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Electronic Supporting Information

Computational studies of DNA base repair mechanisms by nonheme iron dioxygenases: Selective epoxidation and hydroxylation pathways

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QM/MM calculations

Table S1. Absolute energies (Hartree) for the minima structures in the oxygen activation path catalyzed by snapshot Sn_{300} . The superoxo (**Reoo**), peracid (**Re**_{Cyc}) and ferryl oxo (**I**_P) intermediates are all shown alongside the single stable dioxygen species (**Reoo***). The energies are broken down into separate MM and QM contributions as well as the combined total energies.

	$QM_Turbomole$	$MM_{\text{DL}_\text{poly}}$	
³ R _{oo} *	-3306.886852	-103.979836	-3410.866688
³ R _{oo}	-3306.89148	-103.986474	-3410.877954
⁵ Roo	-3306.912856	-103.980312	-3410.893168
⁷ R ₀₀	-3306.913601	-103.980158	-3410.893759
³ Re _{Cyc}	-3306.94486	-103.993571	-3410.938431
⁵ Re _{Cyc}	-3306.965479	-104.001547	-3410.967026
⁷ Re _{Cyc}	-3306.896847	-103.983259	-3410.880106
³ І _Р	-3306.963497	-103.995344	-3410.95884
⁵ l _P	-3306.976063	-103.999213	-3410.975276
7 _P	-3306.98011	-103.978433	-3410.958543

Table S2. Absolute energies (Hartree) for the minima structures in the oxygen activation path catalyzed by snapshot Sn_{400} . The superoxo (**Reoo**), peracid (**Re**_{Cyc}) and ferryl oxo (**I**_P) intermediates are all shown. The energies are broken down into separate MM and QM contributions as well as the combined total energies.

	$QM_{Turbomole}$	$MM_{\text{DL}_\text{poly}}$	
³ R _{oo}	-3306.953522	-103.765796	-3410.719318
⁵Roo	-3306.972756	-103.761067	-3410.733823
⁷ R _{oo}	-3306.973599	-103.761283	-3410.734882
³ Re _{Cyc}	-3307.002789	-103.777868	-3410.780657
⁵ Re _{Cyc}	-3307.021458	-103.77596	-3410.797417
⁷ Re _{Cyc}	-3306.964413	-103.760432	-3410.724846
³ I _P	-3307.050834	-103.754474	-3410.805308
⁵ l _P	-3307.062671	-103.752004	-3410.814675
71 _P	-3306.985073	-103.722586	-3410.707659

Table S3. Absolute energies (Hartree) for the minima structures in the oxygen activation path catalyzed by snapshot Sn_{500} . The superoxo (**Reoo**), peracid (**Re**_{Cyc}) and ferryl oxo (**I**_P) intermediates are all shown. The energies are broken down into separate MM and QM contributions as well as the combined total energies.

	$QM_Turbomole$	MM_{DL_poly}	QMMM _{Total}
³ R ₀₀	-3306.936762	-103.894914	-3410.831676
⁵ R _{oo}	-3306.973147	-103.870705	-3410.843852
⁷ R ₀₀	-3306.962929	-103.880245	-3410.843175
³ Re _{Cyc}	-3306.940199	-103.874952	-3410.815151
⁵ Re _{Cyc}	-3307.039706	-103.878672	-3410.918378
⁷ Re _{Cyc}	-3306.896832	-103.983277	-3410.880109
³ I _P	-3307.041241	-103.870006	-3410.911247
5 1 P	-3307.048999	-103.869045	-3410.918044
71 _P	-3307.047534	-103.868693	-3410.916227

Table S4. Grouped spin populations for the minima structures in the oxygen activation path catalyzed by snapshot Sn₃₀₀. The superoxo (Reoo), peracid (Recyc) and ferryl oxo (I_P) intermediates are all shown alongside the single stable dioxygen species (Reoo*). The coordinating amino acids, iron co-factor as well as the distal (O₁) and proximal (O₂) oxygens of the bound dioxygen are all considered. The spin densities on the α KG co-enzyme is split into two groups for both the succinate (suc) and CO₂ components.

