

Supporting Information

Dialing in on pharmacological features for a therapeutic antioxidant small molecule

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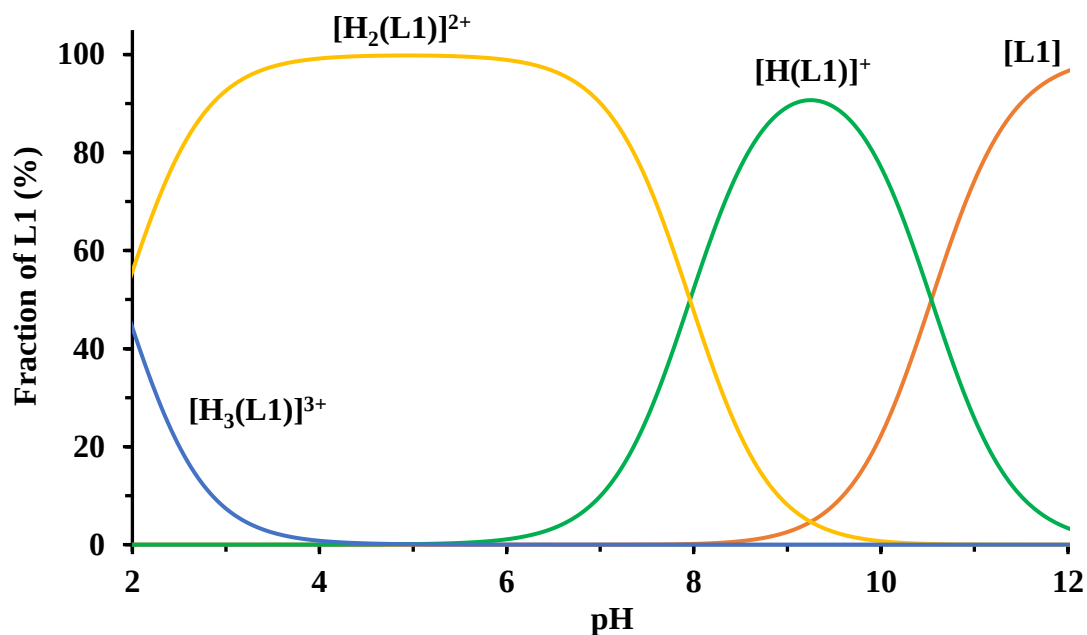


Figure S1. Equilibrium distribution diagram of **L1** based on ref ⁴⁶ ($c_L = 2.5$ mM, $I = 0.1$ M KNO_3 , $T = 298$ K).

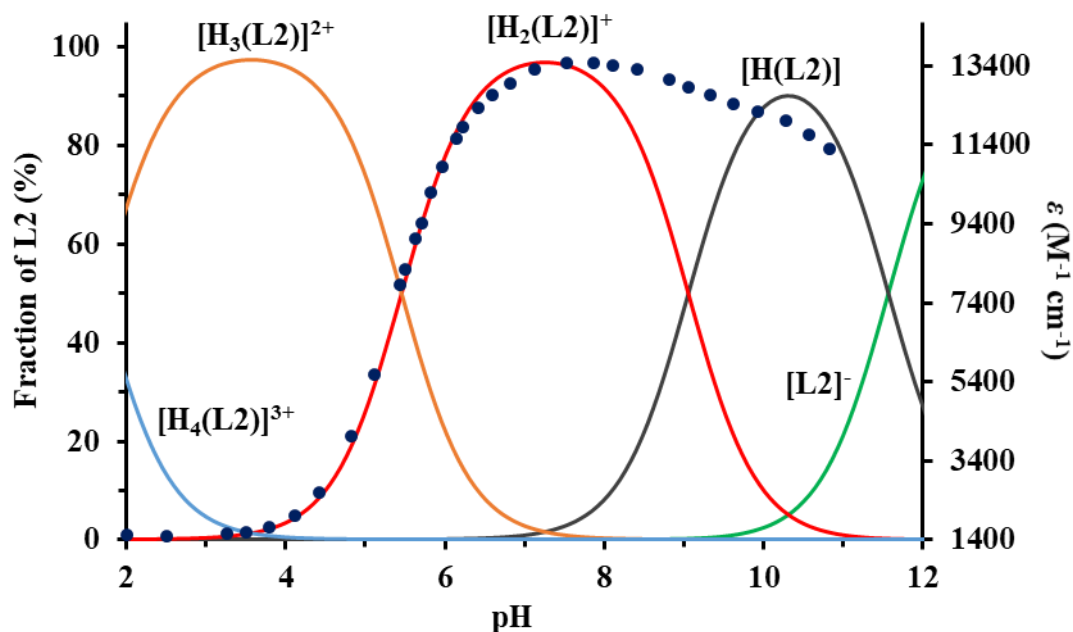


Figure S2. The equilibrium distribution curves for **L2**, calculated using the protonation constants obtained from pH-potentiometric titrations along with the molar extinction coefficients at 255 nm from UV spectrophotometric titrations ($c_L = 0.1$ mM, $I = 0.15$ M NaCl , $T = 298$ K).

