

Electronic Supplementary Information for:

Zn^{II} and Hg^{II} binding to a designed peptide that accommodates different coordination geometries

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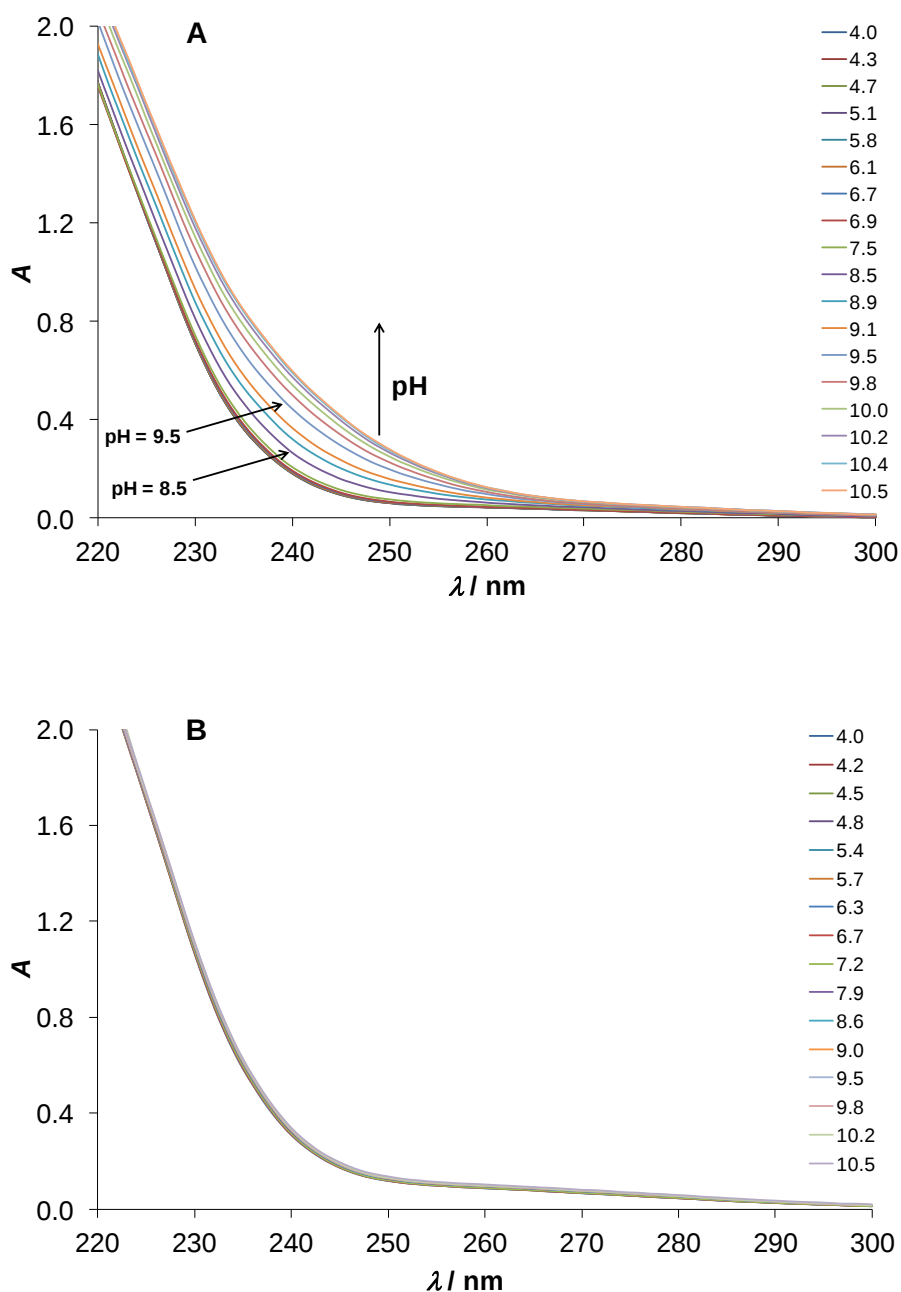


Fig. S1A-B.: UV-spectra recorded for Hg^{II} – HS 0.5:1 (A) and 1:1 (B). ($c_{\text{HS}} = 1.0 \times 10^{-4} \text{ M}$ (A,B), $I = 0.1 \text{ M NaClO}_4$, $T = 298 \text{ K}$, $l = 1.0 \text{ cm}$)

