

# Evaluation of anticancer potential of a binuclear copper (II) complex with 3-(pyridin-3-yl) coumarin ligand: Synthesis, crystal structure, magnetism, DNA interaction and molecular dynamic simulations

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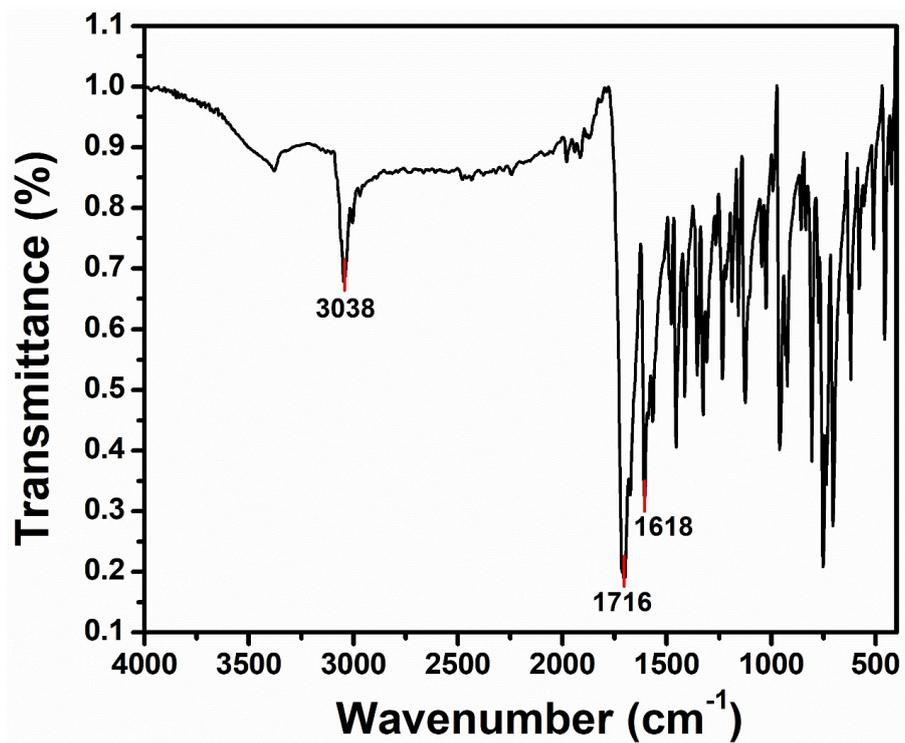


Fig. S1. IR spectrum of 3-(pyridin-3-yl) coumarin (**L**).

