

Supporting Information

Discovery of copper(II) complexes with plumbagin and bipyridine as chemodynamic therapy agents with enhanced antitumor activity by targeting mitochondria

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1. The characterization of Cu1–Cu4

P1_220726104258 #1 RT: 0.01 AV: 1 NL: 7.15E7
T: FTMS + p ESI Full ms [220.00-800.00]

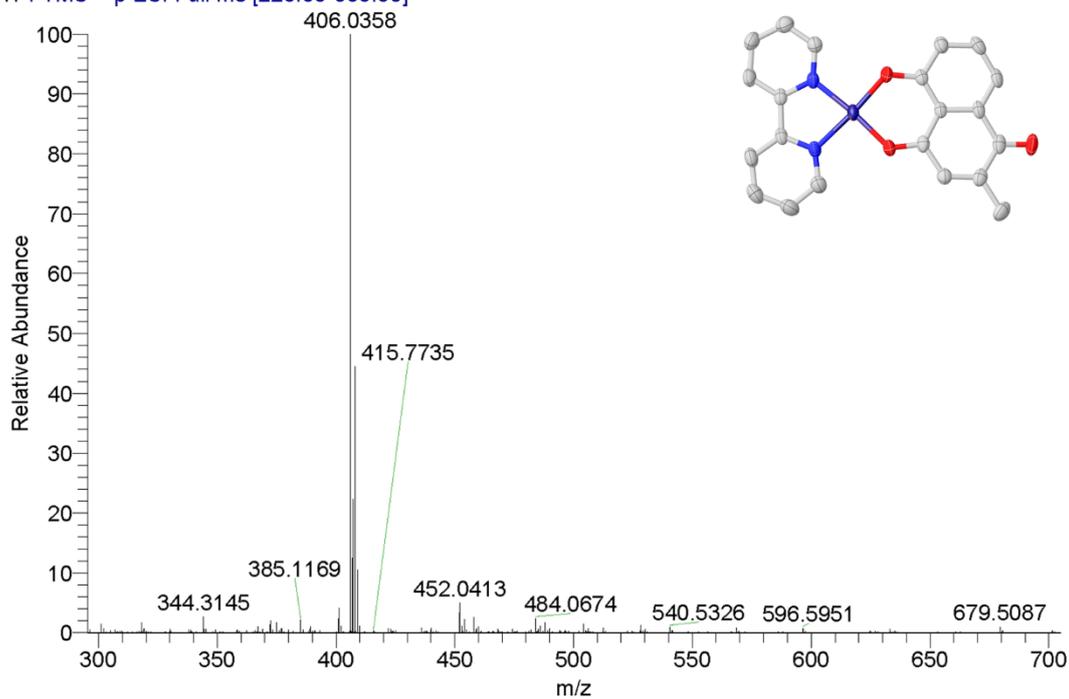


Figure S1. HRMS spectrum of **Cu1**.

P2_220726105159 #1 RT: 0.01 AV: 1 NL: 6.03E8
T: FTMS + p ESI Full ms [220.00-800.00]

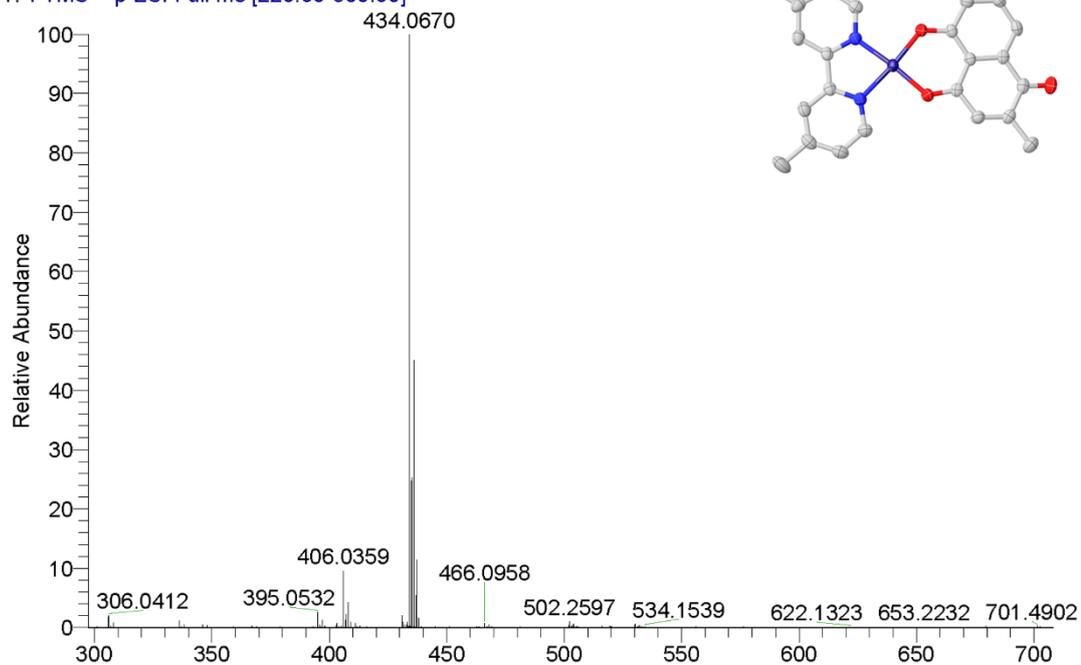


Figure S2. HRMS spectrum of **Cu2**.

F2_220726112230 #1 RT: 0.01 AV: 1 NL: 4.17E8
T: FTMS + p ESI Full ms [200.00-800.00]

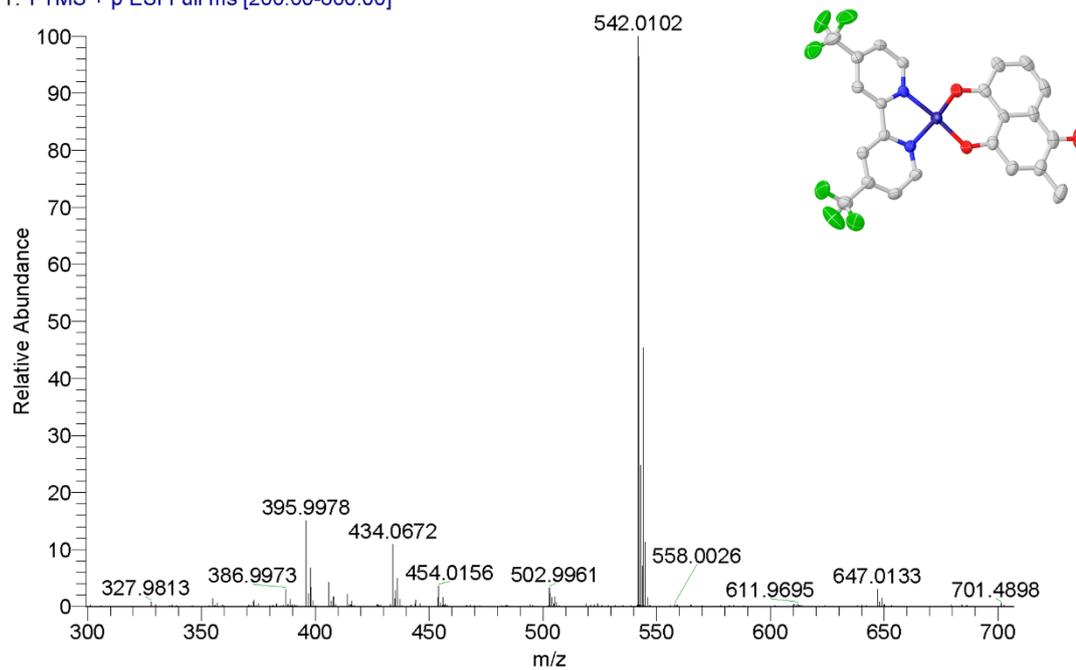


Figure S3. HRMS spectrum of **Cu3**.