	Fe	01	O ₂	CO ₂	suc	HIS ₁₃₁	His ₁₈₇	ARG	ASP
³ R ₀₀ *	3.67	-0.92	-0.93	0.06	0.01	0.04	0.05	-0.01	0.02
³ R ₀₀	2.98	-0.54	-0.62	0.07	0.04	-0.01	0.05	-0.01	0.05
⁵ R _{oo}	4.19	-0.30	-0.53	0.16	0.10	0.14	0.10	-0.01	0.16
⁷ R ₀₀	4.18	0.56	0.61	0.17	0.07	0.13	0.11	0.01	0.18
³ Re _{Cyc}	2.04	0.02	-0.02	0.01	-0.05	0.01	0.04	0.01	-0.06
⁵ Re _{Cyc}	3.75	0.06	0.01	0.00	0.03	0.07	0.03	0.00	0.04
⁷ Re _{Cyc}	4.29	0.20	0.12	0.52	0.49	0.12	0.10	0.00	0.15
³ І _Р	2.84	-0.59	-0.24	0.01	-0.05	0.02	0.09	0.02	-0.10
⁵ l _P	4.21	-0.55	0.06	0.00	0.09	0.11	0.11	0.05	-0.08
7 I P	4.20	1.01	0.32	0.01	0.12	0.08	0.12	0.00	0.15

Table S5. Grouped spin populations for the minima structures in the oxygen activation path catalyzed by snapshot Sn₄₀₀. The superoxo (Reoo), peracid (Rec_{yc}) and ferryl oxo (I_P) intermediates. The coordinating amino acids, iron co-factor as well as the distal (O₁) and proximal (O₂) oxygens of the bound dioxygen are all considered. The spin densities on the α KG co-enzyme is split into two groups for both the succinate (suc) and CO₂ components.

	Fe	01	O ₂	CO ₂	suc	HIS ₁₃₁	His ₁₈₇	ARG	ASP
³ R ₀₀	3.20	-0.66	-0.70	0.07	0.02	0.00	0.04	-0.01	0.03
⁵R _{oo}	4.18	-0.30	-0.50	0.16	0.09	0.13	0.10	0.00	0.14
⁷ R ₀₀	4.17	0.61	0.61	0.14	0.06	0.12	0.11	0.01	0.17
³ Re _{Cyc}	1.99	0.05	-0.02	0.00	-0.05	0.03	0.04	0.00	-0.05
⁵ Re _{Cyc}	3.76	0.07	0.01	0.01	0.03	0.07	0.03	0.00	0.03
⁷ Re _{Cyc}	4.17	0.61	0.61	0.14	0.06	0.12	0.11	0.01	0.17
³І _Р	2.84	-0.59	-0.24	0.01	-0.05	0.02	0.09	0.02	-0.10
⁵ l _P	4.20	-0.54	0.02	0.00	0.10	0.13	0.12	0.05	-0.08
7 P	4.46	0.11	-0.01	0.93	0.25	0.10	0.04	0.04	0.06

Table S5. Grouped spin populations for the minima structures in the oxygen activation path catalyzed by snapshot Sn_{500} . The superoxo (Reoo), peracid (Recyc) and ferryl oxo (IP) intermediates. The coordinating amino acids, iron co-factor as well as the distal (O₁) and proximal (O₂) oxygens of the bound dioxygen are all considered. The spin densities on the α KG co-enzyme is split into two groups for both the succinate (suc) and CO₂ components.

	Fe	O 1	O ₂	CO ₂	suc	HIS ₁₃₁	His ₁₈₇	ARG	ASP
³ R ₀₀	3.32	-0.76	-0.71	0.07	0.01	0.01	0.05	0.00	0.00
⁵R _{oo}	4.19	-0.31	-0.52	0.18	0.07	0.13	0.09	-0.01	0.19
⁷ R ₀₀	4.18	0.52	0.64	0.15	0.07	0.12	0.10	0.01	0.20
³ Re _{Cyc}	1.12	0.05	0.09	0.44	0.35	0.00	-0.03	0.00	-0.02
⁵ Re _{Cyc}	3.76	0.06	0.01	0.01	0.02	0.07	0.03	0.00	0.04
⁷ Re _{Cyc}	4.32	0.16	0.09	0.55	0.49	0.12	0.09	0.00	0.16
³I _Р	1.37	0.63	0.04	0.00	-0.03	0.04	-0.03	0.00	-0.02
⁵ l _P	4.14	-0.29	-0.29	0.02	0.08	0.09	0.11	0.01	0.14
⁷ I _P	4.18	1.06	0.25	0.00	0.12	0.09	0.09	0.00	0.21

Table S6. Relative energies (kcal mol⁻¹) for the reaction landscape in the oxygen activation path catalyzed by snapshot Sn_{300} . The superoxo (**Re**₀₀), peracid (**Re**_{Cyc}) and ferryl oxo (**I**_P) intermediates are all shown alongside the transition state for the initial attack of the superoxo on the beta carbon of the co-enzyme (**TS**_{0,C}) and the barrier for the OO cleavage of the peracid (**TS**_{C,C}). The energies are given for all possible spin states.