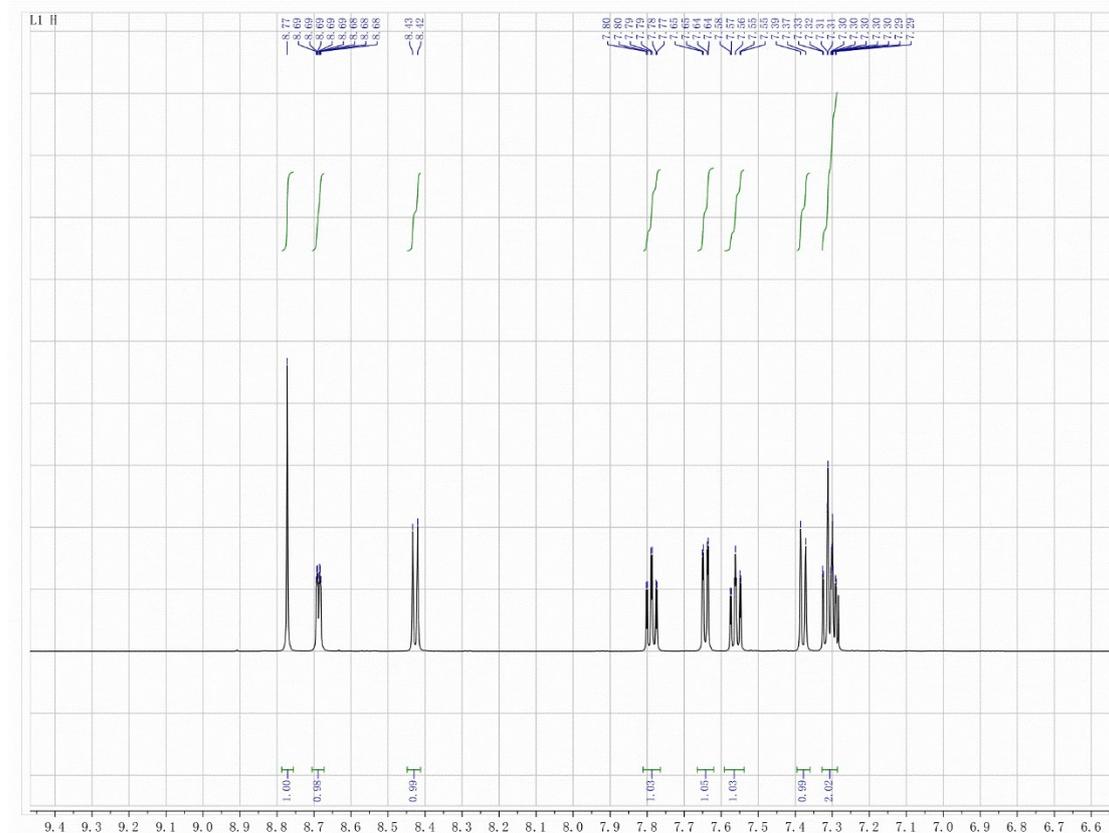
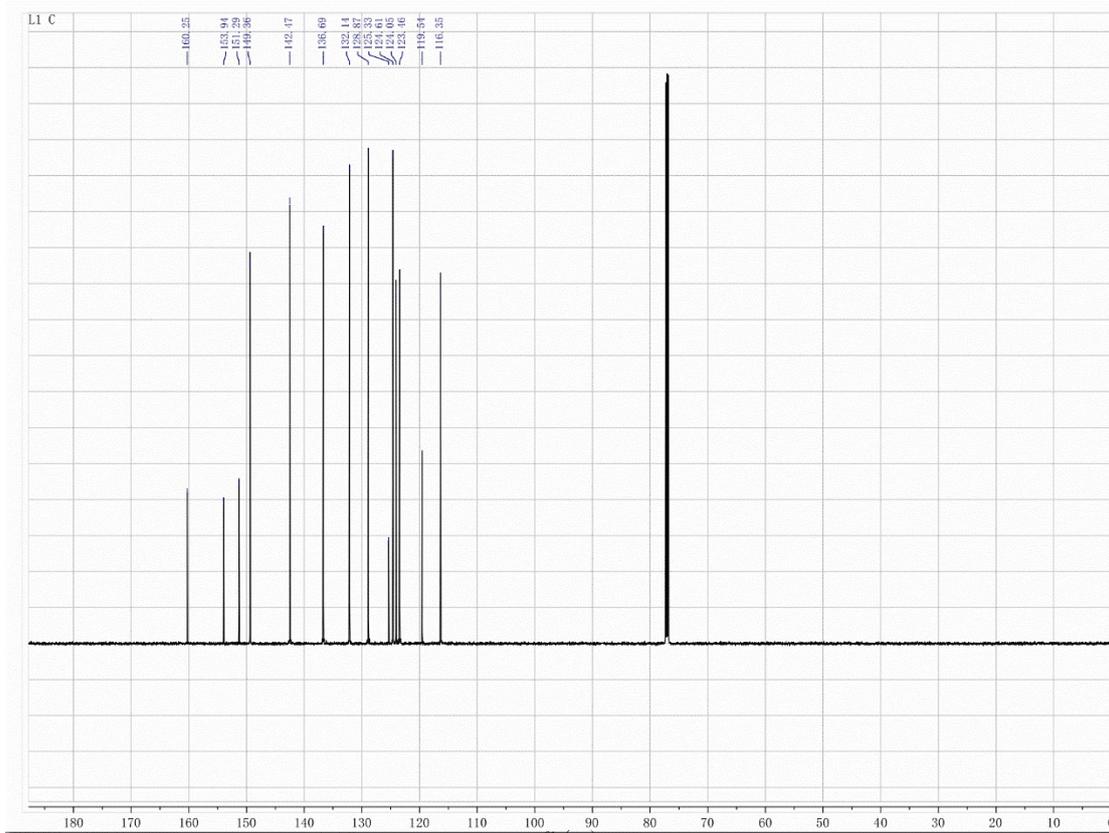
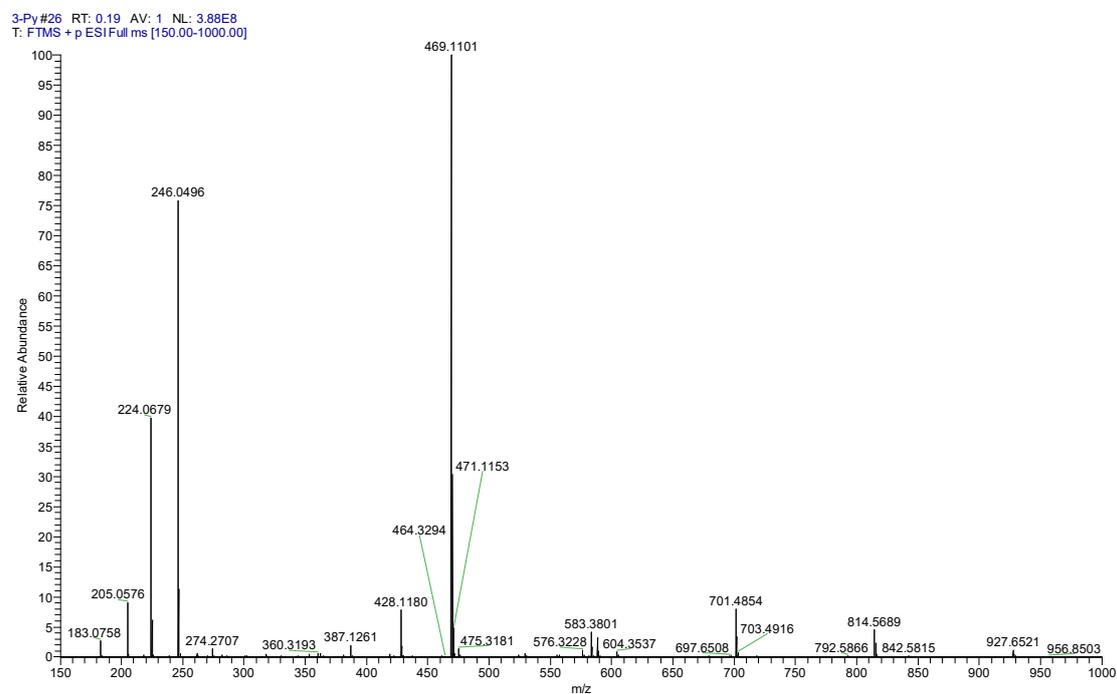


