## Supplementary Information

## Comparison of mononuclear and dinuclear copper(II) biomimetic complexes: Spectroelectrochemical mechanistic study of their catalytic pathways

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**Scheme S1**. Mechanism of tyrosinase catalytic activity toward oxidation of phenol and catechol according to Hamann J.N. et al. [5].



**Fig. S1.** Panel A represents UV-Vis spectra recorded at concentrations 10, 50, 100, 150, 200, 300, 400, 500, 750, 1000 and 1500  $\mu$ mol L<sup>-1</sup> of 3,5-DTBQ obtained in pure MeOH containing 0.1 mol L<sup>-1</sup> LiClO<sub>4</sub>. Panel B shows the corresponding calibration curve for 3,5-DTBQ spectrophotometric detection at 400 nm.



**Fig. S2**. UV-Vis spectroelectrochemistry of compounds **C1** (A), **C2** (B) and **C3** (C) in 0.1M TBAPF<sub>6</sub> and acetonitrile during their irreversible oxidation at 1.7 V.



**Fig. S3**. UV-Vis spectroelectrochemistry of 3,5-DTBC during oxidation at potentials -0.45, -0.30, -0.25 and -0.2 V (A) and during continuous oxidation at potentials -0.20, -0.10, 0.05, 0.15, 0.40 V (B) in  $0.1M \text{ TBAPF}_6$  and acetonitrile.



**Table S1**. Minimal energies calculated for different possible intermediates of catalysts **C1**, **C2** and **C3** proposed according to literature (Spartan 20', Wavefunction, Inc.).

C3			
E <sub>min</sub> / kJ/mol	780.6402	732.8906	
C3			
E <sub>min</sub> / kJ/mol	184.7303	602.6719	
C3			
E <sub>min</sub> / kJ/mol	472.3717	399.0295	



**Fig. S4**. FTIR spectrum of **C1** complex obtained at single-bounce diamond ATR crystal using Nicolet<sup>™</sup> iS50 FTIR Spectrometer from Thermo Fisher Scientific<sup>™</sup> (Waltham, Massachusetts, USA).



**Fig. S5**. FTIR spectrum of **C2** complex obtained at single-bounce diamond ATR crystal using Nicolet<sup>™</sup> iS50 FTIR Spectrometer from Thermo Fisher Scientific<sup>™</sup> (Waltham, Massachusetts, USA).



**Fig. S6**. FTIR spectrum of **C3** complex obtained at single-bounce diamond ATR crystal using Nicolet<sup>™</sup> iS50 FTIR Spectrometer from Thermo Fisher Scientific<sup>™</sup> (Waltham, Massachusetts, USA).



**Fig. S7.** Full scan positive-ion ESI mass spectra of A/ complex **C1**, B/ complex **C2**, and C/ complex **C3** obtained using hybrid quadrupole time of flight mass analyzer (micrOTOF-Q, Bruker Daltonics, Germany) in positive-ion mode in the range of m/z 100 – 1000. Individual m/z values correspond to the most abundant ion peak within the characteristic isotopic distribution of annotated ions.