	S = 1	S = 2	S = 3
Roo	9.92	0.37	0.00
TS _{o,c}	21.15	9.31	10.77
Re _{cyc}	-28.03	-45.98	8.57
TS _{o,o}	-26.80	-41.26	24.25
IP	-40.84	-51.15	-40.65

Table S7. Relative energies (kcal mol⁻¹) for the reaction landscape in the oxygen activation path catalyzed by snapshot Sn_{400} . The superoxo (**Re**_{OO}), peracid (**Re**_{Cyc}) and ferryl oxo (**I**_P) intermediates are all shown alongside the transition state for the initial attack of the superoxo on the beta carbon of the co-enzyme (**TS**_{O,C}) and the barrier for the OO cleavage of the peracid (**TS**_{C,C}). The energies are given for all possible spin states.

Sn400	S = 1	S = 2	S = 3
R _{oo}	9.77	0.66	0.00
TS _{o,c}	19.68	13.53	13.14
Re _{cyc}	-28.72	-39.24	6.30
TS _{o,o}	-19.98	-32.73	28.97
I _P	-44.19	-50.07	17.08

Table S8. Relative energies (kcal mol⁻¹) for the reaction landscape in the oxygen activation path catalyzed by snapshot Sn_{500} . The superoxo (**Reoo**), peracid (**Re**_{Cyc}) and ferryl oxo (**I**_P) intermediates are all shown alongside the transition state for the initial attack of the superoxo on the beta carbon of the co-enzyme (**TS**_{0,C}) and the barrier for the OO cleavage of the peracid (**TS**_{C,C}). The energies are given for all possible spin states.

Sn500	S = 1	S = 2	S = 3
Roo	7.22	-0.42	0.00
TS _{o,c}	21.23	9.33	7.02
Re _{Cyc}	17.59	-47.19	-23.18
TS _{o,o}	24.22	-38.75	-7.49
IP	-42.72	-46.98	-45.84



Fig. S1. Comparative orbital occupations and important bond lengths for the triplet (³Roo), quintet (⁵Roo) and septet (⁷Roo) superoxo QM species.



Fig. S2. Graphical representation of changes in the QM region at different minima point across the low energy quintet reaction pathways for oxygen activation. Important bond lengths for the superoxo (⁵Roo), peracid (⁵R_{Cyc}) and ferryl oxo (⁵I_P) species are all shown.



Fig. S3. Optimized geometries of ${}^{5}\text{Re} \{{}^{3}\text{R}_{e}\}$ for epoxidation reaction by iron(IV)-oxo species with alkylated Cytosine. All bond lengths are given in angstroms and the value of the imaginary frequency in the transition state in cm⁻¹.



Fig. S4. Optimized geometries of ${}^{5}TS_{E} \{ {}^{3}TS_{E} \}$ for epoxidation reaction by iron(IV)-oxo species with alkylated Cytosine. All bond lengths are given in angstroms and the value of the imaginary frequency in the transition state in cm⁻¹.



Fig. S5. Optimized geometries of ${}^{5}IM_{E} \{ {}^{3}IM_{E} \}$ for epoxidation reaction by iron(IV)-oxo species with alkylated Cytosine. All bond lengths are given in angstroms and the value of the imaginary frequency in the transition state in cm⁻¹.



Fig. S6. Optimized geometries of ${}^{5}TS_{rc}$ { ${}^{3}TS_{rc}$ } for epoxidation reaction by iron(IV)-oxo species with alkylated Cytosine. All bond lengths are given in angstroms and the value of the imaginary frequency in the transition state in cm⁻¹.



Fig. S7. Optimized geometries of ${}^{5}P_{E} \{ {}^{3}P_{E} \}$ for epoxidation reaction by iron(IV)-oxo species with alkylated Cytosine. All bond lengths are given in angstroms and the value of the imaginary frequency in the transition state in cm⁻¹.



Fig. S8. Optimized geometries of ⁵Re (left) and ⁵TS_E (right) for epoxidation reaction by iron(IV)oxo species with alkylated Adenine. All bond lengths are given in angstroms and the value of the imaginary frequency in the transition state in cm⁻¹.



Fig. S9. Optimized geometries of ⁵Re (left) and ⁵TS_E (right) for hydroxylation reaction by iron(IV)-oxo species with alkylated Guanine. All bond lengths are given in angstroms and the value of the imaginary frequency in the transition state in cm⁻¹.



Fig. S10. Optimized geometries of ${}^{5}\text{Re} \{{}^{3}\text{Re}\}$ for hydroxylation reaction by iron(IV)-oxo species with alkylated Cytosine. All bond lengths are given in angstroms and the value of the imaginary frequency in the transition state in cm⁻¹.



Fig. S11. Optimized geometries of ${}^{5}TS_{H} \{ {}^{3}TS_{H} \}$ for hydroxylation reaction by iron(IV)-oxo species with alkylated Cytosine. All bond lengths are given in angstroms and the value of the imaginary frequency in the transition state in cm⁻¹.



