

Supporting Information for

**Self-activating nuclease and anticancer activities of
copper(II) complexes with aryl-modified
2,6-di(thiazol-2-yl)pyridine**

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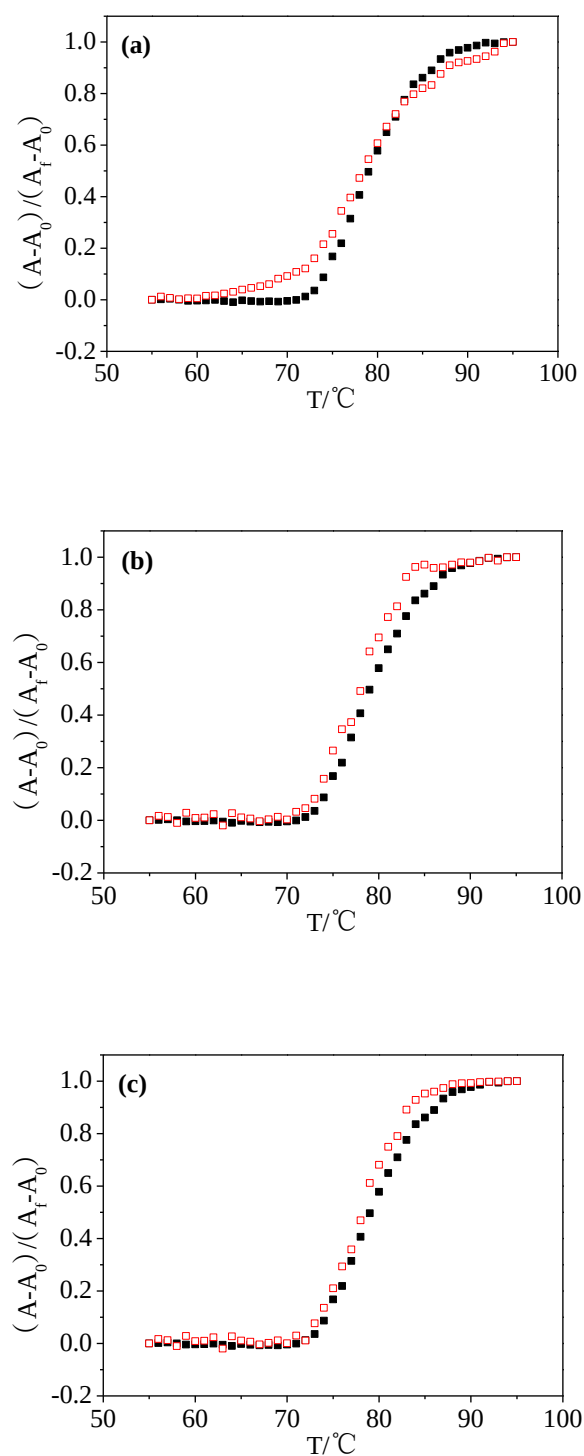


Figure S1. The thermal melt curves of calf thymus DNA (100 μM, ■) incubated with different concentration of complexes **CuPDTP** (a), **CuADTP** (b), **CuBFDTP** (c). The ratio of [Cu]/[DNA] is 0.2 (□). The data was determined by measuring the absorbance of DNA at 260 nm.

