqueous buffer, $\epsilon = 15,000 \text{ M}^{-1} \text{ cm}^{-1}$ at $\lambda = 734 \text{ nm}$ for $\text{ABTS}^{\bullet-}$ vs $\epsilon = 43.6 \text{ M}^{-1} \text{ cm}^{-1}$ at $\lambda = 240 \text{ nm}$ for H_2O_2

Addition of 5 μM **Ru1** to 50 μM $\text{ABTS}^{\bullet-}$ in phosphate buffered saline (PBS, pH 7.4) followed by the addition of 25 mM NAD^+ produced an 87% decrease in radical absorbance within 30 min (Fig. 4.1A, red line) that was 100% complete within 45 min. The UV/vis spectrum after 45 min confirmed a 1:1 correlation between $\text{ABTS}^{\bullet-}$ consumed and ABTS^{2-} produced (Fig. 4.2). Attempts to characterize the NAD^+ oxidation product were unsuccessful, due to the low concentration constraints of the $\text{ABTS}^{\bullet-}$ reduction reaction, but calorimetric and computational studies by others suggest that dehydrogenation of the $-\text{CHOH}-$ moiety at the ribose 2'-position would be thermodynamically the most favorable.^{26,27} No radical reduction occurred in the presence of 5 μM **Ru1** alone (Fig. 4.1A, green line), which revealed that **Ru1** by itself could not reduce $\text{ABTS}^{\bullet-}$. Similarly, no $\text{ABTS}^{\bullet-}$ reduction was observed in the absence of **Ru1**, even with NAD^+ concentrations as high as 50 mM, which demonstrated that NAD^+ by itself could not reduce $\text{ABTS}^{\bullet-}$. However, addition of 5 μM NADH produced a rapid (within mixing time) 18% decrease in radical absorbance (Fig. 4. 1A, blue line), consistent with NADH functioning as a two e^- reductant. After the initial decrease, no additional $\text{ABTS}^{\bullet-}$ reduction was observed beyond normal thermal decay.

To determine if **Ru1** remained catalytically active after the reduction of 10 equiv of $\text{ABTS}^{\bullet-}$, two subsequent aliquots of 50 μM $\text{ABTS}^{\bullet-}$ were added (*) and $[\text{ABTS}^{\bullet-}]$ decreased to zero each time (Fig. 4.1B). The time necessary for complete $\text{ABTS}^{\bullet-}$ reduction increased with each successive aliquot due to the fact that ABTS^{2-} inhibits **Ru1**-catalyzed $\text{ABTS}^{\bullet-}$ reduction.²⁵ After the initial decrease produced by NADH , however, addition of 10 μM $\text{ABTS}^{\bullet-}$ aliquots (*) only increased absorbance proportional

to the [ABTS^{•-}] in each aliquot (Fig. 4.1C), which indicated that the reducing ability of NADH had been exhausted. Treatment of this solution containing 70 μM ABTS^{•-} with 5 μM **Ru1** and 25 mM NAD⁺ (§), produced complete ABTS^{•-} reduction within 1 h.

The ability of **Ru1** to catalyze ABTS^{•-} reduction was assayed with the individual components of NAD⁺: nicotinamide, adenine, and ribose. No ABTS^{•-} reduction occurred upon treatment of 50 μM ABTS^{•-} and 5 μM **Ru1** with either 25 mM nicotinamide or 1.0 mM adenine, which indicated that neither component afforded NAD⁺ its terminal reductant ability. In contrast, the addition of 25 mM D-ribose or 1.0 mM D-ribose phosphate produced complete ABTS^{•-} reduction within 20 min (Figs. 4.3–4.4). The faster reactivity with D-ribose phosphate is consistent with the higher affinity of cationic **I_A** for anionic D-ribose phosphate than neutral D-ribose. Collectively, these results demonstrated that the terminal reductant function of NAD⁺ is derived from its ribose subunits.

The kinetics of **Ru1**-catalyzed ABTS^{•-} reduction with NAD⁺ were analyzed for consistency with the mechanism in Scheme 4.2. Increasing the solution pH led to faster ABTS^{•-} reduction, with no reduction observed in pure H₂O (Fig. 4.5), which indicated that H⁺ dissociation was necessary and was consistent with the conversion of **I_A** to **I_B**. Varying the reaction temperature revealed $\Delta S^\ddagger = 17.1 \pm 4.9 \text{ cal mol}^{-1} \text{ K}^{-1}$ (Fig. 4.6), which demonstrated that disorder was increasing during the rate-determining step and suggested ligand fragmentation and dissociation (i.e., **TS_{B/C}** and R¹–C(=O)–R² elimination, respectively) was occurring. The ΔS^\ddagger value observed with NAD⁺ also fell within the range of values measured for **Ru1**-catalyzed ABTS^{•-} reduction with other non-

tertiary alcohols ($\Delta S^\ddagger = 11.4\text{--}32.8 \text{ cal mol}^{-1} \text{ K}^{-1}$).²⁵ Collectively, these results were consistent with the formation of **I_C** via β -hydride elimination from a ribose subunit coordinated to Ru and dissociation of the oxidized NAD⁺.

