Emanuel Schwarz

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2519919/publications.pdf

Version: 2024-02-01

60 papers

3,021 citations

279798 23 h-index 50 g-index

65 all docs

65 does citations

65 times ranked

6047 citing authors

#	Article	IF	CITATIONS
1	Computational Modeling of the n-Back Task in the ABCD Study: Associations of Drift Diffusion Model Parameters to Polygenic Scores of Mental Disorders and Cardiometabolic Diseases. Biological Psychiatry: Cognitive Neuroscience and Neuroimaging, 2023, 8, 290-299.	1.5	1
2	Brain structural correlates of upward social mobility in ethnic minority individuals. Social Psychiatry and Psychiatric Epidemiology, 2022, 57, 2037-2047.	3.1	1
3	A DEVELOPMENTAL PERSPECTIVE ON FACETS OF IMPULSIVITY AND BRAIN ACTIVITY CORRELATES FROM ADOLESCENCE TO ADULTHOOD. Biological Psychiatry: Cognitive Neuroscience and Neuroimaging, 2022,	1.5	2
4	Oxytocin receptor expression patterns in the human brain across development. Neuropsychopharmacology, 2022, 47, 1550-1560.	5.4	23
5	Advancing Psychiatric Biomarker Discovery Through Multimodal Machine Learning. Biological Psychiatry, 2022, 91, 524-525.	1.3	O
6	From mechanistic insight towards clinical implementation using normative modeling. Nature Computational Science, 2022, 2, 278-280.	8.0	1
7	Generative network models of altered structural brain connectivity in schizophrenia. NeuroImage, 2021, 225, 117510.	4.2	24
8	Multimodal imaging improves brain age prediction and reveals distinct abnormalities in patients with psychiatric and neurological disorders. Human Brain Mapping, 2021, 42, 1714-1726.	3.6	68
9	Identifying multimodal signatures underlying the somatic comorbidity of psychosis: the COMMITMENT roadmap. Molecular Psychiatry, 2021, 26, 722-724.	7.9	7
10	Hyper-Coordinated DNA Methylation is Altered in Schizophrenia and Associated with Brain Function. Schizophrenia Bulletin Open, 2021, 2, .	1.7	0
11	Brain network dynamics during working memory are modulated by dopamine and diminished in schizophrenia. Nature Communications, 2021, 12, 3478.	12.8	69
12	Differential resting-state patterns across networks are spatially associated with Comt and Trmt2a gene expression patterns in a mouse model of 22q11.2 deletion. NeuroImage, 2021, 243, 118520.	4.2	4
13	Translational medicine in psychiatry: challenges and imaging biomarkers. , 2021, , 203-223.		O
14	The genetic architecture of human brainstem structures and their involvement in common brain disorders. Nature Communications, 2020, 11, 4016.	12.8	26
15	Association of a Reproducible Epigenetic Risk Profile for Schizophrenia With Brain Methylation and Function. JAMA Psychiatry, 2020, 77, 628.	11.0	46
16	Identification of Reproducible BCL11A Alterations in Schizophrenia Through Individual-Level Prediction of Coexpression. Schizophrenia Bulletin, 2020, 46, 1165-1171.	4.3	8
17	Neural correlates of individual differences in affective benefit of real-life urban green space exposure. Nature Neuroscience, 2019, 22, 1389-1393.	14.8	125
18	Genome-wide Association Studies in Ancestrally Diverse Populations: Opportunities, Methods, Pitfalls, and Recommendations. Cell, 2019, 179, 589-603.	28.9	428

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19	Common brain disorders are associated with heritable patterns of apparent aging of the brain. Nature Neuroscience, 2019, 22, 1617-1623.	14.8	358
20	Reproducible grey matter patterns index a multivariate, global alteration of brain structure in schizophrenia and bipolar disorder. Translational Psychiatry, 2019, 9, 12.	4.8	35
21	Computational Approaches for Identification of Pleiotropic Biomarker Profiles in Psychiatry. Advances in Experimental Medicine and Biology, 2019, 1134, 111-128.	1.6	0
22	Gimpute: an efficient genetic data imputation pipeline. Bioinformatics, 2019, 35, 1433-1435.	4.1	7
23	RMTL: an R library for multi-task learning. Bioinformatics, 2019, 35, 1797-1798.	4.1	16
24	Resting-state brain network features associated with short-term skill learning ability in humans and the influence of <i>N</i> -methyl- <scp>d</scp> -aspartate receptor antagonism. Network Neuroscience, 2018, 2, 464-480.	2.6	14
25	Analysis of microbiota in first episode psychosis identifies preliminary associations with symptom severity and treatment response. Schizophrenia Research, 2018, 192, 398-403.	2.0	252
26	A geneâ€based review of <i>RGS4</i> as a putative risk gene for psychiatric illness. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 267-273.	1.7	18
27	Comparative Evaluation of Machine Learning Strategies for Analyzing Big Data in Psychiatry. International Journal of Molecular Sciences, 2018, 19, 3387.	4.1	18
28	Male increase in brain gene expression variability is linked to genetic risk for schizophrenia. Translational Psychiatry, 2018, 8, 140.	4.8	9
29	The Utility of Multiplex Assays for Identification of Proteomic Signatures in Psychiatry. Advances in Experimental Medicine and Biology, 2017, 974, 131-138.	1.6	3
30	Multiplex Immunoassay Profiling of Serum in Psychiatric Disorders. Advances in Experimental Medicine and Biology, 2017, 974, 149-156.	1.6	7
31	The role of blood-based biomarkers in advancing personalized therapy of schizophrenia. Expert Review of Precision Medicine and Drug Development, 2017, 2, 363-370.	0.7	0
32	Association of anandamide with altered binocular depth inversion illusion in schizophrenia. World Journal of Biological Psychiatry, 2017, 18, 483-488.	2.6	22
33	A polygenic score for schizophrenia predicts glycemic control. Translational Psychiatry, 2017, 7, 1295.	4.8	16
34	Identification and Clinical Translation of Biomarker Signatures: Statistical Considerations. Methods in Molecular Biology, 2017, 1546, 103-114.	0.9	2
35	Opportunities and Challenges of Multiplex Assays: A Machine Learning Perspective. Methods in Molecular Biology, 2017, 1546, 115-122.	0.9	6
36	Protein Interaction Networks Link Schizophrenia Risk Loci to Synaptic Function. Schizophrenia Bulletin, 2016, 42, 1334-1342.	4.3	16