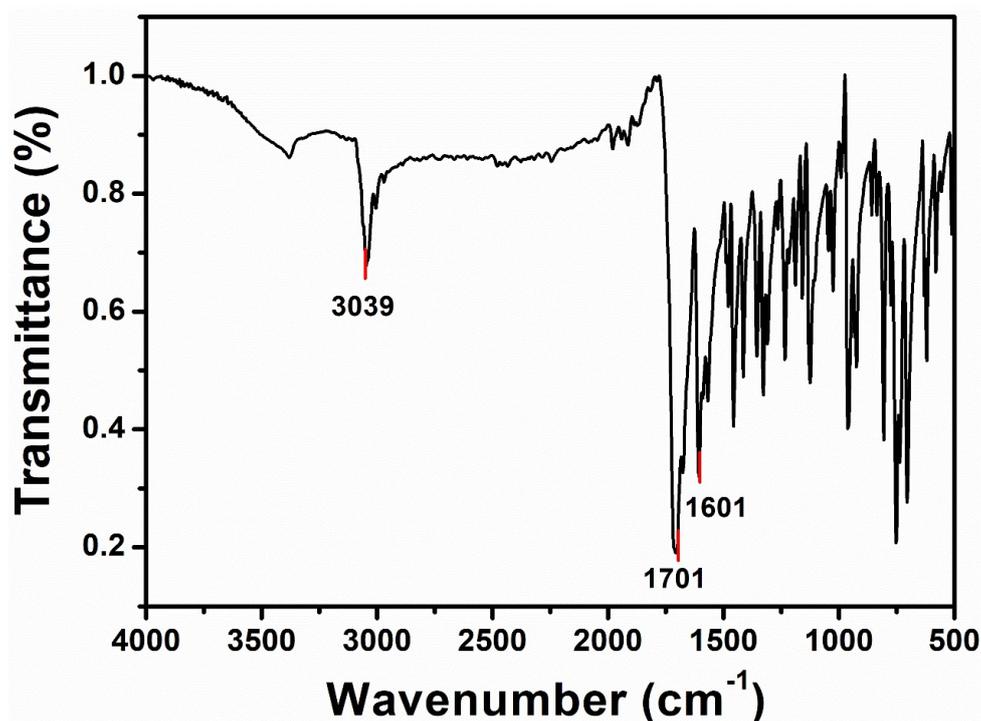
Fig. S2. <sup>1</sup>H NMR spectrum of 3-(pyridin-3-yl) coumarin (**L**) in CDCl<sub>3</sub>.



