

Dimitrios Avramopoulos

List of Publications by Year in descending order

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Version: 2024-02-01

76
papers

5,888
citations

136950

32
h-index

91884

69
g-index

83
all docs

83
docs citations

83
times ranked

11965
citing authors

#	ARTICLE	IF	CITATIONS
1	CRISPR Del/Rei: a simple, flexible, and efficient pipeline for scarless genome editing. <i>Scientific Reports</i> , 2022, 12, .	3.3	1
2	Introduction to epigenetics in psychiatry. , 2021, , 3-24.		1
3	Transcriptome analysis of human induced excitatory neurons supports a strong effect of clozapine on cholesterol biosynthesis. <i>Schizophrenia Research</i> , 2021, 228, 324-326.	2.0	13
4	Schizophrenia risk alleles often affect the expression of many genes and each gene may have a different effect on the risk: A mediation analysis. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2021, 186, 251-258.	1.7	3
5	Transcriptomic data of Clozapine-treated Ngn2-induced Human Excitatory Neurons. <i>Data in Brief</i> , 2021, 35, 106897.	1.0	4
6	Identifying nootropic drug targets via large-scale cognitive GWAS and transcriptomics. <i>Neuropsychopharmacology</i> , 2021, 46, 1788-1801.	5.4	12
7	Polygenic risk scores differentiate schizophrenia patients with toxoplasma gondii compared to toxoplasma seronegative patients. <i>Comprehensive Psychiatry</i> , 2021, 107, 152236.	3.1	5
8	Molecular imaging of beta-amyloid deposition in late-life depression. <i>Neurobiology of Aging</i> , 2021, 101, 85-93.	3.1	15
9	Improving the Utility of Polygenic Risk Scores as a Biomarker for Alzheimer's Disease. <i>Cells</i> , 2021, 10, 1627.	4.1	7
10	Positron emission tomography imaging of serotonin degeneration and beta-amyloid deposition in late-life depression evaluated with multi-modal partial least squares. <i>Translational Psychiatry</i> , 2021, 11, 473.	4.8	18
11	Gene-Environment Interactions in Schizophrenia: A Literature Review. <i>Genes</i> , 2021, 12, 1850.	2.4	40
12	Developing Treatments for Alzheimer's and Related Disorders with Precision Medicine: A Vision. <i>Advances in Experimental Medicine and Biology</i> , 2021, 1339, 395-402.	1.6	2
13	A screen of 1,049 schizophrenia and 30 Alzheimer's-associated variants for regulatory potential. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2020, 183, 61-73.	1.7	31
14	S184. IN SILICO PREDICTION OF T-CELL-MEDIATED MOLECULAR MIMICRY IN TOXOPLASMOSIS AND SCHIZOPHRENIA. <i>Schizophrenia Bulletin</i> , 2020, 46, S108-S108.	4.3	0
15	Allele-specific open chromatin in human iPSC neurons elucidates functional disease variants. <i>Science</i> , 2020, 369, 561-565.	12.6	77
16	Publicly Available hiPSC Lines with Extreme Polygenic Risk Scores for Modeling Schizophrenia. <i>Complex Psychiatry</i> , 2020, 6, 68-82.	0.9	18
17	Modeling Psychiatric Disorder Biology with Stem Cells. <i>Current Psychiatry Reports</i> , 2020, 22, 24.	4.5	25
18	Pleiotropic Meta-Analysis of Cognition, Education, and Schizophrenia Differentiates Roles of Early Neurodevelopmental and Adult Synaptic Pathways. <i>American Journal of Human Genetics</i> , 2019, 105, 334-350.	6.2	86

