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Corrigendum: Analysis of signature genes and association with immune cells infiltration in pediatric septic shock

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KEYWORDS

children, septic shock, weighted gene co-expression network analysis, signature gene, immune cells infiltration

A corrigendum on

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In the published article, there was an error.

A correction has been made to **Method and material**, *Identification of DEGs*. This sentence previously stated:

“Using R software's limma package (19), differentially expressed genes (DEGs) between septic shock cohort and control cohort were analyzed, with the following criterion: $p < 0.05$ and $|\text{fold change (FC)}| > 1$ ”

The corrected sentence appears below:

“Using R software's limma package (19), differentially expressed genes (DEGs) between septic shock cohort and control cohort were analyzed, with the following criterion: $\text{adjust } p \text{ value} < 0.05$ and $|\log \text{ fold change (FC)}| > 1$ ”

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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