The large increase in molar extinction coefficients between pH 4 and 7 aligns well with the deprotonation of the hydroxyl group ($\log K = 5.45$) and the consequent increase of electron delocalization in the 7-center 8-electron π -system. From pH 7 a slow decrease in absorbance can be observed which is caused by the deprotonation of the pyridine N-atom ($\log K = 9.05$). These changes support the proposed protonation order shown in Figure 3.

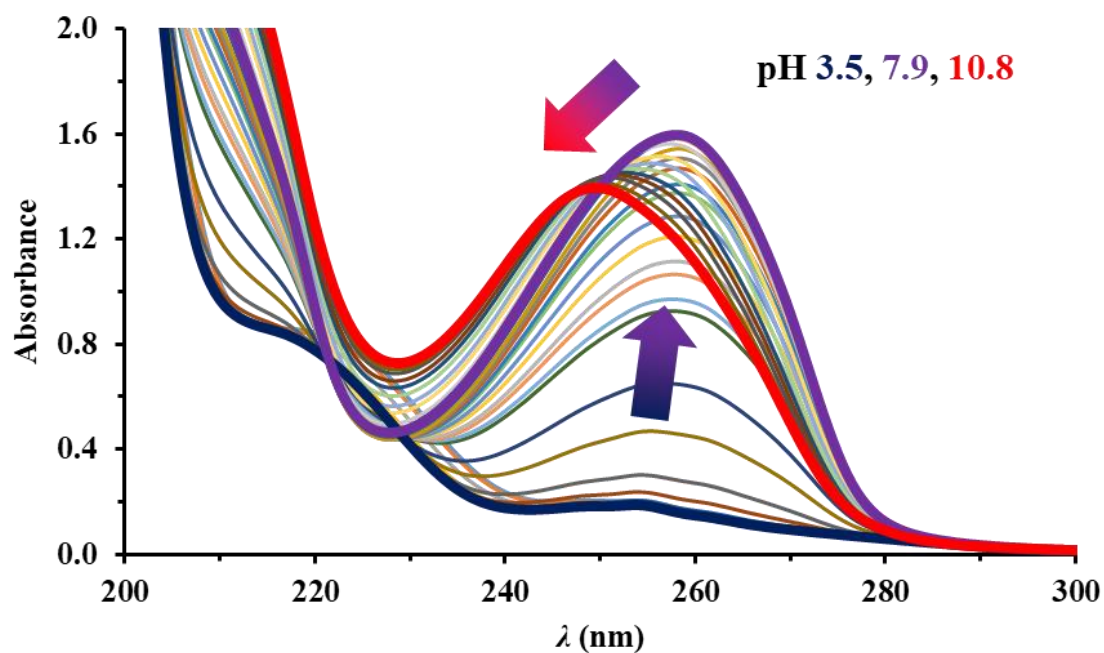


Figure S3. pH dependence of UV spectra in aqueous solution of **L2** ($c_L = 0.1$ mM, pH varied between 1.683 and 10.835).