Fig. S12. Optimized geometries of ${}^{5}IM_{H} \{ {}^{3}IM_{H} \}$ for hydroxylation reaction by iron(IV)-oxo species with alkylated Cytosine. All bond lengths are given in angstroms and the value of the imaginary frequency in the transition state in cm⁻¹.



Fig. S13. Optimized geometries of ${}^{5}TS_{reb}$ { ${}^{3}TS_{reb}$ } for hydroxylation reaction by iron(IV)-oxo species with alkylated Cytosine. All bond lengths are given in angstroms and the value of the imaginary frequency in the transition state in cm⁻¹.



Fig. S14. Optimized geometries of ${}^{5}P_{H} \{ {}^{3}P_{H} \}$ for hydroxylation reaction by iron(IV)-oxo species with alkylated Cytosine. All bond lengths are given in angstroms and the value of the imaginary frequency in the transition state in cm⁻¹.



Fig. S15. Optimized geometries of ⁵Re (left) and ⁵TS_H (right) for hydroxylation reaction by iron(IV)-oxo species with alkylated Guanine. All bond lengths are given in angstroms and the value of the imaginary frequency in the transition state in cm⁻¹.



Fig. S16. Optimized geometries of ${}^{5}\text{Re}$ (left) and ${}^{5}\text{TS}_{\text{H}}$ (right) for hydroxylation reaction by iron(IV)-oxo species with alkylated Adenine. All bond lengths are given in angstroms and the value of the imaginary frequency in the transition state in cm⁻¹.

	E (LACVP)	ZPE	G	E (LACV3P+*)
	(au)	(kcal/mol)	(au)	(au)
¹ Re	-1767.570763	0.378753	-1767.2541	-1768.273169
³ Re	-1767.619247	0.379556	-1767.3014	-1768.323704
⁵Re	-1767.628520	0.378162	-1767.3157	-1768.338381
⁷ Re	-1767.610836	0.376252	-1767.2989	-1768.319167
¹ TS _E	-1767.563873	0.378413	-1767.2471	-1768.262860
³ TS _E	-1767.593207	0.378456	-1767.2764	-1768.293683
⁵TS⊧	-1767.613687	0.376397	-1767.3026	-1768.325039
⁷ TS _E	-1767.600822	0.375990	-1767.2902	-1768.311652
¹ IM _E	-1767.591178	0.378735	-1767.2725	-1768.297585
³ IM _E	-1767.603585	0.377239	-1767.2852	-1768.312414
⁵ IM _E	-1767.614469	0.374670	-1767.3026	-1768.325723
⁷ IM _E	-1767.641175	0.376734	-1767.3297	-1768.358120
¹ TS _{rc}	-1767.597769	0.378468	-1767.2799	-1768.309723
³ TS _{rc}	-1767.578468	0.374595	-1767.2913	-1768.300033
⁵ TS _{rc}	-1767.629027	0.375842	-1767.3190	-1768.349898
⁷ TS _{rc}	-1767.571735	0.374394	-1767.2691	-1768.287445
¹ P _E	-1767.614656	0.380195	-1767.2947	-1768.328372
³ P _E	-1767.612720	0.378152	-1767.2960	-1768.331372
⁵ P _E	-1767.636257	0.376903	-1767.3246	-1768.359487
⁷ P _E	-1767.635387	0.375709	-1767.3220	-1768.356517

 Table S9. Absolute energies of all epoxidation structures described in this work.

