
Computer Modeling (Physical Chemistry) of Enzyme Catalysis, Metalloenzymes

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Outline

Physical Chemistry of Enzyme Catalysis

- (Enzymatic) Reaction Rate and Order
- Michaelis-Menten (and Enzyme) Kinetics

Metals in Enzymology (Theoretical Bioinorganic Chemistry)

- Stability Constants, Selectivity
- Spin-States in Biochemistry
- Crystal Field/Ligand Field Theories
- DFT vs. WFT Methods (Accuracy and Pitfalls)
- Relativistic Effects (Mild Introduction)



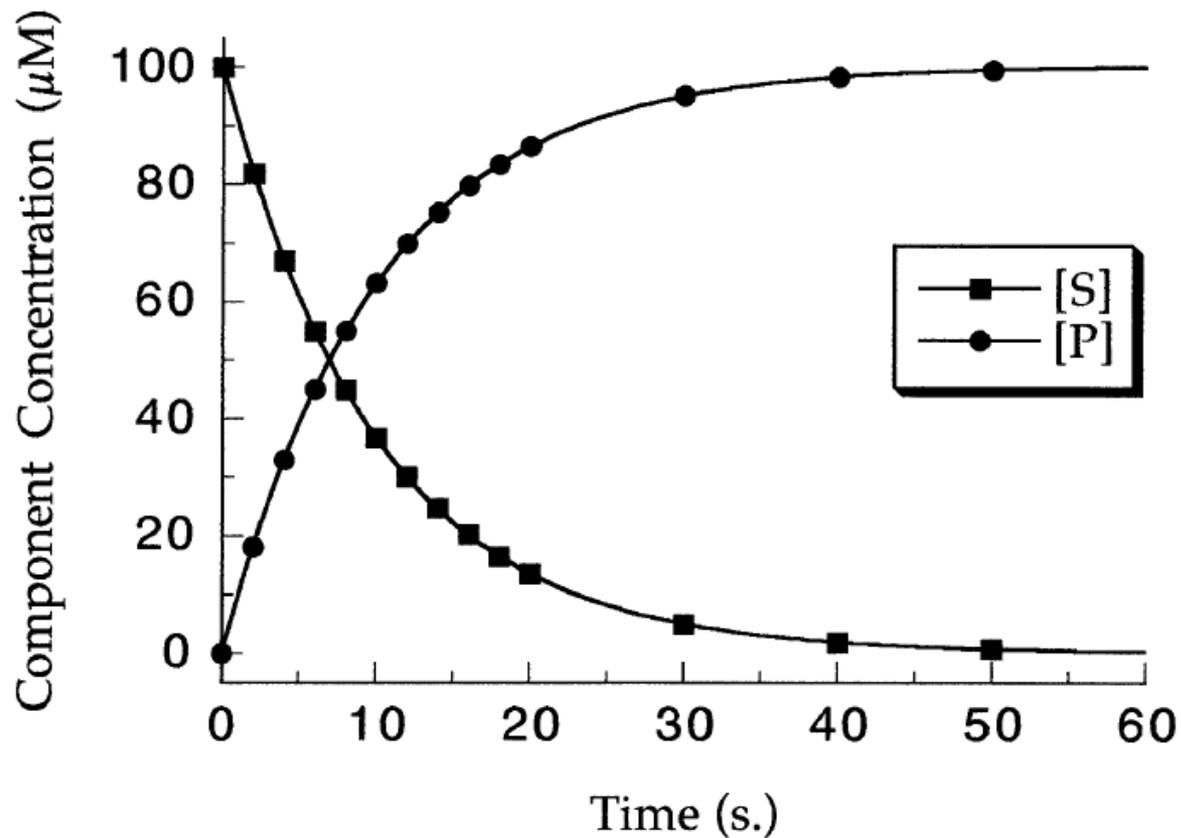


Figure 2.15 Progress curves of product development (circles) and substrate loss (squares) for a first-order reaction.



Chemical reaction: $A + B + C + \dots \rightarrow \{P\}$

Typical rate equation: $v = k[A]^x[B]^y[C]^z\dots$

x... reaction order with respect to **A**

$x + y + z + \dots$ overall reaction order

Reaction order is not always equivalent to stoichiometry and can be determined only experimentally; allows to hypothesize reaction mechanism,

Knowledge of R.O. may suggest the rate-determining step (RDS) (= RLS)

Elementary Reaction: single reaction step, single transition state

Reaction order is then equivalent to stoichiometry



First order reactions

The reaction rate depends on a single reactant and the $x = 1$

Example: S_N1 reaction $\text{ArN}_2^+ + \text{X}^- \rightarrow \text{ArX} + \text{N}_2$, the rate equation is $v = k[\text{ArN}_2^+]$

Second order reactions

$$x + y + \dots = 2$$

$\text{A} + \text{B} + \dots \rightarrow \{\text{P}\}$ can be e.g. $v = k[\text{A}]^2$ or $v = k[\text{A}][\text{B}]$

Example: $\text{NO}_2 + \text{CO} \rightarrow \text{NO} + \text{CO}_2$ is $v = k[\text{NO}_2]^2$

S_N2 : $\text{CH}_3\text{COOC}_2\text{H}_5 + \text{OH}^- \rightarrow \text{CH}_3\text{COO}^- + \text{C}_2\text{H}_5\text{OH}$ is $v = k[\text{CH}_3\text{COOC}_2\text{H}_5][\text{OH}^-]$

Pseudo-first order reactions

If the concentration of one of the reactant stays constant $[\text{B}] = [\text{B}]_0$ (e.g. catalyst or excess concentration)

$$\text{then } v = k[\text{A}][\text{B}] = k'[\text{A}]$$



Determination of reaction order

Method of initial rates

$$\ln v = \ln k + x \ln [A] + y \ln [B] + \dots$$

Integral Method

integrated rate law for a first-order reaction is

$$\ln[A] = -kt + \ln[A]_0$$

Method of Flooding

$v = k [A]^\alpha [B]^\beta$ in excess of reactant B becomes $v = k' [A]^\alpha$



(Henri-)Michaelis-Menten Kinetics



Leonor Michaelis
(1875-1949)

worked together in Berlin



Maud Leonora Menten
(1879-1960)

expanding on the work of Victor Henri who published the first successful mathematical model for describing enzyme kinetics in 1903

Michaelis, L.; Menten, M.L. (1913). "Die Kinetik der Invertinwirkung". *Biochem Z* **49**: 333–369; **recent translation**: *Biochemistry*, **2011**, 50 (39), pp 8264–8269



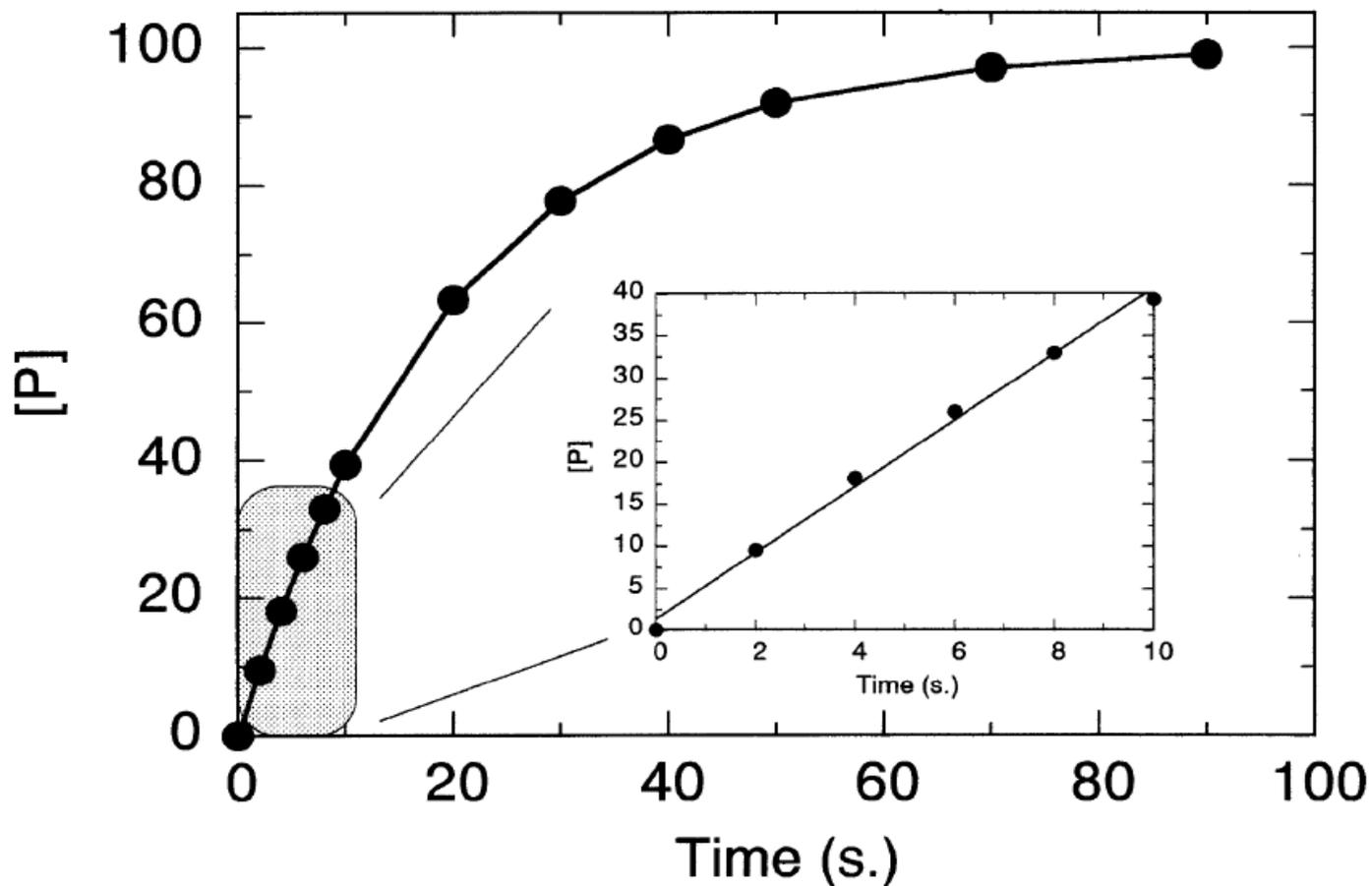


Figure 5.2 Reaction progress curve for the production of product during an enzyme-catalyzed reaction. Inset highlights the early time points at which the initial velocity can be determined from the slope of the linear plot of [P] versus time.



