



Corrigendum: CORO1C is Associated with Poor Prognosis and Promotes Metastasis Through PI3K/AKT Pathway in Colorectal Cancer

Zongxia Wang¹, Lizhou Jia^{1,2}, Yushu sun³, Chunli Li¹, Lingli Zhang⁴, Xiangcheng Wang^{5,6}* and Hao Chen^{2,7}*

¹Cancer Center, Bayannur Hospital, Bayannur, China, ²Department of Pathology, Wannan Medical College, Wuhu, China, ³Department of Oncology, Inner Mongolia Autonomous Region Cancer Hospital, Hohhot, China, ⁴Department of Ophthalmology, Inner Mongolia Autonomous Region People's Hospital, Hohhot, China, ⁵Department of Nuclear Medicine, The Affiliated Hospital of Inner Mongolia Medical University, Hohhot, China, ⁶Key Laboratory of Inner Mongolia Autonomous Region Molecular Imaging, Inner Mongolia Medical University, Hohhot, China, ⁷Faculty of Medical Science, Jinan University, Guangzhou, China

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*Correspondence:

Xiangcheng Wang guyan@nmgfy.com Hao Chen ha0chen@wnmc.edu.cn

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In the original article, there was a mistake in the caption for Figure 4A as published. The incorrect caption stated NCM460 cell lines are normal gastric epithelial cells, however, NCM460 cell lines are normal colorectal epithelial cells. The correct caption for Figure 4 appears below. Additionally, there was a mistake in Figure 1 as published. Figure 1C is immunofluorescence and Figure 1D is immunocomplex by Co-IP. OE-ControlTrop2 should appear on the top row and OE-Trop2 should appear on the bottom row in Figure 1C. Representative images for these parts contain small issues which have been corrected. The corrected Figure 1 appears below.

FIGURE 4. The effects of CORO1C knockdown on CRC growth and metastasis *in vitro* and *in vivo*. (**A**) Levels of CORO1C1 protein expression in CRC cell lines and normal colorectal epithelial cells (NCM460) determined by western blotting. (**B**) COCA2 and HCT116 cells showed a significant decrease in protein level after shCORO1C transfection. (**C**) CORO1C downregulation significantly inhibited the proliferation of both cell lines. (**D**) A significant decrease in cell anchorage-dependent growth was detected after CORO1C knockdown. (**E**, **F**) Decreased CORO1C expression impaired abilities of migration (**E**) and invasion (**F**) of CRC cells (scale bar, 150 µm). All quantitative data of *in vitro* assays were generated from three replicates (**G**). The effects of CORO1C downregulation on the tumor growth in the xenograft mouse model (*n* = 6 mice/per group). **p* < 0.05, ***p* < 0.01, ****p* < 0.001.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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blotting, and (C) Immunofluorescence. (D) Trop2 immunocomplex by Co-IP; (E) Heatmap of proteins interacting with Trop2 in PC and MC. The RNA-seq data were obtained from GSE28702. Red and green colors represent high and low gene expression, respectively. PC: primary CRC; MC: metastatic CRC, **p < 0.01.

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