	1					0	0		0	0
1_	ρ _{Fe}	ρLigands	ρο	PCytosine	ρ _{Total}	QFe	QLigands	Qo	QCytosine	QTotal
⁺ Re	0.69	-0.06	-0.64	0.00	0.00	0.62	-0.27	-0.33	-0.02	0.00
³ Re	1.10	-0.05	0.96	0.00	2.00	0.60	-0.25	-0.33	-0.02	0.00
⁵Re	2.95	0.33	0.72	0.00	4.00	0.74	-0.39	-0.34	-0.02	0.00
⁷ Re	3.98	0.63	1.39	0.00	6.00	0.90	-0.51	-0.36	-0.03	0.00
¹ TS _E	0.81	-0.05	-0.34	-0.41	0.00	0.67	-0.40	-0.45	0.18	0.00
³ TS _E	2.74	0.23	-0.52	-0.45	2.00	0.78	-0.55	-0.44	0.22	0.00
⁵ TS _E	3.64	0.40	0.22	-0.25	4.00	0.82	-0.56	-0.46	0.20	0.00
⁷ TS _E	3.98	0.54	1.16	0.32	6.00	0.86	-0.61	-0.47	0.21	0.00
¹ IM _E	0.99	-0.06	-0.01	-0.93	0.00	0.71	-0.31	-0.57	0.17	0.00
³ IM _E	2.83	0.22	-0.09	-0.96	2.00	0.81	-0.49	-0.52	0.20	0.00
⁵IM _E	4.02	0.56	0.30	-0.89	4.00	0.93	-0.56	-0.61	0.24	0.00
⁷ IM _E	4.04	0.60	0.36	1.01	6.00	0.93	-0.54	-0.60	0.21	0.00
¹ TS _{rc}	-0.06	-0.01	-0.06	0.13	0.00	0.60	-0.70	-0.48	0.57	0.00
³ TS _{rc}	1.98	0.11	0.06	-0.15	2.00	0.69	-0.80	-0.47	0.59	0.00
⁵ TS _{rc}	3.81	0.27	0.11	-0.19	4.00	0.79	-0.84	-0.50	0.55	0.00
⁷ TS _{rc}	4.06	0.66	0.24	1.04	6.00	0.99	-0.49	-0.58	0.08	0.00
	ρ_{Fe}	$\rho_{Ligands}$	$\rho_{Product}$	ρ_{Total}		Q _{Fe}	QLigands	QProduct	Q _{Total}	
¹ P _E	0.01	-0.01	0.00	0.00		0.58	-0.70	0.12	0.00	
³ P _E	1.95	0.04	0.01	2.00		0.67	-0.76	0.09	0.00	
⁵ P _E	3.74	0.25	0.01	4.00		0.83	-0.91	0.08	0.00	
⁷ P _E	4.03	0.59	1.38	6.00		0.93	-0.53	-0.40	0.00	

Table S10. Group spin densities and charges of optimized UB3LYP/B1 geometries in epoxidation reaction.

	ρ _{Fe}	PLigands	ρο	P Cytosine	PTotal	O _{Fe}	OLigands	O ₀	O _{Cytosine}	O _{Total}
¹ Re	0.70	-0.08	-0.61	0.00	0.00	0.62	-0.27	-0.33	-0.02	0.00
³ Re	1.16	-0.07	0.91	0.00	2.00	0.43	-0.21	-0.16	-0.06	0.00
⁵Re	3.10	0.26	0.64	0.00	4.00	0.39	-0.21	-0.13	-0.05	0.00
⁷ Re	4.13	0.55	1.30	0.01	6.00	0.63	-0.42	-0.17	-0.04	0.00
¹ TS _E	0.85	-0.07	-0.34	-0.44	0.00	0.62	-0.40	-0.21	-0.01	0.00
³ TS _E	2.81	0.18	-0.52	-0.47	2.00	0.70	-0.53	-0.19	0.02	0.00
⁵TS₌	3.77	0.31	0.19	-0.27	4.00	0.59	-0.40	-0.21	0.01	0.00
⁷ TS _E	4.15	0.44	1.06	0.34	6.00	0.63	-0.44	-0.24	0.06	0.00
¹ IM _E	1.03	-0.07	-0.01	-0.95	0.00	0.71	-0.34	-0.31	-0.05	0.00
³ IM _E	2.91	0.16	-0.10	-0.98	2.00	0.73	-0.44	-0.21	-0.07	0.00
⁵IM _E	4.18	0.46	0.27	-0.91	4.00	0.77	-0.45	-0.34	0.01	0.00
⁷ IM _E	4.19	0.49	0.30	1.02	6.00	0.78	-0.42	-0.33	-0.03	0.00
¹ TS _{rc}	-0.07	0.00	-0.06	0.13	0.00	0.67	-0.74	-0.22	0.29	0.00
³ TS _{rc}	2.01	0.09	0.05	-0.15	2.00	0.72	-0.83	-0.17	0.29	0.00
⁵ TS _{rc}	3.90	0.21	0.08	-0.19	4.00	0.75	-0.81	-0.21	0.27	0.00
⁷ TS _{rc}	4.22	0.55	0.18	1.05	6.00	0.76	-0.25	-0.19	-0.32	0.00
	ρ_{Fe}	$\rho_{Ligands}$	$\rho_{Product}$	ρ_{Total}		Q _{Fe}	QLigands	QProduct	Q _{Total}	
¹ P _E	0.02	-0.01	0.00	0.00		0.67	-0.96	0.29	0.00	
³ P _E	1.97	0.02	0.01	2.00		0.77	-0.98	0.21	0.00	
⁵PE	3.81	0.20	-0.01	4.00		0.71	-0.81	0.10	0.00	
⁷ P _E	4.19	0.49	1.33	6.00		0.66	-0.35	-0.31	0.00	

Table S11. Group spin densities and charges of single point UB3LYP/B2 calculations onUB3LYP/B1 optimized geometries for epoxidation reaction.



