



Correction to: Evaluation of Forkhead BOX M1 (FOX M1) gene expression in colorectal cancer

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The original version of this article unfortunately contained a mistake. The corrected details are given below for your reading.

In tables 3, 4, 5 and 7, the bold “**0**” denotes significance “**0**”, but “0” alone does not make any sense. Therefore, it needs to be replaced with < 0.001 .

In Results, under heading of “Survival analysis of FOXM1, and various clinicopathological parameters in CRC” Fig. 6E–H should be written as Fig. 6(A–D) and Fig. 7I, J should be written as Fig. 7(A–B) as depicted in figure legends.

After Discussion, under “Conclusion” section the lines” High levels of FOXM1 protein were significantly associated with the most severe pathological characteristics of CRC, demonstrating that FOXM1 “expression” is linked to CRC progression. The word “expression” should be “Overexpression”.

The original article has been corrected.

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