N_220726113050 #1 RT: 0.01 AV: 1 NL: 7.76E7
T: FTMS + p ESI Full ms [200.00-800.00]

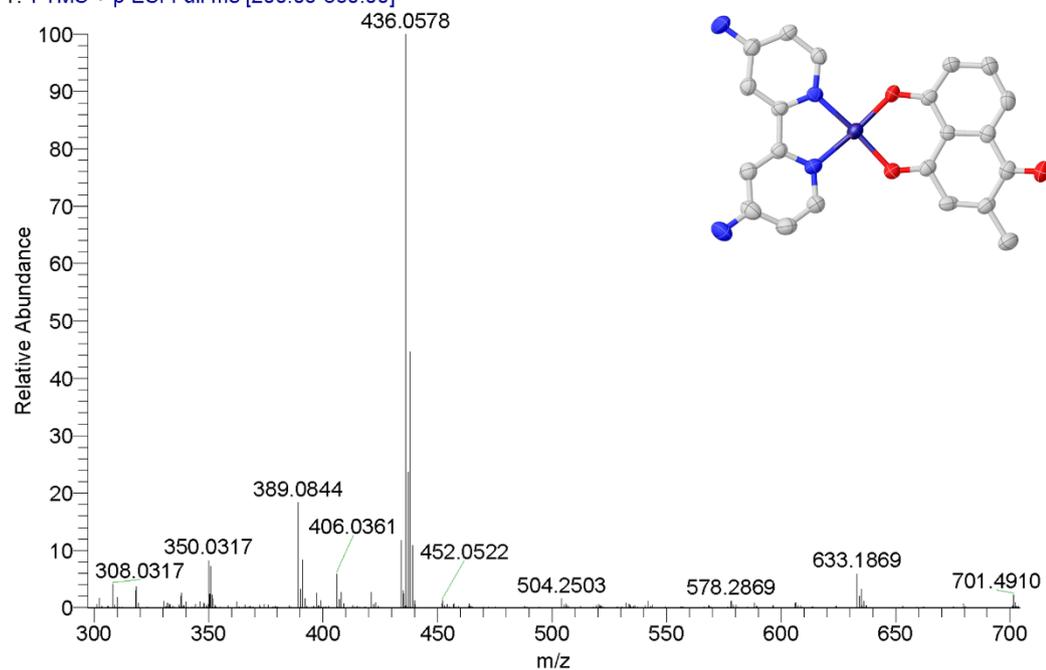


Figure S4. HRMS spectrum of **Cu4**.

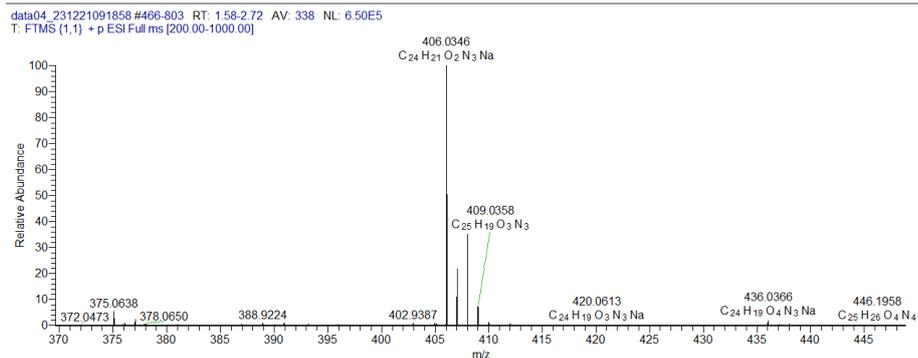
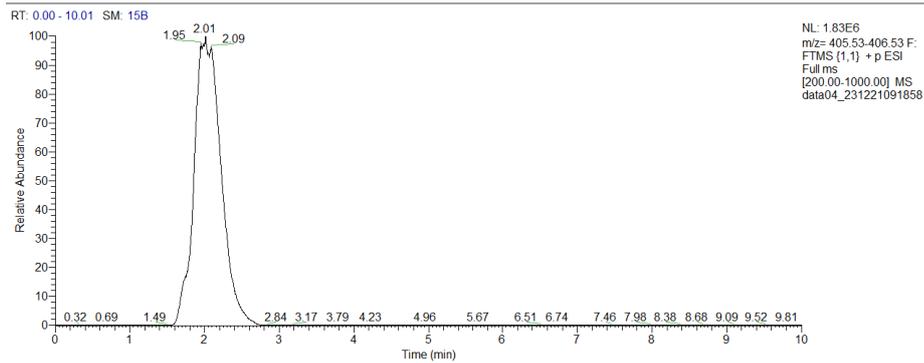


Figure S5. LC-MS spectrum of Cu1.

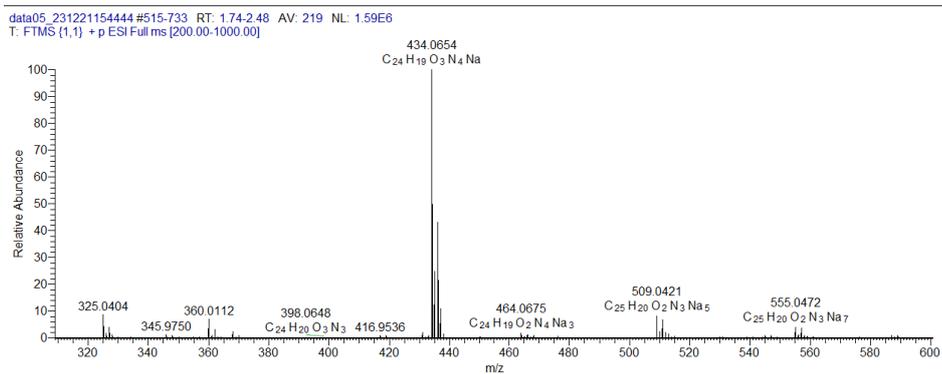
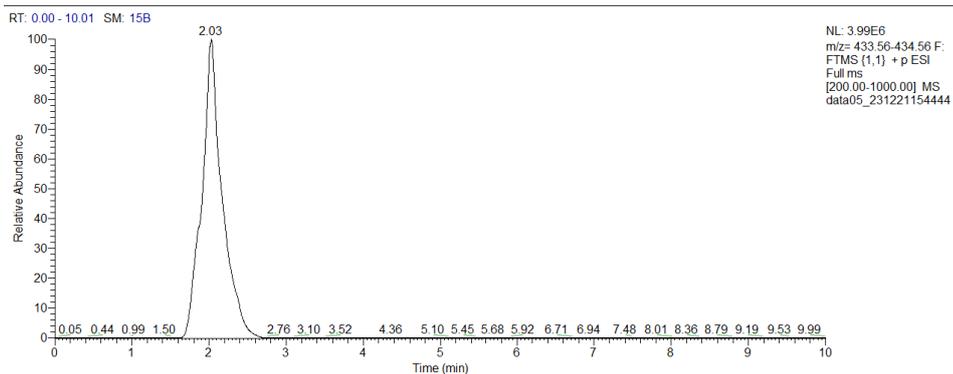


Figure S6. LC-MS spectrum of Cu2.

