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A distinct innate immune signature marks progression from mild to severe COVID-19

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#	Paper	IF	Citations
86	A genetic link between risk for Alzheimer's disease and severe COVID-19 outcomes via the OAS1 gene. <i>Brain</i> , 2021 ,	11.2	14
85	In depth analysis of patients with severe SARS-CoV-2 in sub-Saharan Africa demonstrates distinct clinical and immunological profiles. 2021 ,		2
84	Integrated plasma proteomic and single-cell immune signaling network signatures demarcate mild, moderate, and severe COVID-19. 2021 ,		3
83	Letter in reply to the letter to the editor of Harte JV and Mykytiv V with the title "A panhaemocytometric approach to COVID-19: a retrospective study on the importance of monocyte and neutrophil population data". <i>Clinical Chemistry and Laboratory Medicine</i> , 2021 , 59, e173-e174	5.9	1
82	Blunted Fas signaling favors RIPK1-driven neutrophil necroptosis in critically ill COVID-19 patients.		1
81	Genetic variability associated with OAS1 expression in myeloid cells increases the risk of Alzheimer's disease and severe COVID-19 outcomes.		
80	Systemic Tissue and Cellular Disruption from SARS-CoV-2 Infection revealed in COVID-19 Autopsies and Spatial Omics Tissue Maps. 2021 ,		3
79	Endothelial cell, myeloid, and adaptive immune responses in SARS-CoV-2 infection. <i>FASEB Journal</i> , 2021 , 35, e21577	0.9	6
78	Comparative immune profiling of acute respiratory distress syndrome patients with or without SARS-CoV-2 infection. <i>Cell Reports Medicine</i> , 2021 , 2, 100291	18	4
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71	Neutrophils and COVID-19: Active Participants and Rational Therapeutic Targets. <i>Frontiers in Immunology</i> , 2021 , 12, 680134	8.4	18
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