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Supplementary information

Mixed-ligand copper(II) complexes of guanidine derivatives containing ciprofloxacin: Synthesis, characterization, DFT calculations, DNA interactions and biological activities

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| Table S1 Band assignment of vibrational modes for ciprofloxacin, the starting compounds (1 and 2) and the mixed-ligand | complexes (1Cip | and |
|--|-----------------|-----|
| | | |

2Cip).

| Ciprofloxacin | | Frequency (cm ⁻¹) | | | | |
|---------------|-------------|-------------------------------|--------------|--------------|---|--|
| | 1 | 2 | 1Cip | 2Cip | | |
| 3363br | 3370s-3180s | 3375br-3142w | 3358br | 3238br | v(OH), v(NH), v(NH ₂) | |
| 3044m-2845m | | 2961w, 2856w | 2951w, 2838m | 2947w, 2836w | v(CH), v(=C-H) | |
| | 1689s | 1650s | | | ν(C=C), ν(C=N), δ(NH ₂) | |
| | 1654s | 1552s | | | ν(C=C), ν(C=N), δ(NH ₂) | |
| 1615s | | | 1613s | 1613s | $v_{as}(C=O)_{py}$ | |
| 1586s | | | | | $v_{as}(C=O)_{carb}$ | |
| | | | 1577s | 1579s | $v_{as}(CO_2)_{carb}$ | |
| 1547s | 1547s | 1525s | 1513w | 1523s | v(C=C) _{benzene ring} , v(C=N), | |
| | | | | | δ(NH ₂) | |
| | 1507m | 1494m | | 1479m | ν(C=N), ν(C-NH ₂), δ(NH ₂) | |
| 1474m-1451m | 1473m-1452m | | 1476s-1457s | 1459m | ν(C-N), δ(C-H) | |
| | | | 1376s | 1374m | $v_s(CO_2)_{carb}$ | |
| 1371s | | | | | v(C-H), v(C-O) _{carb} | |
| 1307m | | 1355m-1323m | 1302m | 1352w, 1335w | v(CH) _{ring} | |
| | | | 1254s | 1259s | $\nu(CO_2)_{carb}$ | |
| | 1196s-1117m | 1210m-1056m | | 1192m-1095w | v(ring), p(NH ₂), v(C-O) | |
| 1174m-1075w | 1076w | 1018w | 1196w-1112w | 1050m | ν (C-H) _{ring} | |
| 1035s-1023s | | | 1043w-1027m | 1019w | v(C-F) | |
| 978w | | | 946s | 946m | δ(NH) | |
| 940m-706w | 948 | 929w | 896w-701m | 898w | γ (=C-H), δ (NH ₂) | |
| 652w-621s | 645m | 610m | 655m-629m | | $\delta(ring)_{aromatic}$, NH ₂ wagging | |

Abbreviation: s, strong; m, medium; w, weak; br, broad; v_{as} , asymmetric stretching; v_s , symmetric stretching; δ , bending.

Table S2. Selected bond lengths [Å] and angles [°] for the optimized geometries of 1Cip and 2Cip complexes at B3LYP/def2-TZVP level of theory.

| | 1Cip | | | 2Cip | | |
|----------|-------|-------|-------|-------|-------|-------|
| | 1A | 1B | 1C | 2A | 2B | 2C |
| Cu–O1 | 1.970 | 1.980 | 2.294 | 1.971 | 1.979 | 2.291 |
| Cu–O2 | 1.918 | 1.956 | 1.924 | 1.919 | 1.961 | 1.923 |
| Cu–N1 | 1.954 | 2.004 | 2.076 | 1.954 | 1.998 | 2.082 |
| Cu–N2 | 1.947 | _ | 2.021 | 1.947 | _ | 2.013 |
| Cu–Cl | _ | 2.250 | 2.295 | _ | 2.251 | 2.296 |
| | | | | | | |
| O1–Cu–O2 | 93.42 | 89.78 | 87.43 | 93.31 | 89.79 | 86.92 |
| N1–Cu–N2 | 88.49 | - | 84.15 | 88.68 | - | 84.35 |

| | | Cell cycle distribution (means ± SD)% | | | | | |
|-------|-----------------|---------------------------------------|------------|------------|------------|--|--|
| Cell | Treatment | sub-G1 | G0/G1 | S | G2/M | | |
| | Solvent control | 2.8±0.72 | 60.8±1.23 | 13.2±0.98 | 20.2±1.22 | | |
| | Cisplatin | 24.9±0.88* | 42.2±1.20 | 16.9±0.54* | 13.4±0.28 | | |
| | 1 | 5.1±1.32* | 32.3±1.23 | 19.7±0.87 | 40.9±1.11* | | |
| HeLa | 2 | 3.3±0.89 | 37.0±0.97 | 19.6±0.23 | 39.6±1.41* | | |
| | 1Cip | 10.5±0.65* | 50.9±1.04 | 18.3±0.57* | 16.9±0.99 | | |
| | 2Cip | 10.9±0.99* | 52.2±0.95 | 19.9±0.89* | 17.0±0.36 | | |
| MCF-7 | Solvent control | 6.0±0.86 | 52.7±0.55 | 16.5±0.99 | 20.9±1.14 | | |
| | Cisplatin | 42.4±0.78* | 30.9±0.87 | 18.2±1.17* | 7.5±1.18 | | |
| | 1 | 11.1±0.71* | 50.5±1.12 | 12.9±0.55 | 24.3±1.30* | | |
| | 2 | 13.0±0.54* | 50.7±1.25 | 12.2±0.30 | 22.7±1.15* | | |
| | 1Cip | 16.4±0.32* | 58.0±0.82* | 11.0±0.91 | 14.8±0.98 | | |
| | 2Cip | 15.8±0.98* | 53.9±0.62* | 13.7±0.20 | 15.6±0.99 | | |

Table S3 The cell cycle distribution of HeLa and MCF-7 cells treated with cisplatin (25 μ g mL⁻¹), **1**, **2**, **1Cip** and **2Cip** (50 μ g mL⁻¹ with Trisbuffer containing 0.3% DMSO) on each phase of cell cycle. All data are the mean and standard errors obtained from three independent experiments.

*P < 0.01 significant difference between samples and solvent control.



Fig. S1 Absorption spectra of (a) 1Cip and (b) 2Cip (50 µM) in Tris-buffer containing 0.3% DMSO at room temperature. The absorbance values

at 273 nm for **1Cip** and 274 nm for **2Cip** were used for the calculation of k (slope of the linear fit) and $t_{1/2}(t_{1/2} = \ln(2)/k)$ values.



Fig. S2 Distribution of (a) HeLa and (b) MCF-7 cells treated by **1**, **2**, **1Cip** and **2Cip** (50 μ g mL⁻¹). Cisplatin was used as a positive control and the solvent control was Tris-buffer containing 0.3 % DMSO. The data given are means \pm SD of three independent experiments.



Fig. S3 Antibacterial activity of the complexes in a concentrations of 1.95, 3.9, 7.8, 15.6, 31.2, 62.5, 125 and 250 μ g mL⁻¹ against (a) *E. coli*, (b)

Salmonella, (c) Campylobacter by disc diffusion method.