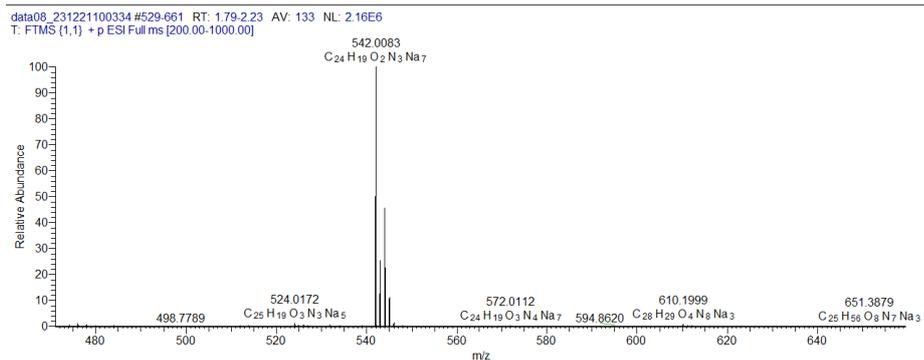
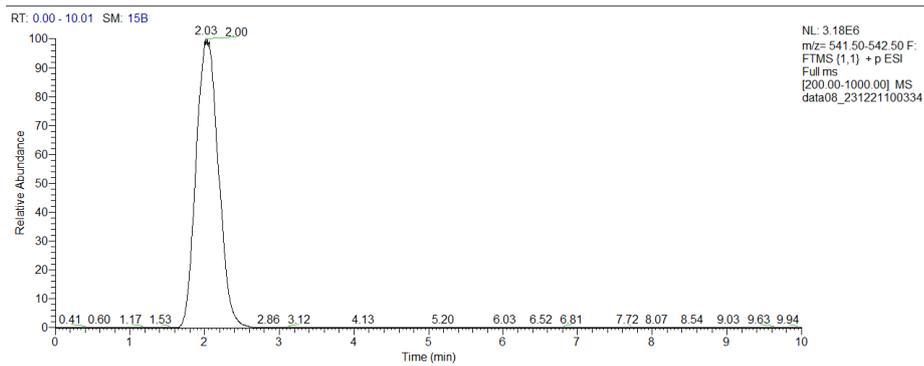


Figure S7. LC-MS spectrum of Cu3.

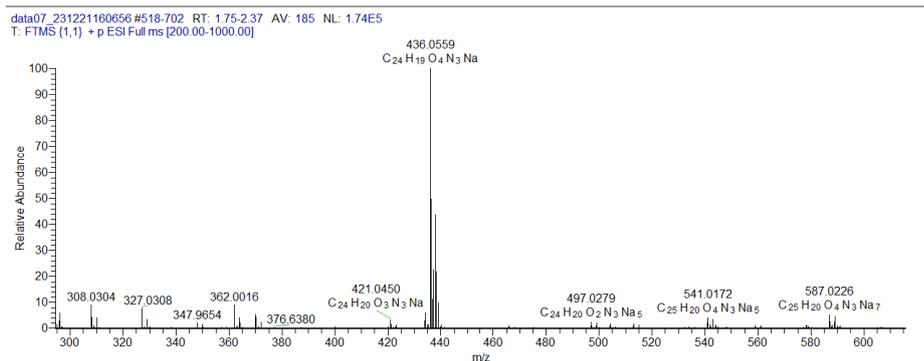
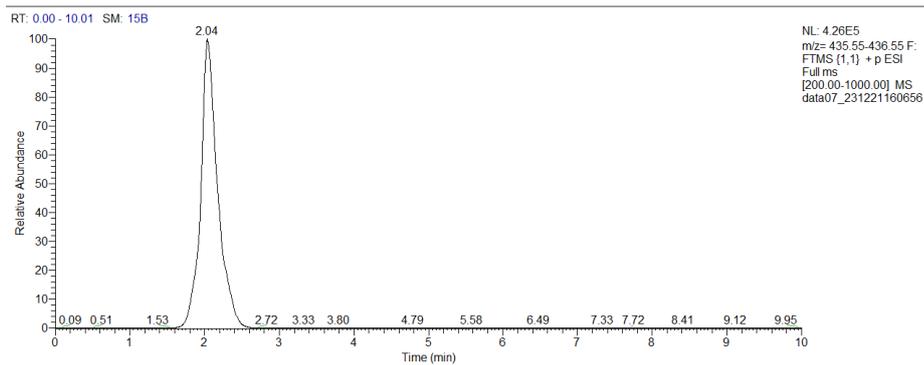


Figure S8. LC-MS spectrum of Cu4.

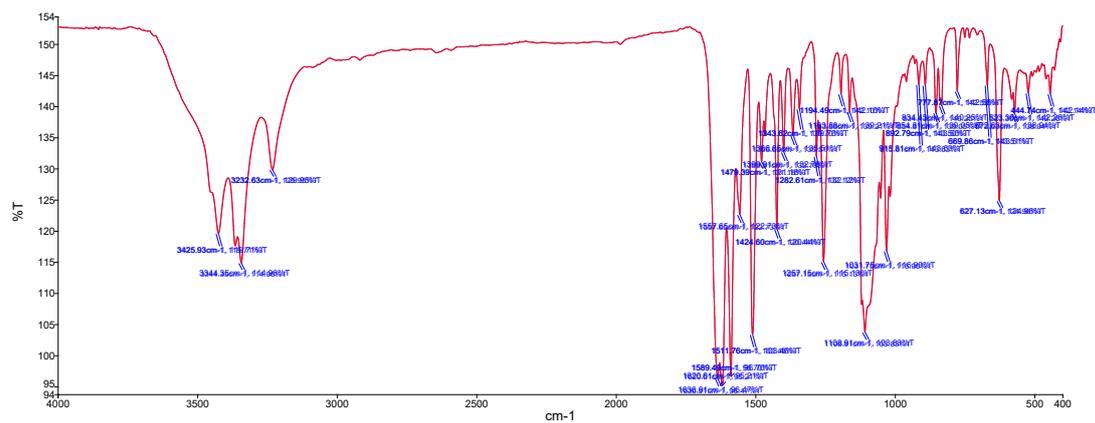


Figure S12. IR spectrum of **Cu4**.

