

# Damjan Vukcevic

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1331834/publications.pdf>

Version: 2024-02-01

39  
papers

21,213  
citations

394421

19  
h-index

414414

32  
g-index

45  
all docs

45  
docs citations

45  
times ranked

30466  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association study of 14,000 cases of seven common diseases and 3,000 shared controls. <i>Nature</i> , 2007, 447, 661-678.	27.8	8,895
2	The UK Biobank resource with deep phenotyping and genomic data. <i>Nature</i> , 2018, 562, 203-209.	27.8	5,221
3	Bayesian Test for Colocalisation between Pairs of Genetic Association Studies Using Summary Statistics. <i>PLoS Genetics</i> , 2014, 10, e1004383.	3.5	2,012
4	Association scan of 14,500 nonsynonymous SNPs in four diseases identifies autoimmunity variants. <i>Nature Genetics</i> , 2007, 39, 1329-1337.	21.4	1,298
5	Identification of 15 new psoriasis susceptibility loci highlights the role of innate immunity. <i>Nature Genetics</i> , 2012, 44, 1341-1348.	21.4	848
6	Genome-wide association study of CNVs in 16,000 cases of eight common diseases and 3,000 shared controls. <i>Nature</i> , 2010, 464, 713-720.	27.8	737
7	Bayesian refinement of association signals for 14 loci in 3 common diseases. <i>Nature Genetics</i> , 2012, 44, 1294-1301.	21.4	469
8	Identification and Stoichiometry of Glycosylphosphatidylinositol-anchored Membrane Proteins of the Human Malaria Parasite <i>Plasmodium falciparum</i> . <i>Molecular and Cellular Proteomics</i> , 2006, 5, 1286-1299.	3.8	243
9	Genome-wide association study of eosinophilic granulomatosis with polyangiitis reveals genomic loci stratified by ANCA status. <i>Nature Communications</i> , 2019, 10, 5120.	12.8	160
10	C5H and Susceptibility to Infectious Diseases. <i>New England Journal of Medicine</i> , 2010, 362, 2092-2101.	27.0	129
11	Strong genetic evidence for a selective influence of GABAA receptors on a component of the bipolar disorder phenotype. <i>Molecular Psychiatry</i> , 2010, 15, 146-153.	7.9	111
12	Common variants in the HLA-DRB1 and HLA-DQA1 HLA class II region are associated with susceptibility to visceral leishmaniasis. <i>Nature Genetics</i> , 2013, 45, 208-213.	21.4	86
13	Genetic utility of broadly defined bipolar schizoaffective disorder as a diagnostic concept. <i>British Journal of Psychiatry</i> , 2009, 195, 23-29.	2.8	83
14	Imputation of KIR Types from SNP Variation Data. <i>American Journal of Human Genetics</i> , 2015, 97, 593-607.	6.2	73
15	The correlation between reading and mathematics ability at age twelve has a substantial genetic component. <i>Nature Communications</i> , 2014, 5, 4204.	12.8	72
16	Bayesian analysis of genetic association across tree-structured routine healthcare data in the UK Biobank. <i>Nature Genetics</i> , 2017, 49, 1311-1318.	21.4	56
17	Polymorphism in a lincRNA Associates with a Doubled Risk of Pneumococcal Bacteremia in Kenyan Children. <i>American Journal of Human Genetics</i> , 2016, 98, 1092-1100.	6.2	39
18	Quantifying the Underestimation of Relative Risks from Genome-Wide Association Studies. <i>PLoS Genetics</i> , 2011, 7, e1001337.	3.5	37

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19	Disease model distortion in association studies. <i>Genetic Epidemiology</i> , 2011, 35, 278-290.	1.3	26
20	Multi-output Gaussian processes for species distribution modelling. <i>Methods in Ecology and Evolution</i> , 2020, 11, 1587-1598.	5.2	19
21	The Effect of ACTN3 Gene Doping on Skeletal Muscle Performance. <i>American Journal of Human Genetics</i> , 2018, 102, 845-857.	6.2	17
22	Genetic variation in VAC14 is associated with bacteremia secondary to diverse pathogens in African children. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E3601-E3603.	7.1	12
23	Epistatic interactions between killer immunoglobulin-like receptors and human leukocyte antigen ligands are associated with ankylosing spondylitis. <i>PLoS Genetics</i> , 2020, 16, e1008906.	3.5	12
24	Temporal clustering of rotational glitches in the Crab pulsar. <i>Monthly Notices of the Royal Astronomical Society</i> , 2019, 482, 3736-3743.	4.4	8
25	Lack of transparency in software used to analyze multiple breath washout data. <i>Pediatric Pulmonology</i> , 2016, 51, 1108-1110.	2.0	7
26	Large-Scale Imputation of KIR Copy Number and HLA Alleles in North American and European Psoriasis Case-Control Cohorts Reveals Association of Inhibitory KIR2DL2 With Psoriasis. <i>Frontiers in Immunology</i> , 2021, 12, 684326.	4.8	7
27	The influence of sighing respirations on infant lung function measured using multiple breath washout gas mixing techniques. <i>Physiological Reports</i> , 2015, 3, e12347.	1.7	6
28	Assertion-Based Approaches to Auditing Complex Elections, with Application to Party-List Proportional Elections. <i>Lecture Notes in Computer Science</i> , 2021, , 47-62.	1.3	5
29	A Unified Evaluation of Two-Candidate Ballot-Polling Election Auditing Methods. <i>Lecture Notes in Computer Science</i> , 2020, , 112-128.	1.3	5
30	Allele imputation for the killer cell immunoglobulin-like receptor KIR3DL1/S1. <i>PLoS Computational Biology</i> , 2022, 18, e1009059.	3.2	5
31	Auditing Hamiltonian Elections. <i>Lecture Notes in Computer Science</i> , 2021, , 235-250.	1.3	4
32	Bayesian modelling of lung function data from multiple breath washout tests. <i>Statistics in Medicine</i> , 2018, 37, 2016-2033.	1.6	3
33	Insights Into the Health of Defence Simulated Workforce Systems Using Data Farming and Analytics. , 2019, , .		1
34	Young Writers Competition: GUESTimation: Breaking the deadlock on wedding guest lists. <i>Significance</i> , 2013, 10, 45-48.	0.4	0
35	Random Errors Are Not Necessarily Politically Neutral. <i>Lecture Notes in Computer Science</i> , 2020, , 19-35.	1.3	0
36	Title is missing!. , 2020, 16, e1008906.		0

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37	Title is missing!. , 2020, 16, e1008906.		0
38	Title is missing!. , 2020, 16, e1008906.		0
39	Title is missing!.. , 2020, 16, e1008906.		0