The observed rate constant (k_{obs}) for **Ru1**-catalyzed ABTS^{•-} reduction with NAD⁺ was 2.53-fold lower in deuterio PBS (pD 7.4) than in proteo PBS. This ABTS^{•-} reduction reaction exhibits a solvent kinetic isotope effect (KIE) of 1.74, which reflects the role of H₂O as an H⁺ acceptor in the conversion of **I_A** to **I_B** and **I_C** back to **I_A**.²⁵ Dividing the proteo/deutero k_{obs} ratio of 2.53 by 1.74 yielded the O–H/D KIE value of 1.45 for NAD⁺. Breakage of an O–H bond in a ribose subunit of NAD⁺ is essential for the formation of **I_B**, whereby H/D substitution causes the activation barrier to increase and the k_{obs} for ABTS^{•-} reduction to decrease. In our previous mechanistic study, the smaller O–H/D KIE value for EtOH (2.92) compared to *i*-PrOH (4.18) reflected the lower pK_a of EtOH (15.9 vs 16.5 for *i*-PrOH).²⁵ Increasing the acidity of the O–H group will increase the O^{δ-}–H^{δ+} bond polarization, which will lower the activation barrier to H⁺ dissociation and thus render the O–H bond less sensitive to H/D isotopic substitution. The substantially lower O–H/D KIE value for NAD⁺ compared to EtOH and *i*-PrOH was thus consistent with the substantially greater acidity of ribose ($pK_a = 11.8$).²⁸

We next sought to demonstrate that NAD⁺ could serve as a reductant under conditions in which the biological supply of NADH had been exhausted. Treatment of a 50 μM ABTS^{•-} solution (Fig. 4.7, *i*) with 18 μM NADH caused a rapid (within mixing time) decrease in radical absorbance corresponding to the reduction of 34 μM ABTS^{•-} (Fig. 4.7, *ii*). This ABTS^{•-}/NADH reaction stoichiometry of 1.9 was consistent with

NADH functioning as a two e^- reductant. Importantly, 16 μM $\text{ABTS}^{\bullet-}$ was not reduced, and no further decreases in $[\text{ABTS}^{\bullet-}]$ occurred. Subsequent addition of 5 μM **Ru1** and 12.5 mM NAD^+ caused the radical absorbance to decrease to zero within 22 min, signifying complete reduction of the remaining $\text{ABTS}^{\bullet-}$ (Fig. 4.7, *iii*). The ratio of NAD^+/NADH used in this experiment (694:1) was consistent with the ratio found in cells,¹²⁻¹⁵ which demonstrates that, under conditions that exhausted the free cellular NADH supply, **Ru1** could utilize the substantially more abundant cellular stores of free NAD^+ to alleviate or prevent oxidative stress.

In the presence of horseradish peroxidase (HRP), addition of H_2O_2 to ABTS^{2-} in PBS results in $\text{ABTS}^{\bullet-}$ formation, and the kinetics of this reaction can be used to evaluate the ability of an antioxidant to prevent or mitigate the onset of oxidative stress.²⁹ Inclusion of 5 μM **Ru1** and 25 mM NAD^+ significantly inhibited $\text{ABTS}^{\bullet-}$ formation, which never exceeded 4.8 μM (Fig. 4.8A, red line). After 15 min, the radical absorbance began to decrease, and complete $\text{ABTS}^{\bullet-}$ reduction was observed 6.6 min later. In contrast, 5 μM NADH completely inhibited $\text{ABTS}^{\bullet-}$ formation for 3.3 min, whereupon the absorbance gradually increased to a maximum of 11 μM (Fig. 4.8A, blue line). This concentration was 7 μM lower than the maximum observed in the control experiment and was consistent with NADH functioning as a two e^- reductant. The subsequent gradual decrease was due to normal $\text{ABTS}^{\bullet-}$ thermal decay.

After complete $\text{ABTS}^{\bullet-}$ reduction was observed following the first H_2O_2 aliquot, two additional 10 μM H_2O_2 aliquots (#) were introduced and $[\text{ABTS}^{\bullet-}]$ peaked at 6.4 μM before being reduced completely each time (Fig. 4.8B), demonstrating that the catalyst

and terminal reductant were both still present and active. Different behavior was observed with NADH after $[\text{ABTS}^{\bullet-}]$ peaked. Adding a second H_2O_2 aliquot (#) caused $[\text{ABTS}^{\bullet-}]$ to increase to 17 μM (Fig. 4.8C), corresponding to 94% ABTS^{2-} oxidation (complete oxidation = 18 μM).²⁴ No change in absorbance was produced by the third H_2O_2 aliquot (#), consistent with all of the ABTS^{2-} having been completely oxidized by the previous H_2O_2 aliquots. Subsequent treatment of this solution with 5 μM **Ru1** and 25 mM NAD^+ (§) resulted in complete $\text{ABTS}^{\bullet-}$ reduction within 45 min.

To demonstrate that the reactivity exhibited by **Ru1** and NAD^+ in Fig. 4.8 derived specifically from $\text{ABTS}^{\bullet-}$ reduction, two 10 μM aliquots of chemically synthesized $\text{ABTS}^{\bullet-}$ (*) were added after the initial reaction with H_2O_2 was complete (Fig. 4.9, red line). The $[\text{ABTS}^{\bullet-}]$ immediately increased by 8.8 μM each time, then decreased to zero 19 and 29 min after addition of the first and second $\text{ABTS}^{\bullet-}$ aliquots, respectively. We had previously shown that ABTS^{2-} is an inhibitor for **Ru1**-catalyzed $\text{ABTS}^{\bullet-}$ reduction with nontertiary alcohols,²⁵ and given that the concentration of ABTS^{2-} increased as each successive $\text{ABTS}^{\bullet-}$ aliquot was reduced, it was unsurprising that the time required for complete $\text{ABTS}^{\bullet-}$ reduction likewise increased. With the NADH experiment, however, the first and second $\text{ABTS}^{\bullet-}$ aliquots produced 9.4 and 9.3 μM increases in $[\text{ABTS}^{\bullet-}]$, respectively, that were stable over time (Fig. 4.9, blue line). Subsequent addition of 5 μM **Ru1** and 25 mM NAD^+ (§) then achieved quantitative $\text{ABTS}^{\bullet-}$ reduction in less than 39 min.