2. Stability of Cu1–Cu4 in PBS

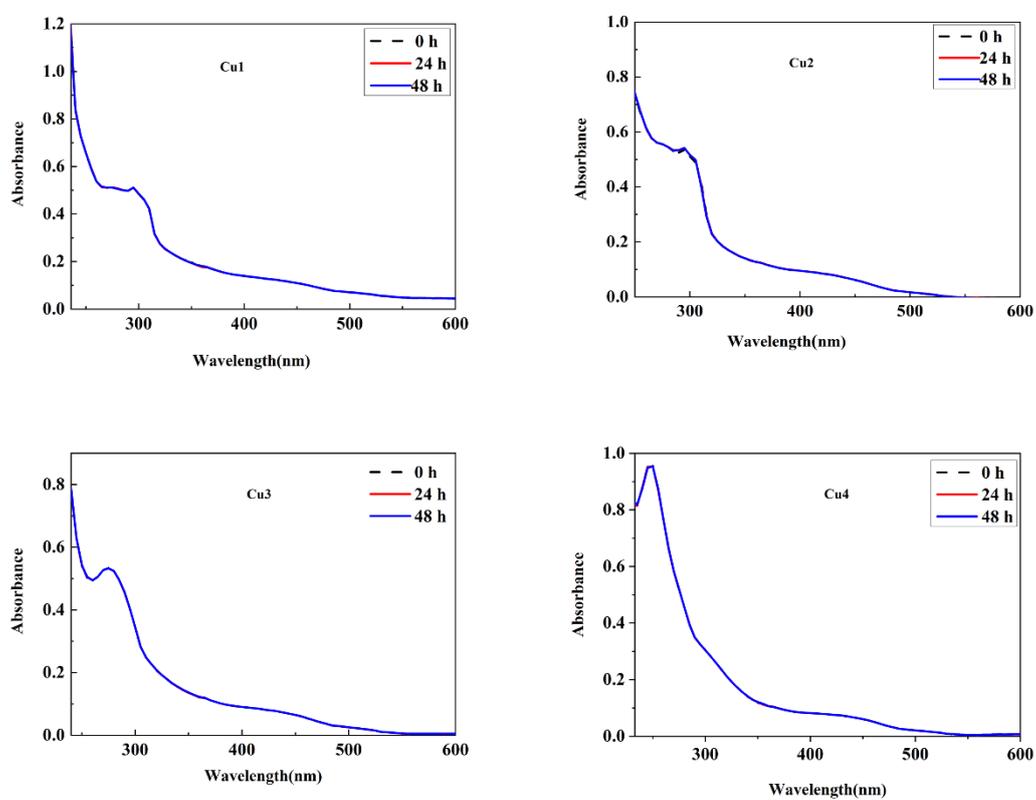


Figure S13. UV-Vis absorption spectra of **Cu1–Cu4** (2.0×10^{-3} M) in PBS buffer containing 1% DMSO solution in the time course 0 h, 24 h, and 48 h.

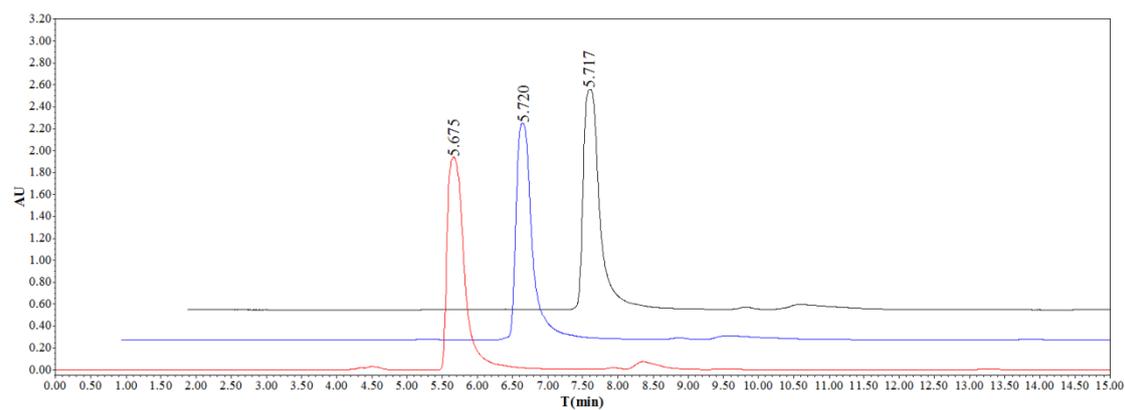


Figure S14. HPLC spectra of **Cu1** (2.0×10^{-3} M) at time 0 h, 24 h, and 48 h in DMSO/CH₃OH.

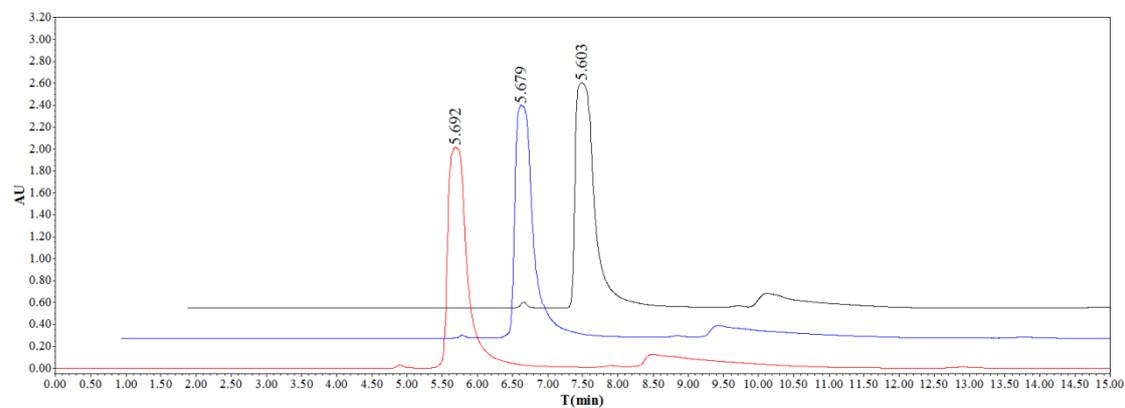


Figure S15. HPLC spectra of **Cu2** (2.0×10^{-3} M) at time 0 h, 24 h, and 48 h in DMSO/CH₃OH.

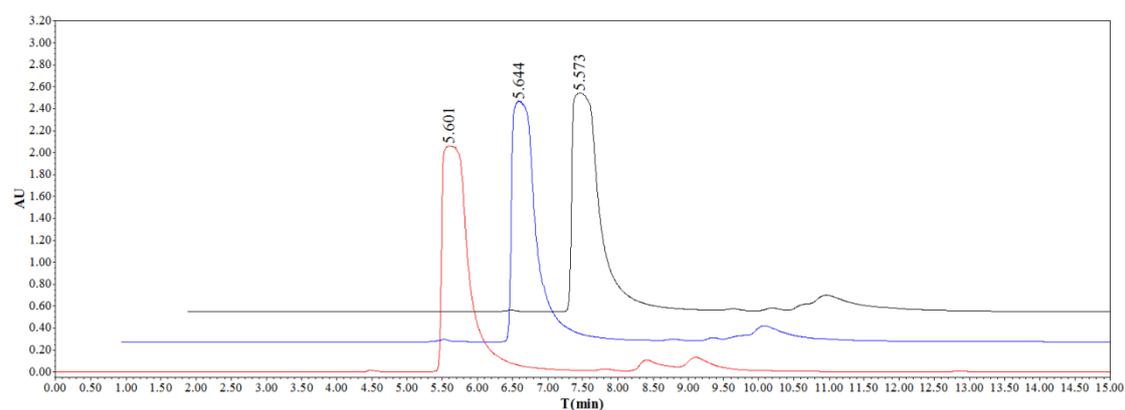


Figure S16. HPLC spectra of **Cu3** (2.0×10^{-3} M) at time 0 h, 24 h, and 48 h in DMSO/CH₃OH.