Substrate concentration

$$[S] = [S_0]e^{-kt}$$

$$v = -\frac{d[S]}{dt} = \frac{d[P]}{dt} = k[S_0]e^{-kt}$$

Initial velocity (~10% of substrate conversion)

$$v_0 = -\frac{\Delta[S]}{\Delta t} = \frac{\Delta[P]}{\Delta t}$$



Effect of substrate concentration on velocity

Brown (1902) – qualitative picture

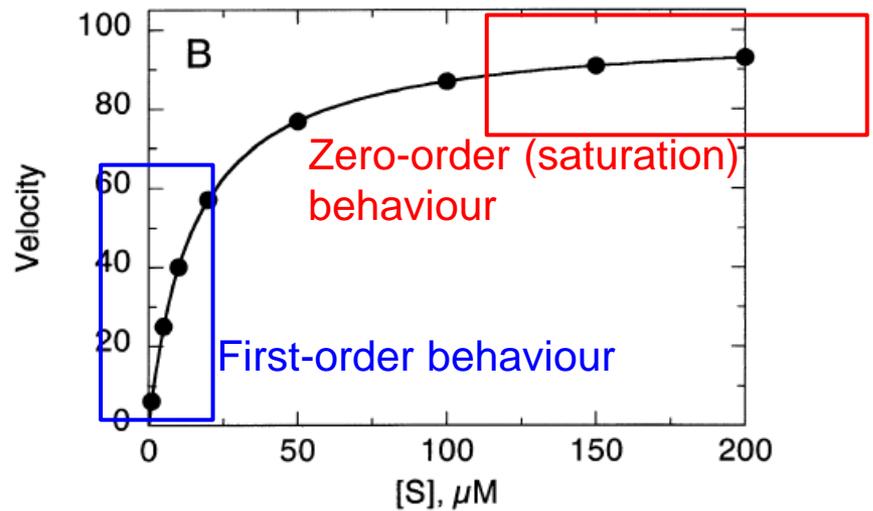
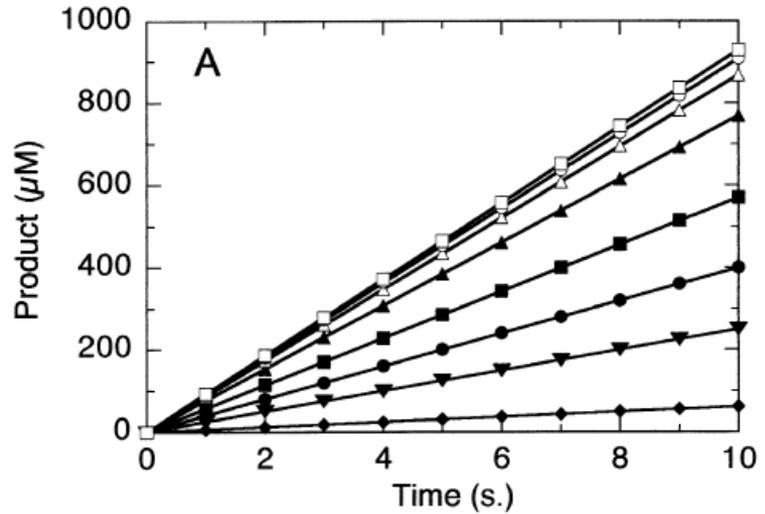
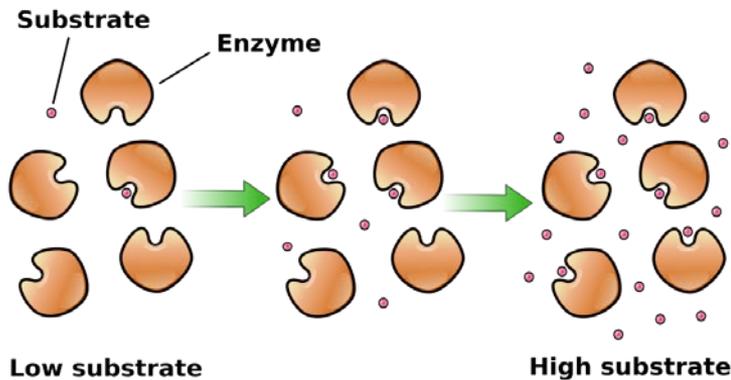
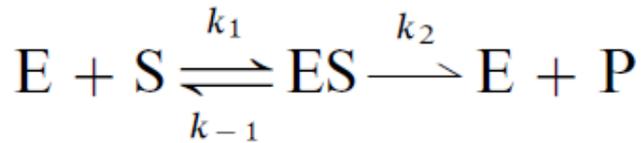
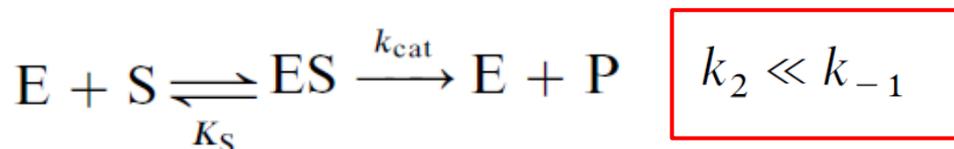


Figure 5.3 (A) Progress curves for a set of enzyme-catalyzed reactions with different starting concentrations of substrate $[S]$. (B) Plot of the reaction velocities, measured as the slopes of the lines from (A), as a function of $[S]$.



THE RAPID EQUILIBRIUM MODEL OF ENZYME KINETICS

Henri (1903) and Michaelis & Menten (1913) put the Brown's model into mathematical framework



$$K_S = \frac{[E]_f [S]}{[ES]}$$

Assuming $[S] = [S_f]$



$$[ES] = \frac{[E][S]}{K_S + [S]}$$

$$v = k_{\text{cat}} [ES]$$

$$v = \frac{k_{\text{cat}} [E][S]}{K_S + [S]}$$

$$V_{\text{max}} = k_{\text{cat}} [E]$$

Original H-M-M equation

$$v = \frac{V_{\text{max}} [S]}{K_S + [S]} = \frac{V_{\text{max}}}{1 + \frac{K_S}{[S]}}$$

However, this original approach is useful in single-turnover (rapid) reactions



THE STEADY STATE MODEL OF ENZYME KINETICS

Briggs and Haldane (1925)

Does not require $k_2 \ll k_{-1}$

Assumptions:

(1) In the initial stage $[E] = [E]_f + [ES]$

(2) $[S] \gg [E]$. $[S]_f \sim [S]$

(3) In the initial stage, depletion of $[S]$ is minimal and $\frac{d[ES]}{dt} = 0$



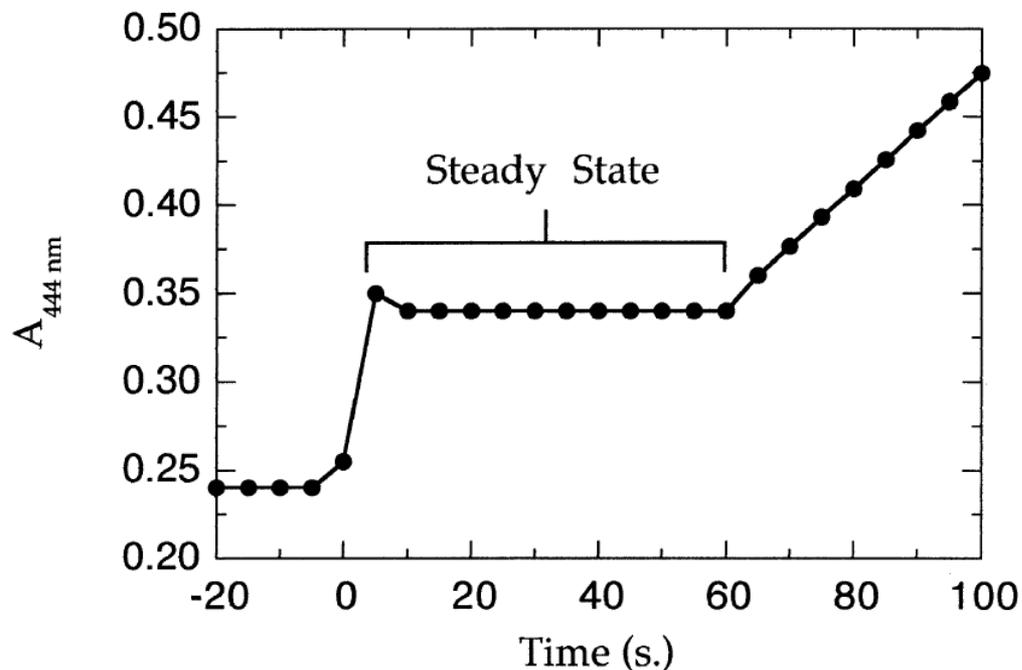


Figure 5.4 Development of the steady state for the reaction of cytochrome *c* oxidase with its substrates, cytochrome *c* and molecular oxygen. The absorbance at 444 nm reflects the ligation state of the active site heme cofactor of the enzyme. Prior to substrate addition (time < 0) the heme group is in the Fe³⁺ oxidation state and is ligated by a histidine group from the enzyme. Upon substrate addition, the active site heme iron is reduced to the Fe²⁺ state and rapidly reaches a steady state phase of substrate utilization in which the iron is ligated by some oxygen species. The steady state phase ends when a significant portion of the molecular oxygen in solution has been used up. At this point the heme iron remains reduced (Fe²⁺) but is no longer bound to a ligand at its sixth coordination site; this heme species has a much larger extinction coefficient at 444 nm; hence the rapid increase in absorbance at this wavelength following the steady state phase. [Data adapted and redrawn from Copeland (1991).]

