

Kristen Brennand

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/7656864/kristen-brennand-publications-by-year.pdf>

Version: 2024-04-25

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

132
papers

8,530
citations

46
h-index

91
g-index

182
ext. papers

10,921
ext. citations

10.8
avg, IF

6.28
L-index

#	Paper	IF	Citations
132	A bidirectional competitive interaction between circHomer1 and Homer1b within the orbitofrontal cortex regulates reversal learning.. <i>Cell Reports</i> , 2022 , 38, 110282	10.6	2
131	Quickly moving too slowly: Interneuron migration in Timothy Syndrome.. <i>Cell Stem Cell</i> , 2022 , 29, 181-188		
130	Chromatin profiling in human neurons reveals aberrant roles for histone acetylation and BET family proteins in schizophrenia.. <i>Nature Communications</i> , 2022 , 13, 2195	17.4	3
129	Using Stem Cell Models to Explore the Genetics Underlying Psychiatric Disorders: Linking Risk Variants, Genes, and Biology in Brain Disease.. <i>American Journal of Psychiatry</i> , 2022 , 179, 322-328	11.9	0
128	Publicly Available hiPSC Lines with Extreme Polygenic Risk Scores for Modeling Schizophrenia. <i>Complex Psychiatry</i> , 2021 , 6, 68-82	2.3	3
127	Common Genetic Variation in Humans Impacts In Vitro Susceptibility to SARS-CoV-2 Infection. <i>Stem Cell Reports</i> , 2021 , 16, 505-518	8	22
126	Circadian rhythms in bipolar disorder patient-derived neurons predict lithium response: preliminary studies. <i>Molecular Psychiatry</i> , 2021 , 26, 3383-3394	15.1	7
125	Xenopus models suggest convergence of gene signatures on neurogenesis in autism. <i>Neuron</i> , 2021 , 109, 743-745	13.9	0
124	Haploinsufficiency of POU4F1 causes an ataxia syndrome with hypotonia and intention tremor. <i>Human Mutation</i> , 2021 , 42, 685-693	4.7	
123	Fitness selection of hyperfusogenic measles virus F proteins associated with neuropathogenic phenotypes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	6
122	Using the dCas9-KRAB system to repress gene expression in hiPSC-derived neurons. <i>STAR Protocols</i> , 2021 , 2, 100580	1.4	0
121	IL10RB as a key regulator of COVID-19 host susceptibility and severity 2021 ,		2
120	Parsing the Functional Impact of Noncoding Genetic Variants in the Brain Epigenome. <i>Biological Psychiatry</i> , 2021 , 89, 65-75	7.9	5
119	Transformative Network Modeling of Multi-omics Data Reveals Detailed Circuits, Key Regulators, and Potential Therapeutics for Alzheimer's Disease. <i>Neuron</i> , 2021 , 109, 257-272.e14	13.9	29
118	Functional genomics of psychiatric disease risk using genome engineering 2021 , 711-734		
117	Molecular subtyping of Alzheimer's disease using RNA sequencing data reveals novel mechanisms and targets. <i>Science Advances</i> , 2021 , 7,	14.3	39
116	Applying stem cells and CRISPR engineering to uncover the etiology of schizophrenia. <i>Current Opinion in Neurobiology</i> , 2021 , 69, 193-201	7.6	6

