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Corrigendum

Corrigendum to "Long Noncoding RNA TUG1/miR-29c Axis Affects Cell Proliferation, Invasion, and Migration in Human Pancreatic Cancer"

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In the article titled "Long Noncoding RNA TUG1/miR-29c Axis Affects Cell Proliferation, Invasion, and Migration in Human Pancreatic Cancer" [1], BLASTn results [2] identified an error in the forward and reverse sequences of the TUG1 shRNA shown in Section 2.2 of the Materials and Methods Section. The reverse sequence of the TUG1 shRNA reported in the original article was identified as targeting the PMM2 gene sequence; however, the authors have confirmed that it was a mistake, and the correct sequences are as follows:

Correct forward sequence should read as follows:

5′-gatccGCTTGGCTTCTATTCTGAATCCTTTCAAG AGAAGGATTCAGAATAGAAGCCAAGCTTTTTTG-3′.

Correct reverse sequence should read as follows:

5′-CAAAAAAGCTTGGCTTCTATTCTGAATCCTT CTCTTGAAAGGATTCAGAATAGAAGCCAAGCG-3′.

References

[1] Y. Lu, L. Tang, Z. Zhang et al., "Long Noncoding RNA TUG1/miR-29c Axis Affects Cell Proliferation, Invasion, and Migration in Human Pancreatic Cancer," *Disease Markers*, vol. 2018, Article ID 6857042, 10 pages, 2018.

[2] C. Labbé, N. Grima, T. Gautier, B. Favier, and J. A. Byrne, "Semi-automated Fact-checking of Nucleotide Sequence Reagents in Biomedical Research Publications: The Seek & Blastn tool," *PLoS One*, vol. 14, no. 3, Article ID e0213266, 2019.