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OPEN Author Correction: Identification of hub genes for adult patients with sepsis via RNA sequencing

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Correction to: Scientific Reports https://doi.org/10.1038/s41598-022-09175-z, published online 24 March 2022

The original version of this Article contained errors in Table 1 and its legend. The "clinical variables" rows, such as ALT (U/L), AST (U/L), Crea (µmol/L), PT-INR, PCT (ng/ml), LAC (mmol/L), CCI and their values were omitted from the Table.

Additionally, the value for Neoplasm was incorrect,

now reads:

In the legend of Table 1,

"SOFA score Sequential Organ Failure Assessment score, WBC white blood count, NEU neutrophile, PCT procalcitonin, PLT platelet, NLR neutrophile/lymphocyte ratio, HGB hemoglobin."

now reads:

"SOFA score, Sequential Organ Failure Assessment score; WBC, White blood count; NEU, neutrophile; PCT, procalcitonin; PLT, platelet; NLR, neutrophile/lymphocyte ratio; HGB, hemoglobin; LAC, lactic acid; ALT, alaninetransaminase; AST, aspartate aminotransferase; Crea, creatinine; PT-INR, prothrombin time/international normalization ratio; CCI, Charlson Comorbidity Index."

The original Table 1 and accompanying legend appear below.

The original Article has been corrected.

nature portfolio

| Clinical variable | Healthy control (n=10) | Sepsis (n = 23) | P value |
|--------------------------------|------------------------|-----------------|---------|
| Age (years) | 50.7 ± 2.181 | 56.65 ± 3.621 | 0.3 |
| Gender (F/M) | 4/6 | 7/16 | 0.59 |
| WBC (×10 ⁹ /L) | 6.843 ± 0.628 | 13.8 ± 1.57 | 0.0076 |
| NEU (×109/L) | 4.141 ± 0.4236 | 12.19 ± 1.557 | 0.0021 |
| PLT (×10 ⁹ /L) | 215.3 ± 15.44 | 172.3 ± 20.13 | 0.19 |
| NLR | 2.04 ± 0.1536 | 23.83 ± 4.385 | 0.0028 |
| HGB (g/L) | 143.6 ± 8.755 | 104.1 ± 5.819 | 0.0007 |
| ALT (U/L) | 19.42 ± 1.912 | 86.22 ± 37.36 | 0.2517 |
| AST (U/L) | 20.55 ± 1.024 | 142.2 ± 56.79 | 0.1716 |
| Crea (µmol/L) | 67.15 ± 3.652 | 121.4 ± 26.79 | 0.1958 |
| PT-INR | N/A | 1.806 ± 0.429 | |
| PCT (ng/ml) | N/A | 29.65 ± 7.549 | |
| LAC (mmol/L) | N/A | 3.236 ± 0.69 | |
| SOFA score | N/A | 6.609 ± 0.8637 | |
| Primary sites of infection (n) | | | |
| Surgical | | 10 | |
| Dermatological infection | | 5 | |
| Trauma | | 3 | |
| Respiratory | | 4 | |
| Urinary tract | | 1 | |
| Pathogens | | | |
| Klebsiella pneumoniae | | 3 | |
| Bacillus coli | | 3 | |
| Baumanii | | 1 | |
| Streptococcus pneumoniae | | 3 | |
| Acinetobacter Jones | | 1 | |
| Pseudomyomyces white | | 1 | |
| Comorbidites (n) | | | |
| Neoplasm | | 5 | |
| Diabetes | | 3 | |
| Hypertension | | 3 | |
| 71 | | 3 | |

Table 1. Clinical information of the subjects. *SOFA score* Sequential Organ Failure Assessment score, *WBC* white blood count, *NEU* neutrophile, *PCT* procalcitonin, *PLT* platelet, *NLR* neutrophile/lymphocyte ratio, *HGB* hemoglobin.

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