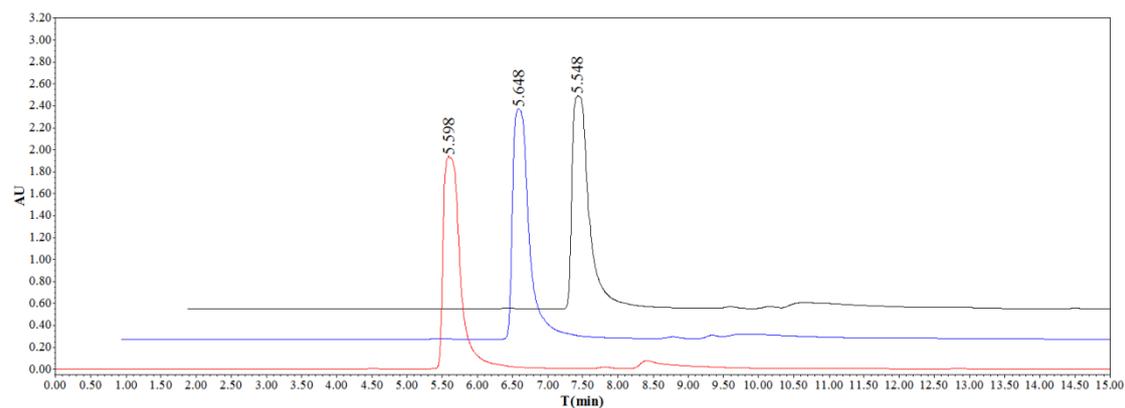


Figure S17. HPLC spectra of **Cu4** (2.0×10^{-3} M) at time 0, 24 h, and 48 h in DMSO/CH₃OH.

3. Solution behavior of Cu1–Cu4

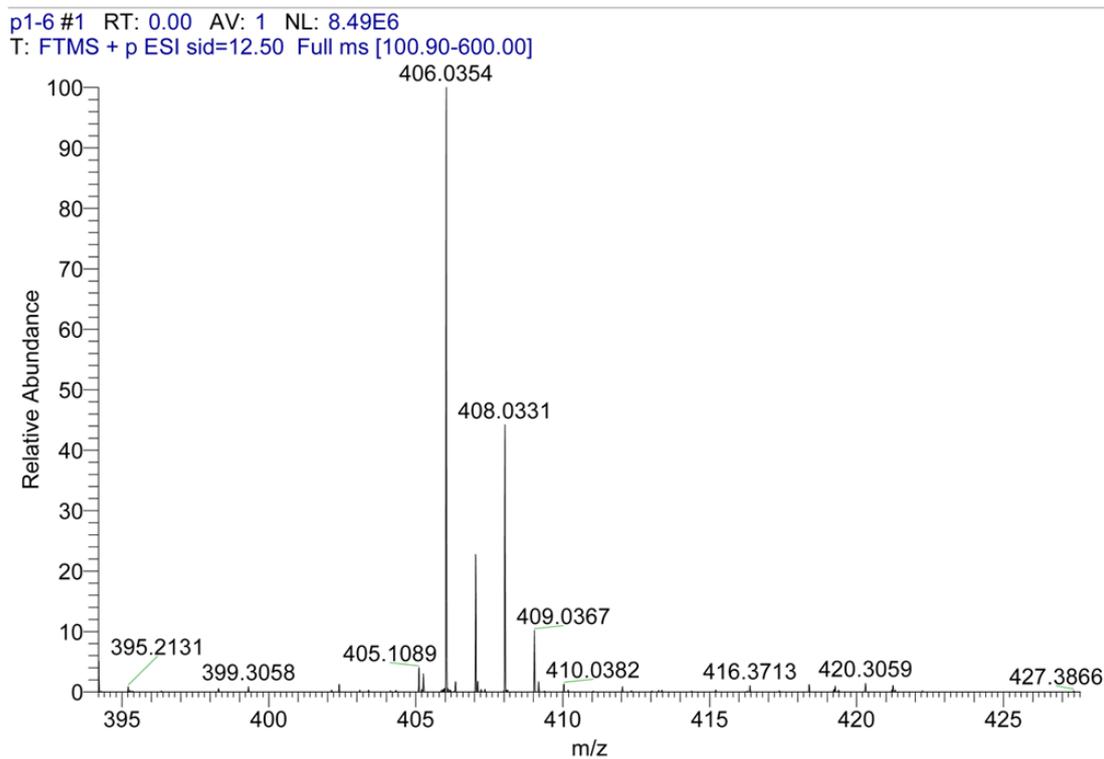


Figure S18. Mass spectrum of **Cu1** in solution.

p2-6 #13-23 RT: 0.05-0.08 AV: 11 NL: 1.64E7
T: FTMS + p ESI sid=12.50 Full ms [200.90-650.00]

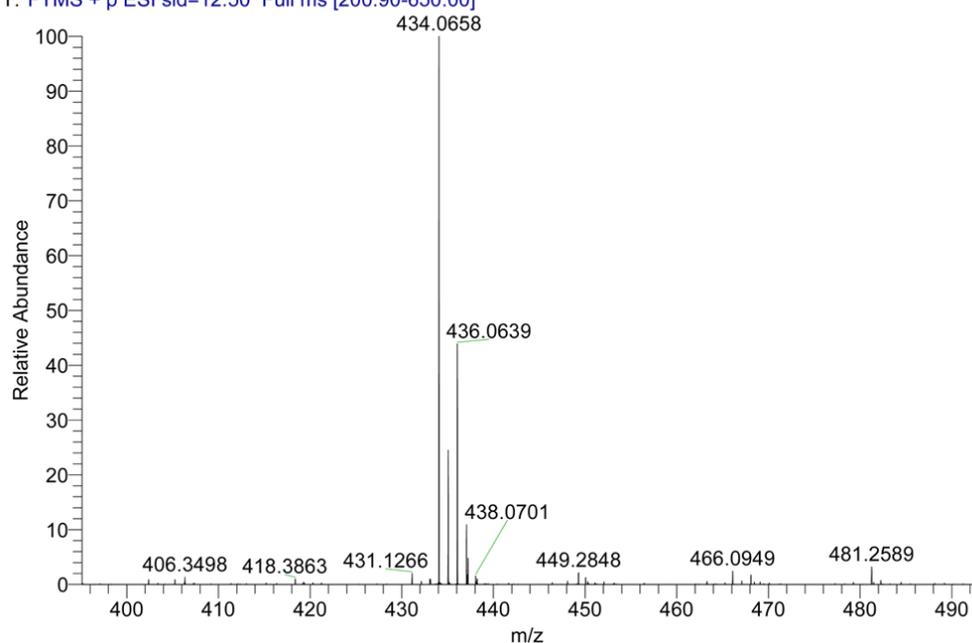


Figure S19. Mass spectrum of Cu_2 in solution.

F2-4 #1 RT: -0.00 AV: 1 NL: 3.87E7
T: FTMS + p ESI sid=12.50 Full ms [300.00-700.00]

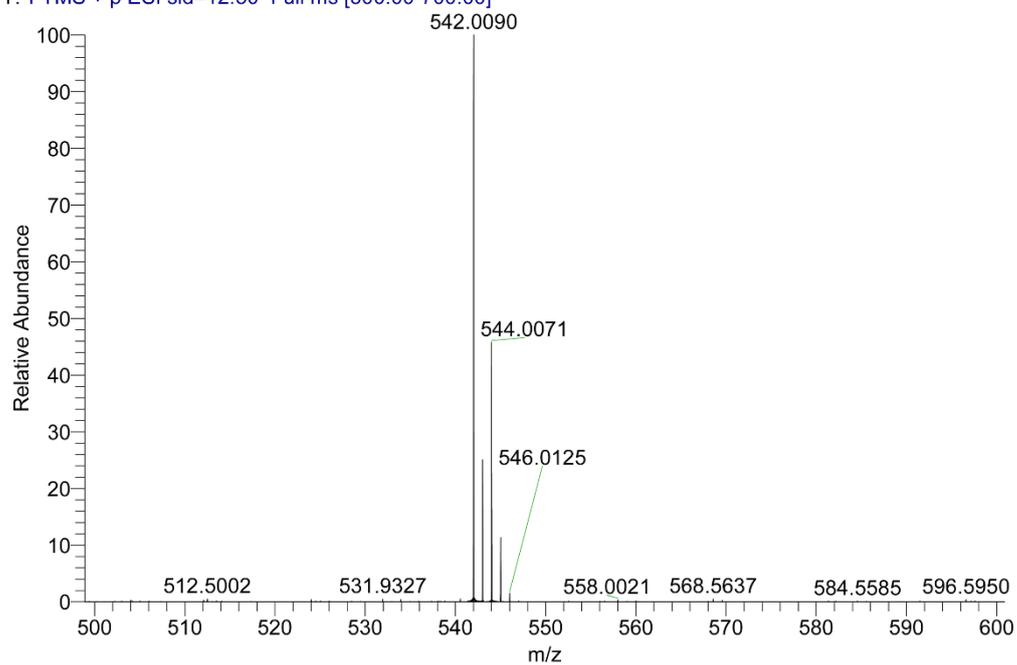


Figure S20. Mass spectrum of Cu_3 in solution.