Synthetic Procedures

General synthetic considerations. The complex **Ru1**^{24,25} and ABTS^{•-} were prepared as previously described.²⁰ All other materials were of reagent quality and used as received. All solvents used were HPLC grade. All manipulations were performed under ambient conditions using standard benchtop techniques.

General spectroscopic considerations. UV-visible absorption spectra were acquired on a Varian Cary 50 Bio spectrometer equipped with a water-cooled Quantum Northwest TC-125 peltier temperature controller. All solution measurements were performed at 25.0 ± 0.1 °C in matched gas-tight quartz cuvettes (Precision Scientific) with 1 cm path lengths and 3.0 mL analyte solution volumes. Absorption spectra were acquired from 900 to 200 nm with a scanning speed of 300 nm min^{-1} and a resolution of 0.5 nm. Each kinetic analysis experiment (5 second intervals) was performed in quadruplicate on four different days. Stock solutions were prepared fresh daily and filtered (0.2 μm PTFE) immediately prior to use.

Kinetic analysis of Ru1-catalyzed ABTS^{•-} reduction with NAD⁺. An aliquot of ABTS^{•-} (40 μL from a 3.75 mM stock solution in H₂O) was added to PBS to afford the ABTS^{•-} working solution and the absorbance spectrum was acquired to confirm ABTS^{•-} concentration. The single wavelength kinetics program was initiated and, after 20 s, an aliquot of **Ru1** (30 μL from 0.5 mM stock solution in CH₃CN) was added, followed by an aliquot of NAD⁺ (30 μL from 2.5 M stock solution in H₂O) 5 min later, and the kinetics program was allowed to continue. After the kinetics program had completed, the

absorbance spectrum was acquired to confirm formation of ABTS^{2-} from the peak at 340 nm.

Inhibition of $\text{ABTS}^{\bullet-}$ formation by HRP in PBS. Derived from a literature procedure.²⁹ Aliquots of ABTS^{2-} (30 μL from a 2.0 mM stock solution in H_2O), HRP (30 μL from a 1.0 μM stock solution in H_2O), and **Ru1**, NAD^+ or NADH were added to PBS (pH 7.4) to afford the $\text{ABTS}^{2-}/\mathbf{Ru1}$ (or NADH)/HRP working solution in PBS. The absorbance spectrum of this working solution was acquired to confirm no $\text{ABTS}^{\bullet-}$ had formed. An aliquot of H_2O_2 (30 μL from 1.0 mM stock solution in H_2O) was added and the single wavelength kinetics program was initiated.

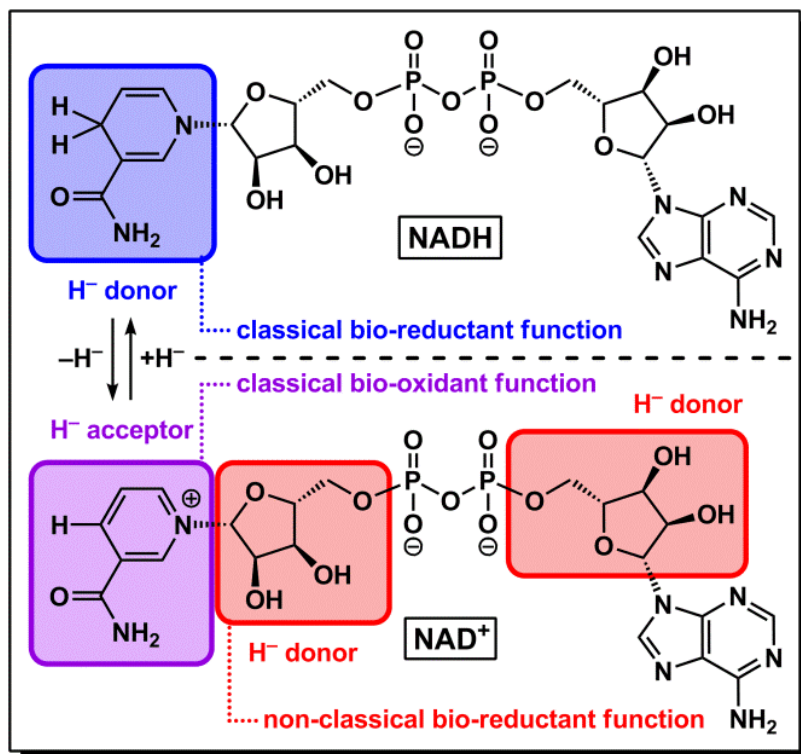
CONCLUSIONS

In summary, NAD^+ is able to function as a terminal reductant for the **Ru1**-catalyzed reduction of $\text{ABTS}^{\bullet-}$ to ABTS^{2-} in aerobic, aqueous solution. Because NAD^+ typically plays the role of H^- acceptor in biological systems, the classical expectation would be that it could not function as an H^- donor. However, the $\text{ABTS}^{\bullet-}$ reduction reactivity observed with NAD^+ and **Ru1** were highly conserved with our previous studies using other nontertiary alcohols as terminal reductants,^{24,25} which suggested that the same mechanism was operative with NAD^+ . The key intermediate responsible for $\text{ABTS}^{\bullet-}$ reduction with NAD^+ and **Ru1** was therefore inferred to be a Ru–H intermediate formed via β -hydride elimination from a ribose subunit coordinated to Ru, whereby this ability of NAD^+ to function as an H^- donor would give rise to its observed ability to function as a reductant. Previous studies by others have revealed that transition metal–hydride complexes formed via H^- transfer from NADH can react with atmospheric O_2 to generate

