CORRECTION

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Correction to: LncRNA-NEAT1 from the competing endogenous RNA network promotes cardioprotective efficacy of mesenchymal stem cell-derived exosomes induced by macrophage migration inhibitory factor via the miR-142-3p/FOXO1 signaling pathway



Correction to: Stem Cell Res Ther 11, 31 (2020) https://doi.org/10.1186/s13287-020-1556-7

Following publication of the original article [1], the authors identified an error in Table 1. The primer of human LncRNA-NEAT1 should be revised to F: 5' -GTGGCTGTTGGAGTCGGTAT - 3' and R: 5' -ACCACGGTCCATGAAGCATT - 3'.

The correct table has been included in this correction.

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the miR-142-3p/FOXO1 signaling pathway. Stem Cell Res Ther. 2020;11:31. https://doi.org/10.1186/s13287-020-1556-7.



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Table 1 Primer sequences

Genes	Sequences
LncRNA-NEAT1	F: 5' - GTGGCTGTTGGAGTCGGTAT – 3'
	R: 5' - ACCACGGTCCATGAAGCATT – 3'
U6	F: 5'- GCTTCGGCAGCACATATACTAAAAT – 3'
	R: 5'- CGCTTCACGAATTTGCGTGTCAT – 3'
miR-142-3p	F: 5' - TGTAGTGTTTCCTACTTTAT – 3'
	R: 5' - GTCGTATCCAGTGCAGGG – 3'
FOXO1	F: 5' - CAGCAAATCAAGTTATGGAGGA – 3'
	R: 5' - TATCATTGTGGGGAGGAGAGTC – 3'
GAPDH	F: 5'- TTGCCATCAATGACCCCTTCA – 3'
	R: 5'- CGCCCCACTTGATTT TGGA – 3'
siRNA-LncRNA-NEAT1	5'- GCCAUCAGCUUUGAAUAAAUU – 3'
siRNA-LncRNA-NT	5'- UUCUCCGAA CGUGUCACGU – 3'
siRNA-FOXO1	5'- CGGAGAAUGUAUACAAGCATT – 3'
siRNA-NT	5'- GGAGUUAUGAGUCAGUAUATT – 3'