Stephen J Glatt

List of Publications by Year
in descending order

52	13,385	230014	206121
papers	citations	27 h-index	51 g-index

59	59	59	36380
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. Biological Psychiatry, 2022, 91, 313-327.	0.7	114
2	Bipolar Disorder, Religion, and Spirituality: A Scoping Review. Journal of Religion and Health, 2022, 61, 3589-3614.	0.8	6
3	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	13.7	929
4	Rare coding variants in ten genes confer substantial risk for schizophrenia. Nature, 2022, 604, 509-516.	13.7	326
5	A polygenic resilience score moderates the genetic risk for schizophrenia. Molecular Psychiatry, 2021, 26, 800-815.	4.1	36
6	Schizophrenia, autism spectrum disorders and developmental disorders share specific disruptive coding mutations. Nature Communications, 2021, 12, 5353.	5.8	44
7	Autophagy, apoptosis, and neurodevelopmental genes might underlie selective brain region vulnerability in attention-deficit/hyperactivity disorder. Molecular Psychiatry, 2021, 26, 6643-6654.	4.1	19
8	Transcriptomic abnormalities in peripheral blood in bipolar disorder, and discrimination of the major psychoses. Schizophrenia Research, 2020, 217, 124-135.	1.1	18
9	Exome sequencing in schizophrenia-affected parentâ€“offspring trios reveals risk conferred by protein-coding de novo mutations. Nature Neuroscience, 2020, 23, 185-193.	7.1	125
10	Methodology in the GBD study of China. Lancet, The, 2020, 396, 25.	6.3	2
11	Gene co-expression networks in peripheral blood capture dimensional measures of emotional and behavioral problems from the Child Behavior Checklist (CBCL). Translational Psychiatry, 2020, 10, 328.	2.4	2
12	Effects of Polygenic Risk and Perceived Friendsâ€™ Drinking and Disruptive Behavior on Development of Alcohol Use Across Adolescence. Journal of Studies on Alcohol and Drugs, 2020, 81, 808-815.	0.6	2
13	An integrated analysis of genes and functional pathways for aggression in human and rodent models. Molecular Psychiatry, 2019, 24, 1655-1667.	4.1	61
14	Effort valuation and psychopathology in children and adults. Psychological Medicine, 2019, 49, 2801-2807.	2.7	5
15	Advanced Paternal Age and Early Onset of Schizophrenia in Sporadic Cases: Not Confounded by Parental Polygenic Risk for Schizophrenia. Biological Psychiatry, 2019, 86, 56-64.	0.7	18
16	Comparative genetic architectures of schizophrenia in East Asian and European populations. Nature Genetics, 2019, 51, 1670-1678.	9.4	440
17	Psychiatric genetics and the structure of psychopathology. Molecular Psychiatry, 2019, 24, 409-420.	4.1	281
18	Nu Support Vector Machine in Prediction of Fluid Intelligence Using MRI Data. Lecture Notes in Computer Science, 2019, , 92-98.	1.0	3