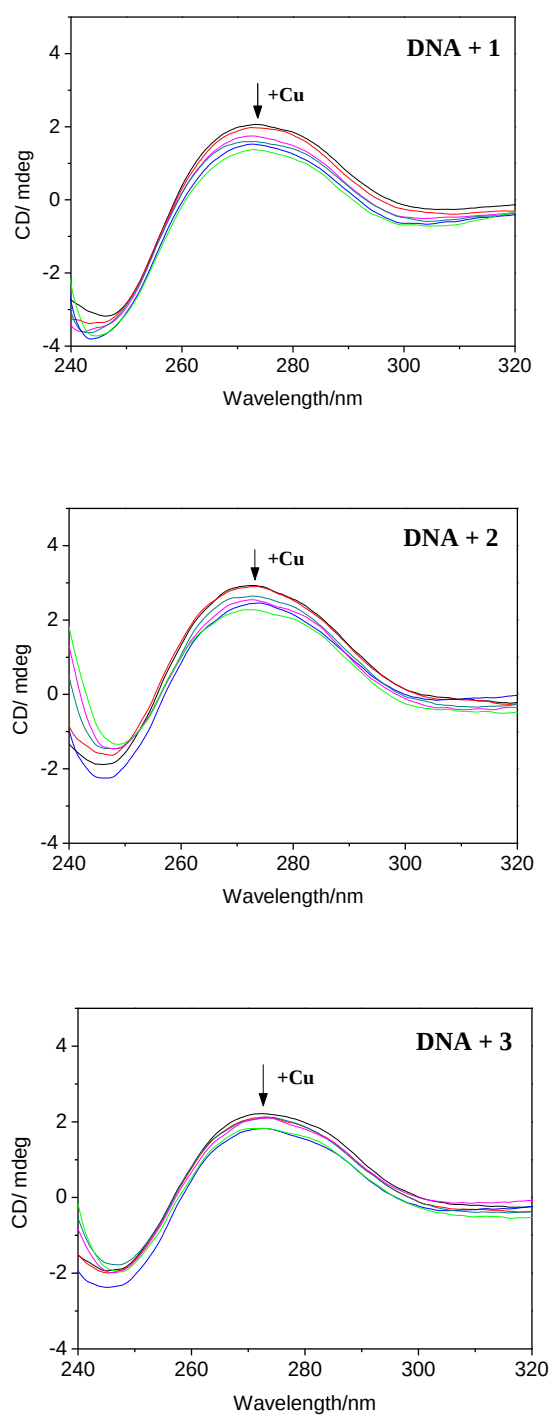


Figure S2. CD spectra of pBR322 DNA (50 μ M bp) in the presence of copper complexes **CuPDTP (1)**, **CuADTP (2)** and **CuBFDTP (3)**. $[\text{Cu}]/[\text{DNA}] = 0, 0.1, 0.2, 0.3, 0.4, 0.5$.

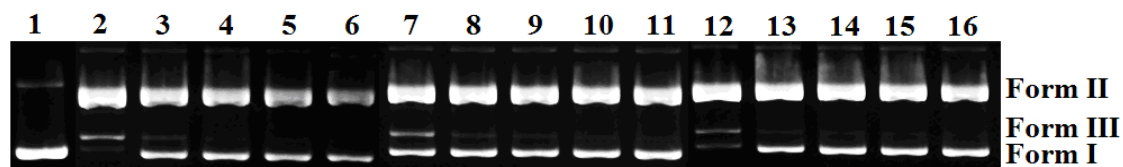


Figure S3. Agarose gel of plasmid DNA cleavage by Cu(II) complexes at different salt concentrations. Lane 1: DNA control (40 μ M), Lane 2-6: **CuPDTP** (100 μ M) + NaCl (18, 50, 100, 150, 200 mM), Lane 7-11: **CuADTP** (100 μ M) + NaCl (18, 50, 100, 150, 200 mM), Lane 12-16: **CuBFDTP** (100 μ M) + NaCl (18, 50, 100, 150, 200 mM).

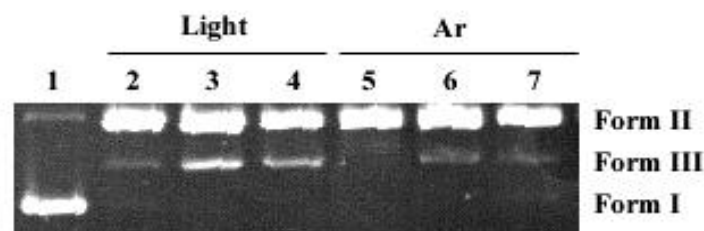


Fig. S4. Gel electrophoresis of pBR 322 DNA cleavage catalyzed by **CuPDTP-CuBFDTP** in phosphate buffer at 37 °C for 1.5 h. (a) Lane 1: DNA control, Lane 2: DNA + **CuPDTP** (100 μ M), Lane 3: DNA + **CuADTP** (100 μ M), Lane 4: DNA + **CuBFDTP** (100 μ M), Lane 5: DNA + **CuPDTP** (100 μ M), Lane 6: DNA + **CuADTP** (100 μ M), Lane 7: DNA + **CuBFDTP** (100 μ M).