Table S12. All bond lengths in angstroms and obtained from geometry optimizations of epoxidation reaction in Gaussian at UB3LYP/B1.

	Fe-O1	Fe-O2	Fe-O3	Fe-O4	Fe-N1	Fe-N2	01-C1	01-C2
¹ Re	1.70326	2.03158	2.11876	1.91060	1.95118	2.06322	3.20627	3.77641
³Re	1.65250	2.04223	2.12473	1.91493	1.94702	2.07739	3.22303	3.85795
⁵Re	1.64927	2.28205	2.13275	1.92770	2.08712	2.06879	3.25171	3.72760
⁷ Re	1.93592	2.36542	2.09256	1.93098	2.10234	2.17470	3.14594	3.85191
¹ TS _E	1.75394	2.02674	2.20290	1.92225	1.94070	2.06782	1.92903	2.64566
³ TS _E	1.74732	2.03843	2.20072	1.93351	1.96169	2.06530	1.94638	2.67336
⁵TS⊧	1.71244	2.34825	2.13294	1.93837	2.12598	2.17696	2.13794	2.69638
⁷ TS _E	1.90480	2.31095	2.15125	1.94079	2.12895	2.22071	2.17446	2.64522
¹ IM _E	1.85849	2.00346	2.12077	1.92395	1.96246	2.02423	1.40860	2.49953
³ IM _E	1.83218	2.36369	2.10034	1.96216	2.04779	2.02685	1.42300	2.50641
⁵IM _E	1.84631	2.27555	2.15312	1.93699	2.12654	2.20339	1.42733	2.42206
⁷ IM _E	1.86934	2.28904	2.14956	1.92761	2.12566	2.19918	1.40732	2.42821
¹ TS _{rc}	1.95220	2.10382	2.20286	2.00338	2.01417	1.98017	1.41033	2.00160
³ TS _{rc}	2.09999	2.24175	2.28532	1.98509	2.14590	2.16792	1.43155	1.94618
⁵ TS _{rc}	2.06541	2.24247	2.29726	1.98792	2.17221	2.17405	1.41456	1.98448
⁷ TS _{rc}	1.95048	2.27279	2.12851	1.91125	2.08059	2.15916	1.52640	1.65565
¹ P _E	2.07269	2.09819	2.23115	2.00605	2.02209	2.00096	1.50436	1.56659
³ P _E	2.23910	2.03463	2.36329	1.98382	2.11388	2.10897	1.49577	1.55259
⁵ P _E	2.47254	3.17129	1.98886	1.95923	2.12496	2.17782	1.48550	1.54077
⁷ P _E	1.85611	2.28069	2.13706	1.92048	2.14957	2.21130	1.40689	2.43387

	E (LACVP)	ZPE	G	E (LACV3P+*)
	(au)	(kcal/mol)	(au)	(au)
¹ Re	-1731.101759	0.408898	-1730.7538	-1731.771229
³ Re	-1731.164160	0.410127	-1730.8191	-1731.833125
⁵Re	-1731.172770	0.408783	-1730.8268	-1731.847229
⁷ Re	-1731.165185	0.407189	-1730.8238	-1731.837261
¹TS _H	-1731.112278	0.403607	-1730.7723	-1731.772673
³ TS _H	-1731.150624	0.403076	-1730.8109	-1731.813370
⁵TS _H	-1731.157975	0.402022	-1730.8192	-1731.829716
⁷ TS _H	-1731.138744	0.400981	-1730.7990	-1731.808883
¹IM _H	-1731.101672	0.407345	-1730.7552	-1731.771548
³ IM _H	-1731.132768	0.405389	-1730.7929	-1731.810758
⁵IM _H	-1731.151604	0.403100	-1730.8098	-1731.825897
⁷ IM _H	-1731.153942	0.403609	-1730.8150	-1731.838528
¹ TS _{reb}	-1731.149271	0.408708	-1730.7989	-1731.814160
³ TS _{reb}	-1731.164602	0.407192	-1730.8266	-1731.827744
⁵ TS _{reb}	-1731.160190	0.403187	-1730.8197	-1731.832007
⁷ TS _{reb}	-1731.164979	0.405386	-1730.8240	-1731.846534
¹ Рн	-1731.214112	0.411702	-1730.8631	-1731.884443
³ Р _Н	-1731.211195	0.409730	-1730.8606	-1731.880578
⁵ P _H	-1731.234052	0.408664	-1730.8951	-1731.915444
⁷ Рн	-1731.169729	0.406679	-1730.8229	-1731.851548

Table S13. Absolute energies of all structures described in this work for hydroxylation reaction.