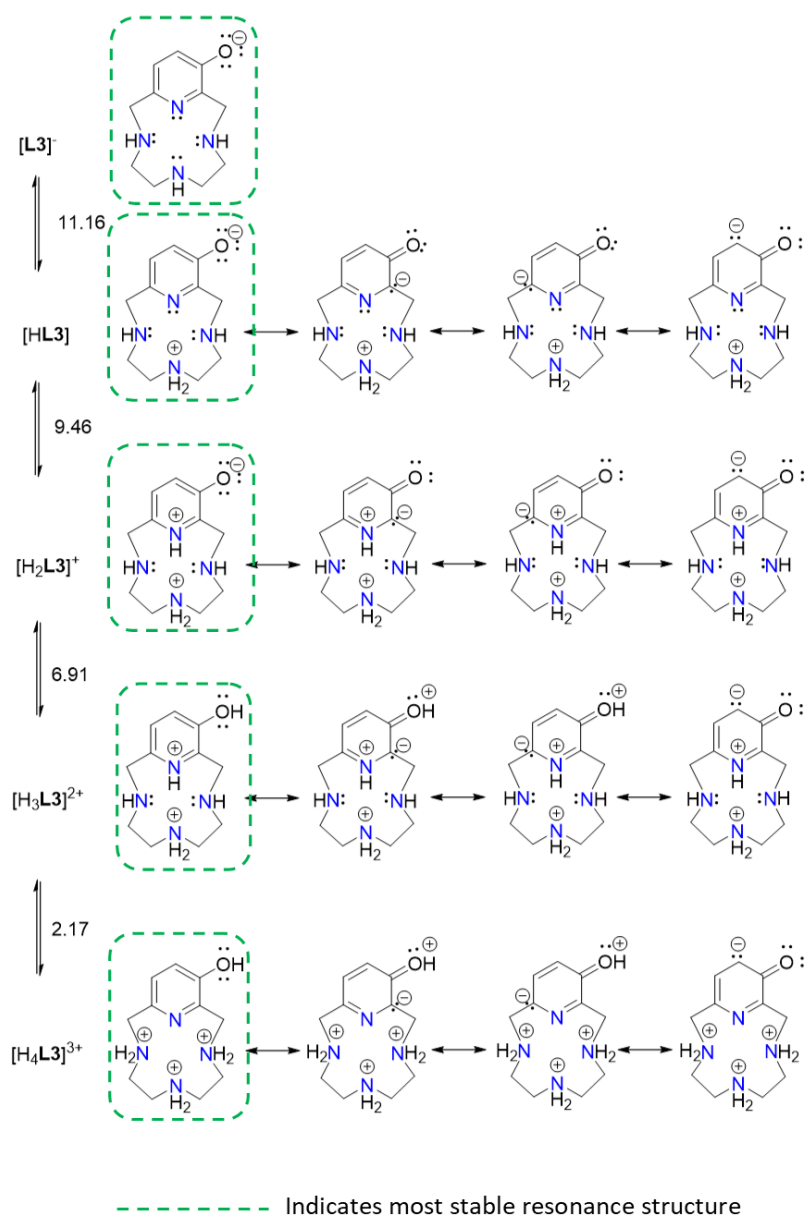


Figure S4. pH dependent resonance structures of **L3**.

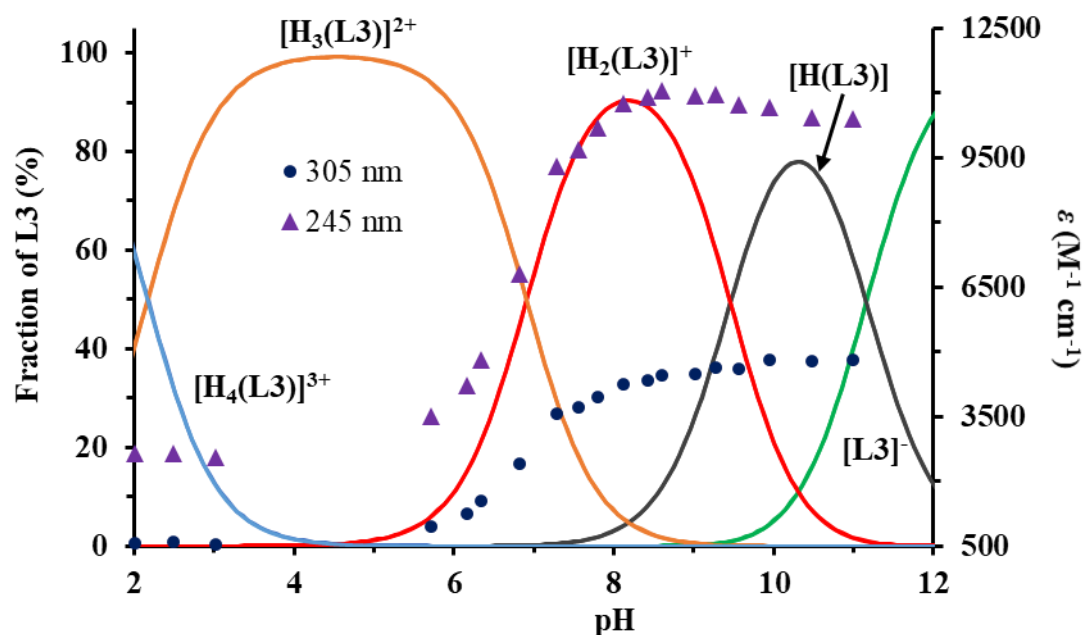


Figure S5. The equilibrium distribution curves for **L3**, calculated using the protonation constants obtained from pH-potentiometric titrations along with the molar extinction coefficients at 245 and 305 nm from UV spectrophotometric titrations ($c_L = 0.03$ mM, $I = 0.15$ M NaCl, $T = 298$ K).