115	Induction of dopaminergic neurons for neuronal subtype-specific modeling of psychiatric disease risk. <i>Molecular Psychiatry</i> , 2021 ,	15.1	2
114	Induced Pluripotent Stem Cells in Psychiatry: An Overview and Critical Perspective. <i>Biological Psychiatry</i> , 2021 , 90, 362-372	7.9	2
113	Analysis framework and experimental design for evaluating synergy-driving gene expression. <i>Nature Protocols</i> , 2021 , 16, 812-840	18.8	2
112	Functional annotation of rare structural variation in the human brain. <i>Nature Communications</i> , 2020 , 11, 2990	17.4	18
111	Transcriptional signatures of participant-derived neural progenitor cells and neurons implicate altered Wnt signaling in Phelan-McDermid syndrome and autism. <i>Molecular Autism</i> , 2020 , 11, 53	6.5	11
110	A computational tool (H-MAGMA) for improved prediction of brain-disorder risk genes by incorporating brain chromatin interaction profiles. <i>Nature Neuroscience</i> , 2020 , 23, 583-593	25.5	81
109	Modeling the complex genetic architectures of brain disease. <i>Nature Genetics</i> , 2020 , 52, 363-369	36.3	21
108	Cell Type-Specific In Vitro Gene Expression Profiling of Stem Cell-Derived Neural Models. <i>Cells</i> , 2020 , 9,	7.9	2
107	ASCL1- and DLX2-induced GABAergic neurons from hiPSC-derived NPCs. <i>Journal of Neuroscience Methods</i> , 2020 , 334, 108548	3	17
106	Integrating CRISPR Engineering and hiPSC-Derived 2D Disease Modeling Systems. <i>Journal of Neuroscience</i> , 2020 , 40, 1176-1185	6.6	9
105	A psychiatric disease-related circular RNA controls synaptic gene expression and cognition. <i>Molecular Psychiatry</i> , 2020 , 25, 2712-2727	15.1	44
104	If there is not one cure for schizophrenia, there may be many. <i>NPJ Schizophrenia</i> , 2020 , 6, 11	5.5	
103	Investigation of Schizophrenia with Human Induced Pluripotent Stem Cells. <i>Advances in Neurobiology</i> , 2020 , 25, 155-206	2.1	3
102	Common genetic variation in humans impacts susceptibility to SARS-CoV-2 infection 2020 ,		4
101	T9. EPIGENETIC PROFILING IN SCHIZOPHRENIA DERIVED HUMAN INDUCED PLURIPOTENT STEM CELLS (HIPSCS) AND NEURONS. <i>Schizophrenia Bulletin</i> , 2020 , 46, S234-S234	1.3	78
100	Massively parallel techniques for cataloguing the regulome of the human brain. <i>Nature Neuroscience</i> , 2020 , 23, 1509-1521	25.5	18
99	Integration of CRISPR-engineering and hiPSC-based models of psychiatric genomics. <i>Molecular and Cellular Neurosciences</i> , 2020 , 107, 103532	4.8	3
98	CRISPR-based functional evaluation of schizophrenia risk variants. <i>Schizophrenia Research</i> , 2020 , 217, 26-36	3.6	4

