

CORRECTION

Correction: Increasing Live Birth Rate by Preimplantation Genetic Screening of Pooled Polar Bodies Using Array Comparative Genomic Hybridization

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There are errors in [Table 1](#). Please see the correct [Table 1](#) here.

Table 1. Characteristics of the total study population.

	non-CGH n = 240	CGH n = 111	p-value
Age (years)	38.4±2.5	39.5±2.3	<0.001
Antagonist protocol	152 (63.3%)	73 (65.8%)	0.394
Agonist protocol	87 (36.3%)	38 (34.2%)	0.394
BMI (kg/m ²)	20.3±9.2	20.5±7.6	0.354
Attempts	2.0 (1.0–3.0)	2.0 (1.0–4.0)	0.002
Attempts >1	142 (59.2%)	78 (70.2%)	0.046
Number of oocytes	6.0 (4.0–10.0)	8.0 (5.0–11.0)	<0.001
Transferred embryos	2.0 (1.0–2.0)	1.0 (0.0–2.0)	<0.001
Transferred embryos >1	154 (65.0%)	48 (43.2%)	<0.001
Day of embryo transfer	3.0 (2.0–3.0)	3.0 (3.0–5.0)	0.070

The data are expressed as the mean ± standard deviation as well as the median (IQR) and counts (%).

* p-value based on the Wilcoxon rank sum test.

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Reference

1. Feichtinger M, Stopp T, Göbl C, Feichtinger E, Vaccari E, Mädler U, et al. (2015) Increasing Live Birth Rate by Preimplantation Genetic Screening of Pooled Polar Bodies Using Array Comparative Genomic Hybridization. PLoS ONE 10(5): e0128317. doi: [10.1371/journal.pone.0128317](https://doi.org/10.1371/journal.pone.0128317) PMID: [26024488](https://pubmed.ncbi.nlm.nih.gov/26024488/)