

Thomas D Als

List of Publications by Year in descending order

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Version: 2024-02-01

50
papers

12,844
citations

147566

31
h-index

174990

52
g-index

69
all docs

69
docs citations

69
times ranked

14774
citing authors

#	ARTICLE	IF	CITATIONS
1	Polygenic liability, stressful life events and risk for secondary-treated depression in early life: a nationwide register-based case-cohort study. <i>Psychological Medicine</i> , 2023, 53, 217-226.	2.7	7
2	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. <i>Biological Psychiatry</i> , 2022, 91, 102-117.	0.7	61
3	Identifying the Common Genetic Basis of Antidepressant Response. <i>Biological Psychiatry Global Open Science</i> , 2022, 2, 115-126.	1.0	31
4	Genome-wide by Environment Interaction Study of Stressful Life Events and Hospital-Treated Depression in the iPSYCH2012 Sample. <i>Biological Psychiatry Global Open Science</i> , 2022, 2, 400-410.	1.0	2
5	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. <i>Nature</i> , 2022, 604, 502-508.	13.7	929
6	A polygenic resilience score moderates the genetic risk for schizophrenia. <i>Molecular Psychiatry</i> , 2021, 26, 800-815.	4.1	36
7	Risk variants and polygenic architecture of disruptive behavior disorders in the context of attention-deficit/hyperactivity disorder. <i>Nature Communications</i> , 2021, 12, 576.	5.8	28
8	A Nationwide Cohort Study of Nonrandom Mating in Schizophrenia and Bipolar Disorder. <i>Schizophrenia Bulletin</i> , 2021, 47, 1342-1350.	2.3	17
9	Identification of genetic loci associated with nocturnal enuresis: a genome-wide association study. <i>The Lancet Child and Adolescent Health</i> , 2021, 5, 201-209.	2.7	27
10	Genetic analyses identify widespread sex-differential participation bias. <i>Nature Genetics</i> , 2021, 53, 663-671.	9.4	124
11	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. <i>Nature Genetics</i> , 2021, 53, 817-829.	9.4	629
12	Polygenic Liability and Recurrence of Depression in Patients With First-Onset Depression Treated in Hospital-Based Settings. <i>JAMA Psychiatry</i> , 2021, 78, 792.	6.0	6
13	Polygenic Heterogeneity Across Obsessive-Compulsive Disorder Subgroups Defined by a Comorbid Diagnosis. <i>Frontiers in Genetics</i> , 2021, 12, 711624.	1.1	7
14	The Genetics of the Mood Disorder Spectrum: Genome-wide Association Analyses of More Than 185,000 Cases and 439,000 Controls. <i>Biological Psychiatry</i> , 2020, 88, 169-184.	0.7	137
15	Genetic liability to ADHD and substance use disorders in individuals with ADHD. <i>Addiction</i> , 2020, 115, 1368-1377.	1.7	47
16	Polygenic risk score, psychosocial environment and the risk of attention-deficit/hyperactivity disorder. <i>Translational Psychiatry</i> , 2020, 10, 335.	2.4	22
17	Large-Scale Exome Sequencing Study Implicates Both Developmental and Functional Changes in the Neurobiology of Autism. <i>Cell</i> , 2020, 180, 568-584.e23.	13.5	1,422
18	Schizophrenia polygenic risk scores, urbanicity and treatment-resistant schizophrenia. <i>Schizophrenia Research</i> , 2019, 212, 79-85.	1.1	19

