

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
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all docs

49
docs citations

49
times ranked

3712
citing authors

#	ARTICLE	IF	CITATIONS
1	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
2	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
3	Genome-Wide Association Study of Ocular Sarcoidosis Confirms HLA Associations and Implicates Barrier Function and Autoimmunity in African Americans. <i>Ocular Immunology and Inflammation</i> , 2021, 29, 244-249.	1.8	21
4	Histone deacetylase 3 controls lung alveolar macrophage development and homeostasis. <i>Nature Communications</i> , 2020, 11, 3822.	12.8	22
5	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. <i>Cancers</i> , 2020, 12, 1220.	3.7	19
6	Single-Cell RNA-Seq Analysis Uncovers Distinct Functional Human NKT Cell Sub-Populations in Peripheral Blood. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 384.	3.7	22
7	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
8	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
9	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
10	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
11	NRG1 variant effects in patients with Hirschsprung disease. <i>BMC Pediatrics</i> , 2018, 18, 292.	1.7	12
12	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
13	Estimating Allele Frequencies. <i>Methods in Molecular Biology</i> , 2017, 1666, 61-81.	0.9	7
14	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
15	Current Developments in Machine Learning Techniques in Biological Data Mining. <i>Bioinformatics and Biology Insights</i> , 2017, 11, 117793221668754.	2.0	6
16	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
17	Cytokine profiles show heterogeneity of interferon- γ response in multiple sclerosis patients. <i>Neurology: Neuroimmunology and NeuroInflammation</i> , 2016, 3, e202.	6.0	34
18	Fine mapping of chromosome 15q25 implicates <i>ZNF592</i> in neurosarcoidosis patients. <i>Annals of Clinical and Translational Neurology</i> , 2015, 2, 972-977.	3.7	17

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19	Detrimental effects of duplicate reads and low complexity regions on RNA- and ChIP-seq data. BMC Bioinformatics, 2015, 16, S10.	2.6	19
20	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
21	Association of <i>HLA-DRB1</i> with Sarcoidosis Susceptibility and Progression in African Americans. American Journal of Respiratory Cell and Molecular Biology, 2015, 53, 206-216.	2.9	42
22	Machine-learning classifiers for imbalanced tornado data. Computational Management Science, 2014, 11, 403-418.	1.3	18
23	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
24	Performance of HLA allele prediction methods in African Americans for class II genes HLA-DRB1, <i>DQB1</i> , and <i>DPB1</i> . BMC Genetics, 2014, 15, 72.	2.7	24
25	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
26	Admixture Fine-Mapping in African Americans Implicates XAF1 as a Possible Sarcoidosis Risk Gene. PLoS ONE, 2014, 9, e92646.	2.5	31
27	Variants at multiple loci implicated in both innate and adaptive immune responses are associated with Sj“gren's syndrome. Nature Genetics, 2013, 45, 1284-1292.	21.4	427
28	ABIN1 Dysfunction as a Genetic Basis for Lupus Nephritis. Journal of the American Society of Nephrology: JASN, 2013, 24, 1743-1754.	6.1	70
29	Role of MYH9 and APOL1 in African and non-African populations with lupus nephritis. Genes and Immunity, 2012, 13, 232-238.	4.1	58
30	A functional haplotype of UBE2L3 confers risk for systemic lupus erythematosus. Genes and Immunity, 2012, 13, 380-387.	4.1	50
31	Association of two independent functional risk haplotypes in <i>TNIP1</i> with systemic lupus erythematosus. Arthritis and Rheumatism, 2012, 64, 3695-3705.	6.7	69
32	Genetics of Sj“gren's syndrome in the genome-wide association era. Journal of Autoimmunity, 2012, 39, 57-63.	6.5	61
33	Genome-Wide Association Study of African and European Americans Implicates Multiple Shared and Ethnic Specific Loci in Sarcoidosis Susceptibility. PLoS ONE, 2012, 7, e43907.	2.5	105
34	Identification of IRF8, TMEM39A, and IKZF3-ZPBP2 as Susceptibility Loci for Systemic Lupus Erythematosus in a Large-Scale Multiracial Replication Study. American Journal of Human Genetics, 2012, 90, 648-660.	6.2	161
35	The genomics of autoimmune disease in the era of genome-wide association studies and beyond. Autoimmunity Reviews, 2012, 11, 267-275.	5.8	58
36	Estimating Allele Frequencies. Methods in Molecular Biology, 2012, 850, 59-76.	0.9	9

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37	Association of a functional variant downstream of TNFAIP3 with systemic lupus erythematosus. Nature Genetics, 2011, 43, 253-258.	21.4	242
38	Identification of a Systemic Lupus Erythematosus Susceptibility Locus at 11p13 between PDHX and CD44 in a Multiethnic Study. American Journal of Human Genetics, 2011, 88, 83-91.	6.2	72
39	Kernel logistic regression using truncated Newton method. Computational Management Science, 2011, 8, 415-428.	1.3	23
40	Confirmation of Linkage to and Localization of Familial Colon Cancer Risk Haplotype on Chromosome 9q22. Cancer Research, 2010, 70, 5409-5418.	0.9	42
41	The p-Centre machine for regression analysis. Optimization Methods and Software, 2010, 25, 171-183.	2.4	0
42	Genome-Wide Association Scan of Dupuytren's Disease. Journal of Hand Surgery, 2010, 35, 2039-2045.	1.6	21
43	Classification and regionalization through kernel principal component analysis. Physics and Chemistry of the Earth, 2010, 35, 316-328.	2.9	21
44	Genetic Epidemiology of Obesity and Cancer. , 2010, , 87-127.		0
45	Support vector machines for spatiotemporal tornado prediction. International Journal of General Systems, 2009, 38, 759-776.	2.5	19
46	Missing Data Imputation Through Machine Learning Algorithms. , 2009, , 153-169.		21
47	Active Learning with Support Vector Machines for Tornado Prediction. Lecture Notes in Computer Science, 2007, , 1130-1137.	1.3	18