

Indra Adrianto

List of Publications by Year in descending order

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Version: 2024-02-01

47
papers

2,138
citations

331670

21
h-index

254184

43
g-index

49
all docs

49
docs citations

49
times ranked

3712
citing authors

#	ARTICLE	IF	CITATIONS
1	Variants at multiple loci implicated in both innate and adaptive immune responses are associated with Sjögren's syndrome. <i>Nature Genetics</i> , 2013, 45, 1284-1292.	21.4	427
2	Association of a functional variant downstream of TNFAIP3 with systemic lupus erythematosus. <i>Nature Genetics</i> , 2011, 43, 253-258.	21.4	242
3	Identification of IRF8, TMEM39A, and IKZF3-ZBP2 as Susceptibility Loci for Systemic Lupus Erythematosus in a Large-Scale Multiracial Replication Study. <i>American Journal of Human Genetics</i> , 2012, 90, 648-660.	6.2	161
4	Genome-Wide Association Study of African and European Americans Implicates Multiple Shared and Ethnic Specific Loci in Sarcoidosis Susceptibility. <i>PLoS ONE</i> , 2012, 7, e43907.	2.5	105
5	Identification of a Systemic Lupus Erythematosus Susceptibility Locus at 11p13 between PDHX and CD44 in a Multiethnic Study. <i>American Journal of Human Genetics</i> , 2011, 88, 83-91.	6.2	72
6	ABIN1 Dysfunction as a Genetic Basis for Lupus Nephritis. <i>Journal of the American Society of Nephrology: JASN</i> , 2013, 24, 1743-1754.	6.1	70
7	Association of two independent functional risk haplotypes in <i>TNIP1</i> with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2012, 64, 3695-3705.	6.7	69
8	High-Density Genetic Mapping Identifies New Susceptibility Variants in Sarcoidosis Phenotypes and Shows Genomic-driven Phenotypic Differences. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2016, 193, 1008-1022.	5.6	68
9	Genetics of Sjögren's syndrome in the genome-wide association era. <i>Journal of Autoimmunity</i> , 2012, 39, 57-63.	6.5	61
10	Identification of a Sjögren's syndrome susceptibility locus at OAS1 that influences isoform switching, protein expression, and responsiveness to type I interferons. <i>PLoS Genetics</i> , 2017, 13, e1006820.	3.5	60
11	Role of MYH9 and APOL1 in African and non-African populations with lupus nephritis. <i>Genes and Immunity</i> , 2012, 13, 232-238.	4.1	58
12	The genomics of autoimmune disease in the era of genome-wide association studies and beyond. <i>Autoimmunity Reviews</i> , 2012, 11, 267-275.	5.8	58
13	A functional haplotype of UBE2L3 confers risk for systemic lupus erythematosus. <i>Genes and Immunity</i> , 2012, 13, 380-387.	4.1	50
14	Confirmation of Linkage to and Localization of Familial Colon Cancer Risk Haplotype on Chromosome 9q22. <i>Cancer Research</i> , 2010, 70, 5409-5418.	0.9	42
15	Association of <i>HLA-DRB1</i> with Sarcoidosis Susceptibility and Progression in African Americans. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2015, 53, 206-216.	2.9	42
16	Suppression of ILC2 differentiation from committed T cell precursors by E protein transcription factors. <i>Journal of Experimental Medicine</i> , 2019, 216, 884-899.	8.5	41
17	Downregulation of E Protein Activity Augments an ILC2 Differentiation Program in the Thymus. <i>Journal of Immunology</i> , 2017, 198, 3149-3156.	0.8	39
18	Cytokine profiles show heterogeneity of interferon- γ response in multiple sclerosis patients. <i>Neurology: Neuroimmunology and NeuroInflammation</i> , 2016, 3, e202.	6.0	34

