Copper depletion inhibits $CoCl_2$-induced aggressive phenotype of MCF-7 cells via downregulation of HIF-1 and inhibition of Snail/Twist-mediated epithelial-mesenchymal transition

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Table S1. Primer sequences for real time-PCR analysis

Gene	Primer sequence (5'→3')
β-actin	Forward: GGCATCCTCACCCTGAAGTA
	Reverse: GGGGTGTTGAAGGTCTCAAA
vimentin	Forward: CTCCGGGAGAAATTGCAGGA
	Reverse: TTCAAGGTCAAGACGTGCCA
VEGFa	Forward: GGCTTTTCTGTGGCAAGAGGTT
	Reverse: CGCTGATGTTTGCTCAAGTGGTC
fibronectin	Forward: GACCTATCCAAGCTCAAGTGGT
	Reverse: ATCCAAGGTTTCTGGGTGGG
Snail	Forward: TGCCCTCAAGATGCACATCCGA
	Reverse: GGGACAGGAGAAGGGCTTCTC
Twist1	Forward: GCCAGGTACATCGACTTCCTCT
	Reverse: TCCATCCTCCAGACCGAGAAGG

Sequences used for *Ctr1* silence

Primer sequence (5'→3') : GGAAGAAGGCAGUGGUAGUdTdT

Quantitative analysis of immunofluorescence data with CoCl2 or TEPA treatment.

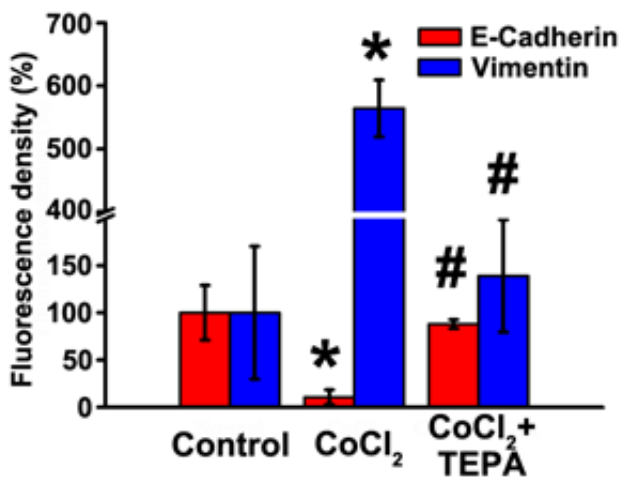

Figure S1. The fluorescence density of E-cadherin and vimentin were measured in each groups. Data was shown as mean ± SD of three independent experiments. $p<0.05$ was considered statistically significant (*compared to the control group and #compared to the $CoCl_2$ group).

Gene silence efficacy validation of *Ctr1* by western blot.

Figure S2. Gene silence efficacy of *Ctr1*. *Ctr1* expression was assayed by western blot after transfection and continued to culture for 24 h.

Morphology of MCF-7 cells under TEPA treatment only.

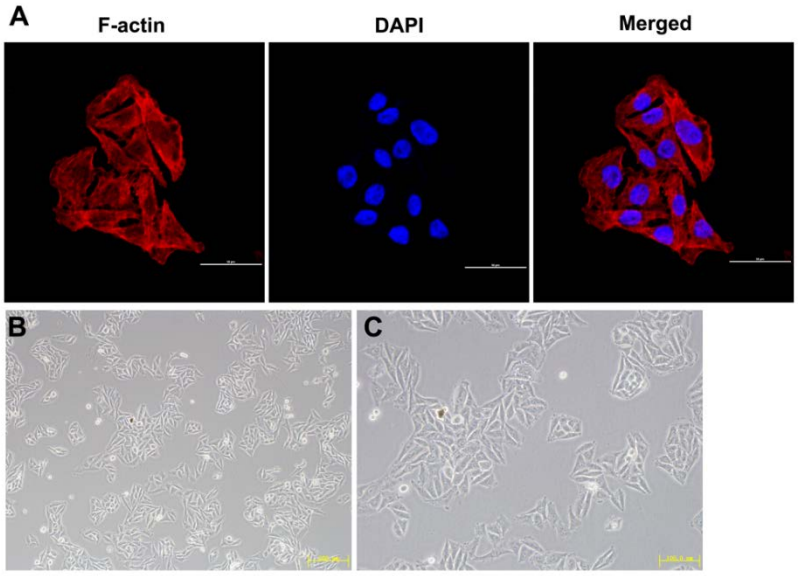

Figure S3. Morphology of MCF-7 cells under TEPA treatment only. (**A**) Spindle-shaped cells were not observed after TEPA treatment for 24 h. F-actin was stained by TRITC-phalloidin (red). Scale bar=50 μm. (**B**) and (**C**) The morphology of MCF-7 cells were visualized by phase-contrast.