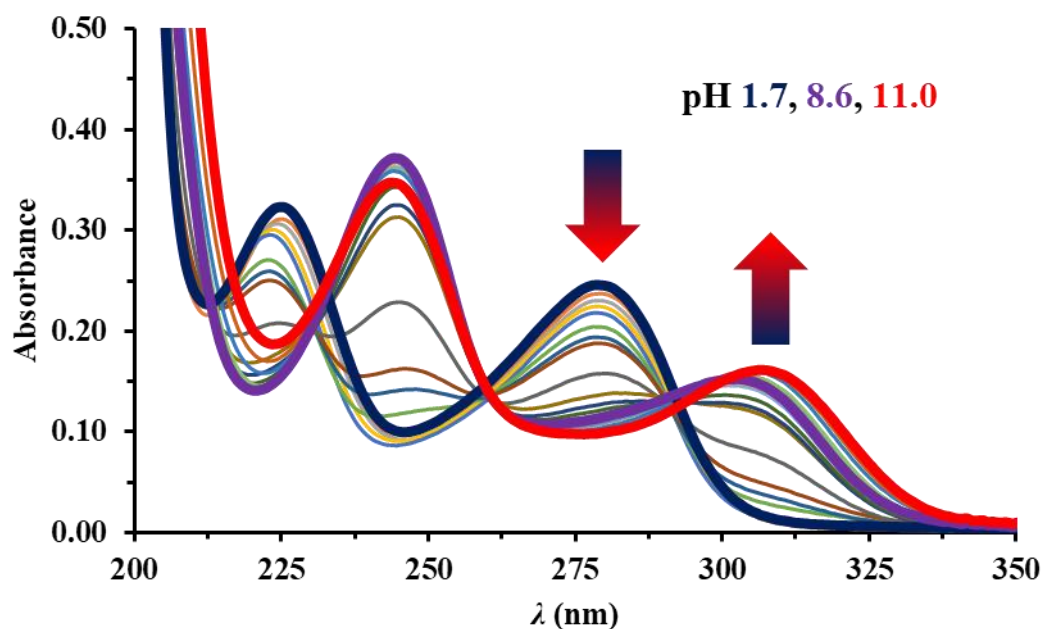


Figure S6. pH dependence of UV spectra in aqueous solution of **L3** ($c_L = 0.03$ mM, pH varied between 1.706 and 11.000).