**Fig. S3.** <sup>13</sup>C NMR spectrum of 3-(pyridin-3-yl) coumarin (L) in CDCl<sub>3</sub>.



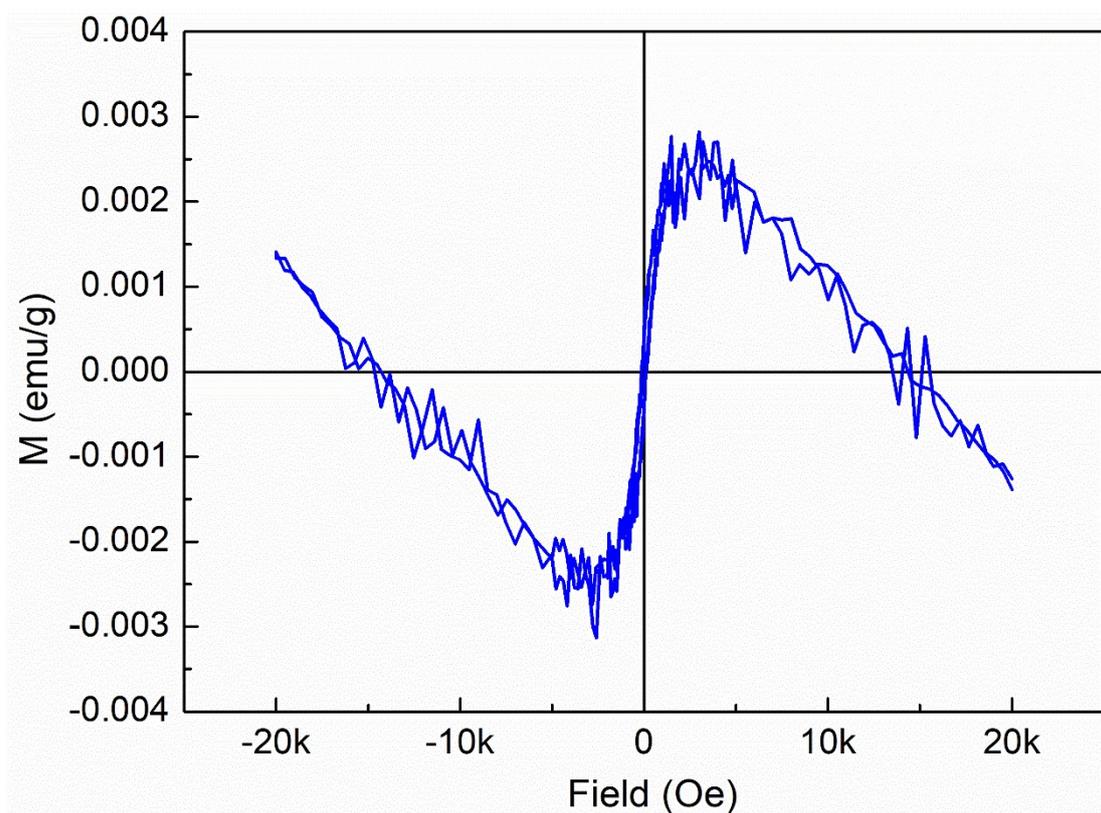
**Fig. S4.** Mass spectrum of 3-(pyridin-3-yl) coumarin (L).