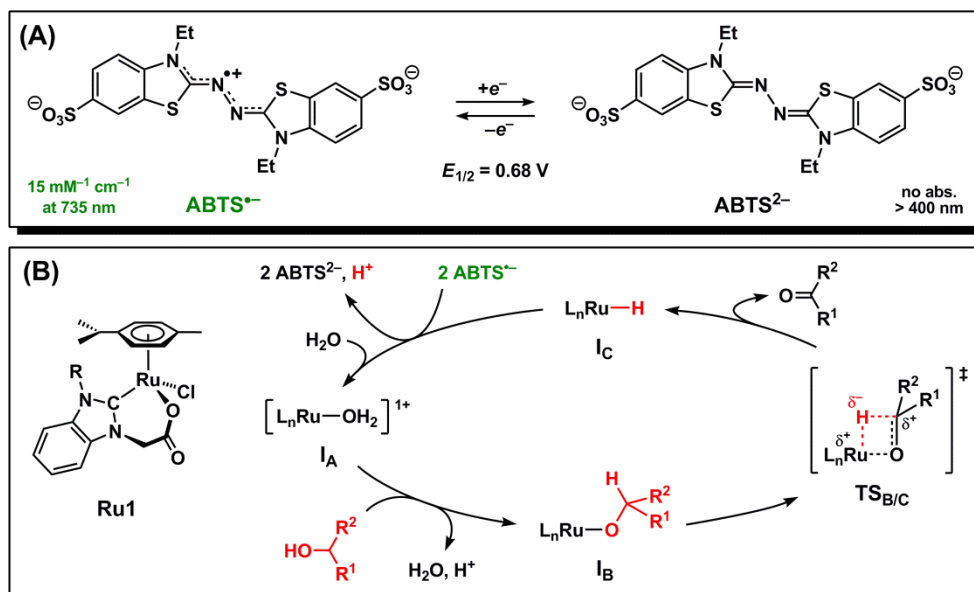
H₂O₂,^{30,31} which in turn can produce cytotoxic effects against cancer cells.^{7-9,32-34} Given that free NAD⁺ is 640–1100 times more abundant in cells than free NADH,¹²⁻¹⁵ we believe that a catalyst that can utilize NAD⁺ as an H⁻ source will be able to generate significantly higher H₂O₂ levels and thus exhibit substantially greater anticancer potency. The biological applications of **Ru1** will be detailed in subsequent reports.

Acknowledgments

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Scheme 4.1. Hydride Transfer with NADH and NAD⁺



Scheme 4.2. Ru1-Catalyzed $\text{ABTS}^{\bullet+}$ Reduction with Alcohols

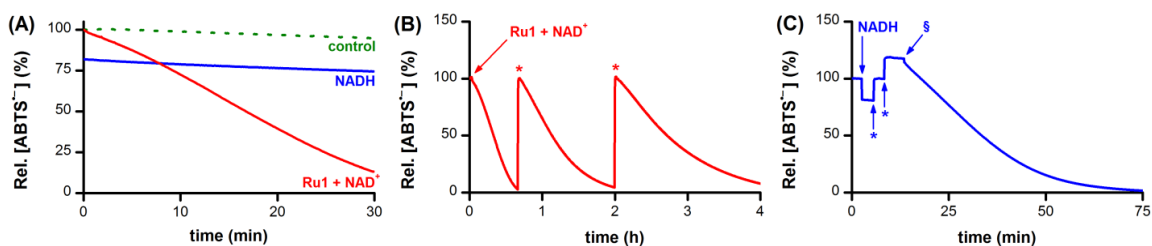


Fig. 4.1. Plot of relative $[\text{ABTS}^{\bullet-}]$ vs time, which shows the reduction of $\text{ABTS}^{\bullet-}$ to ABTS^{2-} following (A) the addition of **Ru1** and NAD^+ (red line), NADH (blue line), or **Ru1** without NAD^+ as a control (dotted green line), (B) two additional $50 \mu\text{M}$ $\text{ABTS}^{\bullet-}$ aliquots (*) after the initial reduction by **Ru1** and NAD^+ , or (C) two additional $10 \mu\text{M}$ $\text{ABTS}^{\bullet-}$ aliquots (*) after the initial reduction by NADH , followed by **Ru1** and NAD^+ (§). Conditions: $[\text{Ru1}]_0$ or $[\text{NADH}]_0 = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[\text{NAD}^+]_0 = 25 \text{mM}$, PBS (pH 7.4), $25 \text{ }^\circ\text{C}$.