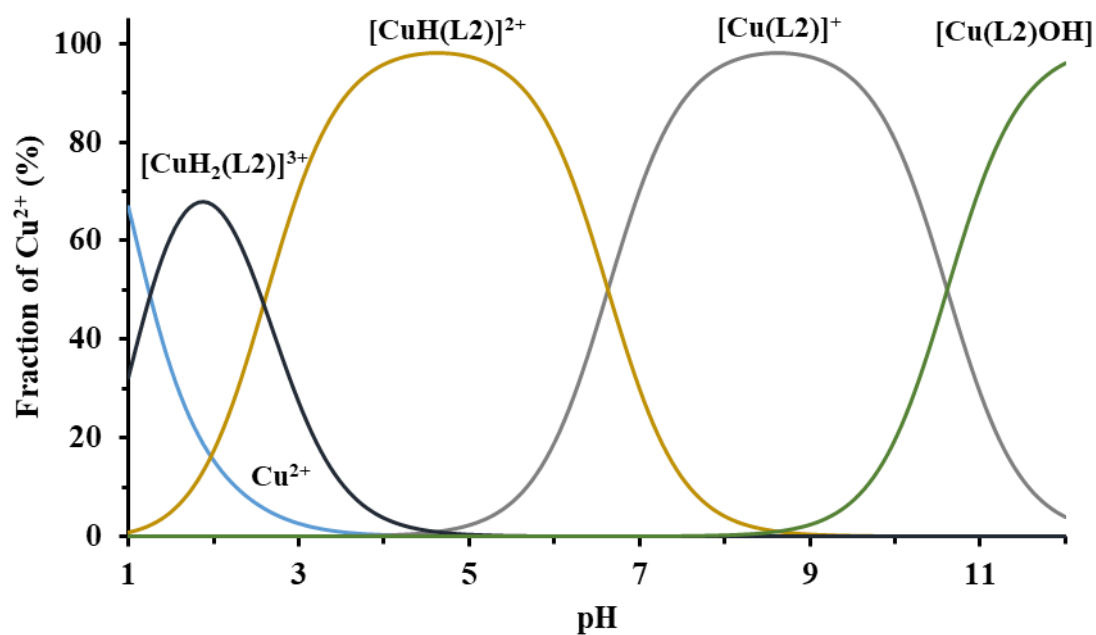


Figure S7. Equilibrium distribution diagram in the Cu²⁺-L2 system calculated by using 2.00 mM concentration of Cu²⁺ ion and the ligand.

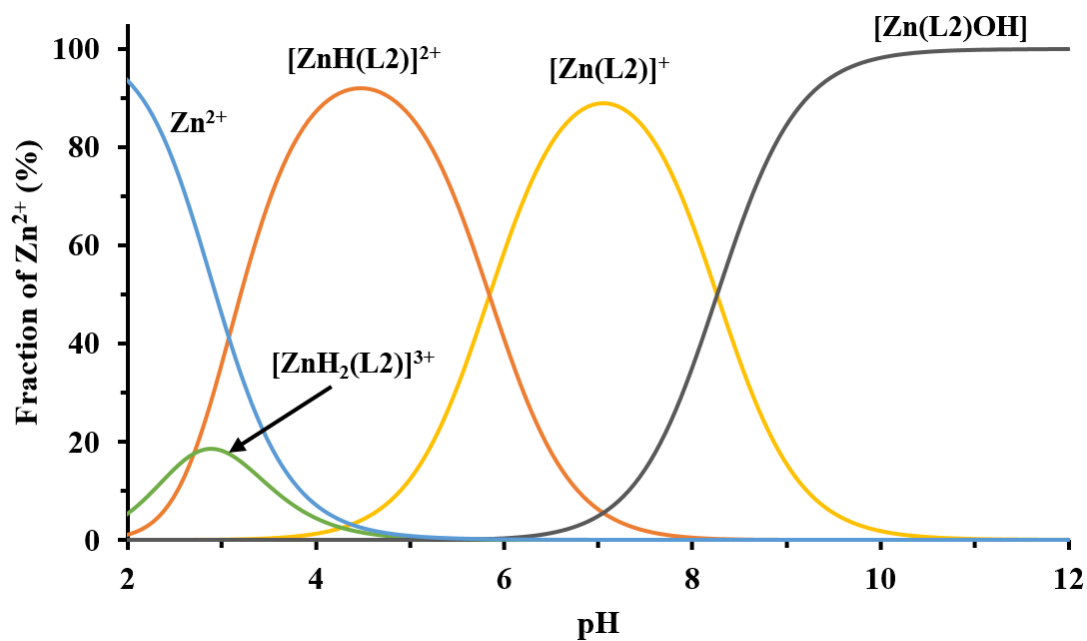


Figure S8. Equilibrium distribution diagram in the Zn²⁺-L2 system calculated by using 2.00 mM concentration of Zn²⁺ ion and the ligand.