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19	Linear models enable powerful differential activity analysis in massively parallel reporter assays. <i>BMC Genomics</i> , 2019, 20, 209.	2.8	322
20	Escitalopram for agitation in Alzheimer's disease (Sâ€CitAD): Methods and design of an investigatorâ€initiated, randomized, controlled, multicenter clinical trial. <i>Alzheimer's and Dementia</i> , 2019, 15, 1427-1436.	0.8	28
21	Preliminary Report on the Feasibility and Efficacy of the Modified Atkins Diet for Treatment of Mild Cognitive Impairment and Early Alzheimerâ€™s Disease. <i>Journal of Alzheimer's Disease</i> , 2019, 68, 969-981.	2.6	66
22	Screening Human Embryos for Polygenic Traits Has Limited Utility. <i>Cell</i> , 2019, 179, 1424-1435.e8.	28.9	78
23	Affected Sib-Pair Analyses Identify Signaling Networks Associated With Social Behavioral Deficits in Autism. <i>Frontiers in Genetics</i> , 2019, 10, 1186.	2.3	2
24	Insights into genetics, human biology and disease gleaned from family based genomic studies. <i>Genetics in Medicine</i> , 2019, 21, 798-812.	2.4	161
25	Genome-wide association study in two populations to determine genetic variants associated with <i>Toxoplasma gondii</i> infection and relationship to schizophrenia risk. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2019, 92, 133-147.	4.8	26
26	Familial monophasic acute transverse myelitis due to the pathogenic variant in <i>VPS37A</i> . <i>Neurology: Genetics</i> , 2018, 4, e213.	1.9	4
27	HLA typing using genome wide data reveals susceptibility types for infections in a psychiatric disease enriched sample. <i>Brain, Behavior, and Immunity</i> , 2018, 70, 203-213.	4.1	10
28	Stress-Dependent Association Between Polygenic Risk for Schizophrenia and Schizotypal Traits in Young Army Recruits. <i>Schizophrenia Bulletin</i> , 2018, 44, 338-347.	4.3	33
29	Neuregulin 3 and its roles in schizophrenia risk and presentation. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018, 177, 257-266.	1.7	28
30	Study of 300,486 individuals identifies 148 independent genetic loci influencing general cognitive function. <i>Nature Communications</i> , 2018, 9, 2098.	12.8	484
31	Recent Advances in the Genetics of Schizophrenia. <i>Molecular Neuropsychiatry</i> , 2018, 4, 35-51.	2.9	81
32	Genome-wide association meta-analysis in 269,867 individuals identifies new genetic and functional links to intelligence. <i>Nature Genetics</i> , 2018, 50, 912-919.	21.4	893
33	Multi-Trait Analysis of GWAS and Biological Insights Into Cognition: A Response to Hill (2018). <i>Twin Research and Human Genetics</i> , 2018, 21, 394-397.	0.6	3
34	Structural variants caused by <i>Alu</i> insertions are associated with risks for many human diseases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E3984-E3992.	7.1	113
35	Thorase variants are associated with defects in glutamatergic neurotransmission that can be rescued by Perampanel. <i>Science Translational Medicine</i> , 2017, 9, .	12.4	20
36	Large-Scale Cognitive GWAS Meta-Analysis Reveals Tissue-Specific Neural Expression and Potential Nootropic Drug Targets. <i>Cell Reports</i> , 2017, 21, 2597-2613.	6.4	103

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37	Neuregulin 3 Knockout Mice Exhibit Behaviors Consistent with Psychotic Disorders. <i>Molecular Neuropsychiatry</i> , 2016, 2, 79-87.	2.9	27
38	New targets for monitoring and therapy in Barth syndrome. <i>Genetics in Medicine</i> , 2016, 18, 1001-1010.	2.4	32
39	A population pharmacokinetic model for R- and S-citalopram and desmethylcitalopram in Alzheimer's disease patients with agitation. <i>Journal of Pharmacokinetics and Pharmacodynamics</i> , 2016, 43, 99-109.	1.8	14
40	Citalopram for the Treatment of Agitation in Alzheimer Dementia. <i>Journal of Geriatric Psychiatry and Neurology</i> , 2016, 29, 59-64.	2.3	8
41	Functional Characterization of Schizophrenia-Associated Variation in CACNA1C. <i>PLoS ONE</i> , 2016, 11, e0157086.	2.5	61
42	Common genetic variation and schizophrenia polygenic risk influence neurocognitive performance in young adulthood. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2015, 168, 392-401.	1.7	52
43	Identification and Functional Studies of Regulatory Variants Responsible for the Association of <i>NRG3</i> with a Delusion Phenotype in Schizophrenia. <i>Molecular Neuropsychiatry</i> , 2015, 1, 36-46.	2.9	14
44	Genome-wide association study of schizophrenia in Ashkenazi Jews. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2015, 168, 649-659.	1.7	203
45	Functional Variants in <i>DPYSL2</i> Sequence Increase Risk of Schizophrenia and Suggest a Link to mTOR Signaling. <i>G3: Genes, Genomes, Genetics</i> , 2015, 5, 61-72.	1.8	39
46	Assessment of incidental findings in 232 whole-exome sequences from the Baylor-Hopkins Center for Mendelian Genomics. <i>Genetics in Medicine</i> , 2015, 17, 782-788.	2.4	41
47	The Genetic Basis of Mendelian Phenotypes: Discoveries, Challenges, and Opportunities. <i>American Journal of Human Genetics</i> , 2015, 97, 199-215.	6.2	574
48	Infection and Inflammation in Schizophrenia and Bipolar Disorder: A Genome Wide Study for Interactions with Genetic Variation. <i>PLoS ONE</i> , 2015, 10, e0116696.	2.5	92
49	Exome Sequencing in 53 Sporadic Cases of Schizophrenia Identifies 18 Putative Candidate Genes. <i>PLoS ONE</i> , 2014, 9, e112745.	2.5	79
50	Bipolar disorder ANK3 risk variant effect on sustained attention is replicated in a large healthy population. <i>Psychiatric Genetics</i> , 2012, 22, 210-213.	1.1	18
51	Gene expression reveals overlap between normal aging and Alzheimer's disease genes. <i>Neurobiology of Aging</i> , 2011, 32, 2319.e27-2319.e34.	3.1	45
52	Linkage and association on 8p21.2-p21.1 in schizophrenia. , 2011, 156, 188-197.		26
53	Schizophrenia-Related Neuregulin-1 Single-Nucleotide Polymorphisms Lead to Deficient Smooth Eye Pursuit in a Large Sample of Young Men. <i>Schizophrenia Bulletin</i> , 2011, 37, 822-831.	4.3	16
54	Fine mapping of the chromosome 10q11-q21 linkage region in Alzheimer's disease cases and controls. <i>Neurogenetics</i> , 2010, 11, 335-348.	1.4	17