97	Sex-Specific Role for the Long Non-coding RNA LINC00473 in Depression. <i>Neuron</i> , 2020 , 106, 912-926.e513,9	46
96	Global landscape and genetic regulation of RNA editing in cortical samples from individuals with schizophrenia. <i>Nature Neuroscience</i> , 2019 , 22, 1402-1412	25.5 32
95	Differential transcriptional response following glucocorticoid activation in cultured blood immune cells: a novel approach to PTSD biomarker development. <i>Translational Psychiatry</i> , 2019 , 9, 201	8.6 12
94	Synergistic effects of common schizophrenia risk variants. <i>Nature Genetics</i> , 2019 , 51, 1475-1485	36.3 90
93	Spatial genome exploration in the context of cognitive and neurological disease. <i>Current Opinion in Neurobiology</i> , 2019 , 59, 112-119	7.6 10
92	9. MODELLING THE IMPACT OF RARE AND COMMON VARIANTS IN SCHIZOPHRENIA USING STEM CELLS. <i>Schizophrenia Bulletin</i> , 2019 , 45, S101-S101	1.3 78
91	Leveraging Human Induced Pluripotent Stem Cell-Based Models Provides Biological Context to Genome-wide Association Study Findings. <i>Biological Psychiatry</i> , 2019 , 85, 532-533	7.9 1
90	Entrainment of Circadian Rhythms to Temperature Reveals Amplitude Deficits in Fibroblasts from Patients with Bipolar Disorder and Possible Links to Calcium Channels. <i>Molecular Neuropsychiatry</i> , 2019 , 5, 115-124	4.9 4
89	Examining the relationship between astrocyte dysfunction and neurodegeneration in ALS using hiPSCs. <i>Neurobiology of Disease</i> , 2019 , 132, 104562	7.5 13
88	CRISPR-based functional evaluation of common SZ risk variants. <i>FASEB Journal</i> , 2019 , 33, 205.2	0.9
87	Neuronal impact of patient-specific aberrant NRXN1 splicing. <i>Nature Genetics</i> , 2019 , 51, 1679-1690	36.3 41
86	Type I interferon response impairs differentiation potential of pluripotent stem cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 1384-1393	11.5 25
85	Chronotype and cellular circadian rhythms predict the clinical response to lithium maintenance treatment in patients with bipolar disorder. <i>Neuropsychopharmacology</i> , 2019 , 44, 620-628	8.7 43
84	New considerations for hiPSC-based models of neuropsychiatric disorders. <i>Molecular Psychiatry</i> , 2019 , 24, 49-66	15.1 45
83	Mapping regulatory variants in hiPSC models. <i>Nature Genetics</i> , 2018 , 50, 1-2	36.3 21
82	Marker chromosome genomic structure and temporal origin implicate a chromoanasythesis event in a family with pleiotropic psychiatric phenotypes. <i>Human Mutation</i> , 2018 , 39, 939-946	4.7 16
81	THC exposure of human iPSC neurons impacts genes associated with neuropsychiatric disorders. <i>Translational Psychiatry</i> , 2018 , 8, 89	8.6 21
80	Inhibition of STEP ameliorates deficits in mouse and hiPSC-based schizophrenia models. <i>Molecular Psychiatry</i> , 2018 , 23, 271-281	15.1 28

79	MEF2C transcription factor is associated with the genetic and epigenetic risk architecture of schizophrenia and improves cognition in mice. <i>Molecular Psychiatry</i> , 2018 , 23, 123-132	15.1	46
78	Modeling the Brain in the Culture Dish: Advancements and Applications of Induced Pluripotent Stem-Cell-Derived Neurons 2018 , 119-157		
77	Modeling Neuropsychiatric and Neurodegenerative Diseases With Induced Pluripotent Stem Cells. <i>Frontiers in Pediatrics</i> , 2018 , 6, 82	3.4	13
76	GJA1 (connexin43) is a key regulator of Alzheimer's disease pathogenesis. <i>Acta Neuropathologica Communications</i> , 2018 , 6, 144	7.3	37
75	Neuron-specific signatures in the chromosomal connectome associated with schizophrenia risk. <i>Science</i> , 2018 , 362,	33.3	91
74	20. THE APPLICATION OF STEM CELL MODELS TO VALIDATE RARE AND COMMON VARIANTS CONTRIBUTING TO SCHIZOPHRENIA. <i>Schizophrenia Bulletin</i> , 2018 , 44, S32-S33	1.3	78
73	20.4 MODELING THE CONTRIBUTION OF COMMON VARIANTS TO SCHIZOPHRENIA RISK. <i>Schizophrenia Bulletin</i> , 2018 , 44, S34-S34	1.3	78
72	Expression-based drug screening of neural progenitor cells from individuals with schizophrenia. <i>Nature Communications</i> , 2018 , 9, 4412	17.4	39
71	Landscape of Conditional eQTL in Dorsolateral Prefrontal Cortex and Co-localization with Schizophrenia GWAS. <i>American Journal of Human Genetics</i> , 2018 , 102, 1169-1184	11	73
70	Using hiPSCs to model neuropsychiatric copy number variations (CNVs) has potential to reveal underlying disease mechanisms. <i>Brain Research</i> , 2017 , 1655, 283-293	3.7	12
69	Altered proliferation and networks in neural cells derived from idiopathic autistic individuals. <i>Molecular Psychiatry</i> , 2017 , 22, 820-835	15.1	224
68	Common developmental genome deprogramming in schizophrenia - Role of Integrative Nuclear FGFR1 Signaling (INFS). <i>Schizophrenia Research</i> , 2017 , 185, 17-32	3.6	37
67	Divergent Levels of Marker Chromosomes in an hiPSC-Based Model of Psychosis. <i>Stem Cell Reports</i> , 2017 , 8, 519-528	8	6
66	In utero exposure to maternal smoking is associated with DNA methylation alterations and reduced neuronal content in the developing fetal brain. <i>Epigenetics and Chromatin</i> , 2017 , 10, 4	5.8	55
65	Variations in brain defects result from cellular mosaicism in the activation of heat shock signalling. <i>Nature Communications</i> , 2017 , 8, 15157	17.4	14
64	Application of CRISPR/Cas9 to the study of brain development and neuropsychiatric disease. <i>Molecular and Cellular Neurosciences</i> , 2017 , 82, 157-166	4.8	19
63	THC Treatment Alters Glutamate Receptor Gene Expression in Human Stem Cell-Derived Neurons. <i>Molecular Neuropsychiatry</i> , 2017 , 3, 73-84	4.9	4
62	Patient-derived hiPSC neurons with heterozygous CNTNAP2 deletions display altered neuronal gene expression and network activity. <i>NPJ Schizophrenia</i> , 2017 , 3, 35	5.5	21

