

# Ina Giegling

## List of Publications by Year in descending order

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Version: 2024-02-01

61  
papers

14,469  
citations

94433

37  
h-index

118850

62  
g-index

65  
all docs

65  
docs citations

65  
times ranked

20479  
citing authors

#	ARTICLE	IF	CITATIONS
1	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. <i>Biological Psychiatry</i> , 2022, 91, 102-117.	1.3	61
2	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , 2022, 91, 313-327.	1.3	114
3	Three genetic“environmental networks for human personality. <i>Molecular Psychiatry</i> , 2021, 26, 3858-3875.	7.9	58
4	Shared genetic risk between eating disorder“and substance“use“related phenotypes: Evidence from genome“wide association studies. <i>Addiction Biology</i> , 2021, 26, e12880.	2.6	28
5	Genome-wide association study identifies 48 common genetic variants associated with handedness. <i>Nature Human Behaviour</i> , 2021, 5, 59-70.	12.0	79
6	HLA-DQB1 6672G>C (rs113332494) is associated with clozapine-induced neutropenia and agranulocytosis in individuals of European ancestry. <i>Translational Psychiatry</i> , 2021, 11, 214.	4.8	12
7	Identifying nootropic drug targets via large-scale cognitive GWAS and transcriptomics. <i>Neuropsychopharmacology</i> , 2021, 46, 1788-1801.	5.4	12
8	Common variants in Alzheimer“s disease and risk stratification by polygenic risk scores. <i>Nature Communications</i> , 2021, 12, 3417.	12.8	140
9	The Relationship Between Polygenic Risk Scores and Cognition in Schizophrenia. <i>Schizophrenia Bulletin</i> , 2020, 46, 336-344.	4.3	60
10	Uncovering the complex genetics of human character. <i>Molecular Psychiatry</i> , 2020, 25, 2295-2312.	7.9	77
11	Uncovering the complex genetics of human temperament. <i>Molecular Psychiatry</i> , 2020, 25, 2275-2294.	7.9	72
12	Genetics of resilience: Implications from genome“wide association studies and candidate genes of the stress response system in posttraumatic stress disorder and depression. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2020, 183, 77-94.	1.7	54
13	High-risk Allele for Herpes Labialis Severity at the IFNL3/4 Locus is Associated With Vestibular Neuritis. <i>Frontiers in Neurology</i> , 2020, 11, 570638.	2.4	10
14	Proton Magnetic Resonance Spectroscopy in Common Dementias“Current Status and Perspectives. <i>Frontiers in Psychiatry</i> , 2020, 11, 769.	2.6	21
15	Polymorphisms in CRYBB2 encoding $\beta$ 2-crystallin are associated with antisaccade performance and memory function. <i>Translational Psychiatry</i> , 2020, 10, 113.	4.8	3
16	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
17	Genome-wide association study identifies eight risk loci and implicates metabo-psychiatric origins for anorexia nervosa. <i>Nature Genetics</i> , 2019, 51, 1207-1214.	21.4	641
18	Genetic meta-analysis of diagnosed Alzheimer“s disease identifies new risk loci and implicates $A\beta$ , tau, immunity and lipid processing. <i>Nature Genetics</i> , 2019, 51, 414-430.	21.4	1,962

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19	Moderation of the relationship between <i>Toxoplasma gondii</i> seropositivity and trait impulsivity in younger men by the phenylalanine-tyrosine ratio. <i>Psychiatry Research</i> , 2018, 270, 992-1000.	3.3	8
20	A genome-wide association study of early gamma-band response in a schizophrenia case-control sample. <i>World Journal of Biological Psychiatry</i> , 2018, 19, 602-609.	2.6	3
21	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. <i>Nature Neuroscience</i> , 2018, 21, 1656-1669.	14.8	490
22	Study of 300,486 individuals identifies 148 independent genetic loci influencing general cognitive function. <i>Nature Communications</i> , 2018, 9, 2098.	12.8	484
23	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
24	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
25	The Genetics of Endophenotypes of Neurofunction to Understand Schizophrenia (GENUS) consortium: A collaborative cognitive and neuroimaging genetics project. <i>Schizophrenia Research</i> , 2018, 195, 306-317.	2.0	17
26	Genetics of schizophrenia: A consensus paper of the WFSBP Task Force on Genetics. <i>World Journal of Biological Psychiatry</i> , 2017, 18, 492-505.	2.6	48
27	Maob and Maob polymorphisms and personality traits in suicide attempters and healthy controls: a preliminary study. <i>Psychiatry Research</i> , 2017, 249, 212-217.	3.3	5
28	Cognitive Characterization of Schizophrenia Risk Variants Involved in Synaptic Transmission: Evidence of CACNA1C's Role in Working Memory. <i>Neuropsychopharmacology</i> , 2017, 42, 2612-2622.	5.4	28
29	Nine differentially expressed genes from a post mortem study and their association with suicidal status in a sample of suicide completers, attempters and controls. <i>Journal of Psychiatric Research</i> , 2017, 91, 98-104.	3.1	6
30	Rare coding variants in <i>PLCG2</i> , <i>ABI3</i> , and <i>TREM2</i> implicate microglial-mediated innate immunity in Alzheimer's disease. <i>Nature Genetics</i> , 2017, 49, 1373-1384.	21.4	783
31	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
32	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. <i>Nature Genetics</i> , 2017, 49, 27-35.	21.4	838
33	Consensus paper of the WFSBP Task Force on Genetics: Genetics, epigenetics and gene expression markers of major depressive disorder and antidepressant response. <i>World Journal of Biological Psychiatry</i> , 2017, 18, 5-28.	2.6	75
34	Genetik und Gen-Umwelt-Interaktionen bei psychischen Erkrankungen. , 2017, , 147-191.		2
35	Parental Origin of Interstitial Duplications at 15q11.2-q13.3 in Schizophrenia and Neurodevelopmental Disorders. <i>PLoS Genetics</i> , 2016, 12, e1005993.	3.5	51
36	Blood Levels of Monoamine Precursors and Smoking in Patients with Schizophrenia. <i>Frontiers in Public Health</i> , 2016, 4, 182.	2.7	5