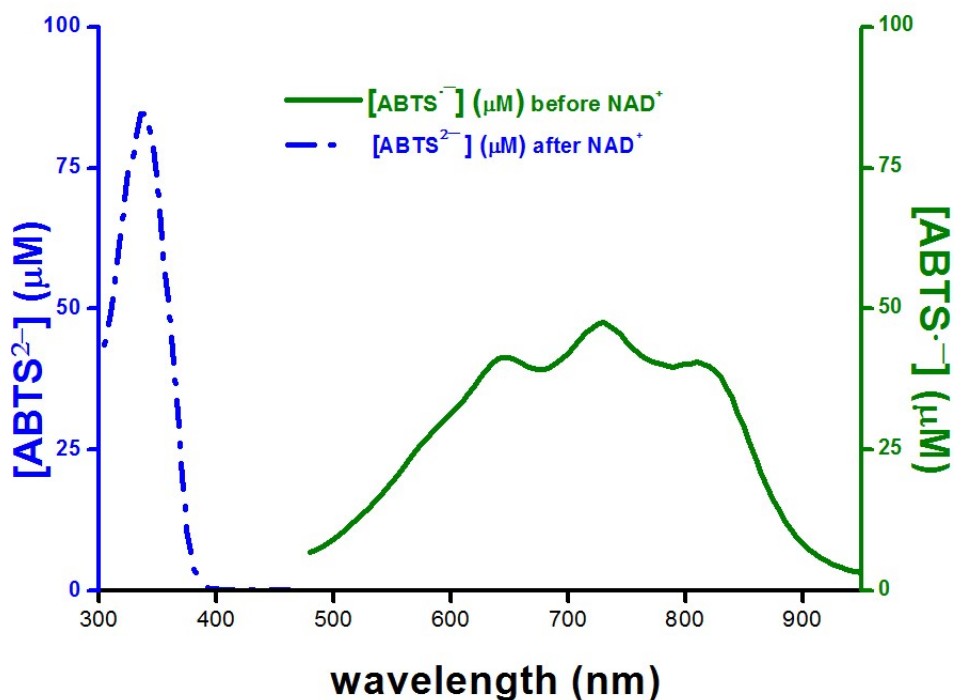


Fig. 4.2. UV/vis of ABTS^{2-} formed from $\text{ABTS}^{\bullet-}$ reduction catalyzed by **Ru1** with NAD^+ as terminal reductant. Green line shows the spectra obtained at $t = 0$ before the addition of NAD^+ , representing the amount of $\text{ABTS}^{\bullet-}$ and blue dash line shows the spectra obtained at $t = 40$ min, representing the amount of the additional ABTS^{2-} produced as well as the ABTS^{2-} from the original stock solution ($30 \mu\text{M}$). $\text{ABTS}^{\bullet-}$ has maximum absorption at 734 nm and ABTS^{2-} has the maximum absorbance at 340 nm (blue dash line). Conditions: $[\text{Ru1}]_0 = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[\text{NAD}^+]_0 = 25 \text{ mM}$; PBS, pH 7.4, $25 \text{ }^\circ\text{C}$, absorbance measured from $300\text{--}950 \text{ nm}$.

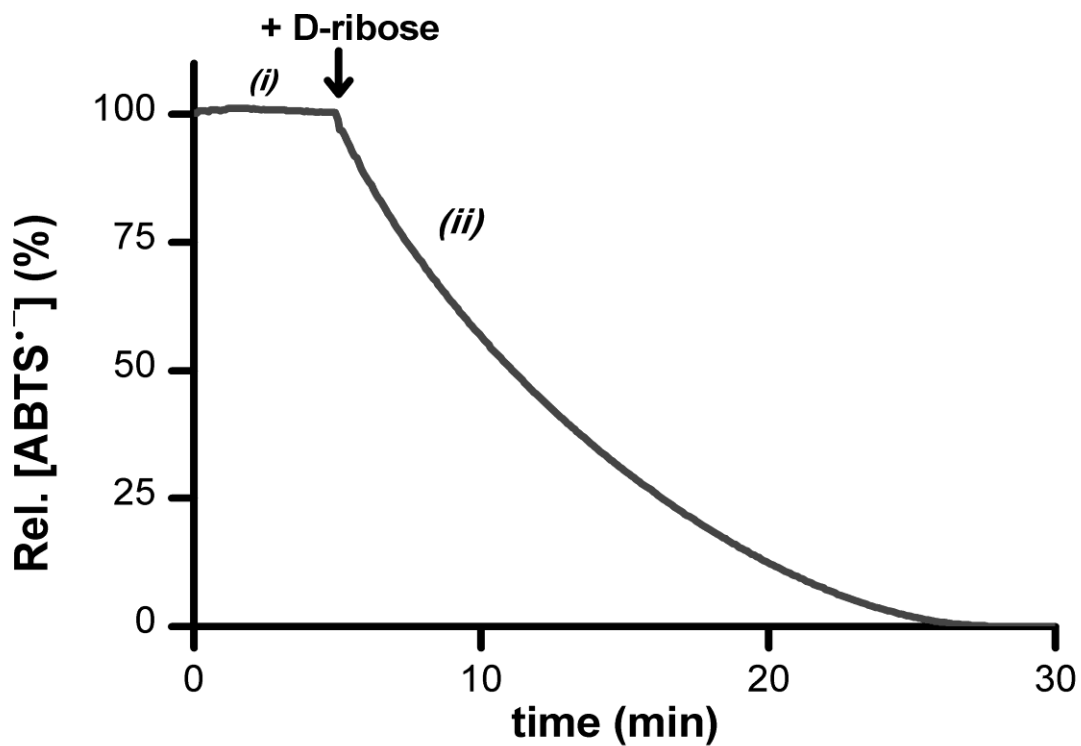


Fig. 4.3. D-Ribose caused complete reduction of ABTS•⁻ in the presence of **Ru1** catalyst. (i) In PBS solution containing only **Ru1**, no reduction of ABTS•⁻ was observed, but (ii) the absorbance decreased gradually after the addition of D-ribose. Conditions: [ABTS•⁻]₀ = 50 μM, [**Ru1**]₀ = 5 μM, [D-ribose]₀ = 25 mM; PBS, pH 7.4, 25 °C, absorbance measured at 734 nm.

