

Chris Cotsapas

List of Publications by Year in descending order

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Version: 2024-02-01

56
papers

13,910
citations

145106

33
h-index

175968

55
g-index

68
all docs

68
docs citations

68
times ranked

28877
citing authors

#	ARTICLE	IF	CITATIONS
1	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. <i>Biological Psychiatry</i> , 2022, 91, 102-117.	0.7	61
2	Seasonal variation and risk of febrile seizures; a Danish nationwide cohort study. <i>Neuroepidemiology</i> , 2022, , .	1.1	2
3	Epilepsy risk in offspring of affected parents; a cohort study of the "maternal effect" in epilepsy. <i>Annals of Clinical and Translational Neurology</i> , 2021, 8, 153-162.	1.7	6
4	Shared associations identify causal relationships between gene expression and immune cell phenotypes. <i>Communications Biology</i> , 2021, 4, 279.	2.0	3
5	Birth characteristics and risk of febrile seizures. <i>Acta Neurologica Scandinavica</i> , 2021, 144, 51-57.	1.0	12
6	Shared genetic basis between genetic generalized epilepsy and background electroencephalographic oscillations. <i>Epilepsia</i> , 2021, 62, 1518-1527.	2.6	5
7	Sub-genic intolerance, ClinVar, and the epilepsies: A whole-exome sequencing study of 29,165 individuals. <i>American Journal of Human Genetics</i> , 2021, 108, 965-982.	2.6	35
8	Do monogenic inborn errors of immunity cause susceptibility to severe COVID-19?. <i>Journal of Clinical Investigation</i> , 2021, 131, .	3.9	3
9	Ultra-Rare Genetic Variation in the Epilepsies: A Whole-Exome Sequencing Study of 17,606 Individuals. <i>American Journal of Human Genetics</i> , 2019, 105, 267-282.	2.6	237
10	Multiple sclerosis genomic map implicates peripheral immune cells and microglia in susceptibility. <i>Science</i> , 2019, 365, .	6.0	710
11	ImmuneRegulation: a web-based tool for identifying human immune regulatory elements. <i>Nucleic Acids Research</i> , 2019, 47, W142-W150.	6.5	4
12	Childhood seizures and risk of psychiatric disorders in adolescence and early adulthood: a Danish nationwide cohort study. <i>The Lancet Child and Adolescent Health</i> , 2019, 3, 99-108.	2.7	31
13	Low-Frequency and Rare-Coding Variation Contributes to Multiple Sclerosis Risk. <i>Cell</i> , 2018, 175, 1679-1687.e7.	13.5	115
14	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , 2018, 360, .	6.0	1,085
15	Genome-wide association studies of multiple sclerosis. <i>Clinical and Translational Immunology</i> , 2018, 7, e1018.	1.7	58
16	Multiple sclerosis. <i>Handbook of Clinical Neurology</i> / Edited By P J Vinken and G W Bruyn, 2018, 148, 723-730.	1.0	50
17	Novel determinants of mammalian primary microRNA processing revealed by systematic evaluation of hairpin-containing transcripts and human genetic variation. <i>Genome Research</i> , 2017, 27, 374-384.	2.4	78
18	Limited statistical evidence for shared genetic effects of eQTLs and autoimmune-disease-associated loci in three major immune-cell types. <i>Nature Genetics</i> , 2017, 49, 600-605.	9.4	205