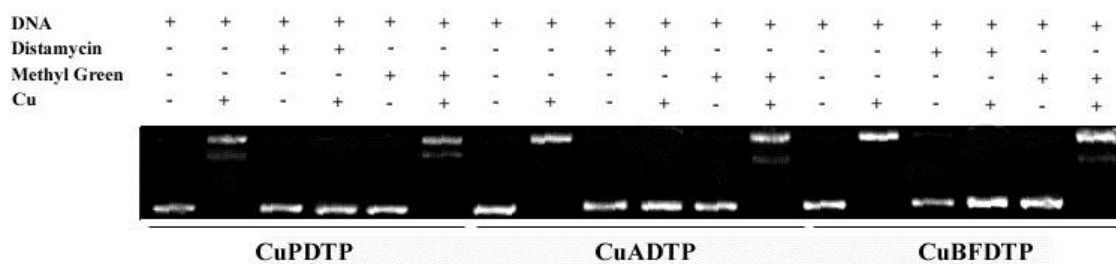


Figure S5. The DNA groove binding preference of the Cu(II) complexes. [DNA] = 40 μ M, [Cu] = 100 μ M, [Distamycin] = 200 μ M, [Methyl Green] = 200 μ M.

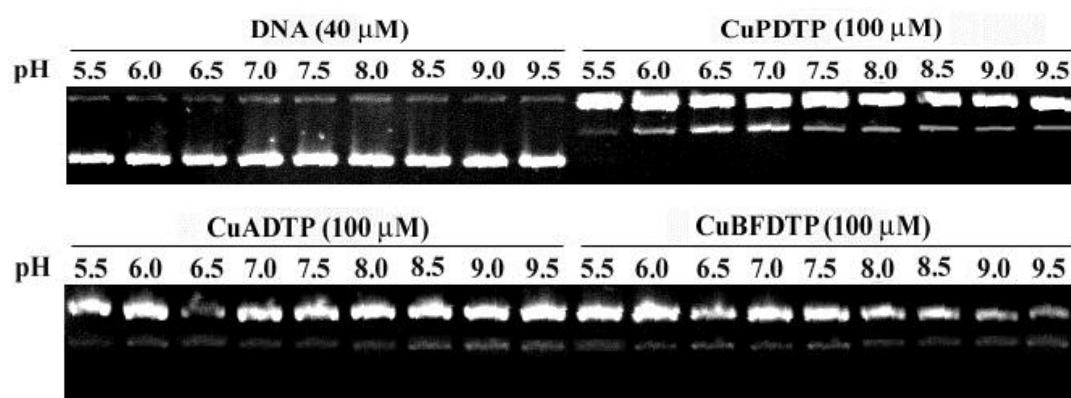


Figure S6. The pH effect on the DNA cleavage activity of Cu(II) complexes.

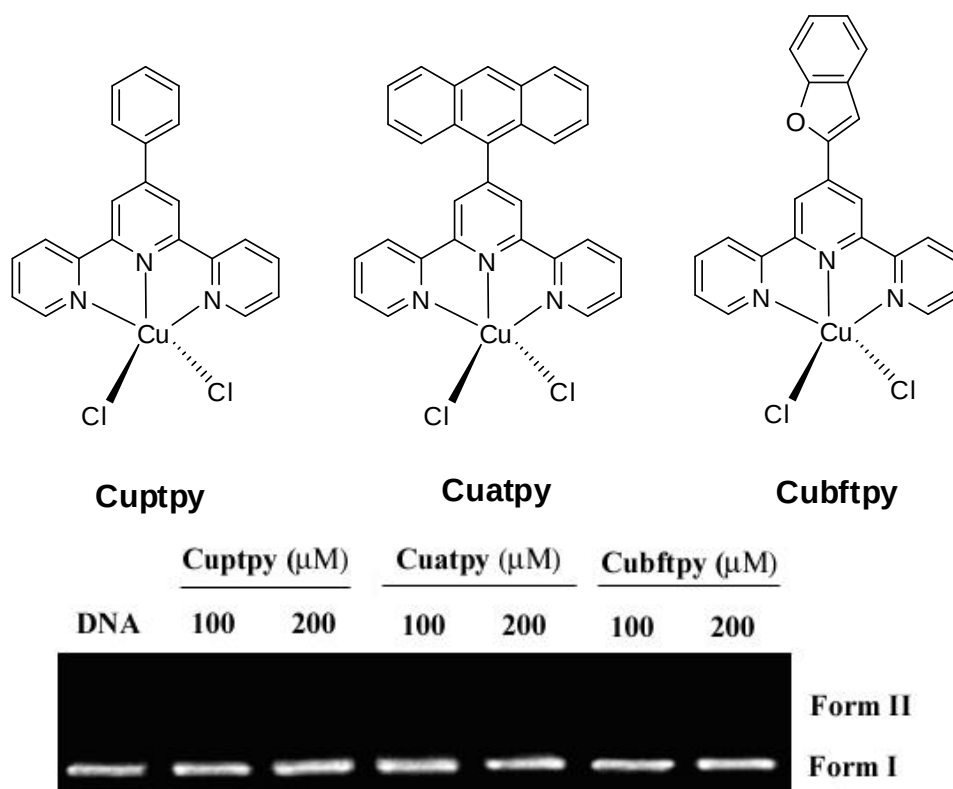


Fig. S7. Gel electrophoresis of pBR 322 DNA cleavage catalyzed by Cupty-Cubftpy in phosphate buffer at 37 °C for 1.5 h. DNA control (40 μM).

