

TRIM28 Down-Regulation on Methylation Imprints in Bovine Preimplantation Embryos – ERRATUM

Erratum

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The publishers apologise for errors in the layout of Table 1. The correct Table is as follows:

Table 1. Sequence of primers and siRNA

Name	Nucleotide sequences(5'-3')	Fragment size(bp)	GenBank Accession Number or Reference
TRIM28	F-CTGGATGGGGGGATAGTA	145	NM_001206809.1
	R-CTGGGGAGAAGGTGGAGTC		
18s rRNA	F-GACTCATTGGCCCTGTAATTGGAATGAGTC		
	R-GCTGCTGGCACCAGACTTG		
H19	F-TTTGGTTTTTGTGGATT	385	NR_003958.2
	R-ATAACTTCAAATTACCTCCTACC		
	F-GGTTTTTGTGGATTTTG	382	
	R-ATAACTTCAAATTACCTCCTACC		
Peg10	F-GTTTGGTATAGGTGTGGGATTT	230	NM_001127210.2
	R-TCAAAACCTAAAACTTAAATTCTC R-ACACCTTACTCAAACCTACC		
Mest	F-GGTATTATTGGGTTTTTATATTGTGA	355	NM_001083368.1(O'Doherty et al. 2012)
	R-CAAACCTCCTAAATCCCTAACTA		
si647	S-GCGUGCAAGUGGAUGUUAATT A-UUAACAUCACUUGCAGCCTT	N/A	
si742	S-GCAGCAAUGUUUCUCAAATT A-UUUGGAGAAACAUUGCUGCTT	N/A	
si1153	S-GCACUAGCUGUGAGGAUAATT	N/A	
	A-UUAUCCUCACAGCUAGUGCTT		
Nonsense	S-UGACCUCAACUACUAGGUUTT	N/A	
	A-AACCAUGUAGUUGAGGUCATT		

Reference

Ma X, Zhang S, Zhang M, Zhu Y, Ma P, Yang S, Su L, Li Z, Lv W and Luan W (2018). TRIM28 down-regulation on methylation imprints in bovine preimplantation embryos. *Zygote*, 26, 449-456. <https://doi.org/10.1017/S0967199418000424>.