	ρ _{Fe}	ρ _{Ligands}	ρο	раснз	ρ_{Total}	Q _{Fe}	QLigands	Qo	Qach3	Q _{Total}
¹ Re	0.80	-0.08	-0.72	0.00	0.00	0.61	-0.18	-0.38	0.94	1.00
³ Re	1.10	-0.06	0.95	0.00	2.00	0.59	-0.10	-0.34	0.85	1.00
⁵Re	2.99	0.32	0.69	0.00	4.00	0.75	-0.25	-0.37	0.87	1.00
⁷ Re	3.99	0.64	1.37	0.00	6.00	0.90	-0.46	-0.37	0.93	1.00
	ρ_{Fe}	$\rho_{Ligands}$	рон	расн2	ρ_{Total}	Q _{Fe}	QLigands	Qoh	Q _{ACH2}	Q _{Total}
¹TSн	0.80	-0.04	-0.37	-0.39	0.00	0.57	-0.09	-0.11	0.63	1.00
^з ТSн	2.77	0.21	-0.51	-0.48	2.00	0.72	-0.28	-0.12	0.68	1.00
⁵TSн	3.70	0.41	0.15	-0.26	4.00	0.83	-0.33	-0.16	0.65	1.00
⁷ TS⊦	4.05	0.56	0.84	0.55	6.00	0.87	-0.40	-0.19	0.72	1.00
¹IM _H	0.94	-0.06	0.10	-0.98	0.00	0.63	-0.14	-0.31	0.82	1.00
³ IM _H	2.02	0.16	0.03	-0.21	2.00	0.72	-0.26	-0.29	0.83	1.00
⁵IM _H	4.05	0.53	0.40	-0.98	4.00	0.91	-0.44	-0.31	0.84	1.00
⁷ IM⊦	4.06	0.58	0.37	0.99	6.00	0.90	-0.40	-0.36	0.86	1.00
¹ TS _{reb}	0.85	-0.05	0.03	-0.84	0.00	0.60	-0.12	-0.23	0.75	1.00
³ TS _{reb}	2.55	0.18	0.00	-0.74	2.00	0.73	-0.32	-0.22	0.81	1.00
⁵ TS _{reb}	4.03	0.48	0.27	-0.78	4.00	0.87	-0.44	-0.30	0.87	1.00
⁷ TS _{reb}	4.11	0.69	0.23	0.96	6.00	0.95	-0.24	-0.25	0.53	1.00
	ρ_{Fe}	$\rho_{Ligands}$	ρ _{ACH2}	ρ_{Total}		Q _{Fe}	QLigands	Q _{ACH2}	Q _{Total}	
¹ Рн	0.00	0.00	0.00	0.00		0.54	-0.44	0.91	1.00	
³Рн	1.98	0.03	-0.01	2.00		0.63	-0.54	0.92	1.00	
⁵P _H	3.96	0.04	0.01	4.00		0.76	-0.63	0.87	1.00	
⁷ Рн	4.09	0.81	1.10	6.00		1.05	-0.24	0.19	1.00	

Table S14. Group spin densities and charges of optimized UB3LYP/B1 geometries forhydroxylation reaction in Gaussian.

