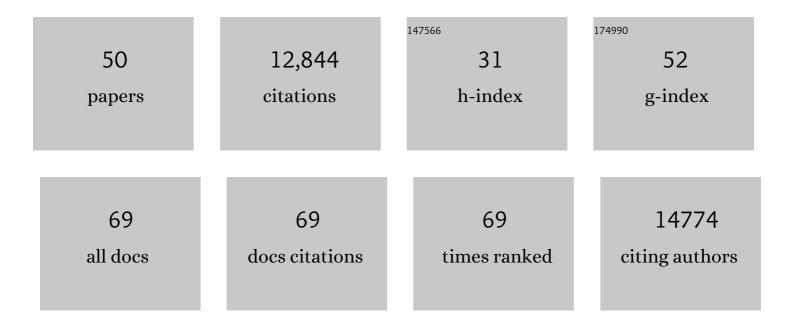
## Thomas D Als

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3814451/publications.pdf Version: 2024-02-01



ΤΗΟΜΛΟ Ο ΔΙΟ

#	Article	IF	CITATIONS
1	Discovery of the first genome-wide significant risk loci for attention deficit/hyperactivity disorder. Nature Genetics, 2019, 51, 63-75.	9.4	1,594
2	ldentification of common genetic risk variants for autism spectrum disorder. Nature Genetics, 2019, 51, 431-444.	9.4	1,538
3	Large-Scale Exome Sequencing Study Implicates Both Developmental and Functional Changes in the Neurobiology of Autism. Cell, 2020, 180, 568-584.e23.	13.5	1,422
4	Common schizophrenia alleles are enriched in mutation-intolerant genes and in regions under strong background selection. Nature Genetics, 2018, 50, 381-389.	9.4	1,332
5	Genome-wide association study identifies 30 loci associated with bipolar disorder. Nature Genetics, 2019, 51, 793-803.	9.4	1,191
6	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. Cell, 2019, 179, 1469-1482.e11.	13.5	935
7	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	13.7	929
8	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. Nature Genetics, 2021, 53, 817-829.	9.4	629
9	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. Cell, 2018, 173, 1705-1715.e16.	13.5	623
10	All roads lead to home: panmixia of European eel in the Sargasso Sea. Molecular Ecology, 2011, 20, 1333-1346.	2.0	176
11	A Mosaic of Chemical Coevolution in a Large Blue Butterfly. Science, 2008, 319, 88-90.	6.0	174
12	The evolution of alternative parasitic life histories in large blue butterflies. Nature, 2004, 432, 386-390.	13.7	163
13	Gene expression imputation across multiple brain regions provides insights into schizophrenia risk. Nature Genetics, 2019, 51, 659-674.	9.4	154
14	A Genetic Investigation of Sex Bias in the Prevalence of Attention-Deficit/Hyperactivity Disorder. Biological Psychiatry, 2018, 83, 1044-1053.	0.7	146
15	Genetic Variants Associated With Anxiety and Stress-Related Disorders. JAMA Psychiatry, 2019, 76, 924.	6.0	140
16	The Genetics of the Mood Disorder Spectrum: Genome-wide Association Analyses of More Than 185,000 Cases and 439,000 Controls. Biological Psychiatry, 2020, 88, 169-184.	0.7	137
17	Sequencing and de novo assembly of 150 genomes from Denmark as a population reference. Nature, 2017, 548, 87-91.	13.7	130
18	Genetic analyses identify widespread sex-differential participation bias. Nature Genetics, 2021, 53, 663-671.	9.4	124

