Binuclear and Tetranuclear Zn(II) Complexes With Thiosemicarbazones: Synthesis, X-ray Crystal Structures, ATPsensing, DNA-binding, Phosphatase activity and theoretical calculations

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4. Figure S4: Mass spectra of H_2L^2



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12. Figure S12: Extended NMR spectra of complex 2



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16. Figure S16: a) Red bars: UV-visible absorbance of complex 1 (1 x 10⁻⁵) in presence of 8 equiv of ATP. Green bars: UV-visible spectrum of complex 1 (1 x 10⁻⁵) in presence of 8 equiv of ATP & 24 equiv of other anions. b) Blue bars: UV-visible absorbance of complex 2 (1 x 10⁻⁵) in presence of 8 equiv of ATP. Orange bars: UV-visible spectrum of complex 2 (1 x 10⁻⁵) in presence of 8 equiv of ATP & 24 equiv of other anions



(a)



17. Figure S17: Benesi-Hildeband plot for determination of Ka values of complexes 1(a) and 2(b) with ATP.

The Benesi-Hildeband equation used is : $1/\Delta A = 1/\Delta A_{max} + (1/K_b[C]) (1/\Delta Amax)$, where ΔA and ΔA_{max} are the change in absorbance at a given concentration of ATP, and when all the metal complex is fully bound to ATP respectively, [C] = [ATP].

The detection limit DL of **Complexes** for ATP was determined from the following equation: $DL = K \times Sb1/S$. Where K = 2 or 3 (we take 2 in this case); Sb1 is the standard deviation; S is the slope of the calibration curve.



18. Figure S18: Detection limit and calibration curves of a) complex **1** and b) complex **2** with ATP.

From the Absorbance vs. [ATP] graphs (S17), we get slope = 12005.49026, and Sb1 value is 0.04066 for complex **1**. Thus using the formula, we get the Detection Limit for Complex $1 = 6.7 \times 10^{-6}$. Similarly for complex **2** Detection Limit = 1.7×10^{-6} .



19. Figure S19: Plots of $[DNA]/(\epsilon_a - \epsilon_f)$ vs. [DNA] for the titration of DNA with the a) complex 1 b)complex 2.



20. Figure S20: Plot of a) I_0/I vs. [Complex 1] b) I_0/I vs. [Complex 2].



21. Figure S21. Time dependent spectra of PNPP (10⁻⁴ M) and PNPP(10⁻⁴ M) in presence of Zn(OAc)₂ solution (10⁻⁵ M).