61	High-Content Screening in hPSC-Neural Progenitors Identifies Drug Candidates that Inhibit Zika Virus Infection in Fetal-like Organoids and Adult Brain. <i>Cell Stem Cell</i> , 2017 , 21, 274-283.e5	18	144
60	An Efficient Platform for Astrocyte Differentiation from Human Induced Pluripotent Stem Cells. <i>Stem Cell Reports</i> , 2017 , 9, 600-614	8	158
59	Evaluating Synthetic Activation and Repression of Neuropsychiatric-Related Genes in hiPSC-Derived NPCs, Neurons, and Astrocytes. <i>Stem Cell Reports</i> , 2017 , 9, 615-628	8	46
58	Cerebral organoids reveal early cortical maldevelopment in schizophrenia-computational anatomy and genomics, role of FGFR1. <i>Translational Psychiatry</i> , 2017 , 7, 6	8.6	90
57	The methyltransferase SETDB1 regulates a large neuron-specific topological chromatin domain. <i>Nature Genetics</i> , 2017 , 49, 1239-1250	36.3	88
56	Neural organoids for disease phenotyping, drug screening and developmental biology studies. <i>Neurochemistry International</i> , 2017 , 106, 85-93	4.4	29
55	Transcriptional signatures of schizophrenia in hiPSC-derived NPCs and neurons are concordant with post-mortem adult brains. <i>Nature Communications</i> , 2017 , 8, 2225	17.4	92
54	The Importance of Non-neuronal Cell Types in hiPSC-Based Disease Modeling and Drug Screening. <i>Frontiers in Cell and Developmental Biology</i> , 2017 , 5, 117	5.7	17
53	Prospects for Modeling Abnormal Neuronal Function in Schizophrenia Using Human Induced Pluripotent Stem Cells. <i>Frontiers in Cellular Neuroscience</i> , 2017 , 11, 360	6.1	15
52	Identification of small-molecule inhibitors of Zika virus infection and induced neural cell death via a drug repurposing screen. <i>Nature Medicine</i> , 2016 , 22, 1101-1107	50.5	458
51	Integrative network analysis of nineteen brain regions identifies molecular signatures and networks underlying selective regional vulnerability to Alzheimer's disease. <i>Genome Medicine</i> , 2016 , 8, 104	14.4	135
50	Dysregulation of miRNA-9 in a Subset of Schizophrenia Patient-Derived Neural Progenitor Cells. <i>Cell Reports</i> , 2016 , 15, 1024-1036	10.6	82
49	Rapid Ngn2-induction of excitatory neurons from hiPSC-derived neural progenitor cells. <i>Methods</i> , 2016 , 101, 113-24	4.6	71
48	Is Huntington's disease a neurodevelopmental disorder?. <i>Science Translational Medicine</i> , 2016 , 8, 320ec1-329ec1	3.7	19
47	hiPSC Models Relevant to Schizophrenia. <i>Handbook of Behavioral Neuroscience</i> , 2016 , 391-406	0.7	
46	P1-021: Exploring Cell Autonomous and Non-Cell Autonomous Effects of APOE Genotype in iPSC-Derived Astrocytes and Neurons 2016 , 12, P407-P407		
45	The Pharmacogenomics of Bipolar Disorder study (PGBD): identification of genes for lithium response in a prospective sample. <i>BMC Psychiatry</i> , 2016 , 16, 129	4.2	42
44	Gene expression elucidates functional impact of polygenic risk for schizophrenia. <i>Nature Neuroscience</i> , 2016 , 19, 1442-1453	25.5	622

