

Lorenzo Bonaguro

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2239078/publications.pdf>

Version: 2024-02-01

13
papers

2,831
citations

1040056

9
h-index

1125743

13
g-index

20
all docs

20
docs citations

20
times ranked

6234
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 1 | Complement activation induces excessive T cell cytotoxicity in severe COVID-19. <i>Cell</i> , 2022, 185, 493-512.e25. | 28.9 | 122 |
| 2 | Disease severity-specific neutrophil signatures in blood transcriptomes stratify COVID-19 patients. <i>Genome Medicine</i> , 2021, 13, 7. | 8.2 | 193 |
| 3 | Neutrophils in COVID-19. <i>Frontiers in Immunology</i> , 2021, 12, 652470. | 4.8 | 206 |
| 4 | Swarm Learning for decentralized and confidential clinical machine learning. <i>Nature</i> , 2021, 594, 265-270. | 27.8 | 375 |
| 5 | Creld1 regulates myocardial development and function. <i>Journal of Molecular and Cellular Cardiology</i> , 2021, 156, 45-56. | 1.9 | 11 |
| 6 | Early IFN- γ signatures and persistent dysfunction are distinguishing features of NK cells in severe COVID-19. <i>Immunity</i> , 2021, 54, 2650-2669.e14. | 14.3 | 145 |
| 7 | Creld2 function during unfolded protein response is essential for liver metabolism homeostasis. <i>FASEB Journal</i> , 2021, 35, e21939. | 0.5 | 15 |
| 8 | Induction of Rosette-to-Lumen stage embryoids using reprogramming paradigms in ESCs. <i>Nature Communications</i> , 2021, 12, 7322. | 12.8 | 6 |
| 9 | Severe COVID-19 Is Marked by a Dysregulated Myeloid Cell Compartment. <i>Cell</i> , 2020, 182, 1419-1440.e23. | 28.9 | 1,162 |
| 10 | Optimized workflow for single-cell transcriptomics on infectious diseases including COVID-19. <i>STAR Protocols</i> , 2020, 1, 100233. | 1.2 | 10 |
| 11 | CRELD1 modulates homeostasis of the immune system in mice and humans. <i>Nature Immunology</i> , 2020, 21, 1517-1527. | 14.5 | 13 |
| 12 | Human Monocyte Subsets and Phenotypes in Major Chronic Inflammatory Diseases. <i>Frontiers in Immunology</i> , 2019, 10, 2035. | 4.8 | 529 |
| 13 | A cross-species approach to identify transcriptional regulators exemplified for Dnajc22 and Hnf4a. <i>Scientific Reports</i> , 2017, 7, 4056. | 3.3 | 3 |