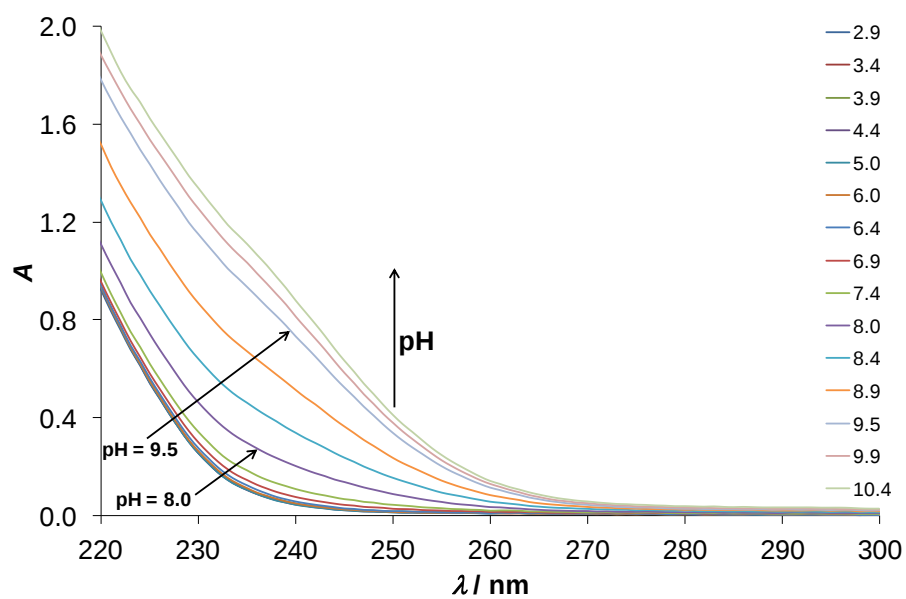


Fig. S2.: UV-spectra recorded for **HS** in the absence of metal ions. ($c_{\text{HS}} = 1.0 \times 10^{-4}$ M, $I = 0.1$ M NaClO₄, $T = 298$ K, $l = 1.0$ cm)

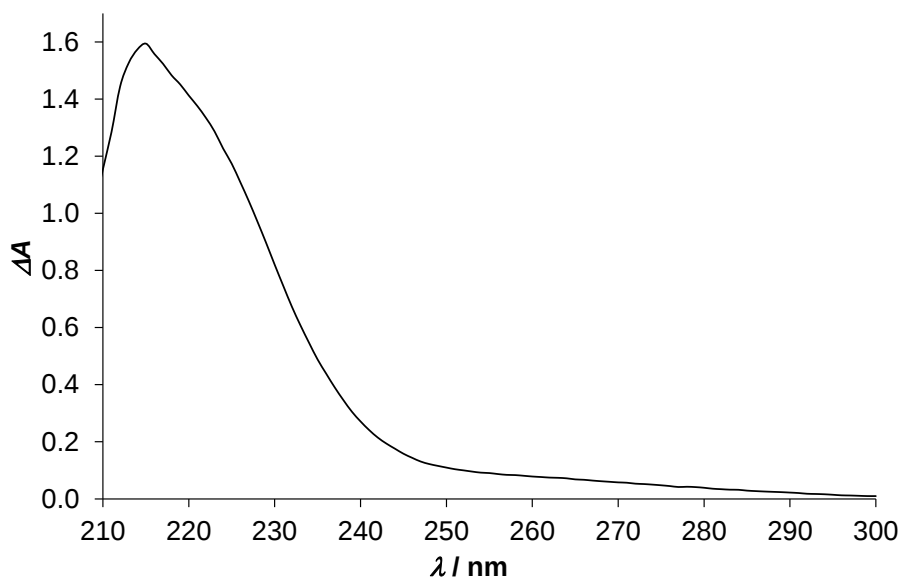


Fig. S3.: Difference of the UV-spectra recorded for Hg^{II} -**HS** 1:1 and **HS** at pH = 4.0. Note that the position of the maximum (λ_{max}) of the observed band has a significant uncertainty due to the large absorbances recorded between 210-220 nm ($c_{\text{HS}} = 1.0 \times 10^{-4}$ M, $I = 0.1$ M NaClO_4 , $T = 298$ K, $l = 1.0$ cm)

