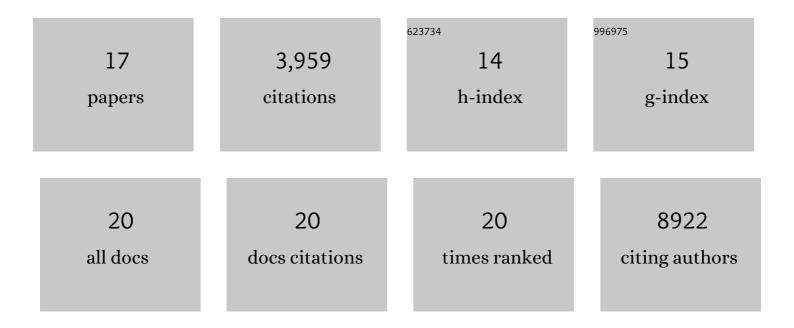
Vladimir Jojic

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11050220/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	The immunoregulatory landscape of human tuberculosis granulomas. Nature Immunology, 2022, 23, 318-329.	14.5	110
2	An inflammatory aging clock (iAge) based on deep learning tracks multimorbidity, immunosenescence, frailty and cardiovascular aging. Nature Aging, 2021, 1, 598-615.	11.6	202
3	ImmGen at 15. Nature Immunology, 2020, 21, 700-703.	14.5	55
4	Modeling Multiplexed Images with <i>Spatial-LDA</i> Reveals Novel Tissue Microenvironments. Journal of Computational Biology, 2020, 27, 1204-1218.	1.6	42
5	Reconstructing Tissue Properties From Medical Images With Application in Cancer Screening. IEEE Transactions on Medical Robotics and Bionics, 2019, 1, 6-13.	3.2	Ο
6	Design of synthetic bacterial communities for predictable plant phenotypes. PLoS Biology, 2018, 16, e2003962.	5.6	182
7	Expression of specific inflammasome gene modules stratifies older individuals into two extreme clinical and immunological states. Nature Medicine, 2017, 23, 174-184.	30.7	304
8	Tradict enables accurate prediction of eukaryotic transcriptional states from 100 marker genes. Nature Communications, 2017, 8, 15309.	12.8	18
9	Learning Microbial Interaction Networks from Metagenomic Count Data. Journal of Computational Biology, 2016, 23, 526-535.	1.6	39
10	Variation in the Human Immune System Is Largely Driven by Non-Heritable Influences. Cell, 2015, 160, 37-47.	28.9	828
11	Cytomegalovirus infection enhances the immune response to influenza. Science Translational Medicine, 2015, 7, 281ra43.	12.4	277
12	Systems analysis of sex differences reveals an immunosuppressive role for testosterone in the response to influenza vaccination. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 869-874.	7.1	542
13	DRUG-INDUCED mRNA SIGNATURES ARE ENRICHED FOR THE MINORITY OF GENES THAT ARE HIGHLY HERITABLE. , 2014, , .		1
14	Apoptosis and other immune biomarkers predict influenza vaccine responsiveness. Molecular Systems Biology, 2013, 9, 659.	7.2	173
15	Identification of transcriptional regulators in the mouse immune system. Nature Immunology, 2013, 14, 633-643.	14.5	179
16	Conservation and divergence in the transcriptional programs of the human and mouse immune systems. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 2946-2951.	7.1	296
17	Deciphering the transcriptional network of the dendritic cell lineage. Nature Immunology, 2012, 13, 888-899.	14.5	688