

## Supporting Information

# Proton-coupled electron transfer at a mis-metalated zinc site detected with protein charge ladders.

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**Table S1.** Metal equivalents of copper per dimeric SOD1, as determined via ICP-MS. Fewer than 0.1 equivalents of metal present per dimer classified protein as “metal free”. All experiments were performed in excess copper. The buffer used for capillary electrophoresis was also tested for copper.

pH	Zn: apo WT SOD1 dimer	Cu: apo WT SOD1 dimer	Cu: WT metallated SOD1 dimer	Buffer
5.6	0.01	0.04	4.74	-0.03
7.4	-0.05	0.02	4.94	-0.10
8.3	0.01	0.04	4.76	0.02

**Table S2. Computed pK<sub>a</sub>s before and after reducing the Cu ion in the canonical Zn-binding site of SOD1.**

Residue ID	Residue Name	Subunit #1			Subunit #2		
		pK <sub>a,ox</sub>	pK <sub>a,red</sub>	ΔpK <sub>a,(red.-ox.)</sub>	pK <sub>a,ox</sub>	pK <sub>a,red</sub>	ΔpK <sub>a,(red.-ox.)</sub>
3	Lys	11.7	11.0	-0.7	11.6	11.5	-0.1
9	Lys	9.4	9.4	0.0	9.5	9.4	-0.1
11	Asp	3.8	3.9	0.0	3.8	3.8	0.0
21	Glu	4.7	4.7	0.0	4.6	4.3	-0.3
23	Lys	10.2	10.3	0.1	10.2	10.3	0.1
24	Glu	4.6	4.6	0.0	4.5	4.6	0.1
30	Lys	10.3	10.3	0.1	10.4	10.3	-0.1
36	Lys	9.9	9.8	-0.1	10.0	9.9	-0.1
40	Glu	3.9	3.9	0.0	3.6	4.4	0.9
43	His	11.0	11.5	0.5	11.4	11.4	0.0
49	Glu	5.1	5.3	0.2	5.1	5.2	0.1
52	Asp	1.5	1.5	0.0	1.4	1.4	-0.1
70	Lys	10.5	10.1	-0.4	9.9	10.2	0.3
75	Lys	10.5	10.5	0.0	10.4	10.5	0.0
76	Asp	3.3	3.3	0.0	3.3	3.4	0.2
77	Glu	4.5	4.4	-0.1	4.4	4.4	0.0
78	Glu	3.7	3.7	0.0	3.8	3.9	0.0
90	Asp	3.0	3.1	0.2	2.7	3.1	0.4
91	Lys	10.9	10.9	0.0	10.9	10.8	0.0
92	Asp	5.1	5.4	0.3	4.9	5.2	0.2
96	Asp	4.3	4.6	0.3	4.6	4.4	-0.2
100	Glu	4.0	4.2	0.2	4.3	4.3	0.0
101	Asp	-1.8	-2.4	-0.7	-3.0	-1.6	1.4
109	Asp	3.4	3.4	0.0	3.3	3.6	0.3
110	His	7.1	6.7	-0.3	6.9	7.0	0.2
121	Glu	3.7	3.9	0.2	3.7	3.8	0.1
122	Lys	10.2	10.2	-0.1	10.2	10.2	0.1
124	Asp	-1.8	0.8	2.6	-2.3	0.6	2.9
125	Asp	2.5	3.5	1.0	2.6	2.2	-0.4
128	Lys	10.5	10.7	0.2	10.6	10.7	0.1
132	Glu	3.6	3.8	0.2	3.8	3.9	0.0
133	Glu	4.6	4.6	0.0	4.7	4.9	0.2
136	Lys	10.3	10.3	0.0	10.3	10.3	0.0

**Table S3.** Summary of classical molecular dynamics simulation conditions and timescales.

Redox State	Zn-Site Bonded Ligands	Zn-Site Non-Bonded Ligands	Simulated pH	Simulated Time (ns)
Cu <sup>+1</sup> Cu <sup>+2</sup>	His-63, His-71, His-80, Asp-83		-4, -2, 0, 2, 4, 6, 8, 10, 12	24 ns at each pH
Cu <sup>+1</sup> Cu <sup>+1</sup>	His-63, His-71, His-80, Asp-83		-4, -2, 0, 2, 4, 6, 8, 10, 12	24 ns at each pH
Cu <sup>+1</sup> Cu <sup>+1</sup>	His-71, His-80, Asp-83	His-63 (N <sub>ε</sub> protonated)		314
Cu <sup>+1</sup> Cu <sup>+1</sup>	His-71, His-80, Asp-83	His-63 (N <sub>ε</sub> and N <sub>δ</sub> protonated)		313
Cu <sup>+1</sup> Cu <sup>+1</sup>	His-63, His-80, Asp-83	His-71 (N <sub>ε</sub> protonated)		308
Cu <sup>+1</sup> Cu <sup>+1</sup>	His-63, His-80, Asp-83	His-71 (N <sub>ε</sub> and N <sub>δ</sub> protonated)		291
Cu <sup>+1</sup> Cu <sup>+1</sup>	His-63, His-71, Asp-83	His-80 (N <sub>ε</sub> protonated)		292
Cu <sup>+1</sup> Cu <sup>+1</sup>	His-63, His-71, Asp-83	His-80 (N <sub>ε</sub> and N <sub>δ</sub> protonated)		382
Cu <sup>+1</sup> Cu <sup>+1</sup>	His-63, His-71, His-80	Asp-83 (unprotonated)		300
Cu <sup>+1</sup> Cu <sup>+1</sup>	His-63, His-71, His-80	Asp-83 (protonated)		331