HMM Equation

$$v = k_2[\text{ES}]$$

$$\frac{d[\text{ES}]}{dt} = k_1[\text{E}]_f[\text{S}]_f \quad \text{and} \quad -\frac{d[\text{ES}]}{dt} = (k_{-1} + k_2)[\text{ES}]$$

$$k_1[\text{E}]_f[\text{S}]_f = (k_{-1} + k_2)[\text{ES}]$$

$$[\text{ES}] = \frac{[\text{E}]_f[\text{S}]_f}{K_m} \quad \text{defining} \quad K_m = \frac{k_{-1} + k_2}{k_1}$$

replace $[\text{E}]_f$ by $([\text{E}] - [\text{ES}])$

$$v = k_{\text{cat}}[\text{E}] \frac{[\text{S}]}{[\text{S}] + K_m}$$

$$v = \frac{V_{\text{max}}[\text{S}]}{K_m + [\text{S}]} = \frac{V_{\text{max}}}{1 + \frac{K_m}{[\text{S}]}}$$



The K_m – **Michaelis constant** - is the substrate concentration that provides a reaction velocity that is half of the maximal velocity obtained under saturating substrate conditions.

K_m is not equal to K_S in rapid equilibrium model, but can be used as the fairly good *relative* estimate of substrate binding affinity

The apparent unimolecular rate constant k_{cat} is also called *turnover number* and denotes the maximum number of enzymatic reactions catalysed per second. [s^{-1}]

In vivo, often $[S] \ll K_m$, (typically $0.1 - 1 K_M$) the overall reaction may be limited by the diffusional rate of encounter of the free enzyme with substrate, which is defined by k_1 . The rate constant for diffusional encounters between molecules like enzymes and substrates is typically in the range of $10^8 - 10^9 M^{-1}s^{-1}$

⇒ **Catalytic perfection**

The **catalytic efficiency** of an enzyme is best defined by the ratio of the kinetic constants, k_{cat}/K_m

$$\Delta G_{ES^\ddagger} = -RT \ln \left(\frac{k_{cat}}{K_m} \right) + RT \ln \left(\frac{k_B T}{h} \right) \quad \Delta \Delta G_{ES^\ddagger} = -RT \ln \left[\frac{(k_{cat}/K_m)^1}{(k_{cat}/K_m)^2} \right]$$

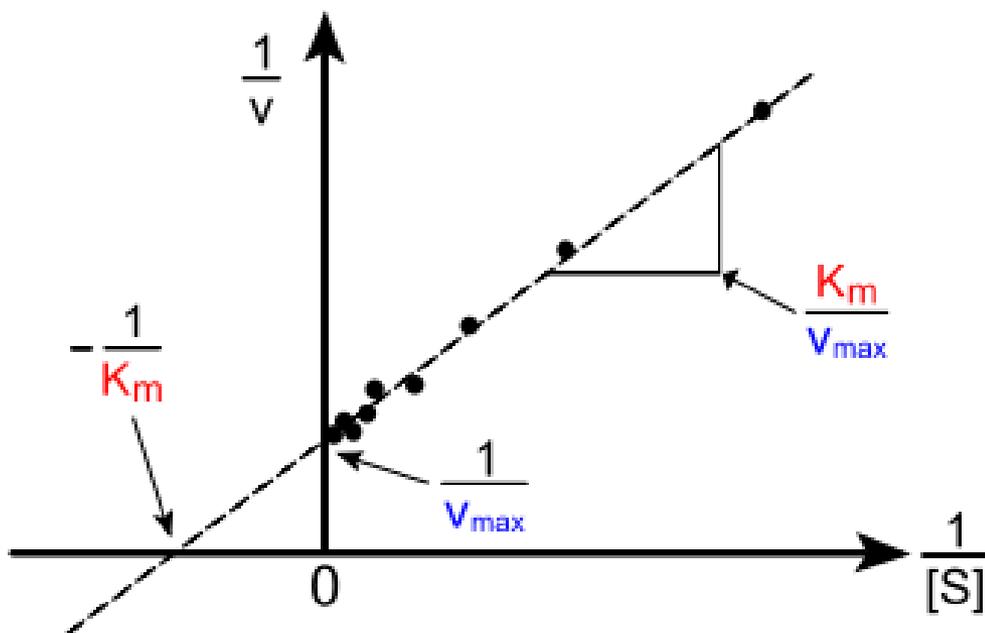


Linear plots of the Michaelis–Menten equation

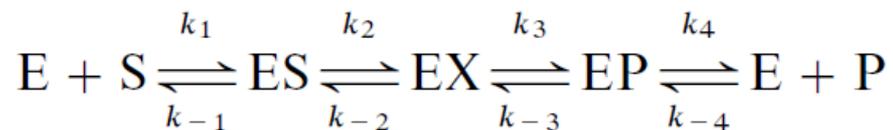
The [Lineweaver–Burk plot](#) or double reciprocal plot is a common way of illustrating kinetic data.

Rearrange HMM to

$$\frac{1}{v} = \left(\frac{K_m}{V_{\max}} \frac{1}{[S]} \right) + \frac{1}{V_{\max}}$$



TRANSIENT STATE KINETIC MEASUREMENTS



pre—steady state kinetics

stopped-flow and rapid reaction quenching (RFQ)

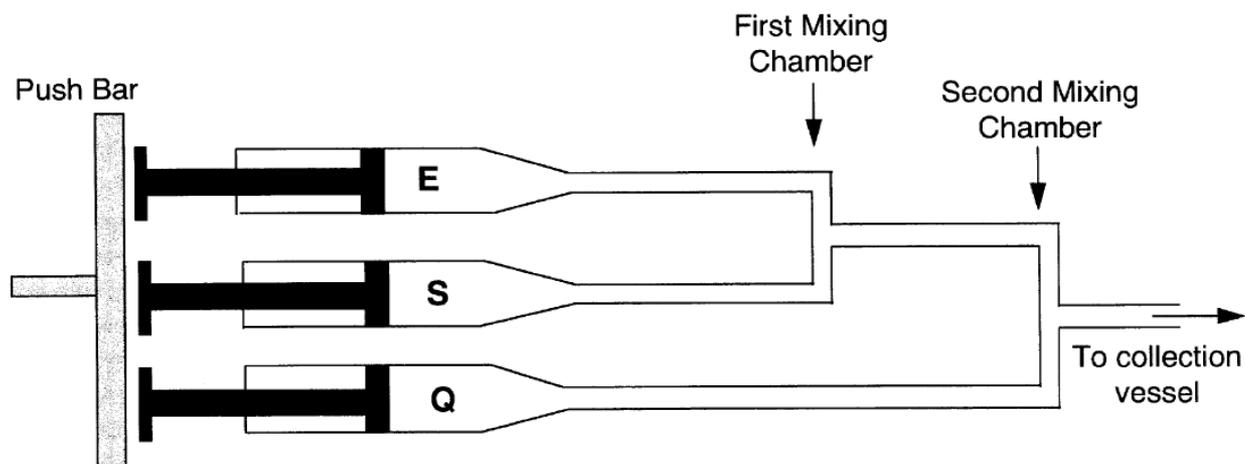
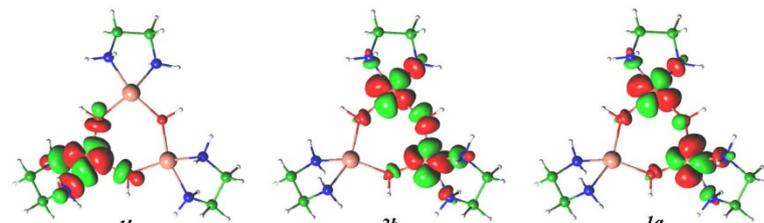
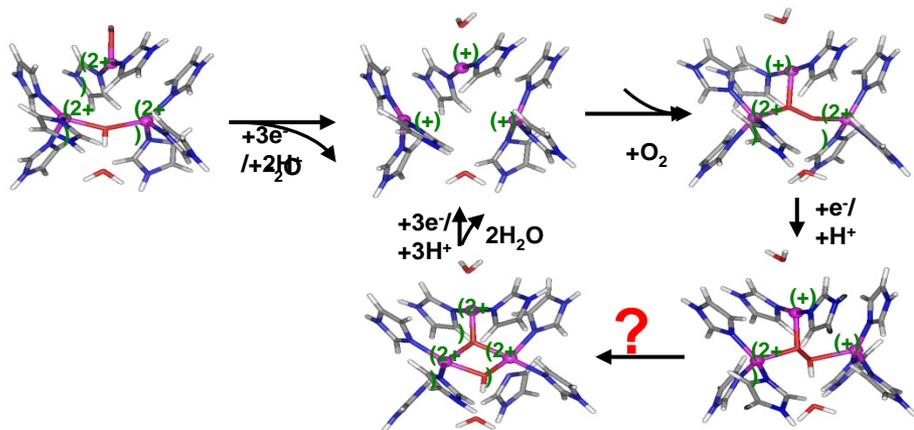
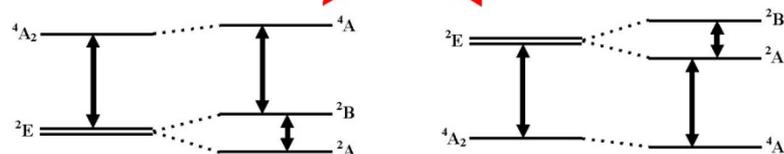


Figure 5.18 Schematic diagram of a typical rapid quench instrument for rapid kinetic measurements.

