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APPROVED BY
Frontiers Editorial Office,
Frontiers Media SA, Switzerland

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RECEIVED 11 April 2024

ACCEPTED 12 April 2024

PUBLISHED 23 April 2024

CITATION

Dai W, Zheng P, Wu J, Chen S, Deng M, Tong X, Liu F, Shang X and Qian K (2024) Corrigendum: Integrated analysis of single-cell RNA-seq and chipset data unravels PANoptosis-related genes in sepsis. *Front. Immunol.* 15:1415915. doi: 10.3389/fimmu.2024.1415915

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Corrigendum: Integrated analysis of single-cell RNA-seq and chipset data unravels PANoptosis-related genes in sepsis

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KEYWORDS

sepsis, Boruta algorithm, PANoptosis, single-cell RNA-seq, ssGSEA

A Corrigendum on

Integrated analysis of single-cell RNA-seq and chipset data unravels PANoptosis-related genes in sepsis

by Dai W, Zheng P, Wu J, Chen S, Deng M, Tong X, Liu F, Shang X and Qian K (2024). *Front. Immunol.* 14:1247131. doi: 10.3389/fimmu.2023.1247131

In the published article, a reference was missing for the section: “We downloaded the sepsis single-cell sequencing sample [GSE167363](https://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE167363) from the GEO database and processed the data using the Seurat package, retaining cells with less than 10% mitochondrial genes, cells with a total number of genes greater than 200, and expression ranging from 200 to 12,000 and Genes expressed in at least 3 cells.”

The reference “Qiu X, Li J, Bonenfant J, Jaroszewski L et al. Dynamic changes in human single-cell transcriptional signatures during fatal sepsis. *J Leukoc Biol* 2021 Dec;110 (6):1253-1268. PMID: 34558746” should be added.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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