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19	Admixture Fine-Mapping in African Americans Implicates XAF1 as a Possible Sarcoidosis Risk Gene. PLoS ONE, 2014, 9, e92646.	2.5	31
20	Performance of HLA allele prediction methods in African Americans for class II genes HLA-DRB1, HLA-DQB1, and HLA-DPB1. BMC Genetics, 2014, 15, 72.	2.7	24
21	Kernel logistic regression using truncated Newton method. Computational Management Science, 2011, 8, 415-428.	1.3	23
22	Histone deacetylase 3 controls lung alveolar macrophage development and homeostasis. Nature Communications, 2020, 11, 3822.	12.8	22
23	Single-Cell RNA-Seq Analysis Uncovers Distinct Functional Human NKT Cell Sub-Populations in Peripheral Blood. Frontiers in Cell and Developmental Biology, 2020, 8, 384.	3.7	22
24	Genome-Wide Association Scan of Dupuytren's Disease. Journal of Hand Surgery, 2010, 35, 2039-2045.	1.6	21
25	Classification and regionalization through kernel principal component analysis. Physics and Chemistry of the Earth, 2010, 35, 316-328.	2.9	21
26	Genome-Wide Association Study of Ocular Sarcoidosis Confirms HLA Associations and Implicates Barrier Function and Autoimmunity in African Americans. Ocular Immunology and Inflammation, 2021, 29, 244-249.	1.8	21
27	Missing Data Imputation Through Machine Learning Algorithms. , 2009, , 153-169.		21
28	Support vector machines for spatiotemporal tornado prediction. International Journal of General Systems, 2009, 38, 759-776.	2.5	19
29	Detrimental effects of duplicate reads and low complexity regions on RNA- and ChIP-seq data. BMC Bioinformatics, 2015, 16, S10.	2.6	19
30	Identification of Distinct Heterogenic Subtypes and Molecular Signatures Associated with African Ancestry in Triple Negative Breast Cancer Using Quantified Genetic Ancestry Models in Admixed Race Populations. Cancers, 2020, 12, 1220.	3.7	19
31	Machine-learning classifiers for imbalanced tornado data. Computational Management Science, 2014, 11, 403-418.	1.3	18
32	Disease Activity in Systemic Lupus Erythematosus Correlates With Expression of the Transcription Factor AT–Interactive Domain 3A. Arthritis and Rheumatology, 2014, 66, 3404-3412.	5.6	18
33	Role of NOD2 Pathway Genes in Sarcoidosis Cases with Clinical Characteristics of Blau Syndrome. American Journal of Respiratory and Critical Care Medicine, 2015, 192, 1133-1135.	5.6	18
34	Active Learning with Support Vector Machines for Tornado Prediction. Lecture Notes in Computer Science, 2007, , 1130-1137.	1.3	18
35	Fine mapping of chromosome 15q25 implicates ZNF592 in neurosarcoidosis patients. Annals of Clinical and Translational Neurology, 2015, 2, 972-977.	3.7	17
36	Efficient Generalized Least Squares Method for Mixed Population and Family-based Samples in Genome-wide Association Studies. Genetic Epidemiology, 2014, 38, 430-438.	1.3	14

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37	NRG1 variant effects in patients with Hirschsprung disease. <i>BMC Pediatrics</i> , 2018, 18, 292.	1.7	12
38	Antibodies to periodontogenic bacteria are associated with higher disease activity in lupus patients. <i>Clinical and Experimental Rheumatology</i> , 2019, 37, 106-111.	0.8	11
39	Extended methods for gene-environment-wide interaction scans in studies of admixed individuals with varying degrees of relationships. <i>Genetic Epidemiology</i> , 2019, 43, 414-426.	1.3	10
40	Estimating Allele Frequencies. <i>Methods in Molecular Biology</i> , 2012, 850, 59-76.	0.9	9
41	A stacked regression ensemble approach for the quantitative determination of biomass feedstock compositions using near infrared spectroscopy. <i>Spectrochimica Acta - Part A: Molecular and Biomolecular Spectroscopy</i> , 2022, 276, 121231.	3.9	8
42	Estimating Allele Frequencies. <i>Methods in Molecular Biology</i> , 2017, 1666, 61-81.	0.9	7
43	Current Developments in Machine Learning Techniques in Biological Data Mining. <i>Bioinformatics and Biology Insights</i> , 2017, 11, 117793221668754.	2.0	6
44	Tribbles 2 pseudokinase confers enzalutamide resistance in prostate cancer by promoting lineage plasticity. <i>Journal of Biological Chemistry</i> , 2022, 298, 101556.	3.4	4
45	The p-Centre machine for regression analysis. <i>Optimization Methods and Software</i> , 2010, 25, 171-183.	2.4	0
46	Effects of Harm Events on the Rate of 30-Day Readmissions in Surgical Patients. <i>Journal of the American College of Surgeons</i> , 2018, 227, S99-S100.	0.5	0
47	Genetic Epidemiology of Obesity and Cancer. , 2010, , 87-127.		0