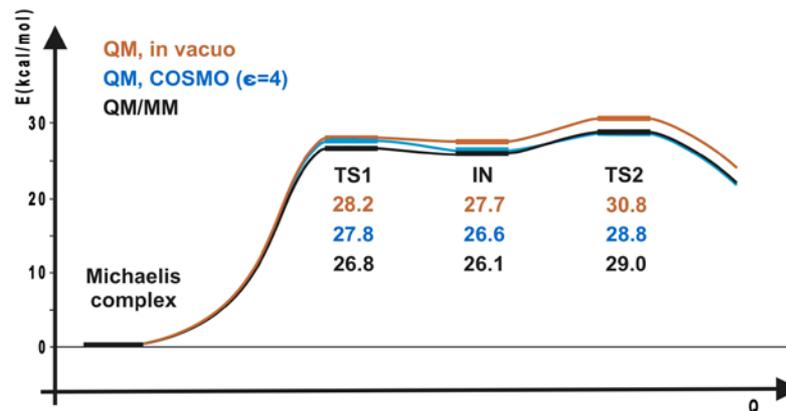
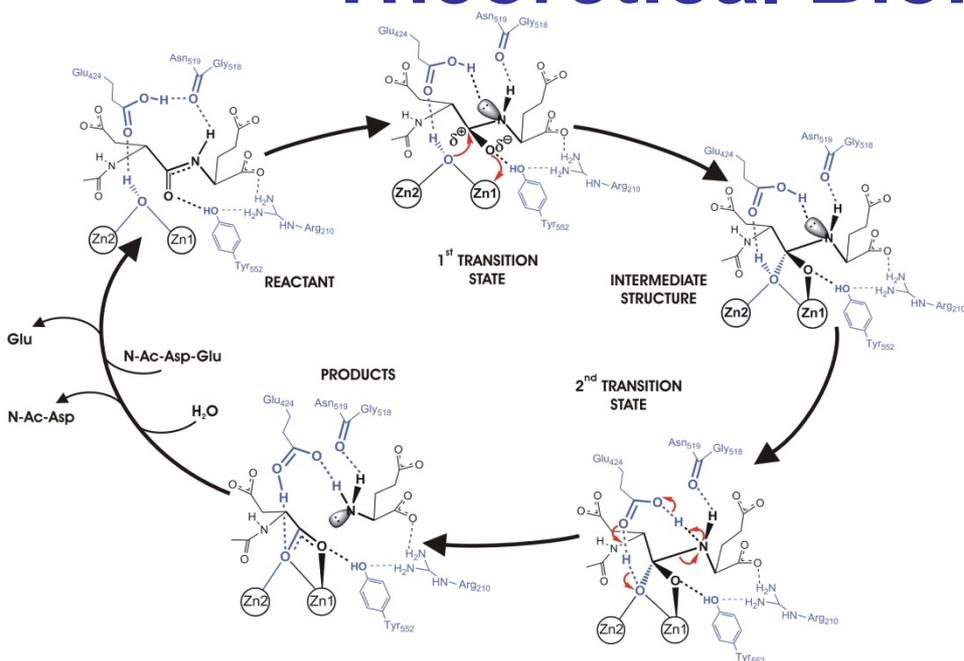


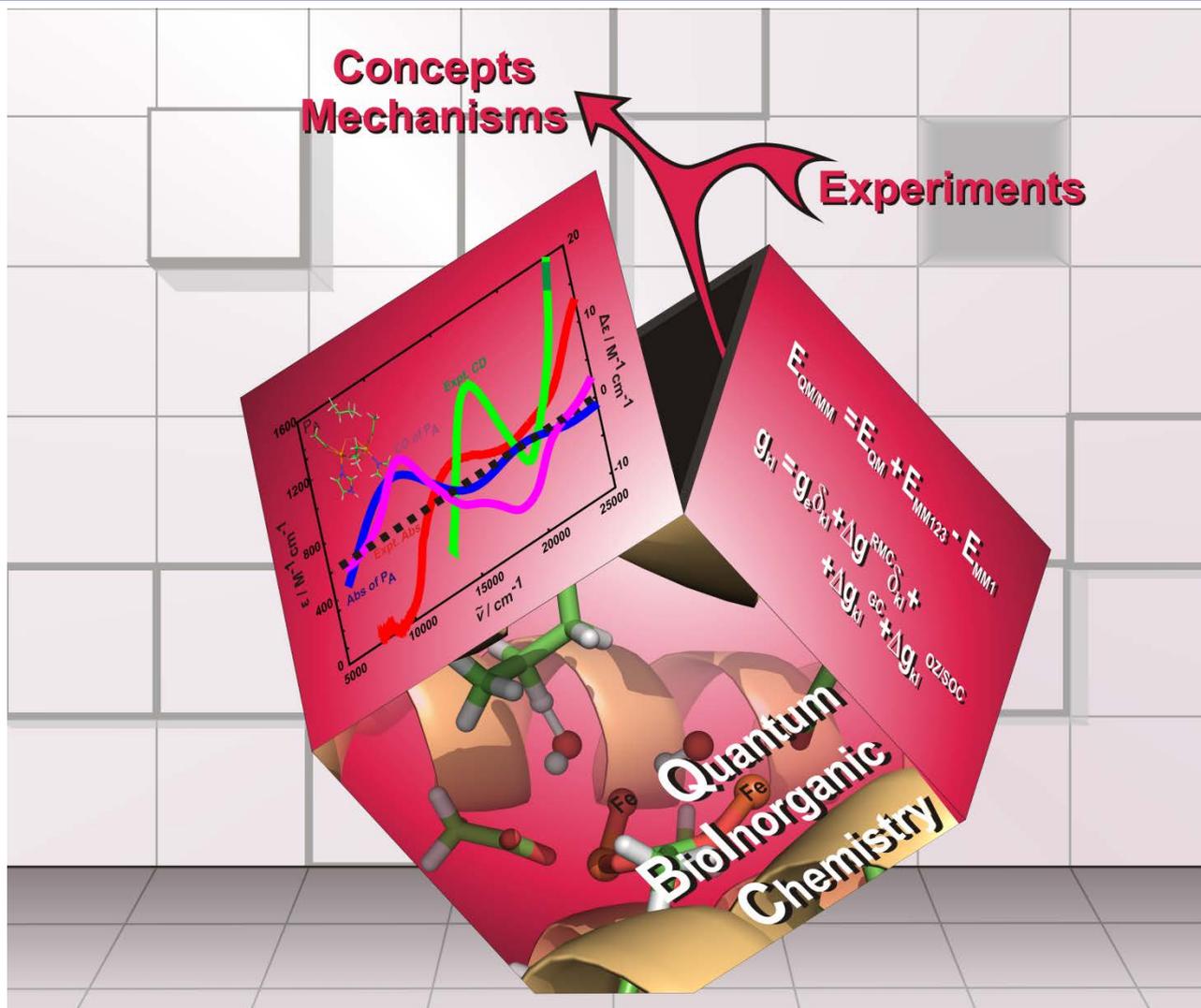


$$\begin{aligned}
 1^2A (D_1) &\sim 0.61((1a)^{\alpha}(1b)^2 - (1a)^{\alpha}(2b)^2) + 0.43(1a)^{\alpha}(1b)^{\beta}(2b)^{\alpha} - 0.25(1a)^{\alpha}(1b)^{\alpha}(2b)^{\beta} \\
 1^2B (D_2) &\sim 0.68((1a)^2(1b)^{\alpha} + (1b)^{\alpha}(2b)^2) + 0.20(1b)^2(2b)^{\alpha} - (1a)^2(2b)^{\alpha} \\
 1^4A (Q_1) &\sim 1.00(1a)^{\alpha}(1b)^{\alpha}(2b)^{\alpha}
 \end{aligned}$$



Theoretical Bioinorganic Chemistry





adapted from Rokob, T. A.; Srnec, M.; Rulíšek, L.: Theoretical Calculations of Physico-Chemical and Spectroscopic Properties of Bioinorganic Systems: Current Limits and Perspectives. *Dalton Trans.* **2012**, 41, 5754-5768.



Protein Structure → Theoretical Model

full protein without conformational sampling

QM/MM

QM/MM/Exp (X-ray, EXAFS, NMR)

full protein with conformational sampling

QM/MD, QM/MM/FEP, QTCP

cluster model (active site only)

QM+solvation (COSMO-RS, SMD, ...)

Calculations vs. Experiment

spectroscopic properties

Absorption, CD, MCD, EPR, IR, Raman, Mössbauer, NRVS, ...