**Fig. S5.** IR spectrum of complex **Coupy-Cu**.

**Table S1.** Selected Bond Lengths (Å) and Angles (°) of the complex **Coupy-Cu**.

Bond	Dist.(Å)	Bond	Dist.(Å)
Cu1-O8	1.936(12)	Cu2-O7	1.941(14)
Cu1-O1	1.939(14)	Cu2-O4	1.966(11)
Cu1-O3	1.945(13)	Cu2-O2	1.972(18)
Cu1-O6	1.964(15)	Cu2-O5	1.983(15)
Cu1-N1	2.159(17)	Cu2-N2	2.126(16)
Angle	(°)	Angle	(°)
O8-Cu1-O1	88.6(6)	O7-Cu2-O4	170.6(5)
O8-Cu1-O3	165.2(5)	O7-Cu2-O2	90.2(6)
O1-Cu1-O3	89.5(6)	O4-Cu2-O2	89.6(6)
O8-Cu1-O6	87.7(6)	O7-Cu2-O5	88.0(6)
O1-Cu1-O6	167.3(7)	O4-Cu2-O5	90.1(6)
O3-Cu1-O6	90.9(6)	O2-Cu2-O5	166.7(7)
O8-Cu1-N1	95.8(5)	O7-Cu2-N2	96.5(6)
O1-Cu1-N1	96.6(6)	O4-Cu2-N2	92.7(6)
O3-Cu1-N1	99.0(6)	O2-Cu2-N2	101.4(6)
O6-Cu1-N1	95.8(6)	O5-Cu2-N2	91.9(7)

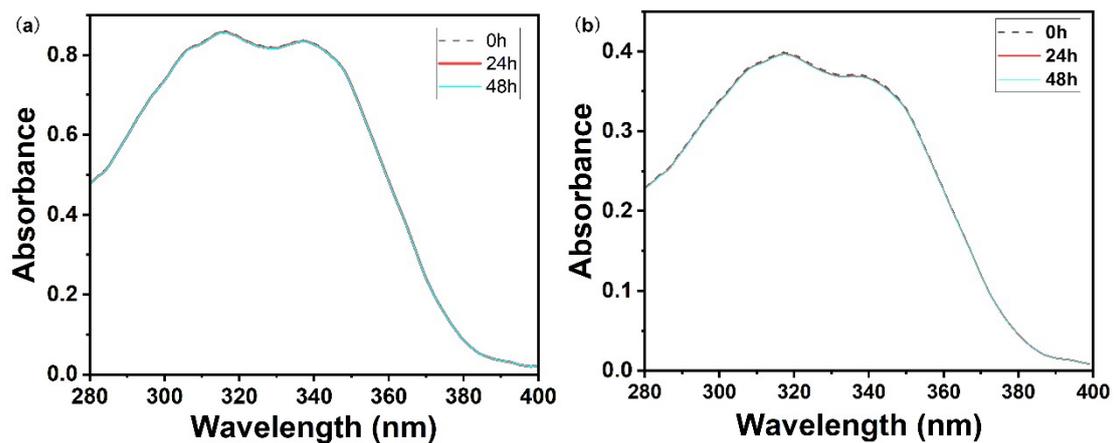


**Fig.S6.** Magnetization ( $M$ , emu/g) as a function of applied magnetic field (Oe) for the complex **Coupy-Cu**, showing the magnetic hysteresis behavior.