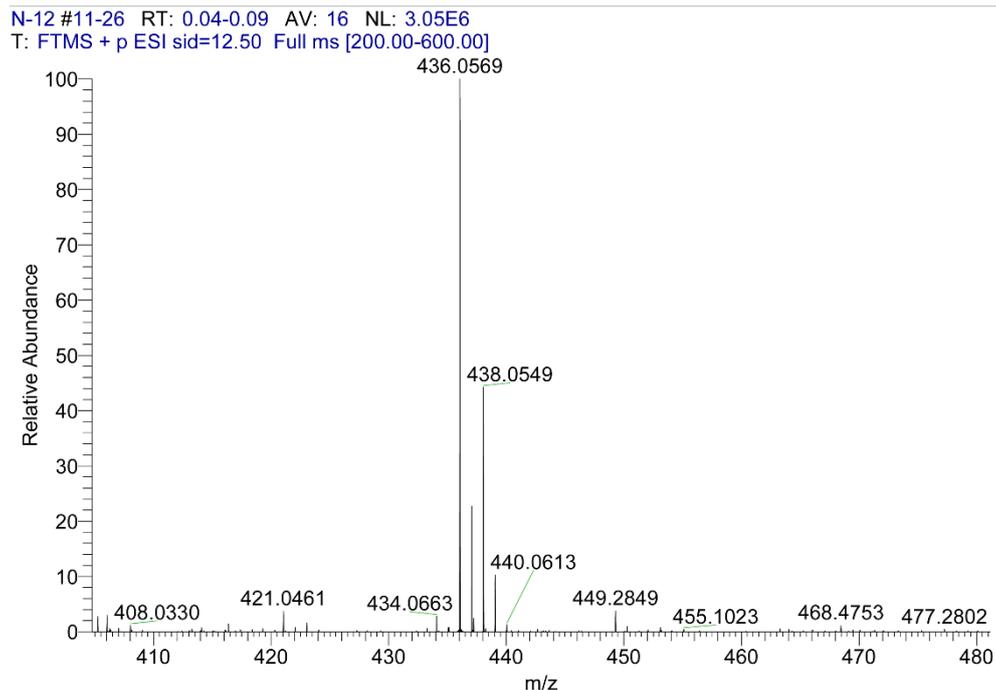


Figure S21. Mass spectrum of **Cu4** in solution.

4. Tables of crystal data of **Cu1–Cu4**

Table S1 Crystal data and structure refinement for **Cu1**

Complex	Cu1
Empirical formula	$C_{21}H_{15}ClCuN_2O_7$
Formula weight (M)	506.34
Crystal system	<i>Triclinic</i>
Space group	<i>P</i> -1
<i>a</i> (Å)	9.2985(5)
<i>b</i> (Å)	10.3543(7)
<i>c</i> (Å)	11.9616(7)
α (°)	112.775(6)
β (°)	107.363(5)
γ (°)	93.359(5)
<i>V</i> /(Å ³)	993.44(11)
<i>Z</i>	2

$D_c(\text{Mg m}^{-3})$	1.693
$F(000)$	514.0
θ range for data collection ($^\circ$)	3.4–26.3 $^\circ$
Reflections collected / unique	12106/4043 [$R_{(\text{int})} = 0.0417$]
Goodness-of-fit on F^2	1.049
R indices [$I > 2\sigma(I)$]	$R_1 = 0.0527$ $\omega R_2 = 0.1248$
R indices (all data)	$R_1 = 0.0791$ $\omega R_2 = 0.1451$

Table S2 Crystal data and structure refinement for **Cu2**

Complex	Cu2
Empirical formula	$\text{C}_{23}\text{H}_{19}\text{ClCuN}_2\text{O}_7$
Formula weight (M)	534.39
Crystal system	<i>Monoclinic</i>
Space group	$P2_1/n$
a (\AA)	9.3791(2)
b (\AA)	10.6291(2)
c (\AA)	22.1108(4)
α ($^\circ$)	90
β ($^\circ$)	92.1043(17)
γ ($^\circ$)	90
V (\AA^3)	2202.78(8)
Z	4
$D_c(\text{Mg m}^{-3})$	1.611
$F(000)$	1092.0
θ range for data collection ($^\circ$)	3.4–26.3 $^\circ$
Reflections collected / unique	13491/4498 [$R_{(\text{int})} = 0.0257$]
Goodness-of-fit on F^2	1.040
R indices [$I > 2\sigma(I)$]	$R_1 = 0.0369$ $\omega R_2 = 0.0922$
R indices (all data)	$R_1 = 0.0453$ $\omega R_2 = 0.0985$

Table S3 Crystal data and structure refinement for **Cu3**

Complex	Cu3
Empirical formula	C ₂₃ H ₁₃ ClCuF ₆ N ₂ O ₇
Formula weight (M)	642.34
Crystal system	<i>monoclinic</i>
Space group	<i>P2₁/c</i>
<i>a</i> (Å)	9.5260(4)
<i>b</i> (Å)	16.5988(5)
<i>c</i> (Å)	15.9965(5)
α (°)	90
β (°)	105.658(4)
γ (°)	90
<i>V</i> /(Å ³)	2435.50(15)
<i>Z</i>	4
<i>D_c</i> (Mg m ⁻³)	1.752
<i>F</i> (000)	1284.0
θ range for data collection (°)	3.3–28.8°
Reflections collected / unique	16165/5645 [<i>R</i> _(int) = 0.0429]
Goodness-of-fit on <i>F</i> ²	1.060
<i>R</i> indices [<i>I</i> > 2σ(<i>I</i>)]	<i>R</i> ₁ = 0.0612 ω <i>R</i> ₂ = 0.1314
<i>R</i> indices (all data)	<i>R</i> ₁ = 0.1023 ω <i>R</i> ₂ = 0.1518

Table S4 Crystal data and structure refinement for **Cu4**

Complex	Cu4
Empirical formula	C ₂₁ H ₁₇ ClCuN ₄ O ₇
Formula weight (M)	536.38
Crystal system	<i>triclinic</i>
Space group	<i>P</i> -1
<i>a</i> (Å)	9.4798(5)
<i>b</i> (Å)	10.4607(8)
<i>c</i> (Å)	12.3592(8)
α (°)	109.072(7)
β (°)	110.542(6)
γ (°)	90.238(5)
<i>V</i> /(Å ³)	1074.86(14)
<i>Z</i>	2
<i>D_c</i> (Mg m ⁻³)	1.657
<i>F</i> (000)	564.0
θ range for data collection (°)	3.2–28.8°
Reflections collected / unique	16062/5050 [<i>R</i> _(int) = 0.0872]
Goodness-of-fit on <i>F</i> ²	1.001
<i>R</i> indices [<i>I</i> > 2σ(<i>I</i>)]	<i>R</i> ₁ = 0.0708 ω <i>R</i> ₂ = 0.1518
<i>R</i> indices (all data)	<i>R</i> ₁ = 0.1551 ω <i>R</i> ₂ = 0.1959