thermodynamic properties

reduction potentials, pK_a values, equilibrium constants

kinetic properties

rate constants, isotope effect

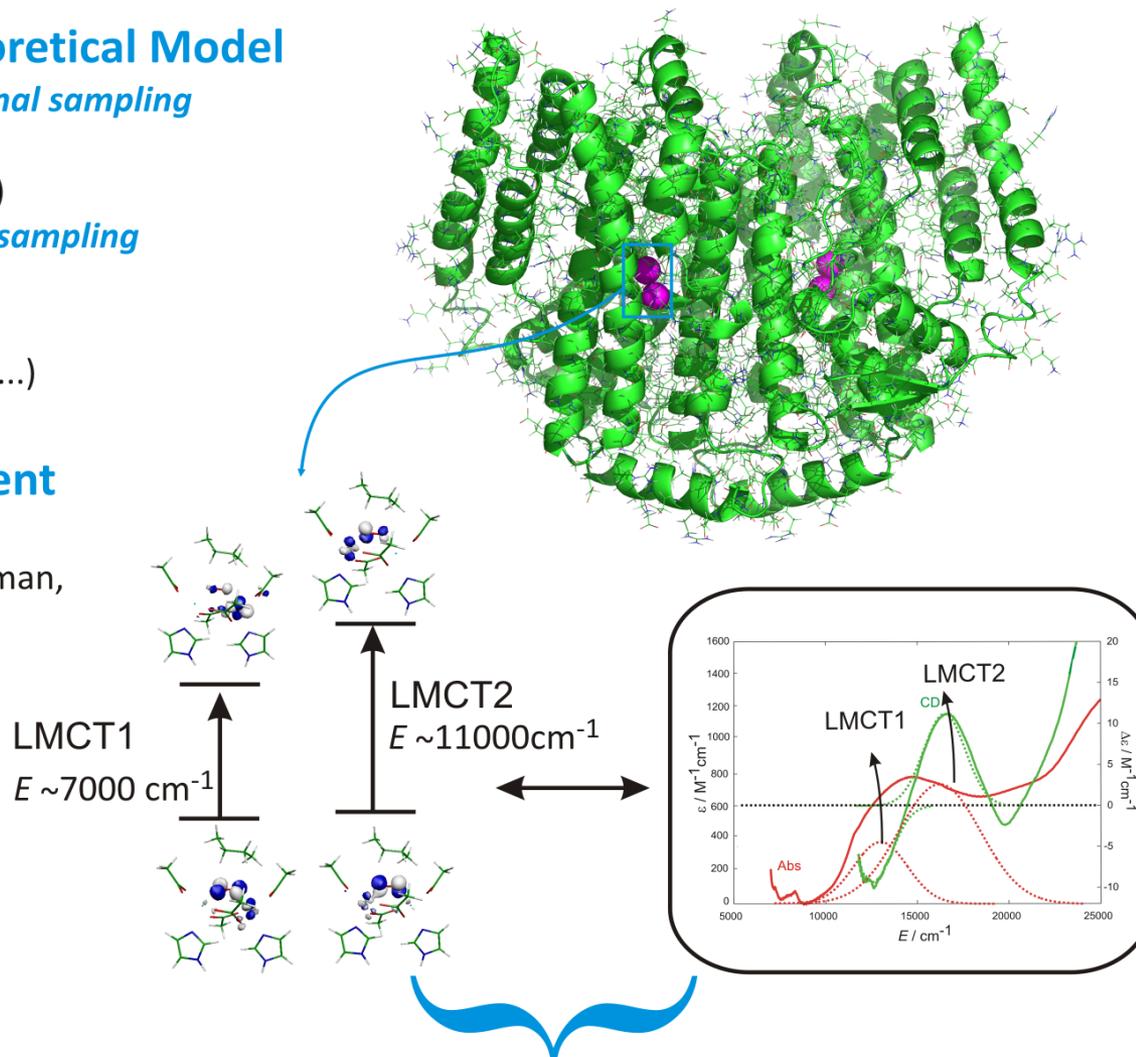
QM Methods

wave function methods

MR-SCF, MR-PT2, MRCI, DMRG, ...

density functional theory (DFT) methods

DFT, DFT+D, ...



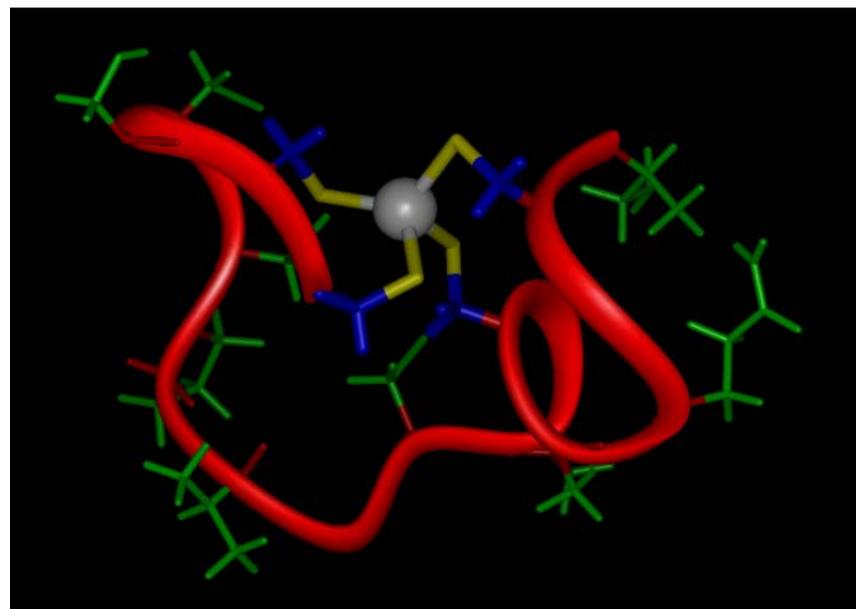
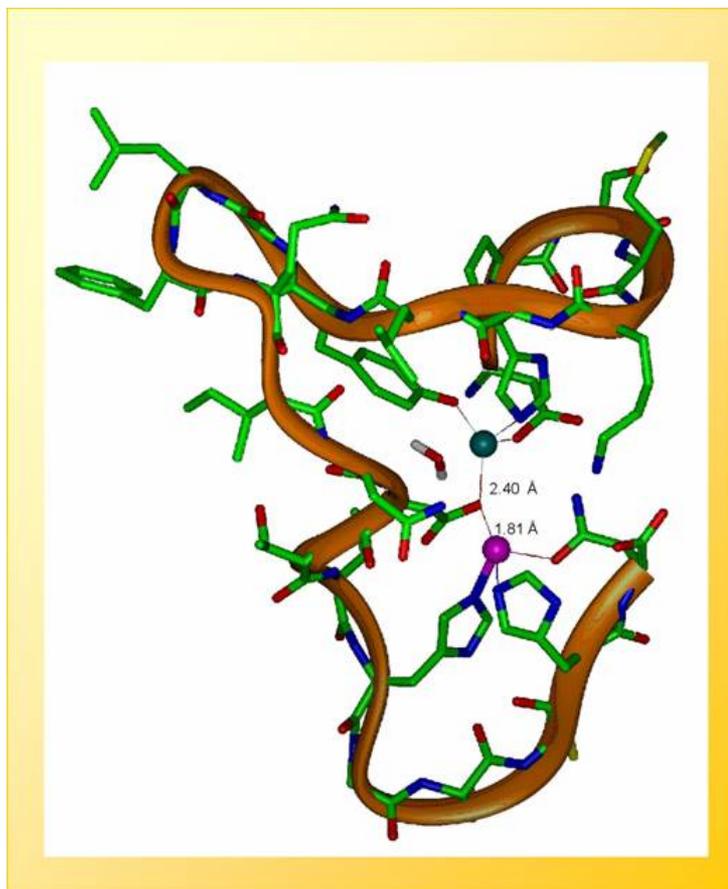
Concepts and Mechanisms

adapted from Rokob, T. A.; Srnec, M.; Rulišek, L.: *Dalton Trans.* **2012**, 41, 5754-5768.



Metal Ion Selectivity

Why Nature selected particular metal ion to perform particular task?



Non-equilibrium concentrations in cells

Cell ~ μM

Sea water ~ nM

- Metalloproteins

- Uphill battle against Irving-Williams series

$\text{Mn(II)} < \text{Fe(II)} < \text{Co(II)} < \text{Ni(II)} < \text{Cu(II)} >$
 Zn(II)

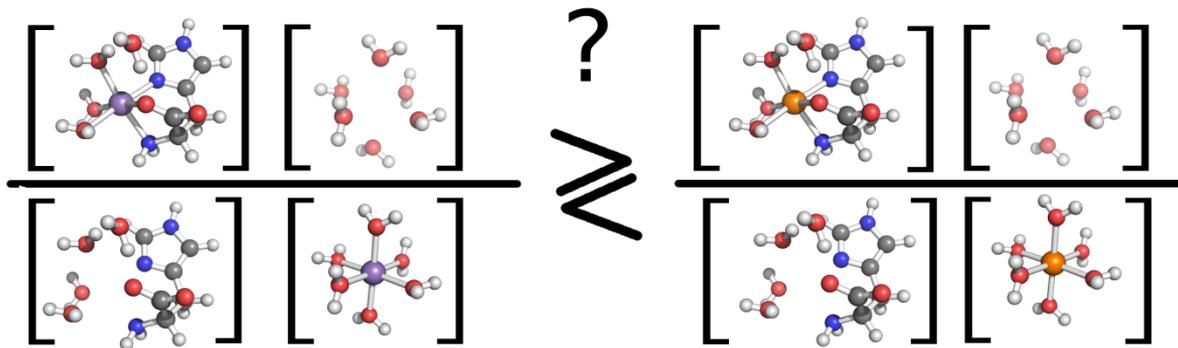


Thermodynamics of Metal Binding

$$K = \frac{[ML_n]}{[M][L]^n}$$

$$\Delta G = -RT \ln K$$

$$G = E_{el} + ZPE - RT \ln Q + G_{solv}$$

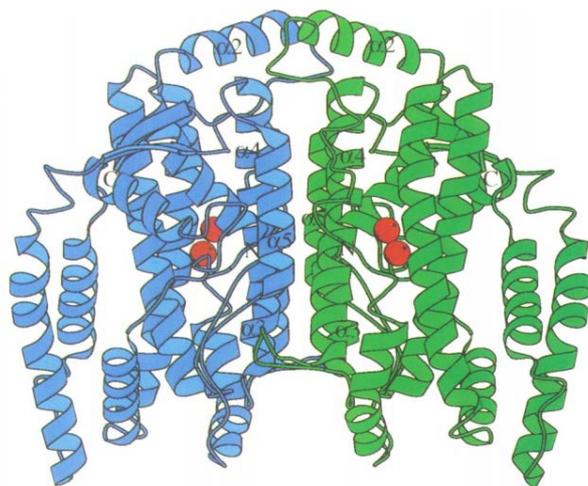


Modelling Metalloproteins

(crystal vs. ligand field theories, spin states in biochemistry, accuracy of QM methods, relativistic effects)