43	Activity-Dependent Changes in Gene Expression in Schizophrenia Human-Induced Pluripotent Stem Cell Neurons. <i>JAMA Psychiatry</i> , 2016 , 73, 1180-1188	14.5	29
42	Spatial genome organization and cognition. <i>Nature Reviews Neuroscience</i> , 2016 , 17, 681-691	13.5	49
41	A guide to generating and using hiPSC derived NPCs for the study of neurological diseases. <i>Journal of Visualized Experiments</i> , 2015 , e52495	1.6	38
40	Noncoding RNAs and neurobehavioral mechanisms in psychiatric disease. <i>Molecular Psychiatry</i> , 2015 , 20, 677-684	15.1	55
39	Differential responses to lithium in hyperexcitable neurons from patients with bipolar disorder. <i>Nature</i> , 2015 , 527, 95-9	50.4	315
38	Increased abundance of translation machinery in stem cell-derived neural progenitor cells from four schizophrenia patients. <i>Translational Psychiatry</i> , 2015 , 5, e662	8.6	34
37	Altered WNT Signaling in Human Induced Pluripotent Stem Cell Neural Progenitor Cells Derived from Four Schizophrenia Patients. <i>Biological Psychiatry</i> , 2015 , 78, e29-34	7.9	62
36	Phenotypic differences in hiPSC NPCs derived from patients with schizophrenia. <i>Molecular Psychiatry</i> , 2015 , 20, 361-8	15.1	272
35	Characterization of molecular and cellular phenotypes associated with a heterozygous deletion using patient-derived hiPSC neural cells. <i>NPJ Schizophrenia</i> , 2015 , 1,	5.5	42
34	From "directed differentiation" to "neuronal induction": modeling neuropsychiatric disease. <i>Biomarker Insights</i> , 2015 , 10, 31-41	3.5	22
33	Creating Patient-Specific Neural Cells for the In Vitro Study of Brain Disorders. <i>Stem Cell Reports</i> , 2015 , 5, 933-945	8	63
32	Dopaminergic differentiation of schizophrenia hiPSCs. <i>Molecular Psychiatry</i> , 2015 , 20, 549-50	15.1	15
31	Modeling Hippocampal Neurogenesis Using Human Pluripotent Stem Cells. <i>Stem Cell Reports</i> , 2014 , 3, 217	8	2
30	A role for noncoding variation in schizophrenia. <i>Cell Reports</i> , 2014 , 9, 1417-29	10.6	174
29	Evolving toward a human-cell based and multiscale approach to drug discovery for CNS disorders. <i>Frontiers in Pharmacology</i> , 2014 , 5, 252	5.6	31
28	Human iPSC neurons display activity-dependent neurotransmitter secretion: aberrant catecholamine levels in schizophrenia neurons. <i>Stem Cell Reports</i> , 2014 , 3, 531-8	8	82
27	Modeling heterogeneous patients with a clinical diagnosis of schizophrenia with induced pluripotent stem cells. <i>Biological Psychiatry</i> , 2014 , 75, 936-44	7.9	46
26	Modeling hippocampal neurogenesis using human pluripotent stem cells. <i>Stem Cell Reports</i> , 2014 , 2, 295-310	8	196