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19	Genome-wide association study implicates CHRNA2 in cannabis use disorder. <i>Nature Neuroscience</i> , 2019, 22, 1066-1074.	7.1	94
20	Genetic Variants Associated With Anxiety and Stress-Related Disorders. <i>JAMA Psychiatry</i> , 2019, 76, 924.	6.0	140
21	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019, 51, 793-803.	9.4	1,191
22	Gene expression imputation across multiple brain regions provides insights into schizophrenia risk. <i>Nature Genetics</i> , 2019, 51, 659-674.	9.4	154
23	Identification of common genetic risk variants for autism spectrum disorder. <i>Nature Genetics</i> , 2019, 51, 431-444.	9.4	1,538
24	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , 2019, 179, 1469-1482.e11.	13.5	935
25	Discovery of the first genome-wide significant risk loci for attention deficit/hyperactivity disorder. <i>Nature Genetics</i> , 2019, 51, 63-75.	9.4	1,594
26	Common schizophrenia alleles are enriched in mutation-intolerant genes and in regions under strong background selection. <i>Nature Genetics</i> , 2018, 50, 381-389.	9.4	1,332
27	A Genetic Investigation of Sex Bias in the Prevalence of Attention-Deficit/Hyperactivity Disorder. <i>Biological Psychiatry</i> , 2018, 83, 1044-1053.	0.7	146
28	Schizophrenia-associated mt-DNA SNPs exhibit highly variable haplogroup affiliation and nuclear ancestry: Bi-genomic dependence raises major concerns for link to disease. <i>PLoS ONE</i> , 2018, 13, e0208828.	1.1	15
29	Complex spatio-temporal distribution and genomic ancestry of mitochondrial DNA haplogroups in 24,216 Danes. <i>PLoS ONE</i> , 2018, 13, e0208829.	1.1	5
30	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. <i>Cell</i> , 2018, 173, 1705-1715.e16.	13.5	623
31	Sequencing and de novo assembly of 150 genomes from Denmark as a population reference. <i>Nature</i> , 2017, 548, 87-91.	13.7	130
32	Nationwide Genomic Study in Denmark Reveals Remarkable Population Homogeneity. <i>Genetics</i> , 2016, 204, 711-722.	1.2	54
33	Systematic Integration of Brain eQTL and GWAS Identifies <i>ZNF323</i> as a Novel Schizophrenia Risk Gene and Suggests Recent Positive Selection Based on Compensatory Advantage on Pulmonary Function. <i>Schizophrenia Bulletin</i> , 2015, 41, 1294-1308.	2.3	48
34	Microevolution in time and space: SNP analysis of historical DNA reveals dynamic signatures of selection in Atlantic cod. <i>Molecular Ecology</i> , 2013, 22, 2424-2440.	2.0	86
35	A genome-wide study of panic disorder suggests the amiloride-sensitive cation channel 1 as a candidate gene. <i>European Journal of Human Genetics</i> , 2012, 20, 84-90.	1.4	45
36	Association of the leucine-7 to proline-7 variation in the signal sequence of neuropeptide Y with major depression. <i>Acta Neuropsychiatrica</i> , 2012, 24, 81-90.	1.0	4

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37	MBL and MASP-2 concentrations in serum and MBL2 promoter polymorphisms are associated to schizophrenia. <i>Acta Neuropsychiatrica</i> , 2012, 24, 199-207.	1.0	11
38	All roads lead to home: panmixia of European eel in the Sargasso Sea. <i>Molecular Ecology</i> , 2011, 20, 1333-1346.	2.0	176
39	Differential timing of gene expression regulation between leptocephali of the two <i>Anguilla</i> eel species in the Sargasso Sea. <i>Ecology and Evolution</i> , 2011, 1, 459-467.	0.8	15
40	Qualitative assessment of the diet of European eel larvae in the Sargasso Sea resolved by DNA barcoding. <i>Biology Letters</i> , 2010, 6, 819-822.	1.0	90
41	Oceanic fronts in the Sargasso Sea control the early life and drift of Atlantic eels. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2010, 277, 3593-3599.	1.2	86
42	Support for a bipolar affective disorder susceptibility locus on chromosome 12q24.3. <i>Psychiatric Genetics</i> , 2010, 20, 93-101.	0.6	6
43	A Mosaic of Chemical Coevolution in a Large Blue Butterfly. <i>Science</i> , 2008, 319, 88-90.	6.0	174
44	Genetics of panic disorder on the Faroe Islands: a replication study of chromosome 9 and panic disorder. <i>Psychiatric Genetics</i> , 2006, 16, 99-104.	0.6	6
45	Highly discrepant proportions of female and male Scandinavian and British Isles ancestry within the isolated population of the Faroe Islands. <i>European Journal of Human Genetics</i> , 2006, 14, 497-504.	1.4	62
46	Microsatellite markers for the large blue butterflies <i>Maculinea nausithous</i> and <i>Maculinea alcon</i> (Lepidoptera: Lycaenidae) and their amplification in other <i>Maculinea</i> species. <i>Molecular Ecology Notes</i> , 2005, 5, 165-168.	1.7	22
47	The evolution of alternative parasitic life histories in large blue butterflies. <i>Nature</i> , 2004, 432, 386-390.	13.7	163
48	The origin of the isolated population of the Faroe Islands investigated using Y chromosomal markers. <i>Human Genetics</i> , 2004, 115, 19-28.	1.8	43
49	Geographical variation in host-ant specificity of the parasitic butterfly <i>Maculinea alcon</i> in Denmark. <i>Ecological Entomology</i> , 2002, 27, 403-414.	1.1	61
50	Adoption of parasitic <i>Maculinea alcon</i> caterpillars (Lepidoptera: Lycaenidae) by three <i>Myrmica</i> ant species. <i>Animal Behaviour</i> , 2001, 62, 99-106.	0.8	45