

# Simone Rizzetto

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9039565/publications.pdf>

Version: 2024-02-01

18  
papers

1,044  
citations

623188

14  
h-index

839053

18  
g-index

25  
all docs

25  
docs citations

25  
times ranked

2249  
citing authors

#	ARTICLE	IF	CITATIONS
1	Pan-Cancer Analysis of Ligandâ€“Receptor Cross-talk in the Tumor Microenvironment. <i>Cancer Research</i> , 2021, 81, 1802-1812.	0.4	41
2	CD8+ T cell landscape in Indigenous and non-Indigenous people restricted by influenza mortality-associated HLA-A*24:02 allomorph. <i>Nature Communications</i> , 2021, 12, 2931.	5.8	20
3	B cell immunodominance in primary hepatitis C virus infection. <i>Journal of Hepatology</i> , 2020, 72, 670-679.	1.8	21
4	Mass cytometry reveals immune signatures associated with cytomegalovirus (CMV) control in recipients of allogeneic haemopoietic stem cell transplant and CMVâ€“specific T cells. <i>Clinical and Translational Immunology</i> , 2020, 9, e1149.	1.7	18
5	Exploring and analysing single cell multi-omics data with VDJView. <i>BMC Medical Genomics</i> , 2020, 13, 29.	0.7	15
6	Lymphoma Driver Mutations in the Pathogenic Evolution of an Iconic Human Autoantibody. <i>Cell</i> , 2020, 180, 878-894.e19.	13.5	82
7	Cytotoxic T cells swarm by homotypic chemokine signalling. <i>ELife</i> , 2020, 9, .	2.8	46
8	Human CD8+ T cell cross-reactivity across influenza A, B and C viruses. <i>Nature Immunology</i> , 2019, 20, 613-625.	7.0	180
9	Single-Cell Transcriptome Analysis of T Cells. <i>Methods in Molecular Biology</i> , 2019, 2048, 155-205.	0.4	3
10	Clonally diverse CD38+HLA-DR+CD8+ T cells persist during fatal H7N9 disease. <i>Nature Communications</i> , 2018, 9, 824.	5.8	107
11	B-cell receptor reconstruction from single-cell RNA-seq with VDJpuzzle. <i>Bioinformatics</i> , 2018, 34, 2846-2847.	1.8	87
12	Toward Large-Scale Computational Prediction of Protein Complexes. <i>Methods in Molecular Biology</i> , 2018, 1819, 271-295.	0.4	2
13	Context-dependent prediction of protein complexes by SiComPre. <i>Npj Systems Biology and Applications</i> , 2018, 4, 37.	1.4	2
14	Mass Cytometry for the Assessment of Immune Reconstitution After Hematopoietic Stem Cell Transplantation. <i>Frontiers in Immunology</i> , 2018, 9, 1672.	2.2	46
15	Impact of sequencing depth and read length on single cell RNA sequencing data of T cells. <i>Scientific Reports</i> , 2017, 7, 12781.	1.6	60
16	A Liver Capsular Network of Monocyte-Derived Macrophages Restricts Hepatic Dissemination of Intraperitoneal Bacteria by Neutrophil Recruitment. <i>Immunity</i> , 2017, 47, 374-388.e6.	6.6	171
17	Linking the T cell receptor to the single cell transcriptome in antigenâ€“specific human T cells. <i>Immunology and Cell Biology</i> , 2016, 94, 604-611.	1.0	118
18	Qualitative and Quantitative Protein Complex Prediction Through Proteome-Wide Simulations. <i>PLoS Computational Biology</i> , 2015, 11, e1004424.	1.5	21