25	Roles of heat shock factor 1 in neuronal response to fetal environmental risks and its relevance to brain disorders. <i>Neuron</i> , 2014 , 82, 560-72	13.9	80
24	Neural Induction of Embryonic Stem/Induced Pluripotent Stem Cells 2013 , 111-129		2
23	Mosaic copy number variation in human neurons. <i>Science</i> , 2013 , 342, 632-7	33.3	404
22	Inducing cellular aging: enabling neurodegeneration-in-a-dish. <i>Cell Stem Cell</i> , 2013 , 13, 635-6	18	9
21	Neural stem and progenitor cells in health and disease. <i>Wiley Interdisciplinary Reviews: Systems Biology and Medicine</i> , 2013 , 5, 701-15	6.6	18
20	Modeling schizophrenia using induced pluripotent stem cell-derived and fibroblast-induced neurons. <i>Schizophrenia Bulletin</i> , 2013 , 39, 4-10	1.3	21
19	Modeling psychiatric disorders at the cellular and network levels. <i>Molecular Psychiatry</i> , 2012 , 17, 1239-53	5.1	92
18	Modeling psychiatric disorders through reprogramming. <i>DMM Disease Models and Mechanisms</i> , 2012 , 5, 26-32	4.1	52
17	Concise review: the promise of human induced pluripotent stem cell-based studies of schizophrenia. <i>Stem Cells</i> , 2011 , 29, 1915-22	5.8	60
16	Rapid cellular turnover in adipose tissue. <i>PLoS ONE</i> , 2011 , 6, e17637	3.7	85
15	Modelling schizophrenia using human induced pluripotent stem cells. <i>Nature</i> , 2011 , 473, 221-5	50.4	1006
14	Induced pluripotent stem cells (iPSCs) and neurological disease modeling: progress and promises. <i>Human Molecular Genetics</i> , 2011 , 20, R109-15	5.6	142
13	Brief report: efficient generation of hematopoietic precursors and progenitors from human pluripotent stem cell lines. <i>Stem Cells</i> , 2011 , 29, 1158-64	5.8	60
12	Investigating synapse formation and function using human pluripotent stem cell-derived neurons. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 3005-10	11.5	113
11	High-efficient generation of induced pluripotent stem cells from human astrocytes. <i>PLoS ONE</i> , 2010 , 5, e15526	3.7	53
10	Slow and steady is the key to beta-cell replication. <i>Journal of Cellular and Molecular Medicine</i> , 2009 , 13, 472-87	5.6	25
9	Reprogramming of pancreatic beta cells into induced pluripotent stem cells. <i>Current Biology</i> , 2008 , 18, 890-4	6.3	343
8	Correction for Szotek et al., Normal ovarian surface epithelial label-retaining cells exhibit stem/progenitor cell characteristics. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008 , 105, 19024-19024	11.5	78

7	Normal ovarian surface epithelial label-retaining cells exhibit stem/progenitor cell characteristics. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008 , 105, 12469-73	11.5	109
6	All beta cells contribute equally to islet growth and maintenance. <i>PLoS Biology</i> , 2007 , 5, e163	9.7	168
5	Gene Expression Elucidates Functional Impact of Polygenic Risk for Schizophrenia		3
4	Transcriptional signatures of schizophrenia in hiPSC-derived NPCs and neurons are concordant with signatures from post mortem adult brains		2
3	Publicly available hiPSC lines with extreme polygenic risk scores for modeling schizophrenia		3
2	Functional annotation of rare structural variation in the human brain		3
1	Molecular Networks and Key Regulators of the Dysregulated Neuronal System in Alzheimer's Disease		1