

CORRECTION

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Epigenetic reprogramming in the porcine germ line

Sara MW Hyldig¹, Nicola Croxall², David A Contreras², Preben D Thomsen¹ and Ramiro Alberio^{2*}

After the publication of this work [1] we became aware that Table one had inadvertently been omitted from the final version of the article. The table is now included in this document as Figure 1.

We regret any inconvenience that this might have caused.

²Division of Animal Sciences, School of Biosciences, University of Nottingham, Loughborough, LE12 5RD, UK.

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Reference

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Author details

¹Department of Basic Animal and Veterinary Sciences, Faculty of Life Sciences, University of Copenhagen, 1870 Frederiksberg C, Denmark.

Sequence	Primers for Bisulfite Analysis	Annealing	Length	# CpGs
<i>IGF2</i> (CTCF3)	5'-GGTTGTGGGTGTGGAGGTAGA-3' 5'-CACACTAAACACCCAACCTTTAACAC-3'	60 °C	207 bp	11
<i>IGF2R</i> (DMR1)	5'-TAGTGATTTTAAGTTGGAAAGTATT-3' 5'-ATAACTCCCTTATTCAAATAATCCC-3'	57 °C	224 bp	29
<i>IGF2R</i> (DMR2)	5'-GGGTAGAGGAAGAAGGTAAGATGGATAG-3' 5'-CCCAAACTCCTAAAACTCCTCATAAC-3'	64 °C	188 bp	16
<i>SINE</i> (SSPRE)	5'-TTTGGTTTTGTTTAGTGGGTTAAG-3' 5'-TATAAAAATCCCAAATAAAAATC-3'	59 °C	~148 bp	~ 8
	Primers for Gender Determination			
<i>AMELX/Y</i>	5'-CRCMTTCATTGAYAATTCAC-3' 5'-CCAGAGGTTGTAACCTTACAG-3'	60 °C	520 bp ♀/♂ 350 ♀	

Figure 1

Figure 1 Table one from Epigenetic reprogramming in the porcine germ line.

* Correspondence: ramiro.alberio@nottingham.ac.uk

²Division of Animal Sciences, School of Biosciences, University of Nottingham, Loughborough, LE12 5RD, UK

Full list of author information is available at the end of the article