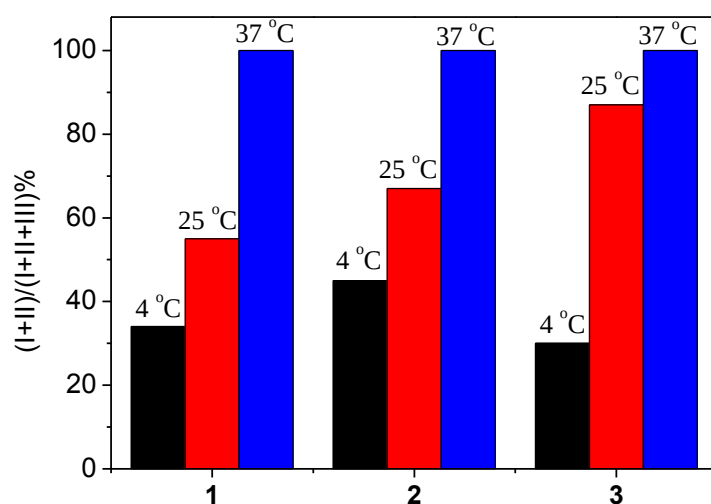


Figure S8. Plasmid DNA cleavage by **CuPDTP** (1), **CuADTP** (2) and **CuBFDTP** (3) (200 μ M) at different temperature incubated 20 min in phosphate buffer. The nuclease activity showing a temperature-dependent manner.

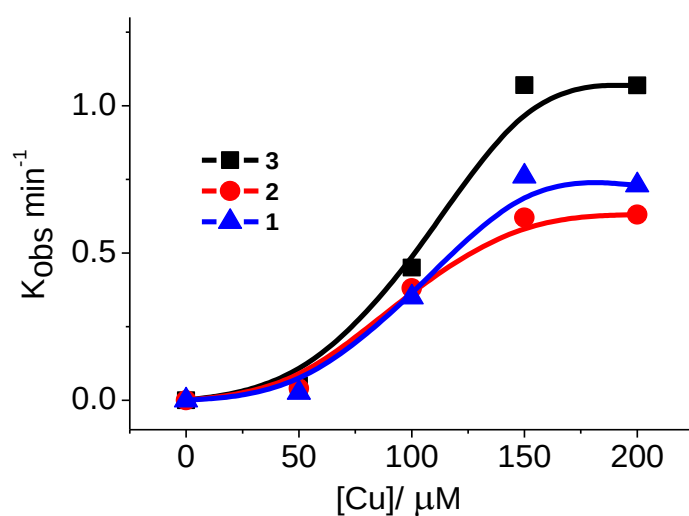


Figure S9. Kinetics for the cleavage of plasmid pBR322 DNA by **CuPDTP** ($\blacktriangle$), **CuADTP** ($\bullet$) and **CuBFDTP** ($\blacksquare$) in a 10 mM phosphate buffer (pH 7.2) at 37 °C.

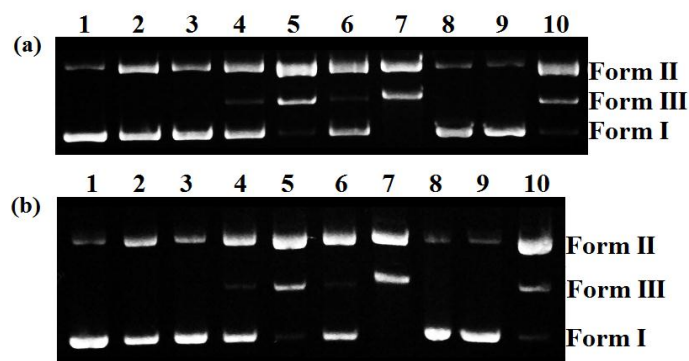


Figure S10. Agarose gel electrophoresis of pBR 322 DNA (40 μ M in bp) with complex **CuPDTP** (a) and **CuADTP** (b) (100 μ M) in the presence of various radical scavengers for 1.5 h at 37 °C. Lane 1: DNA control, Lane 2: DNA + complex (100 μ M); Lane 3: +NaN₃ (100 mM); Lane 4: + Histidine (100 mM); Lane 5: + DMSO (100 mM); Lane 6: + Mannitol (50 mM); Lane 7: + SOD (500 U/mL); Lane 8: + Catalase (1000 U/mL); Lane 9: + EDTA (10 mM); Lane 10: + TEMPO (20 mM).