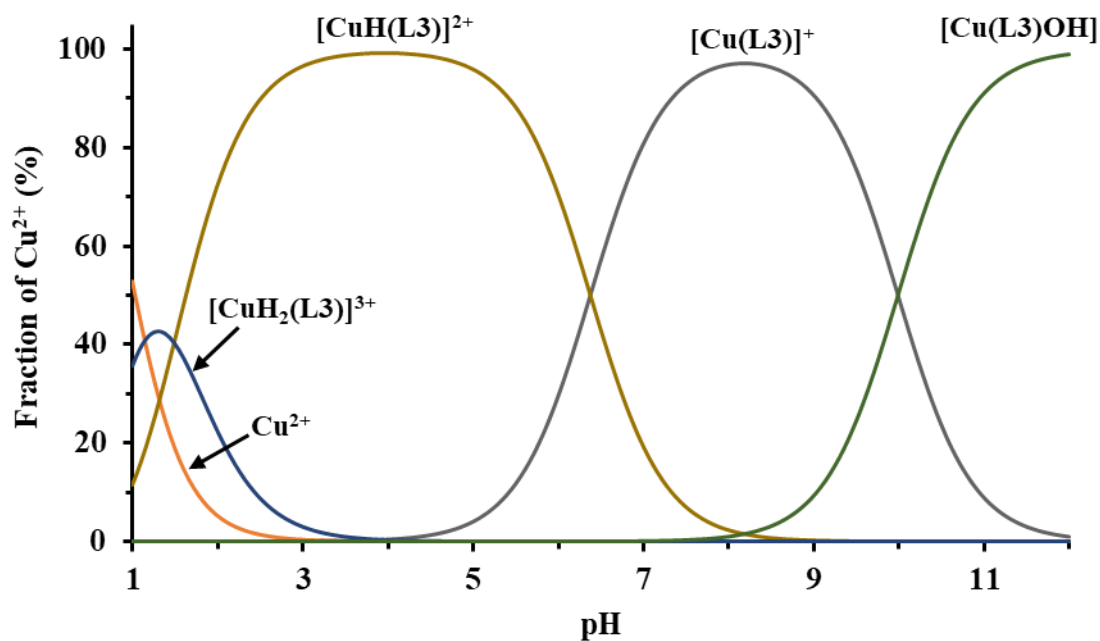


Figure S9. Equilibrium distribution diagram in the Cu²⁺-L3 system calculated by using 2.00 mM concentration of Cu²⁺ ion and the ligand.

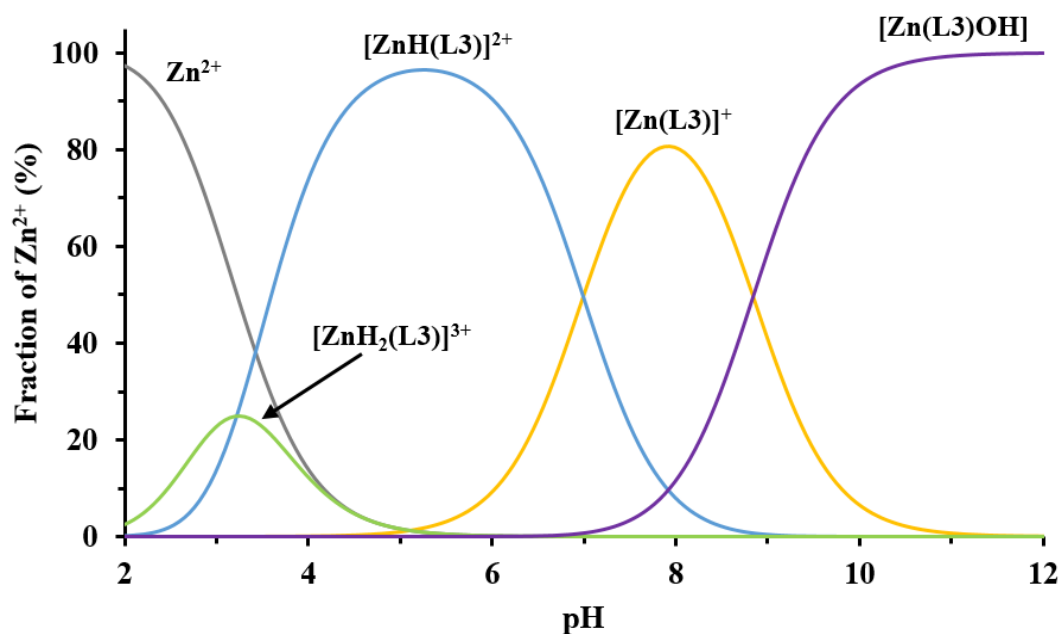


Figure S10. Equilibrium distribution diagram in the Zn²⁺-L3 system calculated by using 2.00 mM concentration of Zn²⁺ ion and the ligand.

Table S1. Fitting parameters for cellular toxicity studies of cyclen, **L1**, and **L2** with HT-22 cells.

	Cyclen	L2	L1
log(inhibitor) vs. normalized response			
Best-fit values			
LogIC ₅₀	1.867	2.474	2.365
IC ₅₀	73.63	298.0	231.8
Std. Error			
LogIC ₅₀	0.05372	0.05707	0.07932
95% Confidence Intervals			
LogIC ₅₀	1.760 to 1.974	2.362 to 2.586	2.207 to 2.524
IC ₅₀	57.50 to 94.29	230.3 to 385.6	160.9 to 334.0
Goodness of Fit			
Degrees of Freedom	63	159	63
R square	0.9422	0.7962	0.9086
Absolute Sum of Squares	5788	33402	11727
Sy.x	9.585	14.49	13.64
Number of points Analyzed	64	160	64