CORRECTION

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Correction: iDESC: identifying differential expression in single-cell RNA sequencing data with multiple subjects

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The original article can be found online at https://doi.org/10.1186/s12859-023-05432-8.

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Correction: BMC Bioinformatics (2023) 24:318
https://doi.org/10.1186/s12859-023-05432-8

Following publication of the original article [1], the authors identified an error in Table 2. The correct table is given below.

The incorrect Table 2 is:

Table 2 Fisher's exact test and Jaccard index measuring the DE genes overlapping between the two chosen scRNA-seq datasets in macrophage and fibroblast

Method	iDESC	MAST-RE	Muscat-MM	Muscat-PB	subT
Macrophage					
P value	1×10^{-5}	1.000	0.269	1×10^{-5}	0.085
Jaccard Index	0.137	0.061	0.108	0.055	0.028
Fibroblast					
P value	4×10^{-21}	4×10^{-6}	3×10^{-5}	2×10^{-7}	0.011
Jaccard Index	0.077	0.067	0.071	0.035	0.024

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Reference
1.　Liu Y, et al. iDESC: identifying differential expression in single-cell RNA sequencing data with multiple subjects. BMC Bioinformatics. 2023;24:318. https://doi.org/10.1186/s12859-023-05432-8.

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