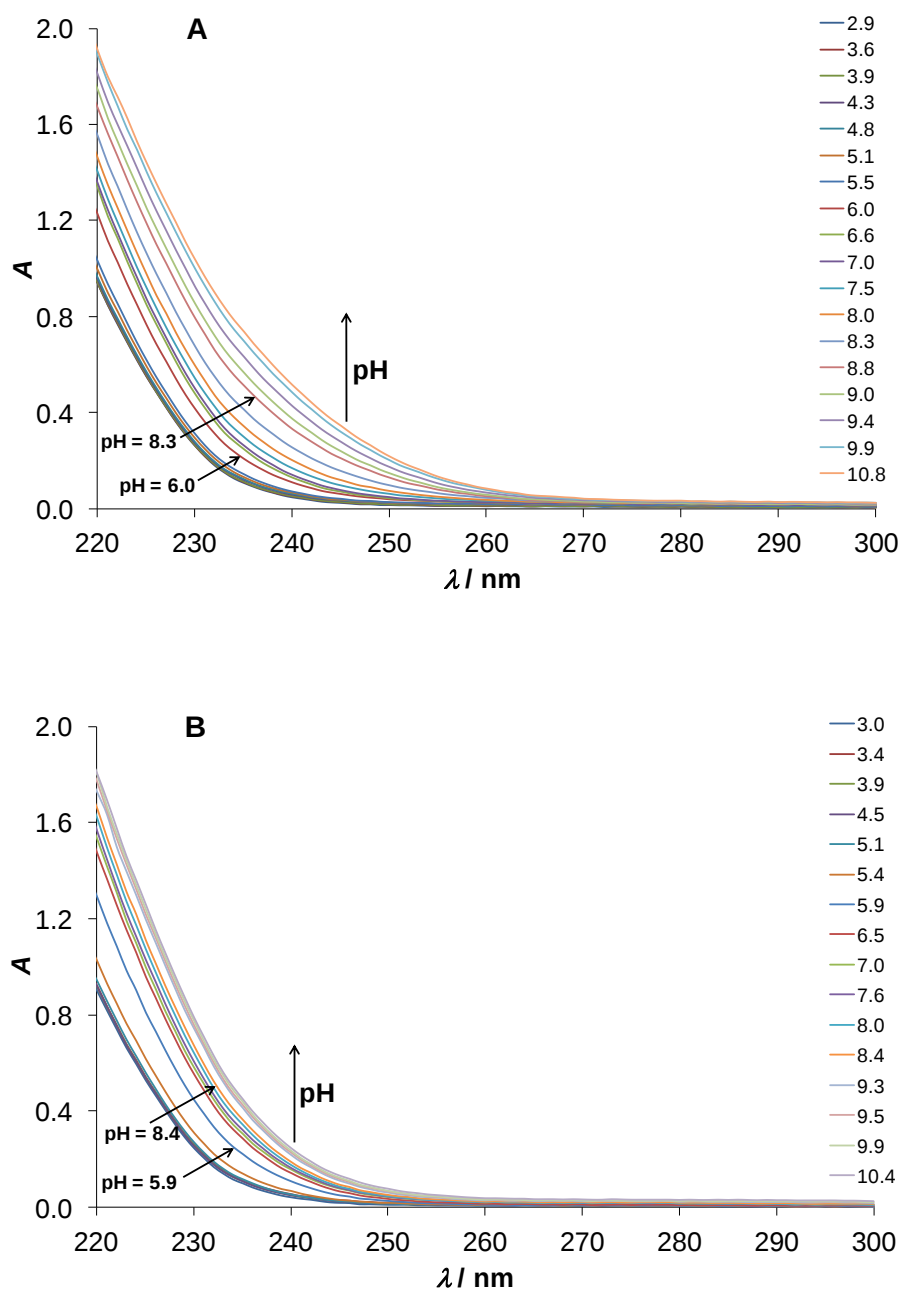


Fig. S4A-B.: UV-spectra recorded for $\text{Zn}^{\text{II}} - \text{HS}$ 0.5:1 (A) and 1:1 (B) ($c_{\text{HS}} = 1.0 \times 10^{-4} \text{ M}$ (A,B), $I = 0.1 \text{ M NaClO}_4$, $T = 298 \text{ K}$, $l = 1.0 \text{ cm}$)