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19	Psychiatric Genetics, Epigenetics, and Cellular Models in Coming Years. Journal of Psychiatry and Brain Science, 2019, 4, .	0.3	1
20	PTSD Blood Transcriptome Mega-Analysis: Shared Inflammatory Pathways across Biological Sex and Modes of Trauma. Neuropsychopharmacology, 2018, 43, 469-481.	2.8	92
21	Genetic correlations among psychiatric and immuneâ€related phenotypes based on genomeâ€wide association data. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 641-657.	1.1	158
22	Predictors of current functioning and functional decline in schizophrenia. Schizophrenia Research, 2017, 188, 158-164.	1.1	37
23	The interaction between the dopamine receptor D4 (DRD4) variable number tandem repeat polymorphism and perceived peer drinking norms in adolescent alcohol use and misuse. Development and Psychopathology, 2017, 29, 173-183.	1.4	5
24	Optimizing the chances of success in the search for epigenetic biomarkers: Embracing genetic variation. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2017, 174, 589-594.	1.1	2
25	Adaptive combination of Bayes factors as a powerful method for the joint analysis of rare and common variants. Scientific Reports, 2017, 7, 13858.	1.6	4
26	Machine-learning classification of 22q11.2 deletion syndrome: A diffusion tensor imaging study. NeuroImage: Clinical, 2017, 15, 832-842.	1.4	22
27	Blood transcriptomic comparison of individuals with and without autism spectrum disorder: A combinedâ€samples megaâ€analysis. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2017, 174, 181-201.	1.1	54
28	RNA sequencing of transformed lymphoblastoid cells from siblings discordant for autism spectrum disorders reveals transcriptomic and functional alterations: Evidence for sexâ€specific effects. Autism Research, 2017, 10, 439-455.	2.1	21
29	SLC9A9 Coâ€expression modules in autismâ€associated brain regions. Autism Research, 2017, 10, 414-429.	2.1	12
30	Interaction Between the Î¼â€Opioid Receptor Gene and the Number of Heavyâ€Drinking Peers on Alcohol Use. Alcoholism: Clinical and Experimental Research, 2017, 41, 2041-2050.	1.4	4
31	Evaluation of the reconsolidation of traumatic memories protocol for the treatment of PTSD: a randomized, wait-list-controlled trial. Journal of Military, Veteran and Family Health, 2017, 3, 21-33.	0.3	19
32	The potential of genetic and gene expression analysis in the diagnosis of neuropsychiatric disorders. Expert Review of Molecular Diagnostics, 2016, 16, 677-695.	1.5	5
33	Transcriptome-wide mega-analyses reveal joint dysregulation of immunologic genes and transcription regulators in brain and blood in schizophrenia. Schizophrenia Research, 2016, 176, 114-124.	1.1	74
34	Analysis of protein-coding genetic variation in 60,706 humans. Nature, 2016, 536, 285-291.	13.7	9,051
35	Minireview: Multiomic candidate biomarkers for clinical manifestations of sickle cell severity: Early steps to precision medicine. Experimental Biology and Medicine, 2016, 241, 772-781.	1.1	16
36	Association of the OPRM1 Variant rs1799971 (A118G) with Non-Specific Liability to Substance Dependence in a Collaborative de novo Meta-Analysis of European-Ancestry Cohorts. Behavior Genetics, 2016, 46, 151-169.	1.4	98

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37	The influence of genes on â€œpositive valence systemsâ€•constructs: A systematic review. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 92-110.	1.1	13
38	Assessment of Lifespan Functioning Attainment (ALFA) scale: A quantitative interview for self-reported current and functional decline in schizophrenia. Journal of Psychiatric Research, 2015, 65, 102-107.	1.5	5
39	Blood-based gene-expression biomarkers of post-traumatic stress disorder among deployed marines: A pilot study. Psychoneuroendocrinology, 2015, 51, 472-494.	1.3	54
40	Meta-analysis of data from the Psychiatric Genomics Consortium and additional samples supports association of CACNA1C with risk for schizophrenia. Schizophrenia Research, 2015, 168, 429-433.	1.1	19
41	Genetic liability, prenatal health, stress and family environment: Risk factors in the Harvard Adolescent Family High Risk for Schizophrenia Study. Schizophrenia Research, 2014, 157, 142-148.	1.1	42
42	On the outside, looking in: A review and evaluation of the comparability of blood and brain â€œâ€•omesâ€•. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2013, 162, 595-603.	1.1	208
43	Bloodâ€•based geneâ€•expression predictors of PTSD risk and resilience among deployed marines: A pilot study. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2013, 162, 313-326.	1.1	63
44	Blood-Based Gene Expression Signatures of Infants and Toddlers With Autism. Journal of the American Academy of Child and Adolescent Psychiatry, 2012, 51, 934-944.e2.	0.3	98
45	Dysfunctional gene splicing as a potential contributor to neuropsychiatric disorders. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 382-392.	1.1	57
46	Similarities and differences in peripheral blood geneâ€•expression signatures of individuals with schizophrenia and their firstâ€•degree biological relatives. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 869-887.	1.1	53
47	Preliminary evidence of ubiquitin proteasome system dysregulation in schizophrenia and bipolar disorder: Convergent pathway analysis findings from two independent samples. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 494-502.	1.1	97
48	Genome-wide linkage analysis of heroin dependence in Han Chinese: Results from Wave Two of a multi-stage study. Drug and Alcohol Dependence, 2008, 98, 30-34.	1.6	17
49	Evaluation of OPRM1 variants in heroin dependence by family-based association testing and meta-analysis. Drug and Alcohol Dependence, 2007, 90, 159-165.	1.6	54
50	Successful Aging: From Phenotype to Genotype. Biological Psychiatry, 2007, 62, 282-293.	0.7	81
51	Meta-analyses of ALDH2 and ADH1B with alcohol dependence in Asians.. Psychological Bulletin, 2006, 132, 607-621.	5.5	203
52	Assessing the validity of blood-based gene expression profiles for the classification of schizophrenia and bipolar disorder: A preliminary report. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2005, 133B, 1-5.	1.1	205