	ρ _{Fe}	ρ _{Ligands}	ρο	расн3	ρ_{Total}	Q _{Fe}	QLigands	Qo	Q _{ACH3}	Q _{Total}
¹ Re	0.80	-0.10	-0.70	0.00	0.00	0.51	-0.21	-0.24	0.95	1.00
³ Re	1.17	-0.07	0.90	0.00	2.00	0.45	-0.14	-0.23	0.92	1.00
⁵Re	3.13	0.24	0.63	0.00	4.00	0.48	-0.18	-0.22	0.92	1.00
⁷ Re	4.15	0.56	1.29	0.01	6.00	0.62	-0.37	-0.19	0.94	1.00
	ρ _{Fe}	ρLigands	рон	Расн2	ρ_{Total}	Q _{Fe}	QLigands	Qoh	Qach2	Q _{Total}
¹ TSн	0.83	-0.06	-0.38	-0.40	0.00	0.51	-0.20	0.18	0.52	1.00
³ TSн	2.84	0.16	-0.51	-0.49	2.00	0.61	-0.37	0.19	0.58	1.00
⁵ TS _H	3.83	0.32	0.10	-0.25	4.00	0.62	-0.32	0.18	0.52	1.00
⁷ TS _H	4.21	0.48	0.77	0.55	6.00	0.70	-0.41	0.10	0.62	1.00
	4.21	0.47	0.75	0.57	6.00	0.69	-0.41	0.10	0.63	1.00
¹ IMн	0.97	-0.07	0.10	-0.99	0.00	0.56	-0.17	-0.24	0.85	1.00
³ IM _H	2.80	0.16	0.03	-0.99	2.00	0.72	-0.26	-0.29	0.83	1.00
⁵IM _H	4.21	0.44	0.34	-0.99	4.00	0.86	-0.40	-0.29	0.83	1.00
⁷ IM _H	4.21	0.48	0.32	0.99	6.00	0.75	-0.39	-0.25	0.88	1.00
¹ TS _{reb}	0.89	-0.06	0.04	-0.86	0.00	0.59	-0.27	-0.01	0.69	1.00
³ TS _{reb}	2.65	0.15	-0.01	-0.79	2.00	0.72	-0.44	0.01	0.70	1.00
⁵ TS _{reb}	4.17	0.40	0.24	-0.81	4.00	0.76	-0.44	-0.01	0.69	1.00
⁷ TS _{reb}	4.26	0.58	0.18	0.98	6.00	0.75	-0.19	0.01	0.43	1.00
	ρ_{Fe}	$\rho_{Ligands}$	p _{ACH2}	ρ_{Total}		Q _{Fe}	QLigands	Q _{ACH2}	Q _{Total}	
¹ Рн	0.00	0.00	0.00	0.00		0.54	-0.48	0.96	1.00	
³ Рн	2.03	-0.01	-0.02	2.00		0.53	-0.49	0.97	1.00	
⁵Рн	3.84	0.17	-0.01	4.00		0.55	-0.50	0.95	1.00	
⁷ P _H	4.24	0.69	1.07	6.00		0.83	-0.24	0.41	1.00	

Table S15. Group spin densities and charges of single point UB3LYP/B2 calculations onUB3LYP/B1 optimized geometries for hydroxylation reaction in Gaussian.



Table S16. All bond lengths in angstroms and obtained from geometry optimizations for

	Fe-O1	Fe-O2	Fe-O3	Fe-O4	Fe-N1	Fe-N2	01-H1	H1-C1	01-C1
¹ Re	1.74346	2.04614	2.04915	1.89882	1.95509	2.04803	3.61496	1.09283	4.33017
³ Re	1.65568	2.01445	2.07153	1.93670	1.94867	2.05294	1.99734	1.09495	3.08993
⁵Re	1.64995	2.24744	2.09589	1.95733	2.10113	2.04743	1.95870	1.09415	3.02665
⁷ Re	1.94090	2.33657	2.07411	1.94135	2.12006	2.16562	2.15901	1.09244	3.09906
¹ TSн	1.75326	2.01858	2.05638	1.91737	1.95000	2.02665	1.24603	1.34004	2.58562
³ TSн	1.78251	2.07040	2.09459	1.95991	2.00576	2.02952	1.22450	1.35370	2.57795
⁵ TSн	1.74556	2.23539	2.18663	1.97444	2.08353	2.14612	1.29417	1.22925	2.51828
⁷ TSн	1.85239	2.08668	2.10592	1.95619	2.01861	2.08723	1.22147	1.34324	2.56216
¹ IMн	1.82031	2.03036	2.03170	1.93958	1.97078	2.02285	0.98036	3.15281	2.96202
³ IM _H	1.81880	2.05400	2.27944	1.92551	2.20514	2.01088	0.98228	3.13086	2.92549
⁵IMн	1.84838	2.24262	2.21821	2.00466	2.11811	2.17147	0.97954	2.82545	2.97008
⁷ IM _H	1.86324	2.33383	2.14215	1.94260	2.21081	2.15754	0.97596	3.02486	2.99143
¹ TS _{reb}	1.85981	2.00575	2.04344	1.94203	1.95857	2.00445	0.98062	2.54284	2.12298
³ TS _{reb}	1.88607	1.99570	2.04713	1.93758	1.95466	2.01450	0.98210	2.50770	2.07661
⁵ TS _{reb}	1.90647	2.20376	2.16335	1.99757	2.14106	2.16942	0.97729	2.68542	2.26828
⁷ TS _{reb}	1.98228	2.16661	2.18238	1.96483	2.08898	2.14868	0.99887	2.24911	1.80415
¹ P _H	2.03055	2.07331	2.09276	2.09422	1.99724	1.98202	1.06698	2.10221	1.39730
³ Рн	2.02596	2.47652	2.01485	2.29901	2.05438	1.98025	1.08950	2.13458	1.40241
⁵Рн	2.50446	2.35400	2.07597	2.11548	2.14701	2.14738	1.02155	2.02381	1.41016
⁷ Рн	2.07470	2.24636	2.15561	1.89127	2.08590	2.11389	0.98337	2.04671	1.50046

hydroxylation reaction in Gaussian at UB3LYP/B1.