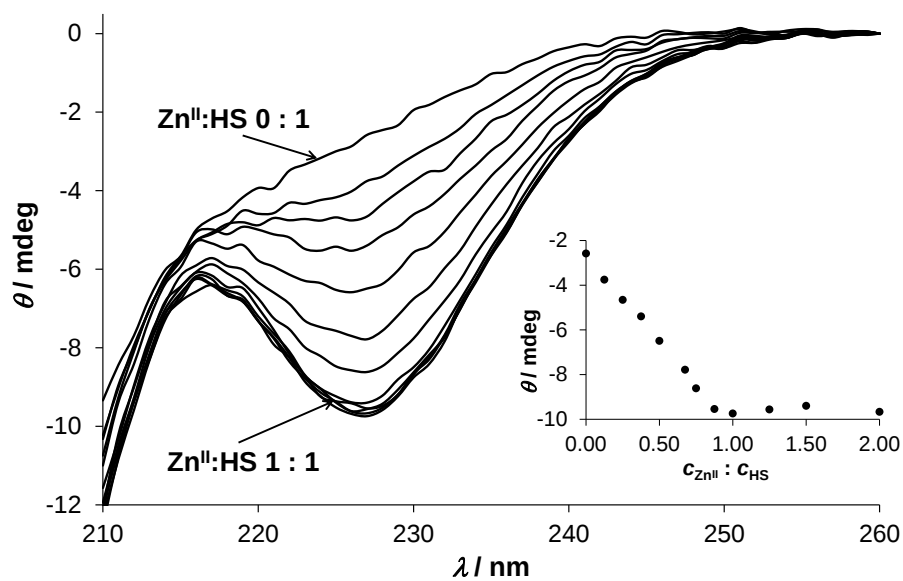


Fig. S5.: SRCD spectra of **HS** as a function of metal ion to ligand ratio in the range of 0:1 to 2:1 at pH = 10.5 ($c_{\text{HS}} = 2.0 \times 10^{-4}$ M, $I = 0.1$ M NaClO₄, $T = 298$ K, $l = 1$ mm). The insert shows the trace of ellipticities at $\lambda = 227$ nm.

