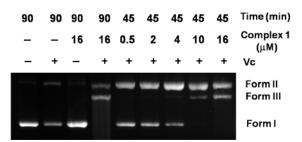
## DNA cleavage behavior of a new *p*-xylyl spaced bisCu(BPA)Cl<sub>2</sub> complex: the steric effect of bulky *p*-xylyl-derived spacer

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**Figure S1.** Agarose gel electrophoresis patterns of plasmid pUC19 DNA (0.02 mgmL<sup>-1</sup>, 30  $\mu$ M base pair) obtained after at different incubation time when incubated with complex 1 of different concentration in Tris-HCl buffer (50 mM, pH 7.4) at 37 °C in the presence of 100-fold excess of ascorbic acid.

## Structural optimization of complex 1

The structure of complex 1 was optimized by density functional theory (DFT) calculations with 6-31G\* basis sets using a suite of Gaussian 03 programs.<sup>i</sup> Different initial structures of complex 1 were constructed based on the crystal structural data of complex  $Cu_2(pTPXA)Cl_4$ . Two stable conformations, *anti*-form and *cis*-form, were obtained (Figure S2). The Cu-Cu distance in *cis*-form (11.84 Å) is remarkably larger than the Cu-Cu distance in  $Cu_2(pTPXA)Cl_4$  (7.98 Å),<sup>ii</sup> whose dicopper centers display the internuclear synergic effect in activating O<sub>2</sub> to form ROS, and the two copper centers in *cis*-form are disable to react with the same O<sub>2</sub> molecule in a synergic manner. As to the *anti*-form, the two copper centers located above and below the benzoxadiazole plane respectively, can't react with the same O<sub>2</sub> molecules either, although the Cu-Cu distance is only 8.53 Å. Therefore, the two Cu(BPA)Cl<sub>2</sub> motifs in complex 1 may function solely as a mononuclear copper complex.

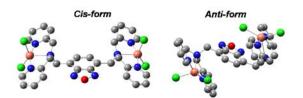


Figure S2. Cis- and anti-form of complex 1 obtained by theoretical study.

<sup>&</sup>lt;sup>i</sup> M. J. Frisch, G. W. Trucks, H. B. Schlegel, et al., GAUSSIAN 03 (Revision D.02), Gaussian Inc., Pittsburg, PA, 2006.

<sup>&</sup>lt;sup>II</sup> Y. Zhao, J. Zhu, W. He, Z. Yang, Y. Zhu, Y. Li, J. Zhang, Z. Guo, Chem. Eur. J. 2006, 12, 6621-6629.