Supporting Information File

Anion directed structural tuning of azomethine derived two Zn²⁺complexes with optoelectronic recognition of Cu²⁺ from aqueous medium with Anti-cancer activities: Expedition from micromolar to femtomolar sensitivity with DFT revelation

Nithun Ranjan Pandit, ^{¥ [a]} Sourav Bej, ^{¥ [bc]} Riyanka Das, ^{¥ [bc]} Nirajan Ghosal ^[d], Ananya Mondal^{[a][e]}, Ranjana Pal^[d], Meenakshi Ghosh^[e], Priyabrata Banerjee, ^{* [bc]} Biplab Biswas^{* [a]}

a. Department of Chemistry, Presidency University, 86/1, College Street, Kolkata 700073, India.

b. Electric Mobility and tribology research Group, CSIR-Central Mechanical Engineering Research Institute, Mahatma Gandhi Avenue, Durgapur, 713209, India.

c. Academy of Scientific & Innovative Research (AcSIR), Ghaziabad-201002, Uttar Pradesh, India

d. Department of Life Sciences, Presidency University, 86/1, College Street, Kolkata 700073, India.

e. Department of Chemistry, Vidyasagar College for Women, 39 Sankar Ghosh Lane, Kolkata-6, India.

± Present Address : School of Science, Harbin Institute of Technology (Shenzhen), Shenzhen 518055, China

Corresponding authors: Dr Biplab Biswas: E-mail: <u>biplab.chem@presiuniv.ac.in</u>, <u>http://presiuniv.ac.in/web/</u>; Tel: +919734246721&Dr Priyabrata Banerjee: E-mail: <u>pr_banerjee@cmeri.res.in</u>, <u>priyabrata_banerjee@yahoo.co.in</u>;

Web: www.cmeri.res.in, www.priyabratabanerjee.in; Tel: +91-9433814081

[¥] NRP, SB and RD contributed equally as first author to this paper



Figure S1: Mass spectrometric data of NS-1



m/z= 264

 H_2O

Scheme S1: Probable Mass spectrometric analysis of NS-1



Figure S2: (A) ¹H-NMR spectra of NS-1

¹H-NMR(400 MHz, D₂O): 1H, s, 8.74 ppm (H_d), 1H, d, 8.57 ppm(H_h), 1H, dd, 8.10-8.14 ppm (H_g), J= 16Hz, 1H, d, 7.88-7.90 ppm, J= 8 Hz, (H_e), 1H, dd, 7.70-7.72 ppm, J= 8 Hz (H_f), 2H, s, 3.50 ppm(H_b), 3H, s, 1.79 ppm (H_i), 6H, s, 1.32 ppm (H_c).



Figure S2: (B) ¹³C-NMR spectra of NS-1



Figure S3: HR-MS data of NS-2



Scheme S2: Probable Mass spectrometric analysis of NS-2



Figure S4: packing of the complex NS-1 in different directions.

Table S1: Bond length associated with the h	nydrogen bonding	of the complex NS-1
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Bond	Distance
O(1)O(3)	2.571(2)

Table S2: Selected bond lengths around the coordination environment of the complex NS-1

Bonds	Distance (in A°)
Zn(1)-N(1)	2.20(1)
Zn(1)-N(2)	2.07(1)
Zn(1)-N(3)	1.95(2)
Zn(1)-O(1)	2.22(1)
Zn(1)-O(2)	1.95(1)

Table S3: Bond length associated with the Schiff base of the complex NS-1

Bond	Distance
C(6)-N(2)	1.26(2)



Figure S5: Selected bond lengths around the coordination environment of the complex NS-1



Figure S6: Selected bond angles around the coordination environment of the complex NS-1

Bond	Angles in degree
N(2)- Zn (1)-N(1)	76.29
N(1)- Zn (1)-O(2)	94.13
O(1)-Zn(1)-N(2)	77.96
N(3)- Zn (1)-O(2)	91.86
O(2)-Zn(1)-O(1)	96.71

Table S4: Selected bond angles around the coordination environment of complex NS-1



Figure S7: (A) X-ray structure of the complex NS-2 (B) Hydrogen bonding network of NS-2Table S7: Bond lengths associated with the hydrogen bonding of the complex NS-2

Bond	Distance
O(2)O(3)	2.824(9)



Figure S8: **(A)** Selected bond lengths around the coordination environment of the complex NS-2 (B) Enlarged coordination view of **NS-2**.