5. The HRMS spectra of Cu1–Cu4 reacted with GSH, separately

Cu1 GSH#1 RT: 0.00 AV: 1 NL: 1.09E5
T: FTMS - p ESI Full ms [100.00-800.00]

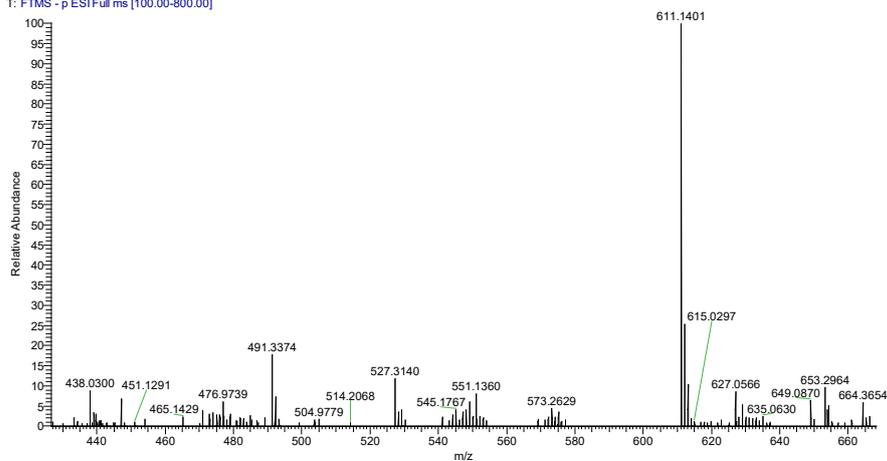


Figure S22. HRMS spectrum of Cu1 with GSH.

Cu2 GSH#1 RT: 0.00 AV: 1 NL: 1.40E5
T: FTMS - p ESI Full ms [100.00-800.00]

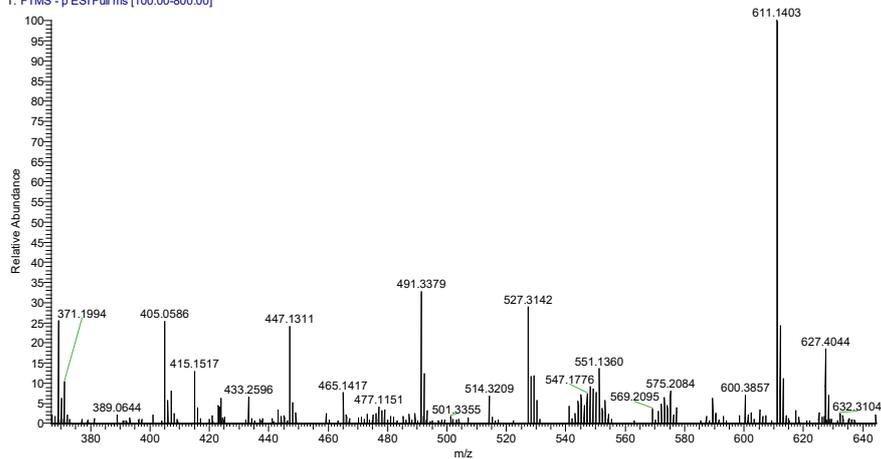


Figure S23. HRMS spectrum of Cu2 with GSH.

Cu3 GSH#1 RT: 0.00 AV: 1 NL: 1.07E4
T: FTMS - p ESI Full ms [100.00-800.00]

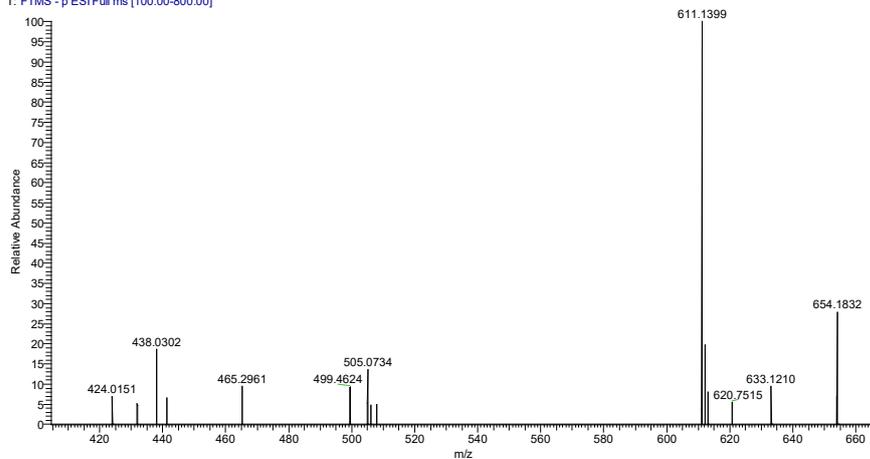


Figure S24. HRMS spectrum of Cu3 with GSH.

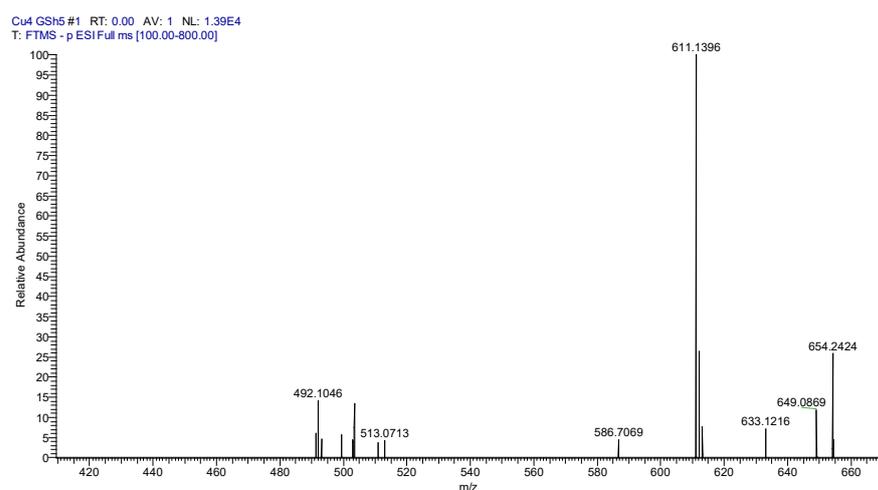


Figure S25. HRMS spectrum of Cu4 with GSH.

