

# Marianna Orlova

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6275844/publications.pdf>

Version: 2024-02-01

23  
papers

924  
citations

623734

14  
h-index

642732

23  
g-index

26  
all docs

26  
docs citations

26  
times ranked

1222  
citing authors

#	ARTICLE	IF	CITATIONS
1	A systems biology approach identifies candidate drugs to reduce mortality in severely ill patients with COVID-19. <i>Science Advances</i> , 2022, 8, .	10.3	14
2	Peculiar Phenotypic and Cytotoxic Features of Pulmonary Mucosal CD8 T Cells in People Living with HIV Receiving Long-Term Antiretroviral Therapy. <i>Journal of Immunology</i> , 2021, 206, 641-651.	0.8	5
3	Genome-wide association study of resistance to <i>Mycobacterium tuberculosis</i> infection identifies a locus at 10q26.2 in three distinct populations. <i>PLoS Genetics</i> , 2021, 17, e1009392.	3.5	17
4	Alveolar macrophages from persons living with HIV show impaired epigenetic response to <i>Mycobacterium tuberculosis</i> . <i>Journal of Clinical Investigation</i> , 2021, 131, .	8.2	19
5	Deep resequencing identifies candidate functional genes in leprosy GWAS loci. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0010029.	3.0	5
6	An observational study identifying highly tuberculosis-exposed, HIV-1-positive but persistently TB, tuberculin and IGRA negative persons with <i>M. tuberculosis</i> specific antibodies in Cape Town, South Africa. <i>EBioMedicine</i> , 2020, 61, 103053.	6.1	22
7	The complex pattern of genetic associations of leprosy with HLA class I and class II alleles can be reduced to four amino acid positions. <i>PLoS Pathogens</i> , 2020, 16, e1008818.	4.7	14
8	Family-based genome-wide association study of leprosy in Vietnam. <i>PLoS Pathogens</i> , 2020, 16, e1008565.	4.7	8
9	Pleiotropic effects for Parkin and LRRK2 in leprosy type-1 reactions and Parkinson's disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 15616-15624.	7.1	50
10	HIV persistence in mucosal CD4+ T cells within the lungs of adults receiving long-term suppressive antiretroviral therapy. <i>Aids</i> , 2018, 32, 2279-2289.	2.2	44
11	An eQTL variant of ZXDC is associated with IFN- $\gamma$ production following <i>Mycobacterium tuberculosis</i> antigen-specific stimulation. <i>Scientific Reports</i> , 2017, 7, 12800.	3.3	5
12	A genome wide association study identifies a lncRNA as risk factor for pathological inflammatory responses in leprosy. <i>PLoS Genetics</i> , 2017, 13, e1006637.	3.5	29
13	Deciphering the genetic control of gene expression following <i>Mycobacterium leprae</i> antigen stimulation. <i>PLoS Genetics</i> , 2017, 13, e1006952.	3.5	37
14	Pauci- and Multibacillary Leprosy: Two Distinct, Genetically Neglected Diseases. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004345.	3.0	57
15	A Missense LRRK2 Variant Is a Risk Factor for Excessive Inflammatory Responses in Leprosy. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004412.	3.0	181
16	Association of TNFSF8 Regulatory Variants With Excessive Inflammatory Responses but not Leprosy Per Se. <i>Journal of Infectious Diseases</i> , 2015, 211, 968-977.	4.0	29
17	CUBN and NEBL common variants in the chromosome 10p13 linkage region are associated with multibacillary leprosy in Vietnam. <i>Human Genetics</i> , 2014, 133, 883-93.	3.8	12
18	Linkage disequilibrium pattern and age-at-diagnosis are critical for replicating genetic associations across ethnic groups in leprosy. <i>Human Genetics</i> , 2013, 132, 107-116.	3.8	32

#	ARTICLE	IF	CITATIONS
19	Gene Set Signature of Reversal Reaction Type I in Leprosy Patients. <i>PLoS Genetics</i> , 2013, 9, e1003624.	3.5	32
20	Crohn's Disease Susceptibility Genes are Associated With Leprosy in the Vietnamese Population. <i>Journal of Infectious Diseases</i> , 2012, 206, 1763-1767.	4.0	70
21	Genetics of leprosy reactions: an overview. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2012, 107, 132-142.	1.6	41
22	Human Leukocyte Antigen Class I Region Single-Nucleotide Polymorphisms are Associated with Leprosy Susceptibility in Vietnam and India. <i>Journal of Infectious Diseases</i> , 2011, 203, 1274-1281.	4.0	49
23	Stepwise replication identifies a low-producing lymphotoxin-1 allele as a major risk factor for early-onset leprosy. <i>Nature Genetics</i> , 2007, 39, 517-522.	21.4	152