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37	Reciprocal moderation by <i>Toxoplasma gondii</i> seropositivity and blood phenylalanine ð€“ tyrosine ratio of their associations with trait aggression. <i>Pteridines</i> , 2016, 27, 77-85.	0.5	8
38	A sequence variant associating with educational attainment also affects childhood cognition. <i>Scientific Reports</i> , 2016, 6, 36189.	3.3	2
39	Meta-analysis of Genome-Wide Association Studies for Extraversion: Findings from the Genetics of Personality Consortium. <i>Behavior Genetics</i> , 2016, 46, 170-182.	2.1	178
40	Combined <i>Toxoplasma gondii</i> seropositivity and high blood kynurenine ð€“ Linked with nonfatal suicidal self-directed violence in patients with schizophrenia. <i>Journal of Psychiatric Research</i> , 2016, 72, 74-81.	3.1	29
41	In psychiatrically healthy individuals, overweight women but not men have lower tryptophan levels. <i>Pteridines</i> , 2015, 26, 79-84.	0.5	9
42	Independent evidence for an association between general cognitive ability and a genetic locus for educational attainment. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2015, 168, 363-373.	1.7	25
43	Meta-analysis of Genome-wide Association Studies for Neuroticism, and the Polygenic Association With Major Depressive Disorder. <i>JAMA Psychiatry</i> , 2015, 72, 642.	11.0	289
44	Loss-of-function variants in <i>ABCA7</i> confer risk of Alzheimer's disease. <i>Nature Genetics</i> , 2015, 47, 445-447.	21.4	283
45	ð€œLatentð€•infection with <i>Toxoplasma gondii</i> : Association with trait aggression and impulsivity in healthy adults. <i>Journal of Psychiatric Research</i> , 2015, 60, 87-94.	3.1	92
46	Integrated Pathway-Based Approach Identifies Association between Genomic Regions at CTCF and <i>CACNB2</i> and Schizophrenia. <i>PLoS Genetics</i> , 2014, 10, e1004345.	3.5	44
47	Pleckstrin homology domain containing 6 protein ( <i>PLEKHA6</i> ) polymorphisms are associated with psychopathology and response to treatment in schizophrenic patients. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2014, 51, 190-195.	4.8	10
48	Partitioning Heritability of Regulatory and Cell-Type-Specific Variants across 11 Common Diseases. <i>American Journal of Human Genetics</i> , 2014, 95, 535-552.	6.2	569
49	Phadiatop Seropositivity in Schizophrenia Patients and Controls: A Preliminary Study. <i>AIMS Public Health</i> , 2014, 1, 43-50.	2.6	4
50	Glutamatergic Dysbalance and Oxidative Stress in In Vivo and In Vitro Models of Psychosis Based on Chronic NMDA Receptor Antagonism. <i>PLoS ONE</i> , 2013, 8, e59395.	2.5	24
51	A Positive Association between <i>T. gondii</i> Seropositivity and Obesity. <i>Frontiers in Public Health</i> , 2013, 1, 73.	2.7	32
52	Dysequilibrium of neuronal proliferation and apoptosis in a pharmacological animal model of psychosis. <i>Methods</i> , 2012, 56, 519-527.	3.8	11
53	Alterations of the early auditory evoked gamma-band response in first-degree relatives of patients with schizophrenia: Hints to a new intermediate phenotype. <i>Journal of Psychiatric Research</i> , 2011, 45, 699-705.	3.1	57
54	Maternally Derived Microduplications at 15q11-q13: Implication of Imprinted Genes in Psychotic Illness. <i>American Journal of Psychiatry</i> , 2011, 168, 408-417.	7.2	95

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55	Reduced Early Auditory Evoked Gamma-Band Response in Patients with Schizophrenia. <i>Biological Psychiatry</i> , 2010, 67, 224-231.	1.3	110
56	A Genome-Wide Investigation of SNPs and CNVs in Schizophrenia. <i>PLoS Genetics</i> , 2009, 5, e1000373.	3.5	383
57	Disruption of the neurexin 1 gene is associated with schizophrenia. <i>Human Molecular Genetics</i> , 2009, 18, 988-996.	2.9	424
58	Common variants conferring risk of schizophrenia. <i>Nature</i> , 2009, 460, 744-747.	27.8	1,572
59	Large recurrent microdeletions associated with schizophrenia. <i>Nature</i> , 2008, 455, 232-236.	27.8	1,619
60	Identification of loci associated with schizophrenia by genome-wide association and follow-up. <i>Nature Genetics</i> , 2008, 40, 1053-1055.	21.4	977
61	A Pharmacological Model for Psychosis Based on N-methyl-D-aspartate Receptor Hypofunction: Molecular, Cellular, Functional and Behavioral Abnormalities. <i>Biological Psychiatry</i> , 2006, 59, 721-729.	1.3	219