6. Intracellular GSH / GSSG ratio

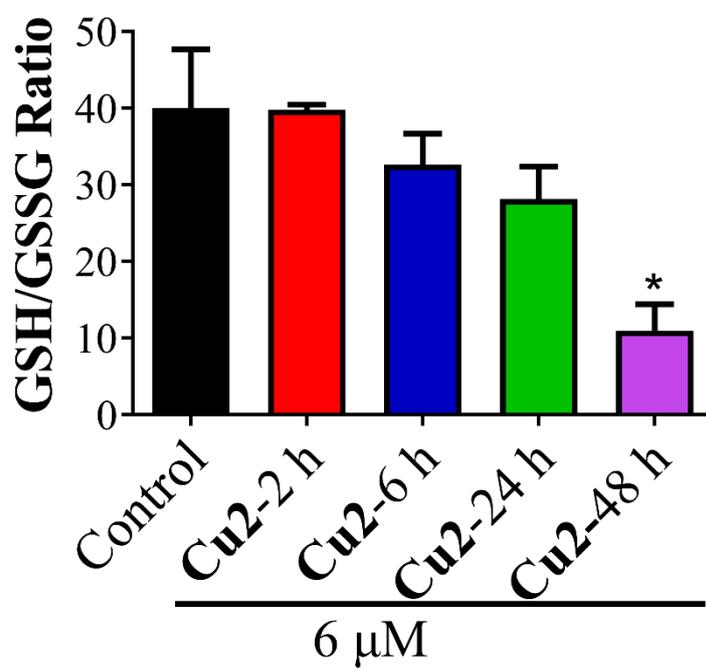


Figure S26. GSH/GSSG ratios in HeLa cells after 2 h, 6 h, 24 h and 48 h of exposure to Cu2 (6 μM, *P < 0.05, ** P < 0.01, *** P < 0.001) .

7. Cell cycle arrest of Cu1–Cu4

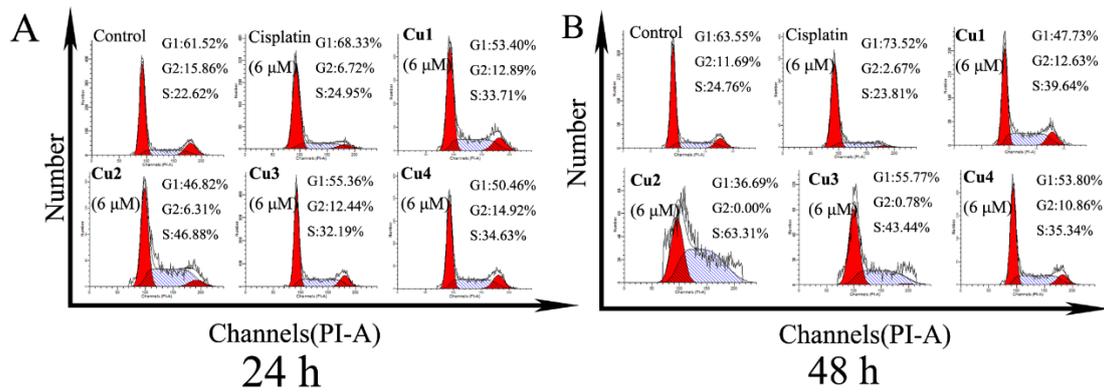


Figure S27. (A)(B) Cell cycle analysis of HeLa cells that treated with **Cu1–Cu4** (6 μ M) and cisplatin (6 μ M) for 24 and 48 h.

8. Expression of apoptosis-related proteins induced by Cu1-Cu4

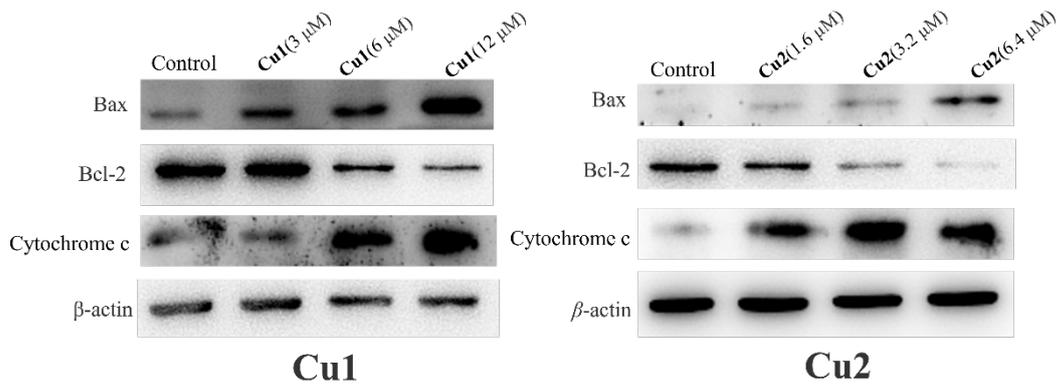


Figure S28. The expression level of apoptosis related proteins following **Cu1** and **Cu2** after exposure to 0.5x IC_{50} , 1x IC_{50} , and 2x IC_{50} .

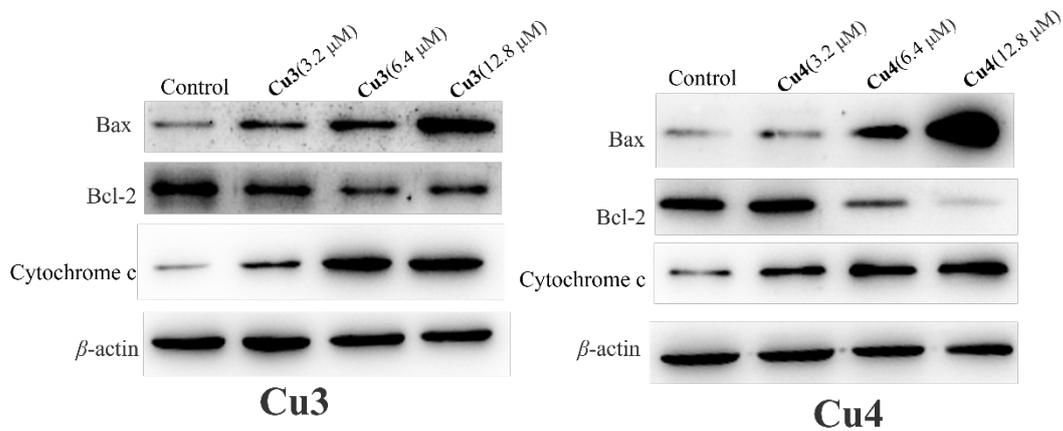


Figure S29. The expression level of apoptosis related proteins following **Cu3** and **Cu4** after exposure to 0.5x IC_{50} , 1x IC_{50} , and 2x IC_{50} .

9. pharmacokinetic evaluation of Cu2

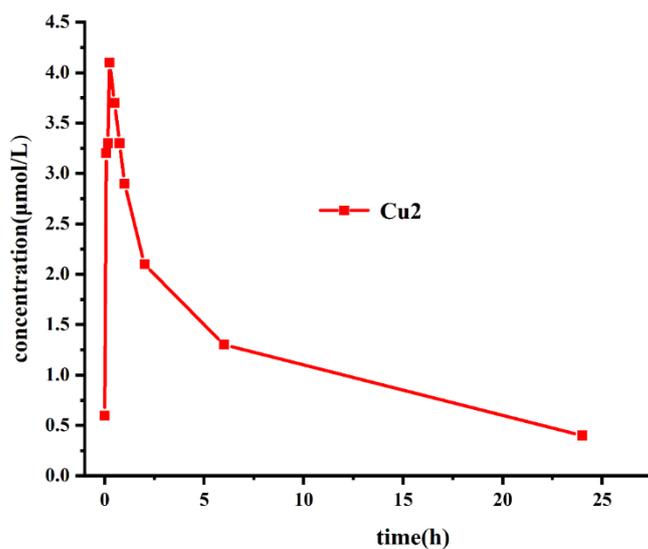


Figure S30. Pharmacokinetic evaluation in mice: concentration-time profile of **Cu2** in plasma following a single dose (20 mg/kg)

10. Acute toxicity test of Cu2

Table S5 acute toxicity of **Cu2** (alive/total)

	18.18 mg/kg	22.725 mg/kg	27.27 mg/kg
Cu2	6/6	6/6	4/6