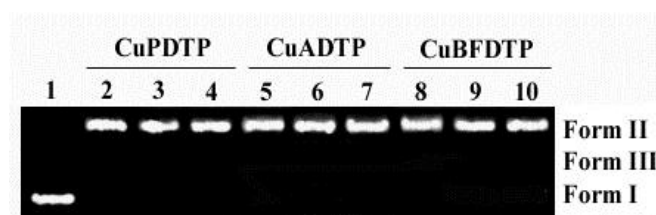


Figure S11. Agarose gel electrophoresis of pBR 322 DNA (40 μ M in bp) with Cu(II) complexes in the presence of various radical scavengers for 1.5 h in the dark at 37 $^{\circ}$ C. Lane 1: DNA control, Lane 2: DNA + **CuPDTP** (100 μ M); Lane 3: DNA + **CuPDTP** (100 μ M) + NaN_3 (100 mM); Lane 4: DNA + **CuPDTP** (100 μ M) + Histidine (100 mM); Lane 5: DNA + **CuADTP** (100 μ M); Lane 6: DNA + **CuADTP** (100 μ M) + NaN_3 (100 mM); Lane 7: DNA + **CuADTP** (100 μ M) + Histidine (100 mM); Lane 8: DNA + **CuBFDTP** (100 μ M); Lane 9: DNA + **CuBFDTP** (100 μ M) + NaN_3 (100 mM); Lane 10: DNA + **CuBFDTP** (100 μ M) + Histidine (100 mM).

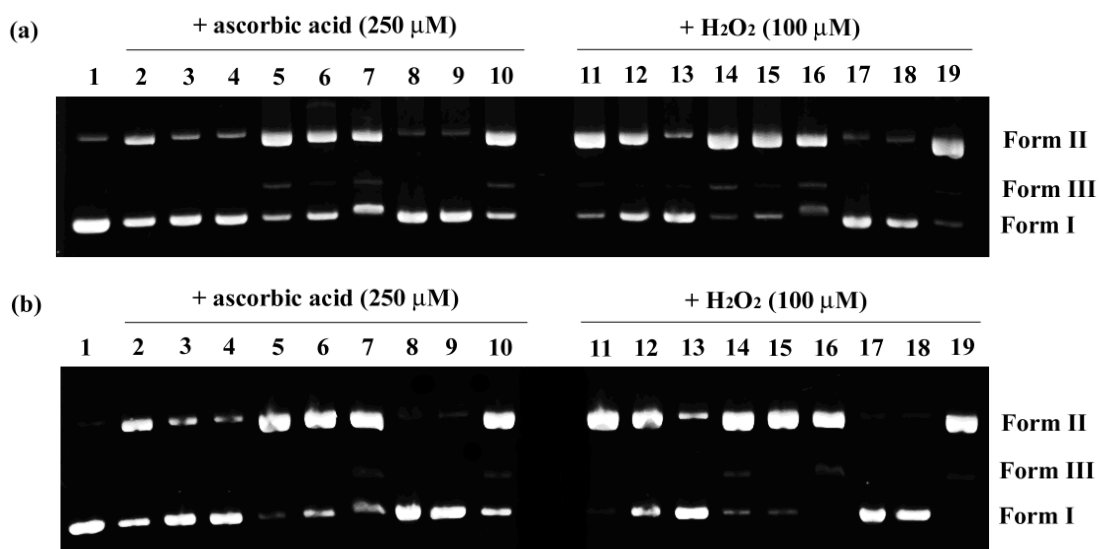


Figure S12. Agarose gel electrophoresis of **CuPDTP** (a) and **CuADTP** (b) (10 μ M) oxidative cleaving DNA in the presence of radical inhibitors under redox condition. Lane 1: DNA control, Lane 2 and 11: DNA + Cu (100 μ M) ; Lane 3 and 12: Cu + NaN_3 (100 mM); Lane 4 and 13: Cu + Histidine (100 mM); Lane 5 and 14: Cu + DMSO (100 mM); Lane 6 and 15: Cu + Mannitol (50 mM); Lane 7 and 16: Cu + SOD (500 U/mL); Lane 8 and 17: Cu + Catalase (1000 U/mL); Lane 9 and 18: Cu + EDTA (10 mM); Lane 10 and 19: Cu + TEMPO (20 mM).