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#	Article	IF	CITATIONS
19	Genome-wide association study implicates CHRNA2 in cannabis use disorder. Nature Neuroscience, 2019, 22, 1066-1074.	7.1	94
20	Qualitative assessment of the diet of European eel larvae in the Sargasso Sea resolved by DNA barcoding. Biology Letters, 2010, 6, 819-822.	1.0	90
21	Oceanic fronts in the Sargasso Sea control the early life and drift of Atlantic eels. Proceedings of the Royal Society B: Biological Sciences, 2010, 277, 3593-3599.	1.2	86
22	Microevolution in time and space: <scp>SNP</scp> analysis of historical <scp>DNA</scp> reveals dynamic signatures of selection in <scp>A</scp> tlantic cod. Molecular Ecology, 2013, 22, 2424-2440.	2.0	86
23	Highly discrepant proportions of female and male Scandinavian and British Isles ancestry within the isolated population of the Faroe Islands. European Journal of Human Genetics, 2006, 14, 497-504.	1.4	62
24	Geographical variation in host-ant specificity of the parasitic butterfly Maculinea alcon in Denmark. Ecological Entomology, 2002, 27, 403-414.	1.1	61
25	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. Biological Psychiatry, 2022, 91, 102-117.	0.7	61
26	Nationwide Genomic Study in Denmark Reveals Remarkable Population Homogeneity. Genetics, 2016, 204, 711-722.	1.2	54
27	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. Schizophrenia Bulletin, 2015, 41, 1294-1308.	2.3	48
28	Genetic liability to ADHD and substance use disorders in individuals with ADHD. Addiction, 2020, 115, 1368-1377.	1.7	47
29	Adoption of parasitic Maculinea alcon caterpillars (Lepidoptera: Lycaenidae) by three Myrmica ant species. Animal Behaviour, 2001, 62, 99-106.	0.8	45
30	A genome-wide study of panic disorder suggests the amiloride-sensitive cation channel 1 as a candidate gene. European Journal of Human Genetics, 2012, 20, 84-90.	1.4	45
31	The origin of the isolated population of the Faroe Islands investigated using Y chromosomal markers. Human Genetics, 2004, 115, 19-28.	1.8	43
32	A polygenic resilience score moderates the genetic risk for schizophrenia. Molecular Psychiatry, 2021, 26, 800-815.	4.1	36
33	Identifying the Common Genetic Basis of Antidepressant Response. Biological Psychiatry Global Open Science, 2022, 2, 115-126.	1.0	31
34	Risk variants and polygenic architecture of disruptive behavior disorders in the context of attention-deficit/hyperactivity disorder. Nature Communications, 2021, 12, 576.	5.8	28
35	Identification of genetic loci associated with nocturnal enuresis: a genome-wide association study. The Lancet Child and Adolescent Health, 2021, 5, 201-209.	2.7	27
36	Microsatellite markers for the large blue butterflies Maculinea nausithous and Maculinea alcon (Lepidoptera: Lycaenidae) and their amplification in other Maculinea species. Molecular Ecology Notes, 2005, 5, 165-168.	1.7	22

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#	Article	IF	CITATIONS
37	Polygenic risk score, psychosocial environment and the risk of attention-deficit/hyperactivity disorder. Translational Psychiatry, 2020, 10, 335.	2.4	22
38	Schizophrenia polygenic risk scores, urbanicity and treatment-resistant schizophrenia. Schizophrenia Research, 2019, 212, 79-85.	1.1	19
39	A Nationwide Cohort Study of Nonrandom Mating in Schizophrenia and Bipolar Disorder. Schizophrenia Bulletin, 2021, 47, 1342-1350.	2.3	17
40	Differential timing of gene expression regulation between leptocephali of the twoâ€, <i>Anguilla</i> â€,eel species in the Sargasso Sea. Ecology and Evolution, 2011, 1, 459-467.	0.8	15
41	Schizophrenia-associated mt-DNA SNPs exhibit highly variable haplogroup affiliation and nuclear ancestry: Bi-genomic dependence raises major concerns for link to disease. PLoS ONE, 2018, 13, e0208828.	1.1	15
42	MBL and MASP-2 concentrations in serum andMBL2promoter polymorphisms are associated to schizophrenia. Acta Neuropsychiatrica, 2012, 24, 199-207.	1.0	11
43	Polygenic liability, stressful life events and risk for secondary-treated depression in early life: a nationwide register-based case-cohort study. Psychological Medicine, 2023, 53, 217-226.	2.7	7
44	Polygenic Heterogeneity Across Obsessive-Compulsive Disorder Subgroups Defined by a Comorbid Diagnosis. Frontiers in Genetics, 2021, 12, 711624.	1.1	7
45	Genetics of panic disorder on the Faroe Islands: a replication study of chromosome 9 and panic disorder. Psychiatric Genetics, 2006, 16, 99-104.	0.6	6
46	Polygenic Liability and Recurrence of Depression in Patients With First-Onset Depression Treated in Hospital-Based Settings. JAMA Psychiatry, 2021, 78, 792.	6.0	6
47	Support for a bipolar affective disorder susceptibility locus on chromosome 12q24.3. Psychiatric Genetics, 2010, 20, 93-101.	0.6	6
48	Complex spatio-temporal distribution and genomic ancestry of mitochondrial DNA haplogroups in 24,216 Danes. PLoS ONE, 2018, 13, e0208829.	1.1	5
49	Association of the leucine-7 to proline-7 variation in the signal sequence of neuropeptide Y with major depression. Acta Neuropsychiatrica, 2012, 24, 81-90.	1.0	4
50	Genome-wide by Environment Interaction Study of Stressful Life Events and Hospital-Treated Depression in the iPSYCH2012 Sample. Biological Psychiatry Global Open Science, 2022, 2, 400-410.	1.0	2