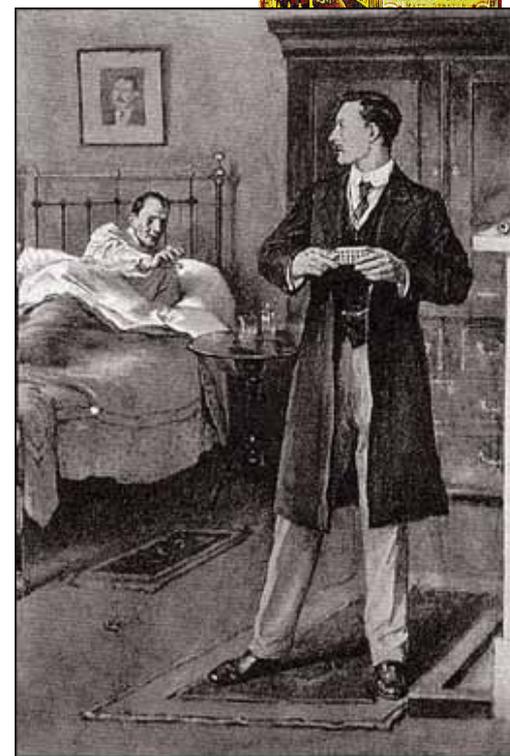
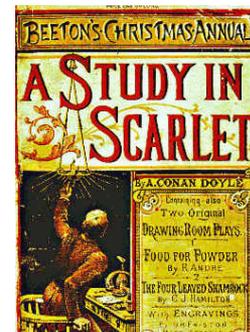
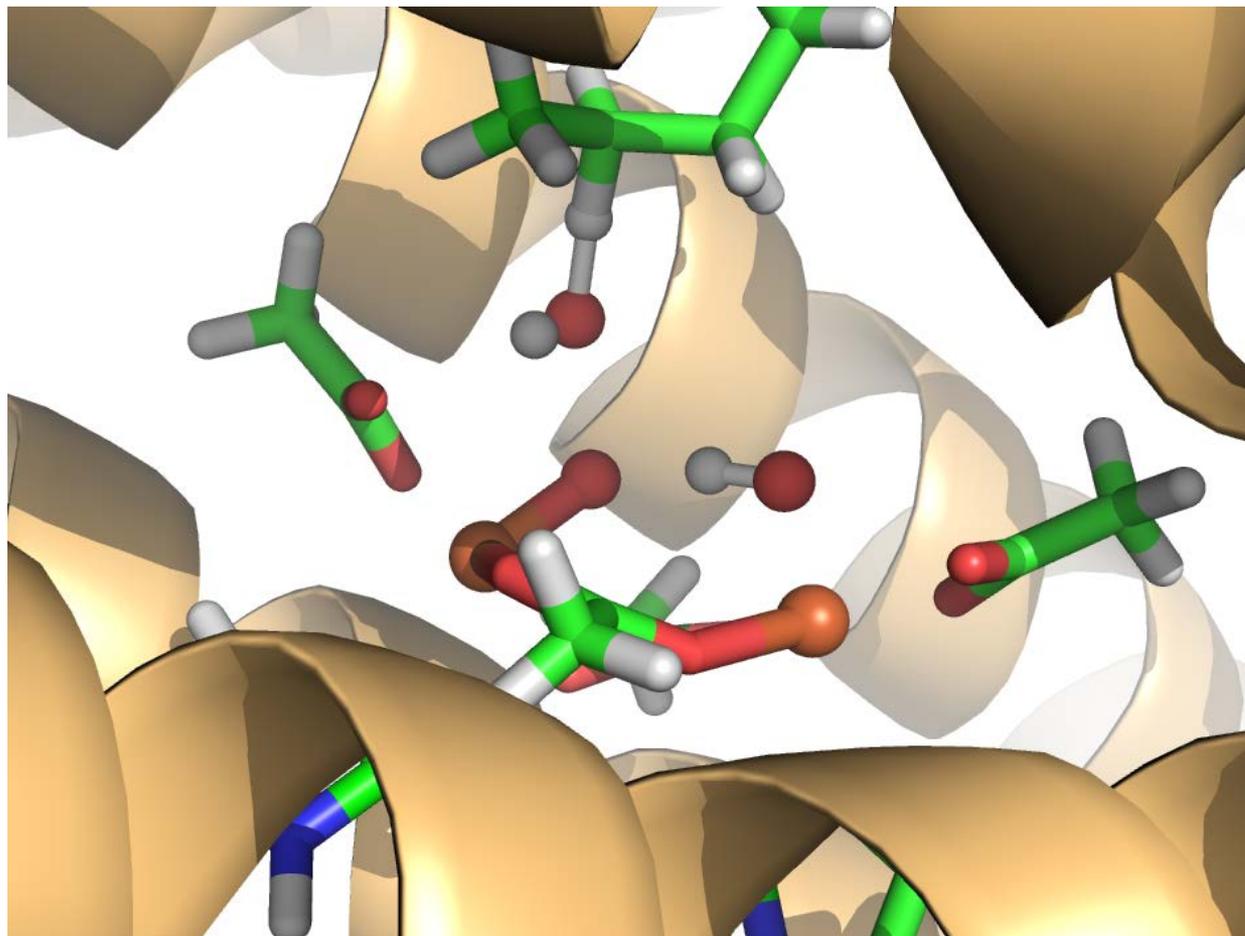
On the example of

Δ^9 Desaturase Reaction Mechanism



(Mystery of) Δ^9 Desaturase Reaction Mechanism

When you have eliminated all which is impossible, then whatever remains, however improbable, must be the truth



NHFe₂ Enzymes

Binuclear Non-Heme Iron Proteins

| reaction type | representative enzyme | catalytic reaction |
|-------------------------------|---|---|
| reversible dioxygen binding | hemerythrin | $[\text{Fe}^{\text{II}}\text{Fe}^{\text{II}}] \xrightleftharpoons{+\text{O}_2} [\text{Fe}^{\text{III}}\text{Fe}^{\text{III}}]-\text{OOH}$ |
| hydroxylation ^a | methane monooxygenase | $[\text{Fe}^{\text{II}}\text{Fe}^{\text{II}}] + \text{CH}_4 \xrightarrow{+\text{O}_2} [\text{Fe}^{\text{III}}\text{Fe}^{\text{III}}] + \text{CH}_3\text{OH}/\text{H}_2\text{O}$ |
| 1-e ⁻ oxidation | ribonucleotide diphosphate reductase | $[\text{Fe}^{\text{II}}\text{Fe}^{\text{II}}] + \text{Tyr} \xrightarrow{+\text{O}_2} [\text{Fe}^{\text{III}}\text{Fe}^{\text{III}}] + \text{Tyr}^\bullet$ |
| desaturation | stearoyl-acyl carrier protein Δ^9 -desaturase | $[\text{Fe}^{\text{II}}\text{Fe}^{\text{II}}] + \text{stearoyl ACP} \xrightarrow{+\text{O}_2} [\text{Fe}^{\text{III}}\text{Fe}^{\text{III}}] + \text{oleoyl ACP}$ |
| hydrolysis of phosphate ester | purple acid phosphatase | $[\text{Fe}^{\text{II}}\text{Fe}^{\text{III}}] + \text{ROHPO}_3 \xrightarrow{+\text{H}_2\text{O}} [\text{Fe}^{\text{II}}\text{Fe}^{\text{III}}] + \text{H}_3\text{PO}_4$ |
| NADH peroxidation | rubrerythrin | $[\text{Fe}^{\text{II}}\text{Fe}^{\text{II}}] + \text{H}_2\text{O}_2 \xrightarrow{+\text{O}_2} [\text{Fe}^{\text{III}}\text{Fe}^{\text{III}}]$ |
| ferroxidation | ferritin | $[\text{Fe}^{\text{II}}\text{Fe}^{\text{II}}] \xrightarrow{+\text{O}_2} [\text{Fe}^{\text{III}}\text{Fe}^{\text{III}}]$ |

taken from: Solomon, E. I.; Brunold, T. C.; Davis, M. I.; Kemsley, J. N.; Lee, S. K.; Lehnert, N.; Neese, F.; Skulan, A. J.; Yang, Y. S.; Zhou, J. *Chem. Rev.* **2000**, *100*, 235-349.

Oxygen intermediates **P**, **P'**, **Q**, and **X** observed in ribonucleotide reductase (RR), Δ^9 desaturase (D9D) and methane monooxygenase (MMO), toluene/o-xylene monooxygenase, toluene 4-monooxygenase

S.J. Lippard, R. A. Friesner, E. I. Solomon, J. D. Lipscomb, L. Que, Jr., ...



Δ^9 Desaturase

- one of the most important enzymes in the fatty-acid metabolism of plants
- catalyzes the oxidation of stearic acid



- the enzyme function is restored by a two-electron reduction mediated by ferredoxin