Figure S9: packing of the complex NS-2.



Figure S10: Selected bond angles around the coordination environment of the complex NS-2.

Table S8: Selected bond lengths around the coordination environment of the complex NS-2

Bonds	Distance (in A°)
Zn(1)-N(2)	2.062
Zn(1)-N(1)	2.150
Zn(1)-O(3)	2.246

Table S9: Bond angle associated with the Schiff base of the complex NS-2

Bond	Distance
C(6)-N(2)	1.259(5)

Table S10: Selected bond angles around the coordination environment of complex NS-2

Bond	Angles in degree
N(1)-Zn(1)-N(1)	93.06
O(3)-Zn(1)-O(1)	90.79
N(1)-Zn(1)-O(3)	90.22

Table S11: crystallographic parameters of NS-1 and NS-2

Complex	NS-1	NS-2
Formula	C ₁₂ H ₁₇ N ₅ ZnO ₃	C ₂₀ H ₂₈ N ₄ O ₂ Zn, 2(ClO ₄)
Formula Weight	343.67	620.75
Temperature (K)	293	273
Crystal System	Orthorhombic	Monoclinic
Space group	P 2 ₁ 2 ₁ 2 ₁	P 2/n
<i>a</i> (Å)	9.508(7)	10.7236(8)
<i>b</i> (Å)	9.719(7)	7.8448(6)
<i>c</i> (Å)	16.852(13)	15.7025(12)
α (°)	90	90
β (°)	90	96.744(3)°
γ (°)	90	90
Z	4	2
d _{cal} (g cm ⁻³)	1.466	1.571
μ(mm ⁻¹)	1.593	1.199
F(000)	708.0	640.0
Total reflection	10775	19282

Unique Reflections	2736	2910
Observe data[l>2σ(l)]	2736	2549
R(int)	0.1766	0.0336
R1, wR2 (all data)	0.1382, 0.1948	0.0695, 0.1883
R1, wR2 [I>2 <i>o</i> (I)]	0.0867, 0.1738	0.0621, 0.1796



Figure S11: TGA data of (a) NS-1 and (b) NS-2



Figure S12: Comparison between P-XRD pattern of experimental data and simulated data of (A) NS-1 (B) NS-2.

The sharp band at 2066 cm⁻¹ corresponds to the azide ion. The C-H bands are at 3088 cm⁻¹ and 2973 cm⁻¹, broad band at 2816 cm⁻¹ corresponds to -OH of the protonated Schiff base ligand. Band at 1752 cm⁻¹ corresponds to the C-N bond of the Schiff base, another band at 1603 cm⁻¹ corresponds to the C-O of acetate ion which is directly attached with Zn in **NS-1**.

On the other hand, the broad band at 3373 cm⁻¹ corresponds to the protonated -OH of the Schiff base in complex-2 (**NS-2**). Medium band in the region 2988-2909 cm⁻¹ correspond to the C-H stretching frequency. The band in the region 1074-1191 cm⁻¹ corresponds to the ClO_4^- . Strong band at 1660 cm⁻¹ corresponds to the C-N of the Schiff base.



Figure S13: FT-IR spectra of the (A) NS-1 (B) NS-2



Figure S14: Chromogenic changes of (A) NS-1 and (B) NS-2 in presence of (i) nitrate, (ii) sulfate, (iii) chloride and(iv) perchlorate salt of copper.