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19	Large-Scale trans-eQTLs Affect Hundreds of Transcripts and Mediate Patterns of Transcriptional Co-regulation. <i>American Journal of Human Genetics</i> , 2017, 100, 581-591.	2.6	86
20	Integrative Genetic and Epigenetic Analysis Uncovers Regulatory Mechanisms of Autoimmune Disease. <i>American Journal of Human Genetics</i> , 2017, 101, 75-86.	2.6	29
21	Microbiota control immune regulation in humanized mice. <i>JCI Insight</i> , 2017, 2, .	2.3	23
22	Regulatory polymorphisms modulate the expression of HLA class II molecules and promote autoimmunity. <i>ELife</i> , 2016, 5, .	2.8	113
23	Progress and challenges for treating Type 1 diabetes. <i>Journal of Autoimmunity</i> , 2016, 71, 1-9.	3.0	23
24	Changes in T _H 1 cell subsets identify responders to Fc γ R nonbinding anti-CD3 mAb (teplizumab) in patients with type 1 diabetes. <i>European Journal of Immunology</i> , 2016, 46, 230-241.	1.6	48
25	NR1H3 p.Arg415Gln Is Not Associated to Multiple Sclerosis Risk. <i>Neuron</i> , 2016, 92, 333-335.	3.8	24
26	Survey of variation in human transcription factors reveals prevalent DNA binding changes. <i>Science</i> , 2016, 351, 1450-1454.	6.0	114
27	Network Analysis of Genome-Wide Selective Constraint Reveals a Gene Network Active in Early Fetal Brain Intolerant of Mutation. <i>PLoS Genetics</i> , 2016, 12, e1006121.	1.5	24
28	Shared genetic basis for migraine and ischemic stroke. <i>Neurology</i> , 2015, 84, 2132-2145.	1.5	91
29	Genetic variants associated with autoimmunity drive NF κ B signaling and responses to inflammatory stimuli. <i>Science Translational Medicine</i> , 2015, 7, 291ra93.	5.8	81
30	Genetic analysis for a shared biological basis between migraine and coronary artery disease. <i>Neurology: Genetics</i> , 2015, 1, e10.	0.9	61
31	Class II HLA interactions modulate genetic risk for multiple sclerosis. <i>Nature Genetics</i> , 2015, 47, 1107-1113.	9.4	312
32	Weight Loss after Gastric Bypass Is Associated with a Variant at 15q26.1. <i>American Journal of Human Genetics</i> , 2013, 92, 827-834.	2.6	65
33	Analysis of immune-related loci identifies 48 new susceptibility variants for multiple sclerosis. <i>Nature Genetics</i> , 2013, 45, 1353-1360.	9.4	1,213
34	Immune-mediated disease genetics: the shared basis of pathogenesis. <i>Trends in Immunology</i> , 2013, 34, 22-26.	2.9	88
35	Network-Based Multiple Sclerosis Pathway Analysis with GWAS Data from 15,000 Cases and 30,000 Controls. <i>American Journal of Human Genetics</i> , 2013, 92, 854-865.	2.6	164
36	Pleiotropy in complex traits: challenges and strategies. <i>Nature Reviews Genetics</i> , 2013, 14, 483-495.	7.7	958

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37	Systematic Localization of Common Disease-Associated Variation in Regulatory DNA. <i>Science</i> , 2012, 337, 1190-1195.	6.0	3,129
38	Unraveling Multiple MHC Gene Associations with Systemic Lupus Erythematosus: Model Choice Indicates a Role for HLA Alleles and Non-HLA Genes in Europeans. <i>American Journal of Human Genetics</i> , 2012, 91, 778-793.	2.6	140
39	Human genetics offers an emerging picture of common pathways and mechanisms in autoimmunity. <i>Current Opinion in Immunology</i> , 2012, 24, 552-557.	2.4	29
40	Heritability of the Weight Loss Response to Gastric Bypass Surgery. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2011, 96, E1630-E1633.	1.8	76
41	Pervasive Sharing of Genetic Effects in Autoimmune Disease. <i>PLoS Genetics</i> , 2011, 7, e1002254.	1.5	540
42	Proteins Encoded in Genomic Regions Associated with Immune-Mediated Disease Physically Interact and Suggest Underlying Biology. <i>PLoS Genetics</i> , 2011, 7, e1001273.	1.5	450
43	Fine Mapping in 94 Inbred Mouse Strains Using a High-Density Haplotype Resource. <i>Genetics</i> , 2010, 185, 1081-1095.	1.2	95
44	Common body mass index-associated variants confer risk of extreme obesity. <i>Human Molecular Genetics</i> , 2009, 18, 3502-3507.	1.4	106
45	Intra- and inter-individual genetic differences in gene expression. <i>Mammalian Genome</i> , 2009, 20, 281-295.	1.0	21
46	Genetic variants near TNFAIP3 on 6q23 are associated with systemic lupus erythematosus. <i>Nature Genetics</i> , 2008, 40, 1059-1061.	9.4	534
47	Genetic Analysis of Human Traits In Vitro: Drug Response and Gene Expression in Lymphoblastoid Cell Lines. <i>PLoS Genetics</i> , 2008, 4, e1000287.	1.5	200
48	Identifying genetic components of drug response in mice. <i>Pharmacogenomics</i> , 2008, 9, 1323-1330.	0.6	2
49	Intra- and inter-individual genetic differences in gene expression. <i>Nature Precedings</i> , 2008, , .	0.1	2
50	Two independent alleles at 6q23 associated with risk of rheumatoid arthritis. <i>Nature Genetics</i> , 2007, 39, 1477-1482.	9.4	497
51	Genome-wide detection and characterization of positive selection in human populations. <i>Nature</i> , 2007, 449, 913-918.	13.7	1,788
52	Hierarchical Bayes variable selection and microarray experiments. <i>Journal of Multivariate Analysis</i> , 2007, 98, 852-872.	0.5	4
53	Normalization procedures and detection of linkage signal in genetical-genomics experiments. <i>Nature Genetics</i> , 2006, 38, 855-856.	9.4	28
54	Genetic dissection of gene regulation in multiple mouse tissues. <i>Mammalian Genome</i> , 2006, 17, 490-495.	1.0	13

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55	Genetic Variation and the Control of Transcription. Cold Spring Harbor Symposia on Quantitative Biology, 2003, 68, 109-114.	2.0	6
56	Pleiotropy in complex traits: challenges and strategies. , 0, .		1