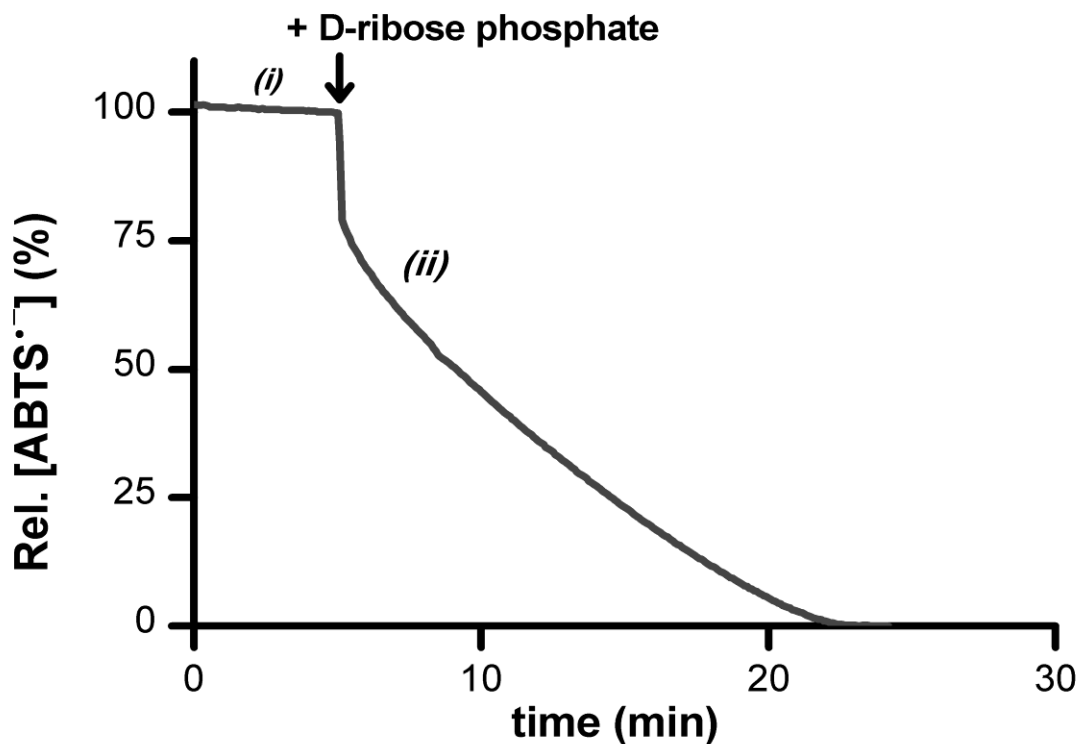


Fig. 4.4. D-Ribose phosphate caused complete reduction of $\text{ABTS}^{\bullet-}$ in the presence of **Ru1** catalyst. (i) In PBS solution containing only **Ru1**, no reduction of $\text{ABTS}^{\bullet-}$ was observed, but (ii) the absorbance decreased rapidly after the addition of ribose phosphate. Conditions: $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[\text{Ru1}]_0 = 5 \mu\text{M}$, $[\text{D-ribose phosphate}]_0 = 1 \text{mM}$; PBS, pH 7.4, 25 °C, absorbance measured at 734 nm.

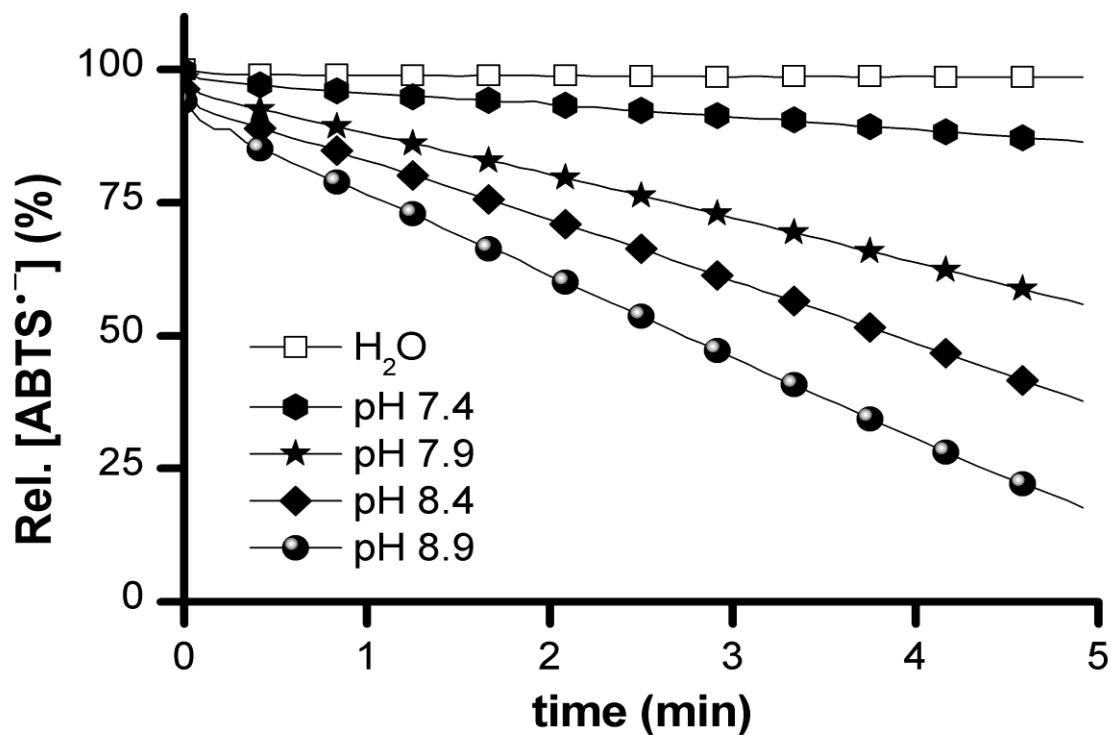


Fig. 4.5. In pure water, NAD^+ caused no reduction of $\text{ABTS}^{\bullet-}$ in the presence of **Ru1** catalyst whereas faster rates of reduction by NAD^+ were observed in PBS solutions with increasing pH. Conditions: $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[\text{Ru1}]_0 = 5 \mu\text{M}$, $[\text{NAD}^+]_0 = 25 \text{mM}$, pH = 7.4, 7.9, 8.4 or 8.9; PBS or H_2O , 25°C , absorbance measured at 734 nm.