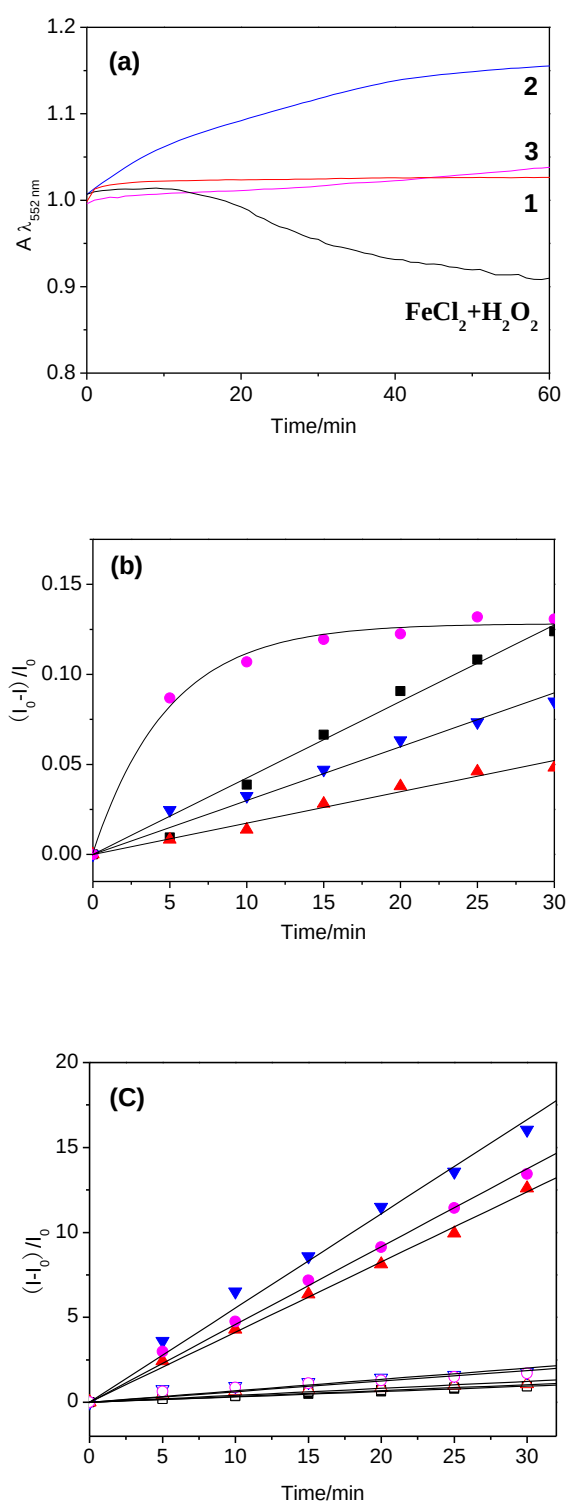


Figure S13. (a) UV absorbance of Rhodamine B (10 μM) in the presence of Cu(II) complexes (100 μM) in phosphate buffer under aerobic condition. FeCl_2 (100 μM) and H_2O_2 mixture (Fenton conditions) as a control experiment. (b) The DPBF (10 μM) consumption percentage as a function of time in the CH_3OH solution of $[\text{Ru}(\text{bpy})_3]^{2+}$ (■, ▲, ▼, ◆).

30 μM), **CuPDTP** ($\blacktriangle$, 30 μM), **CuADTP** ($\blacktriangledown$, 30 μM), **CuBFDTP** ($\bullet$, 30 μM). (c) Dye degradation of DCFH (0.2 μM) incubated with $[\text{Ru}(\text{bpy})_3]^{2+}$ ($\blacksquare$, 30 μM), **CuPDTP** ($\blacktriangle$, 30 μM), **CuADTP** ($\blacktriangledown$, 30 μM), **CuBFDTP** ($\bullet$, 30 μM), and with $[\text{Ru}(\text{bpy})_3]^{2+}$ ($\square$, 30 μM), **CuPDTP** (Δ , 30 μM), **CuBDTP** (∇ , 30 μM), **CuBFDTP** ($\circ$, 30 μM) in the presence of Catalase (1000 U/mL).