

Supporting Information for

**Magnetic Circular Dichroism Studies of Iron(II) Binding to Human
Calprotectin**

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1. Supplementary Data

1.1 Sample Concentrations

Table S1. Protein and Metal Concentrations of MCD Samples

Protein	[Protein] (mM)	[Fe(II)] (mM)	[Ca(II)] (mM)	Glassing agent ^a
CP-Ser	1.36	1.22	27.2	Glycerol
CP-Ser	1.68	1.51	33.6	Sucrose
H103A	1.44	1.30	28.8	Glycerol
H103A	1.84	1.66	36.8	Sucrose
AHA	1.81	1.63	36.2	Glycerol
AHA	1.74	1.57	34.8	Sucrose
AAA	1.36	1.22	27.2	Glycerol
AAA	2.24	2.01	44.8	Sucrose
ΔHis ₃ Asp	1.11	1.00	22.2	Glycerol
ΔHis ₃ Asp	1.54	1.39	30.8	Sucrose
ΔHis ₃ Asp-H103A	1.61	1.45	32.2	Sucrose
CP-Ser	1.68	0.50	33.6	Sucrose
CP-Ser	1.68	1.01	33.6	Sucrose
CP-Ser	1.68	2.02	33.6	Sucrose
ΔHis ₄	1.59	1.43	29.8	Glycerol
ΔHis ₄	1.58	1.42	31.2	Sucrose
ΔHis ₃ Asp	2.26	3.39	45.2	Sucrose
CP-Ser	1.85	1.11	0	Sucrose
ΔHis ₃ Asp	1.42	0.85	0	Sucrose

^aGlycerol samples were prepared with 40% (v/v) 150 mM HEPES, 200 mM NaCl, pH 7.4 60% (v/v) glycerol-d₈ in D₂O. Sucrose samples were prepared with 150 mM HEPES, 200 mM NaCl, pH 7.4 saturated with sucrose in D₂O.

1.2 MCD Spectra

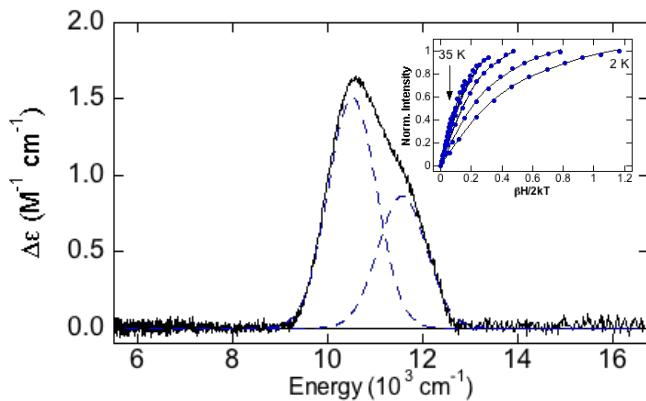


Figure S1. 5 K, 7 T NIR MCD spectra of $\Delta\text{His}_3\text{Asp}/\text{Ca}(\text{II})/\text{Fe}(\text{II})$ in a glycerol glass. Gaussian fits are shown as dashed blue lines. Inset: Saturation magnetization data (dots) and best fit (lines) for $\Delta\text{His}_3\text{Asp}/\text{Ca}(\text{II})/\text{Fe}(\text{II})$ in glycerol collected at 10310 cm^{-1} .

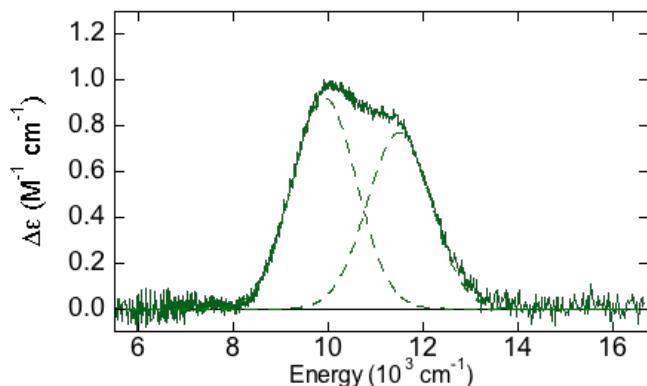


Figure S2. 5 K, 7 T NIR MCD spectra of $\Delta\text{His}_3\text{Asp}(\text{H}103\text{A})/\text{Ca}(\text{II})/\text{Fe}(\text{II})$ in a saturated sucrose buffer solution. Gaussian fits are shown as dashed green lines.