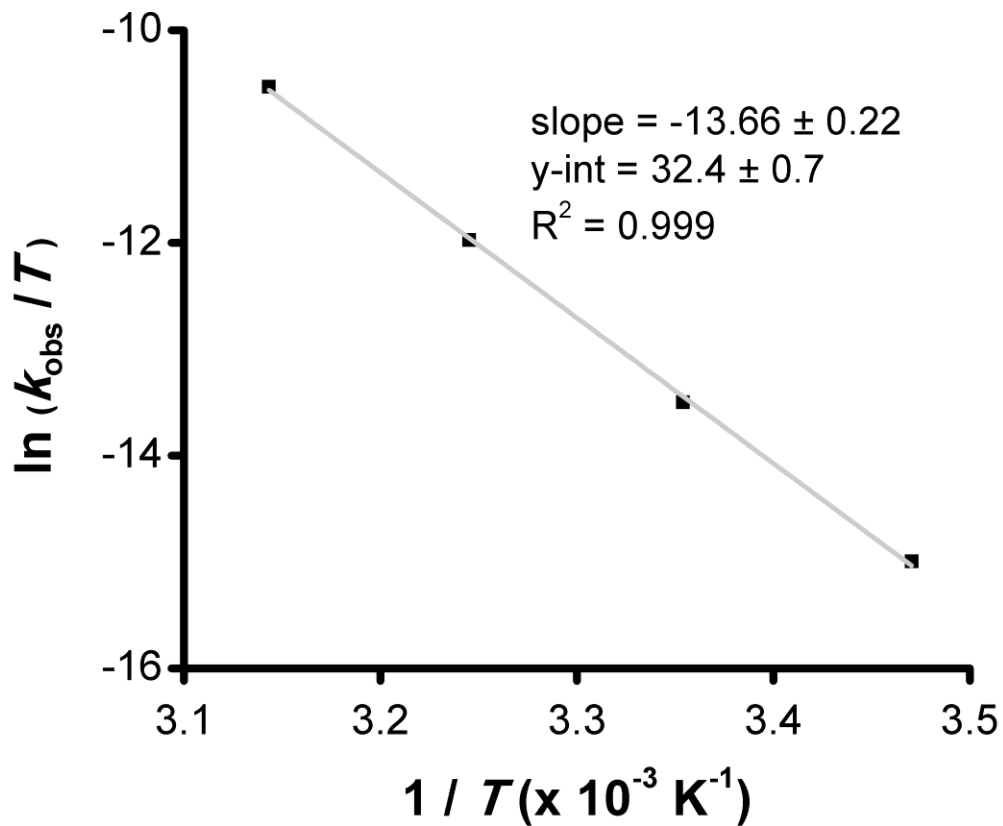


Fig. 4.6. Eyring plot for $\text{ABTS}^{\bullet-}$ reduction catalyzed by **Ru1** with NAD^+ as terminal reductant. Conditions: $[\text{Ru1}]_0 = 5 \mu\text{M}$, $[\text{ABTS}^{\bullet-}]_0 = 50 \mu\text{M}$, $[\text{ABTS}^{2-}]_0 = 100 \mu\text{M}$, $[\text{NAD}^+]_0 = 25 \text{ mM}$, $T = 15, 25, 35$ or $45 \text{ }^\circ\text{C}$; PBS, pH 7.4, absorbance measured at 734 nm.

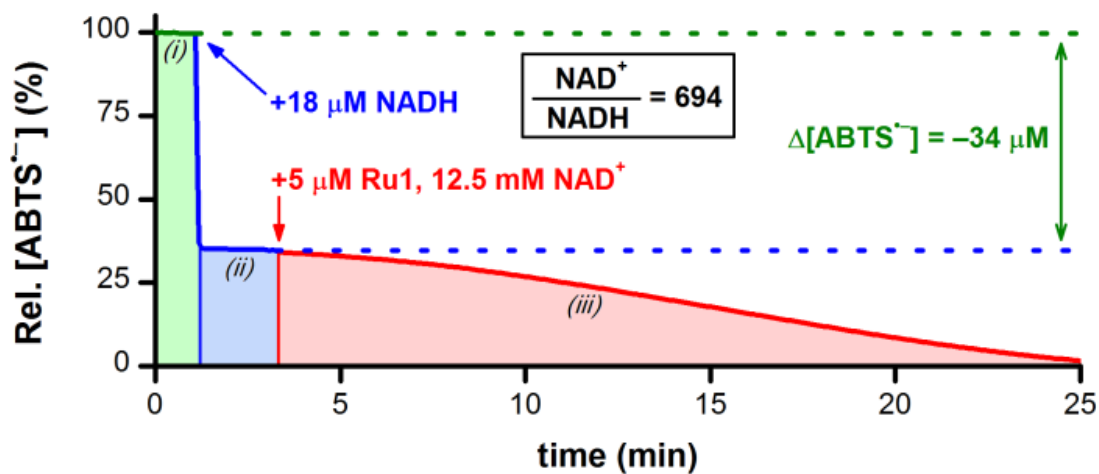


Fig. 4.7. Plot of relative $[ABTS^{\bullet-}]$ vs time (i), which shows the 2:1 stoichiometric reduction of $ABTS^{\bullet-}$ by NADH (ii), followed by the catalytic reduction of $ABTS^{\bullet-}$ by **Ru1** and NAD^+ (iii). Conditions: $[Ru1]_0 = 5 \mu M$, or $[NADH]_0 = 18 \mu M$, $[ABTS^{\bullet-}]_0 = 50 \mu M$, $[NAD^+]_0 = 12.5 \text{ mM}$, PBS (pH 7.4), 25 °C.

