Correction to “Genome-wide CRISPR-Cas9 screening identifies that hypoxia-inducible factor-1a-induced CBX8 transcription promotes pancreatic cancer progression via IRS1/AKT axis”

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Abstract
Correction to “Genome-wide CRISPR-Cas9 screening identifies that hypoxia-inducible factor-1a-induced CBX8 transcription promotes pancreatic cancer progression via IRS1/AKT axis” (PMID: 34853645 PMCID: PMC8603463 DOI: 10.4251/wjgo.v13.i11.1709). In this article, the picture of Figure 6C was misused due to our carelessness while typesetting. We corrected this mistake, and replaced the incorrect image with the correct one.

Key Words: Correction; Error; Figure; CRISPR-Cas9

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TO THE EDITOR

After confirming the figures in our manuscript, we were surprised to find a mistake in Figure 6C [1]. It was an unintentional error that occurred when we typeset the images. We have replaced the incorrect images with the correct Figure 6C (Figure 1). Figure 6D-E was based on the correct image and does not need to be changed. We assure you that this mistake does not change the meaning of the picture or the conclusion of the manuscript. We apologize for our careless mistake, which has caused great inconvenience.

Figure 1 We replaced the incorrect images with the correct Figure 6C in “Genome-wide CRISPR-Cas9 screening identifies that hypoxia-inducible factor-1a-induced CBX8 transcription promotes pancreatic cancer progression via IRS1/AKT axis.”

FOOTNOTES

Author contributions: Teng BW found and corrected the misused figure; Zhang KD, Yang YH, Guo ZY, Chen WW, and Qiu ZJ checked the correction.

Conflict-of-interest statement: There are no conflicts of interest to report.

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