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Supporting information

Mixed-ligand copper(II) Schiff base complexes: the vital role of co-ligands in DNA/protein interactions and cytotoxicity

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Figure captions

Fig S1. ¹H NMR spectra of L and Cu(II) complexes.

Fig S2. ESI-MS spectrum of L.

- Fig S3. ESI-MS spectrum of complex 1.
- Fig S4. ESI-MS spectrum of complex 2.
- Fig S5. ESI-MS spectrum of complex 3.
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Fig S7. Stability properties of L and complexes 1-4.

- Fig S8. DNA studies of CV in complexes 2 and 3.
- Fig S9. DNA studies of DPV in complexes 2 and 3.
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- Fig S11. Cell viability of AGS gastric cancer cells after exposure with L for 24 h.
- **Table S1.** Calculate binding energy in molecular docking studies.
- Table S2. Means of inhibition zones diameter obtained by agar well diffusion method.
- **Table S3.** Means of inhibition zones diameter in *Staphylococcus aureus*.
- **Table S4.** Means of inhibition zones diameter in *Pseudomonas aeruginosa*.



Fig. S1 ¹H NMR spectra of (a) ligand using DMSO-d₆ solvent (b) Cu(II) complexes of **1** and (c) **2** using in CD₃CN solvent.



Fig. S2 ESI-MS spectrum of L.



Fig. S3 ESI-MS spectrum of complex 1.



Fig. S4 ESI-MS spectrum of complex 2.



Fig. S5 ESI-MS spectrum of complex 3.



Fig. S6 ESI-MS spectrum of complex 4.



Fig. S7 Stability of L and complexes **1-4** were monitored for 0-24 h in MeOH/Tris-HCl buffer solution at room temperature.



Fig. S8 CV of complexes **2** and **3** in the absence and presence of HS-DNA at scan rate of 100 mV/s.



Fig. S9 DPV of complexes **2** and **3** in the absence and presence of HS-DNA in buffer/CH₃CN solution.



Fig. S10 Antibacterial activity of compound with *Pseudomonas aeruginosa* [(a) L (c) **1** (e) **2** (g) **3** and (i) **4**] and *S. aureus* [(b) L (d) **1** (f) **2** (h) **3** and (j) **4**].



Fig. S11 Cell viability of AGS gastric cancer cells after exposure with L for 24 h.

Table S1 The calculated free energy, inhibitory constant and binding region of the Cu(II)

 complexes with B-DNA (1BNA)

Complexes	Free energy of binding (FEB) (kcal/mol)	Inhibitory constant (µM)	Binding region
1	-4.66	383.0	Guanine, Cytosine, Adenine
2	-4.19	845.4	Guanine, Cytosine
3	-4.34	658.3	Guanine, Cytosine, Adenine
4	-3.99	1190.0	Guanine, Cytosine, Adenine, Thymine

Table S2 Means of inhibition zones diameter obtained by agar well diffusion method against

 Staphylococcus aureus and Pseudomonas aeruginosa.

Complex	Means of zones of bacterial growth inhibition (mm)				
	S. aureus	P. aeruginosa			
1	14	17			
2	12	14			
3	13	15			
4	12	11			

Table S3: Means of inhibition zones diameter obtained by agar well diffusion method against

 Staphylococcus aureus.

	Zones of growth inhibition (mm)							
Compound	10	25	50	100	250	500	1000	2500
	(µg/mL)	(µg/mL)	(µg/mL)	(µg/mL)	(µg/mL)	(µg/mL)	(µg/mL)	(µg/mL)
L	7	7	7	8	9	9	10	10
1	10	10	11	13	15	16	18	22
2	10	10	10	11	13	14	16	17
3	10	10	10	12	14	16	19	17
4	10	10	10	10	12	14	15	20

	Zones of growth inhibition (mm)							
Compound	10	25	50	100	250	500	1000	2500
	(µg/mL)	(µg/mL)	(µg/mL)	(µg/mL)	(µg/mL)	(µg/mL)	(µg/mL)	(µg/mL)
L	6	6	6	6	8	8	11	11
1	11	11	16	18	19	20	21	23
2	10	10	14	15	14	17	18	20
3	10	11	12	16	17	18	20	22
4	10	10	10	10	12	13	14	15

Table S4: Means of inhibition zones diameter obtained by agar well diffusion method against*Pseudomonas aeruginosa*.