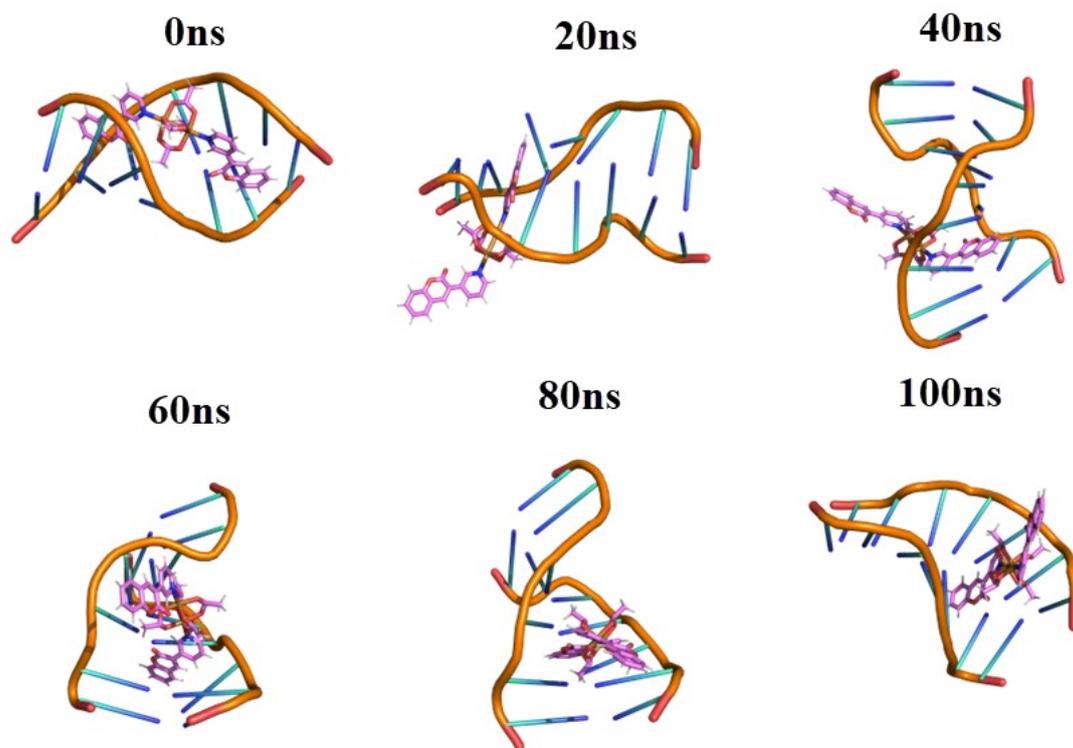
**Table S2.** Summary of simulation parameters for EPR spectra (complex Coupy-Cu).

<b><math>g</math>-value</b>	<b><math>D/\text{cm}^{-1}</math></b>	<b><math>E/\text{cm}^{-1}</math></b>	<b><math>lw/\text{mT}</math></b>
2.08 2.41	0.290	0.265	10, 35

\*  $lw$  is the homogeneous Lorentzian linewidth; the linewidth defines the FWHM (full width at half height, in mT) of the absorption Gaussian (first number) and Lorentzian (second number) broadening of the lines in the spectrum.



**Fig.S7.** UV-Vis spectra of ligand **L** (a) and complex **Coupy-Cu** (b) in Tris-HCl/NaCl buffer (pH = 7.2, with DMSO content < 1% v/v).



**Fig.S8.** Snapshot structures during the simulation.