

The Role of Transcriptomics in Redefining Critical Illness

Critical Care

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Multi-Omics Endotypes in ICU Sepsis-Induced Immunosuppression. <i>Microorganisms</i> , 2023, 11, 1119.	3.6	3
2	Biomarkers to guide immunomodulatory treatment: where do we stand?. <i>Expert Review of Molecular Diagnostics</i> , 2023, 23, 945-958.	3.1	0
3	A machine learning model derived from analysis of time-course gene-expression datasets reveal temporally stable gene markers predictive of sepsis mortality. <i>Shock</i> , 0, , .	2.1	0
4	Machine learning-driven identification of the gene-expression signature associated with a persistent multiple organ dysfunction trajectory in critical illness. <i>EBioMedicine</i> , 2024, 99, 104938.	6.1	0
5	The Omics Complexity in Sepsis: The Limits of the Personalized Medicine Approach. <i>Journal of Personalized Medicine</i> , 2024, 14, 225.	2.5	0