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55	Genetics of Psychiatric Disorders Methods: Molecular Approaches. Clinics in Laboratory Medicine, 2010, 30, 815-827.	1.4	2
56	Genetics of Psychiatric Disorders Methods: Molecular Approaches. Psychiatric Clinics of North America, 2010, 33, 1-13.	1.3	14
57	Neuroglobin and Alzheimer's dementia: Genetic association and gene expression changes. Neurobiology of Aging, 2010, 31, 1835-1842.	3.1	51
58	Familiality of Novel Factorial Dimensions of Schizophrenia. Archives of General Psychiatry, 2009, 66, 591.	12.3	71
59	Fine Mapping on Chromosome 10q22-q23 Implicates Neuregulin 3 in Schizophrenia. American Journal of Human Genetics, 2009, 84, 21-34.	6.2	81
60	Genetics of Alzheimer's disease: recent advances. Genome Medicine, 2009, 1, 34.	8.2	112
61	Genome-wide linkage analysis of 723 affected relative pairs with late-onset Alzheimer's disease. Human Molecular Genetics, 2007, 16, 2703-2712.	2.9	52
62	Stage II follow-up on a linkage scan for bipolar disorder in the Ashkenazim provides suggestive evidence for chromosome 12p and the GRIN2B gene. Genetics in Medicine, 2007, 9, 745-751.	2.4	31
63	Recurrent 10q22-q23 Deletions: A Genomic Disorder on 10q Associated with Cognitive and Behavioral Abnormalities. American Journal of Human Genetics, 2007, 80, 938-947.	6.2	101
64	Impact of Schizophrenia Candidate Genes on Schizotypy and Cognitive Endophenotypes at the Population Level. Biological Psychiatry, 2007, 62, 784-792.	1.3	124
65	Effect of Schizotypy on Cognitive Performance and Its Tuning by COMT val158 Met Genotype Variations in a Large Population of Young Men. Biological Psychiatry, 2007, 61, 845-853.	1.3	66
66	A novel gene derived from a segmental duplication shows perturbed expression in Alzheimer's disease. Neurogenetics, 2007, 8, 111-120.	1.4	8
67	Testing groups of genomic locations for enrichment in disease loci using linkage scan data: A method for hypothesis testing. Human Genomics, 2006, 2, 345-52.	2.9	6
68	Further evidence of a maternal parent-of-origin effect on chromosome 10 in late-onset Alzheimer's disease. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2006, 141B, 537-540.	1.7	26
69	Linkage to chromosome 14q in Alzheimer's disease (AD) patients without psychotic symptoms. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2005, 132B, 9-13.	1.7	26
70	Brain activation in offspring of AD cases corresponds to 10q linkage. Annals of Neurology, 2005, 58, 142-146.	5.3	7
71	Bipolar I Disorder and Schizophrenia: A Single-Nucleotide Polymorphism Screen of 64 Candidate Genes among Ashkenazi Jewish Case-Parent Trios. American Journal of Human Genetics, 2005, 77, 918-936.	6.2	358
72	Results of a high-resolution genome screen of 437 Alzheimer's Disease families. Human Molecular Genetics, 2003, 12, 23-32.	2.9	304

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73	Evidence for parent of origin effect in late-onset Alzheimer disease. American Journal of Medical Genetics Part A, 2002, 114, 679-686.	2.4	61
74	Homologous loci DXYS156X and DXYS156Y contain a polymorphic pentanucleotide repeat (TAAAA)n and map to human X and Y chromosomes. Human Mutation, 1994, 4, 208-211.	2.5	42
75	Linkage mapping of the cystathionine γ -synthase (CBS) gene on human chromosome 21 using a DNA polymorphism in the 3' untranslated region. Human Genetics, 1993, 90, 566-8.	3.8	19
76	Mitotic errors in somatic cells cause trisomy 21 in about 4.5% of cases and are not associated with advanced maternal age. Nature Genetics, 1993, 3, 146-150.	21.4	138