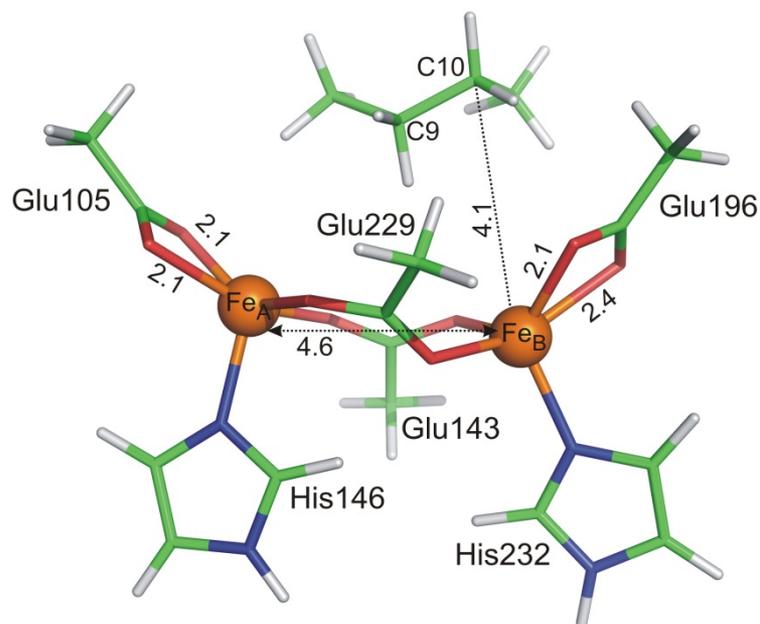
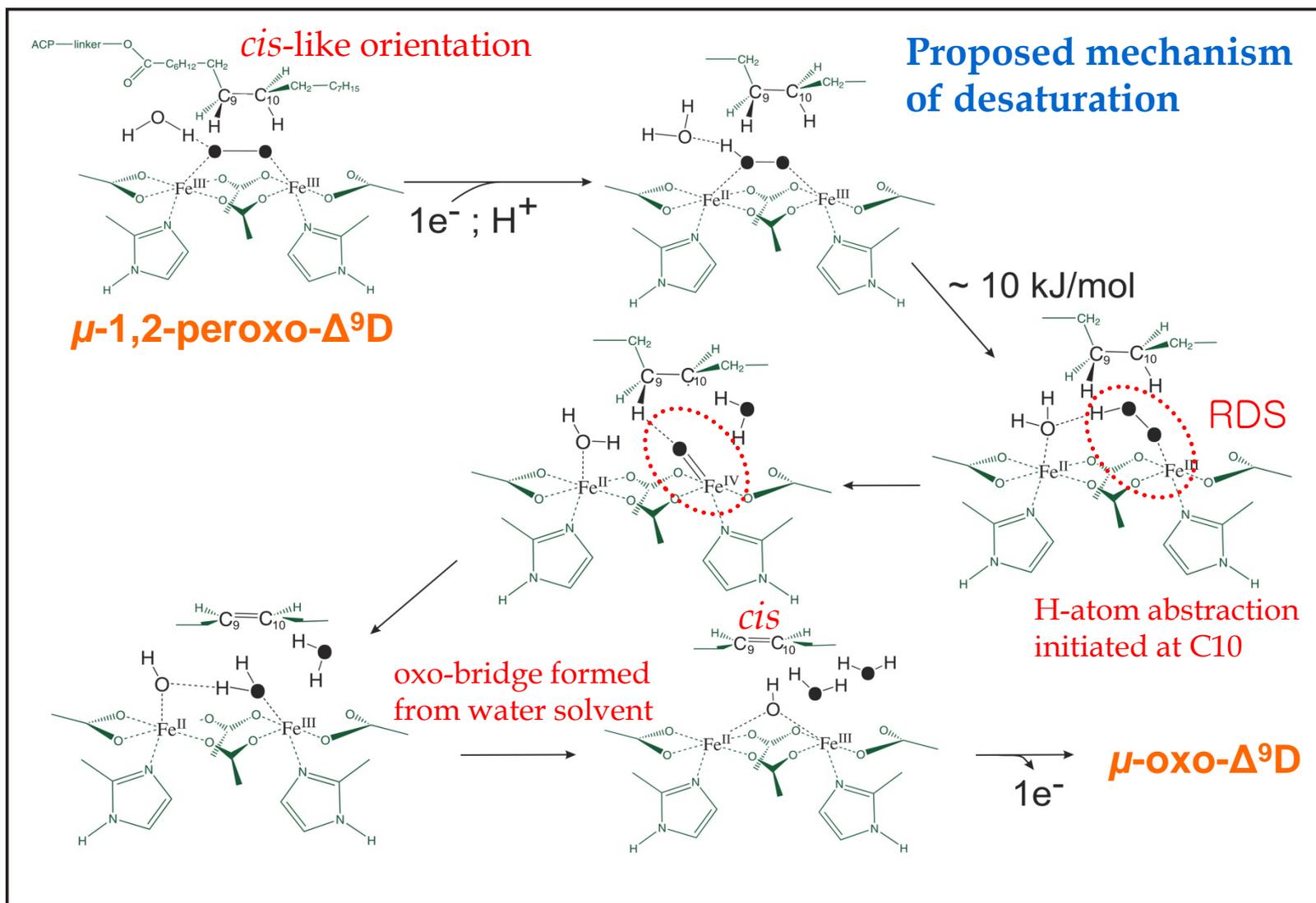


Figure: Quantum system for the reduced [Δ^9 D...substrate] complex (distances in Å).

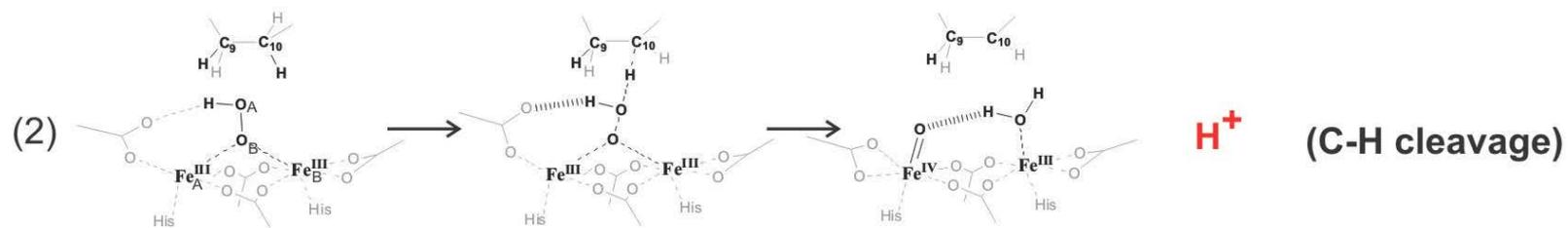
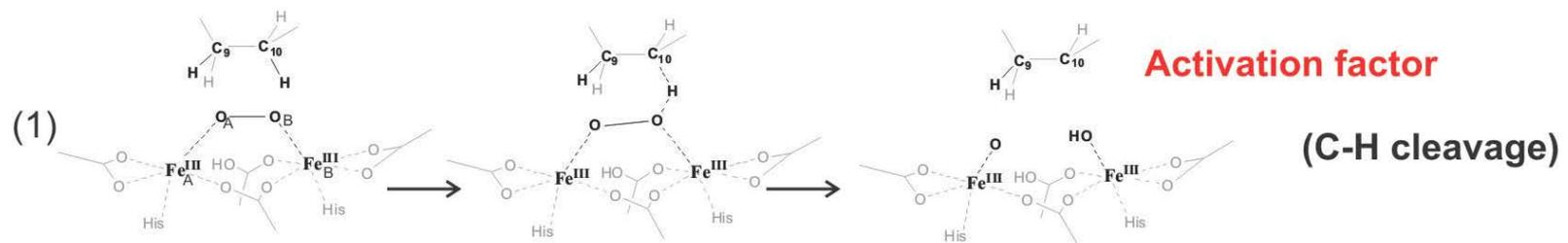


Reaction mechanism of Δ^9 -Desaturase (working +e⁻ version)