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37	Working memory genetics in schizophrenia and related disorders: An RDoC perspective. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 121-131.	1.7	36
38	Dynamic brain network reconfiguration as a potential schizophrenia genetic risk mechanism modulated by NMDA receptor function. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 12568-12573.	7.1	161
39	Identifying multimodal signatures associated with symptom clusters: the example of the IMAGEMEND project. World Psychiatry, 2016, 15, 179-180.	10.4	14
40	Pretreatment levels of the fatty acid handling proteins H-FABP and CD36 predict response to olanzapine in recent-onset schizophrenia patients. Brain, Behavior, and Immunity, 2016, 52, 178-186.	4.1	26
41	Genetic influences on schizophrenia and subcortical brain volumes: large-scale proof of concept. Nature Neuroscience, 2016, 19, 420-431.	14.8	204
42	Investigation of molecular serum profiles associated with predisposition to antipsychotic-induced weight gain. World Journal of Biological Psychiatry, 2015, 16, 22-30.	2.6	20
43	IgG dynamics of dietary antigens point to cerebrospinal fluid barrier or flow dysfunction in first-episode schizophrenia. Brain, Behavior, and Immunity, 2015, 44, 148-158.	4.1	48
44	Sleep deprivation in humans: Effects on melatonin in cerebrospinal fluid and serum. Sleep and Biological Rhythms, 2014, 12, 69-72.	1.0	7
45	Oleoylethanolamide and Human Neural Responses to Food Stimuli in Obesity. JAMA Psychiatry, 2014, 71, 1254.	11.0	31
46	Identification of a Molecular Profile Associated with Immune Status in First-Onset Schizophrenia Patients. Clinical Schizophrenia and Related Psychoses, 2014, 7, 207-215.	1.4	30
47	Identification of Subgroups of Schizophrenia Patients With Changes in Either Immune or Growth Factor and Hormonal Pathways. Schizophrenia Bulletin, 2014, 40, 787-795.	4.3	84
48	Cytokine alterations in first-episode schizophrenia patients before and after antipsychotic treatment. Schizophrenia Research, 2014, 154, 23-29.	2.0	171
49	Proteomic profiling in schizophrenia: enabling stratification for more effective treatment. Genome Medicine, 2013, 5, 25.	8.2	19
50	Distinct Molecular Phenotypes in Male and Female Schizophrenia Patients. PLoS ONE, 2013, 8, e78729.	2.5	48
51	Biomarker discovery in human cerebrospinal fluid: the need for integrative metabolome and proteome databases. Genome Medicine, 2012, 4, 39.	8.2	5
52	Identification of a blood-based biological signature in subjects with psychiatric disorders prior to clinical manifestation. World Journal of Biological Psychiatry, 2012, 13, 627-632.	2.6	50
53	Alterations of primary fatty acid amides in serum of patients with severe mental illness. Frontiers in Bioscience - Elite, 2011, E3, 308-314.	1.8	12
54	The Application of Multiplexed Assay Systems for Molecular Diagnostics. International Review of Neurobiology, 2011, 101, 259-278.	2.0	0

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55	Validation of a Blood-Based Laboratory Test to Aid in the Confirmation of a Diagnosis of Schizophrenia. Biomarker Insights, 2010, 5, BMI.S4877.	2.5	137
56	Clinical bioinformatics for complex disorders: a schizophrenia case study. BMC Bioinformatics, 2009, 10, S6.	2.6	12
57	Biomarker discovery in psychiatric disorders. Electrophoresis, 2008, 29, 2884-2890.	2.4	33
58	High Throughput Lipidomic Profiling of Schizophrenia and Bipolar Disorder Brain Tissue Reveals Alterations of Free Fatty Acids, Phosphatidylcholines, and Ceramides. Journal of Proteome Research, 2008, 7, 4266-4277.	3.7	171
59	Cerebrospinal fluid: identification of diagnostic markers for schizophrenia. Expert Review of Molecular Diagnostics, 2008, 8, 209-216.	3.1	23
60	Peptide correlation: A means to identify high quality quantitative information in largeâ€scale proteomic studies. Journal of Separation Science, 2007, 30, 2190-2197.	2.5	22