Figure S15: Change of absorbance at **(A)** 395 nm and **(B)** 295 nm as the linear function of the concentration of Cu²⁺ for the calculation of LOD for Cu²⁺ by **NS-1** and **NS-2** respectively.



Figure S16: (A) Job's plot of **NS-1** with Cu²⁺ for the stoichiometry determination monitoring the change of absorbance at 395 nm **(B)** Benesi–Hildebrand plot for absorbance of Cu²⁺ with **NS-1** for the association constant determination.

Figure S17: Job's plot of **NS-2** with Cu²⁺ for the stoichiometry determination monitoring the change of absorbance at 295 nm.

Figure S18: Selectivity of (A) NS-1 and (B) NS-2 toward Cu²⁺.

Figure S19: Cross-interference studies of **(A) NS-1** and **(B) NS-2** toward Cu²⁺in presence of other competing cations in similar environment.

Figure S20: Naked eye chromogenic changes of **(A) (1)** pyridine-2-aldehyde, **(2)** pyridine-2-aldehyde+Cu²⁺, **(B) (1)** 2-amino-2 methyl-1-propanol, **(2)** 2-amino-2 methyl-1-propanol +Cu²⁺ and **(C) (1)** in-situ generated ligand **(2)** in-situ generated ligand+Cu²⁺.

Figure S21: UV-Vis spectral change of pyridine-2-aldehyde in presence of Cu²⁺

Figure S22: ESI-MS data of NS-2 after interaction with Cu²⁺.

Figure S23: ESI-MS data of NS-1 after interaction with Cu²⁺.

Figure S24: (A) UV-Vis absorption spectrum of **NS-2** (100 μ M) in acetonitrile (B) UV-Vis spectral alteration of 10 μ M **NS-2** in acetonitrile in presence of 10 μ M Cu²⁺spiked human urine sample.

Figure S28: Change of absorbance at 295 nm as the linear function of the concentration of Cu^{2+} for the calculation of LOD of **NS-2** toward Cu^{2+} in human urine sample.

Table S12: Comparative literature survey of the recently developed complex based chemosensors for the detection of Cu²⁺

Entry	Solvent Medium	LOD	Solid state detection	Detection from real field sample	Cu ²⁺ detection from human urine: A potential biomarker sensor for Wilson's disease diagnosis	Ref.
1.	H ₂ O/DMSO =9/1 (v/v)	4.6 nM	No	No	No	X1
2.	Buffer solution	77 nM	No	Yes	No	X2
3.	HEPES- buffer	2.77 μM	No	No	No	Х3
4.	H ₂ O with 0.5% DMSO	6.1 μM	Yes	Yes	No	X4
5.	H ₂ O/EtOH (7: 3 v/v)	-	No	Yes	No	X5
6.	HEPES buffered (pH 7.0, 10 mM)	-	No	No	No	X6

	water					
7.	THF/H ₂ O, (1/1	8.95×10 ⁻⁸	Yes	Yes	No	X7
	v/v)	М				
8.	CH ₃ CN:H ₂ O	1.53 μM	No	No	No	X8
	= 7:3 v/v					
9.	H ₂ O	20 µM	Yes	Yes	No	X9
10.	Acetonitrile	0.03 μM	Yes	Yes	No	X10
	-water (1:1)					
11.	CH₃CN	0.08 µM	No	No	No	X11
12.	(HEPES:CH3	5 nM	No	Yes	No	X12
	CN (1:1), pH					
	7.4					
13.	CH ₃ CN–H ₂ O	-	No	No	No	X13
	solution					
	(4:1 <i>,</i> v/v)					
14.	MeCN/ H ₂ O	-	No	No	No	X14
	(9/1)					
15.	H₂O	NS-1: 48.6	Yes	Yes	Yes	Prese
		fM				nt
		NS-2: 2.4				work
		μM				

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