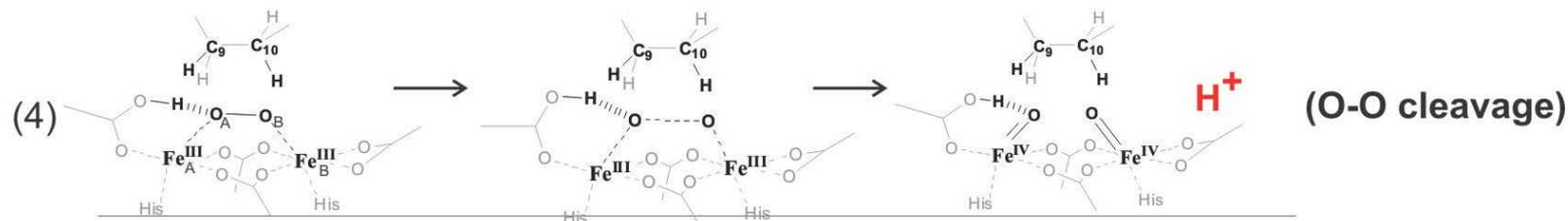


Reaction Pathways Studied

O-O vs. C-H cleavage; activation factors: -, H^+ , H_2O



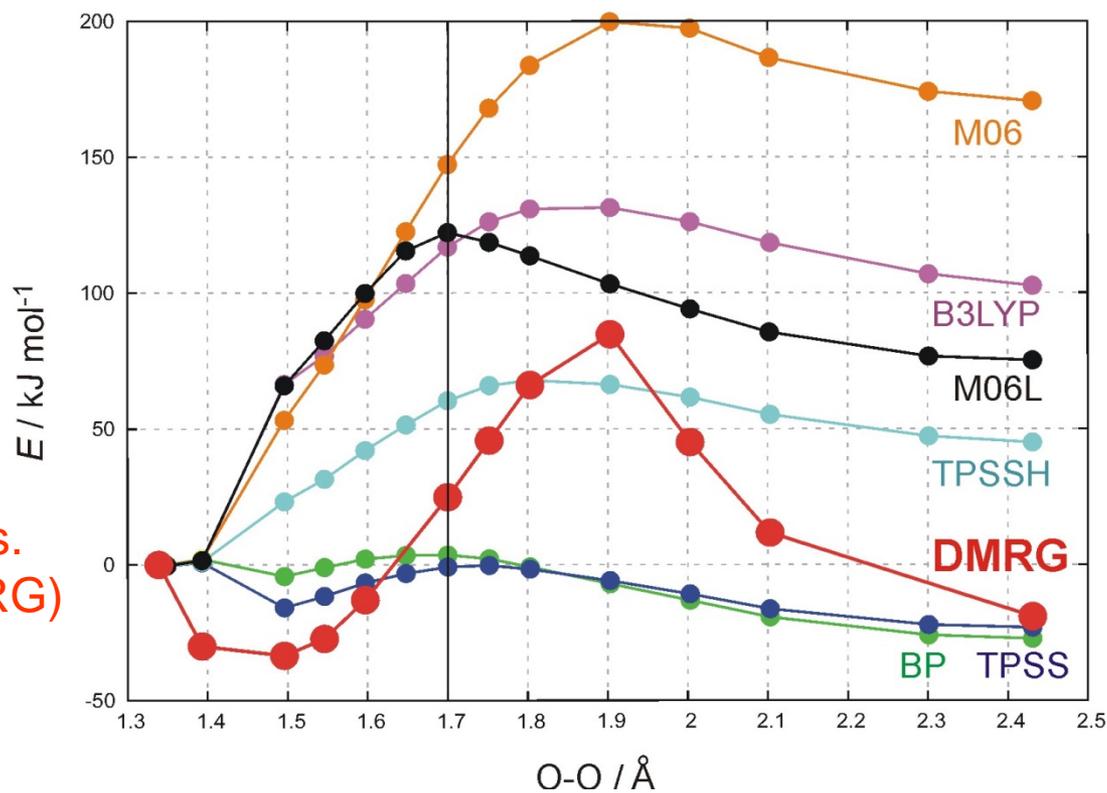
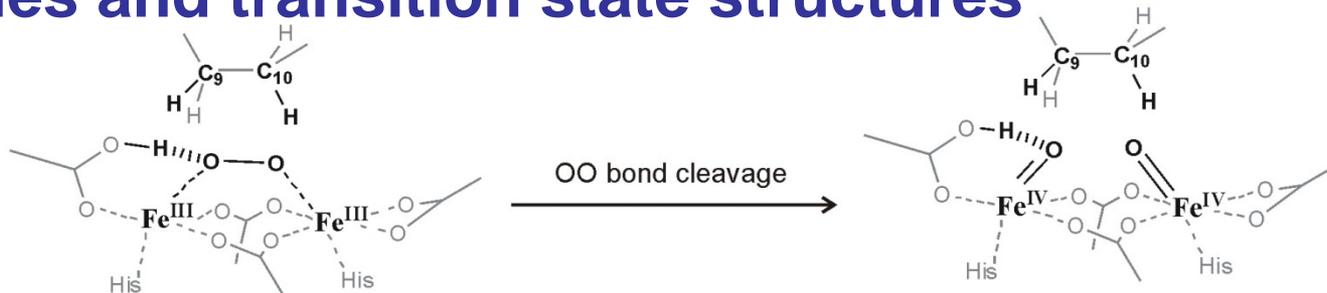
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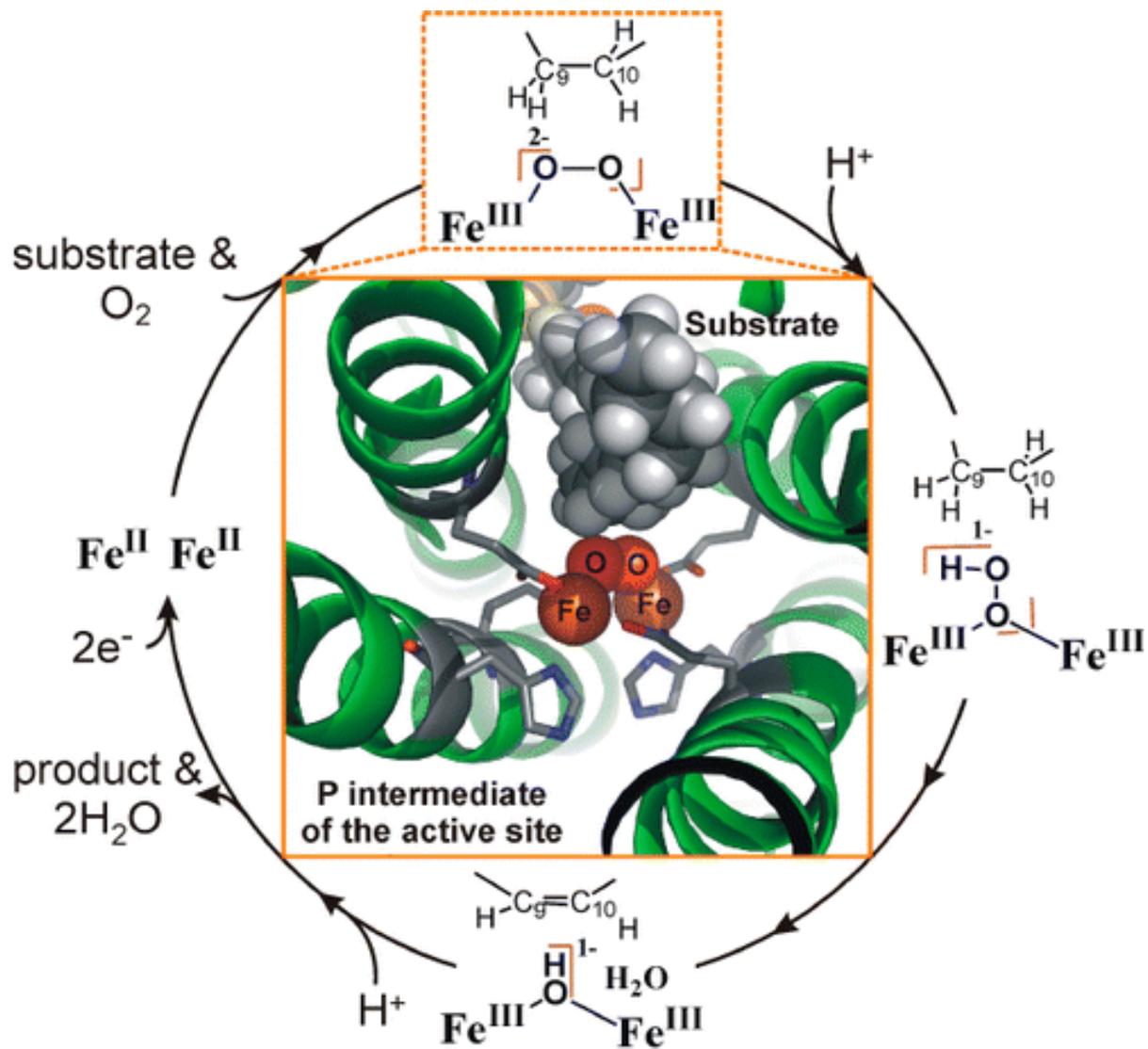


Difference between the DMRG-CASPT2 and DFT equilibrium geometries and transition state structures



Peroxo (DFT) vs. superoxo- (DMRG) structures





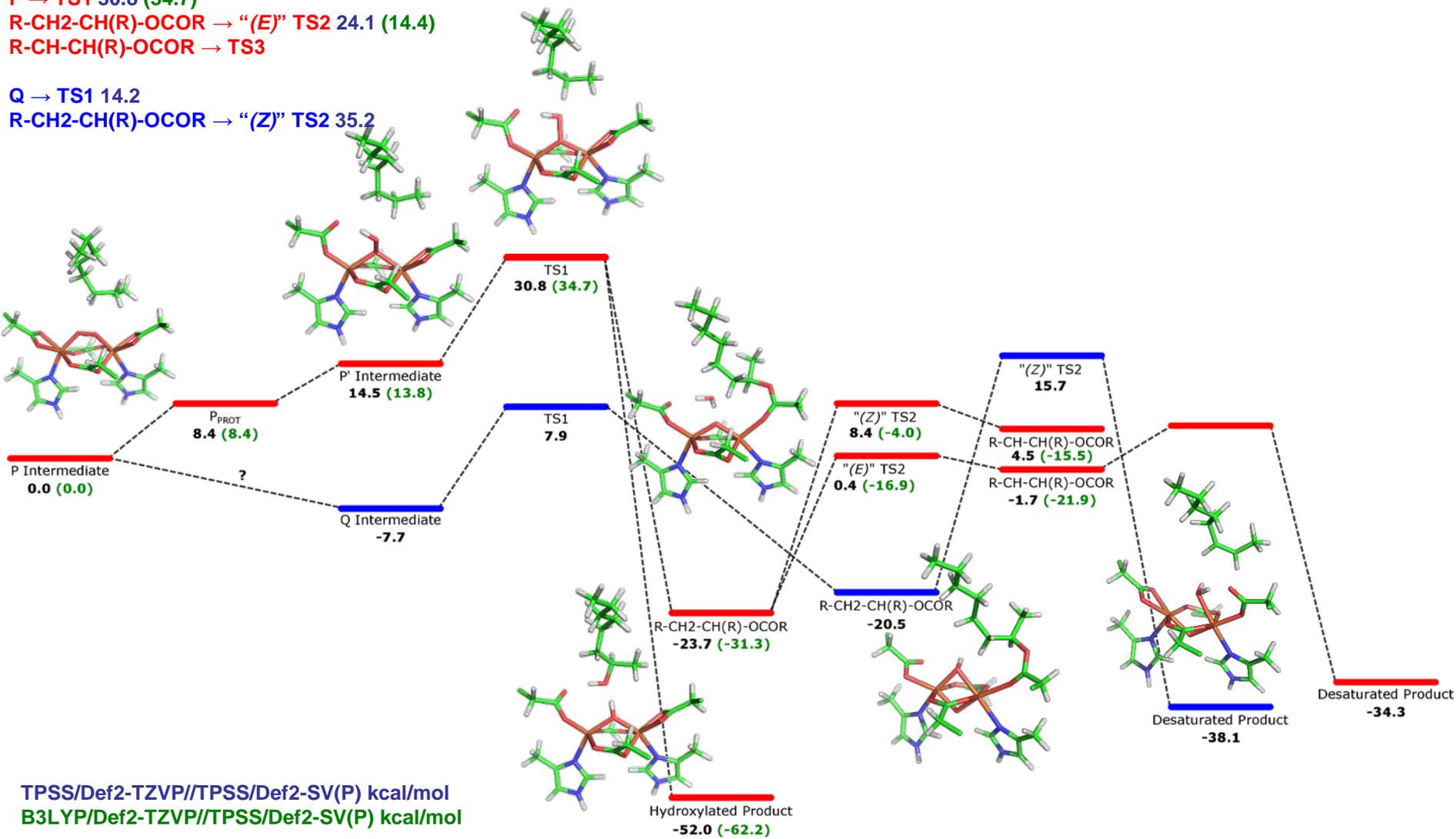
P → TS1 30.8 (34.7)

R-CH₂-CH(R)-OCOR → "(E)" TS2 24.1 (14.4)

R-CH-CH(R)-OCOR → TS3

Q → TS1 14.2

R-CH₂-CH(R)-OCOR → "(Z)" TS2 35.2



Towards 'Functional Metallopeptides': En Route to Disentangle the Catalytic Power of Metalloproteins