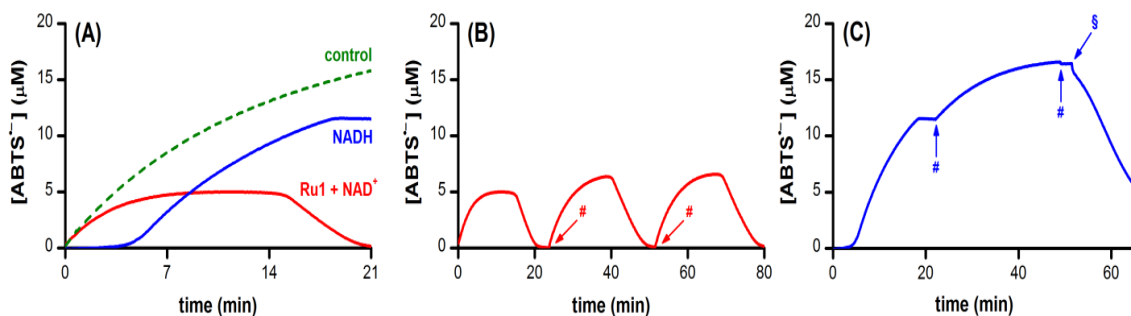


Fig. 4.8. (A) Plot of $[\text{ABTS}^{\bullet-}]$ vs time, which shows the oxidation of ABTS^{2-} to $\text{ABTS}^{\bullet-}$ *in situ* by HRP and H_2O_2 in the presence of **Ru1** and NAD^+ (red line), NADH (blue line), or **Ru1** without NAD^+ as a control (dotted green line). Plot of $[\text{ABTS}^{\bullet-}]$ vs time, which shows $\text{ABTS}^{\bullet-}$ formation following two additional aliquots of $10 \mu\text{M}$ H_2O_2 (#) in the presence of (B) **Ru1** and NAD^+ or (C) NADH. For the NADH experiment shown in (C), **Ru1** and NAD^+ were added (§) after the final aliquot of $10 \mu\text{M}$ H_2O_2 . Conditions: $[\text{HRP}]_0 = 10 \text{ nM}$, $[\text{Ru1}]_0$ or $[\text{NADH}]_0 = 5 \mu\text{M}$, $[\text{H}_2\text{O}_2]_0 = 10 \mu\text{M}$, $[\text{ABTS}^{2-}]_0 = 20 \mu\text{M}$, $[\text{NAD}^+]_0 = 25 \text{ mM}$, PBS (pH 7.4) at $25 \text{ }^\circ\text{C}$.

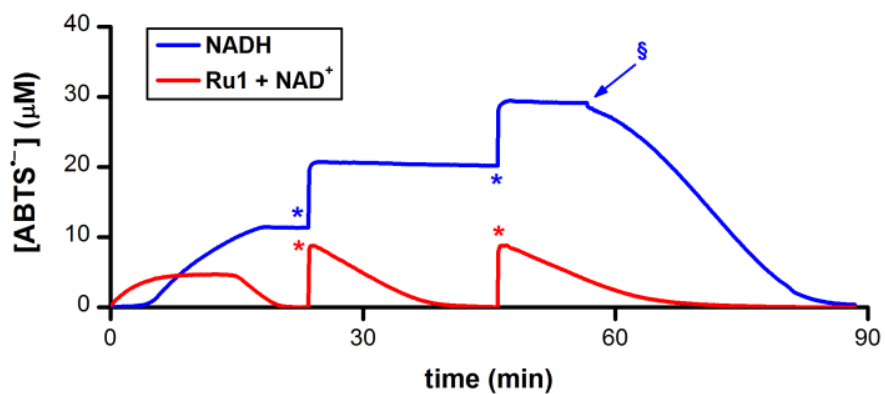


Fig. 4.9. Plot of $[\text{ABTS}^{\bullet-}]$ vs time, which shows the oxidation of ABTS^{2-} to $\text{ABTS}^{\bullet-}$ *in situ* by HRP and H_2O_2 followed by subsequent $\text{ABTS}^{\bullet-}$ reactivity in the presence of **Ru1** and NAD^+ (red line) or NADH (blue line). After the initial reaction of **Ru1** and NAD^+ or NADH had completed, two additional aliquots of $10 \mu\text{M}$ $\text{ABTS}^{\bullet-}$ (*) were introduced. For the NADH experiment (blue line), $5 \mu\text{M}$ **Ru1** and 25mM NAD^+ were added (§) after the final aliquot of $10 \mu\text{M}$ $\text{ABTS}^{\bullet-}$. Conditions: $[\text{HRP}]_0 = 10 \text{ nM}$, $[\text{Ru1}]_0$ or $[\text{NADH}]_0 = 5 \mu\text{M}$, $[\text{H}_2\text{O}_2]_0 = 10 \mu\text{M}$, $[\text{ABTS}^{2-}]_0 = 20 \mu\text{M}$, $[\text{NAD}^+]_0 = 25 \text{mM}$, PBS (pH 7.4) at $25 \text{ }^\circ\text{C}$.

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