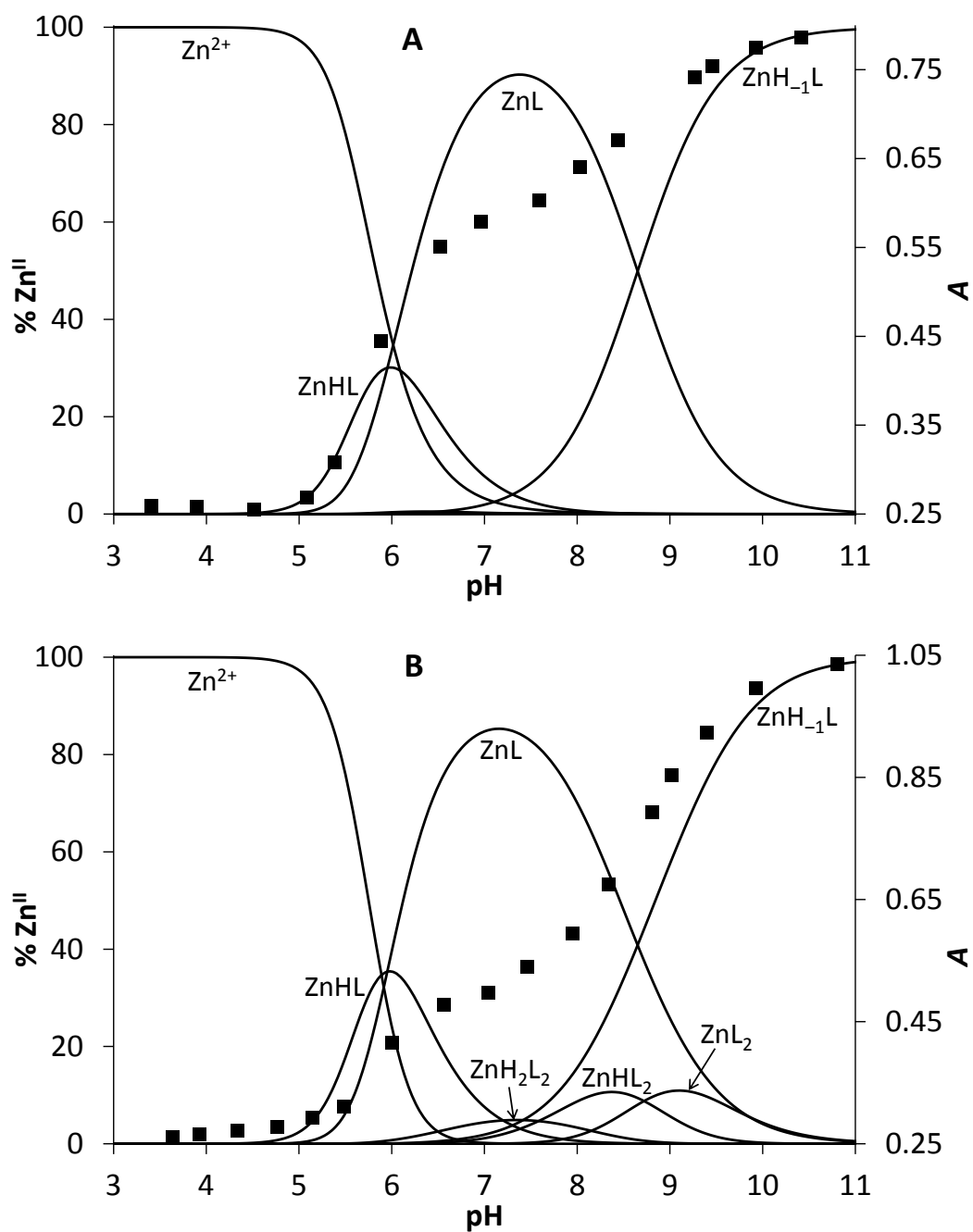


Fig. S6A-B.: Species distribution diagram and for the Zn^{II} : HS 1:1 (A) and 0.5:1 (B) systems ($c_{\text{HS}} = 1.0 \times 10^{-4}$ M). The full squares represent the measured absorbances at 230 nm.

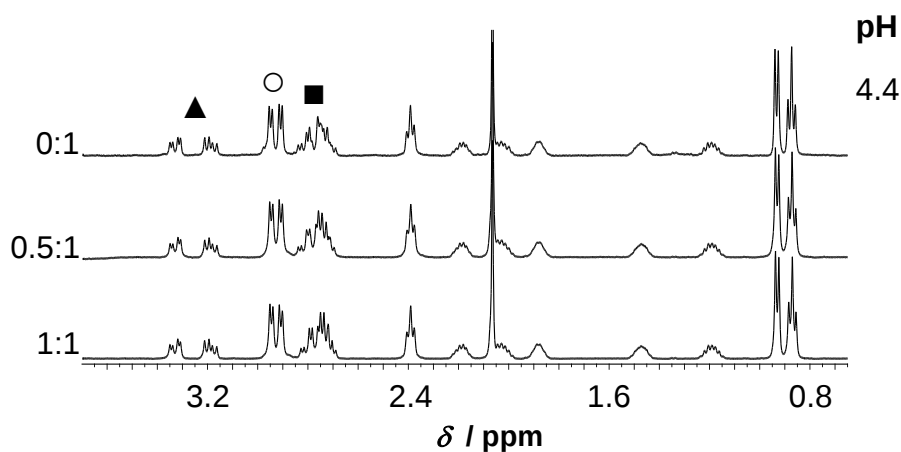


Fig. S7.: Aliphatic region of the ^1H NMR spectra of **HS** recorded in the absence of Zn^{II} and in the $\text{Zn}^{\text{II}}:\text{HS}$ 0.5:1 and 1:1 systems at $\text{pH} = 4.4$ ($\text{H}_2\text{O}/\text{D}_2\text{O} = 90:10$ %v/v, $c_{\text{HS}} = 1.3 \times 10^{-3}$ M, $T = 298$ K). C_βH_2 resonances of the residues with potential donor groups are indicated by the following symbols: His C_βH_2 : ▲; Cys C_βH_2 : ○; Asp C_βH_2 : ■.