

Bettina Konte

List of Publications by Year in descending order

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Version: 2024-02-01

78
papers

20,217
citations

81743

39
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64668

79
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87
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docs citations

87
times ranked

24583
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.	0.7	61
2	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , 2022, 91, 313-327.	0.7	114
3	Three geneticâ€environmental networks for human personality. <i>Molecular Psychiatry</i> , 2021, 26, 3858-3875.	4.1	58
4	Shared genetic risk between eating disorderâ€and substanceâ€useâ€related phenotypes: Evidence from genomeâ€wide association studies. <i>Addiction Biology</i> , 2021, 26, e12880.	1.4	28
5	Genome-wide association study identifies 48 common genetic variants associated with handedness. <i>Nature Human Behaviour</i> , 2021, 5, 59-70.	6.2	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.	2.4	12
7	Genome-wide analyses of smoking behaviors in schizophrenia: Findings from the Psychiatric Genomics Consortium. <i>Journal of Psychiatric Research</i> , 2021, 137, 215-224.	1.5	10
8	Identifying nootropic drug targets via large-scale cognitive GWAS and transcriptomics. <i>Neuropsychopharmacology</i> , 2021, 46, 1788-1801.	2.8	12
9	The Relationship Between Polygenic Risk Scores and Cognition in Schizophrenia. <i>Schizophrenia Bulletin</i> , 2020, 46, 336-344.	2.3	60
10	Uncovering the complex genetics of human character. <i>Molecular Psychiatry</i> , 2020, 25, 2295-2312.	4.1	77
11	Uncovering the complex genetics of human temperament. <i>Molecular Psychiatry</i> , 2020, 25, 2275-2294.	4.1	72
12	A large-scale genome-wide association study meta-analysis of cannabis use disorder. <i>Lancet Psychiatry</i> , 2020, 7, 1032-1045.	3.7	200
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.	1.1	10
14	A Variation in FGF14 Is Associated with Downbeat Nystagmus in a Genome-Wide Association Study. <i>Cerebellum</i> , 2020, 19, 348-357.	1.4	16
15	Polymorphisms in CRYBB2 encoding Î²2-crystallin are associated with antisaccade performance and memory function. <i>Translational Psychiatry</i> , 2020, 10, 113.	2.4	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.	2.6	86
17	GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. <i>American Journal of Psychiatry</i> , 2019, 176, 651-660.	4.0	186
18	Populationâ€based identityâ€byâ€descent mapping combined with exome sequencing to detect rare risk variants for schizophrenia. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019, 180, 223-231.	1.1	2

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19	Common schizophrenia alleles are enriched in mutation-intolerant genes and in regions under strong background selection. <i>Nature Genetics</i> , 2018, 50, 381-389.	9.4	1,332
20	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.	1.7	8
21	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.	1.3	3
22	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. <i>Nature Neuroscience</i> , 2018, 21, 1656-1669.	7.1	490
23	Study of 300,486 individuals identifies 148 independent genetic loci influencing general cognitive function. <i>Nature Communications</i> , 2018, 9, 2098.	5.8	484
24	Estimation of Genetic Correlation via Linkage Disequilibrium Score Regression and Genomic Restricted Maximum Likelihood. <i>American Journal of Human Genetics</i> , 2018, 102, 1185-1194.	2.6	119
25	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.	9.4	893
26	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.	0.3	3
27	Genome-Wide Association Study in Vestibular Neuritis: Involvement of the Host Factor for HSV-1 Replication. <i>Frontiers in Neurology</i> , 2018, 9, 591.	1.1	44
28	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. <i>Cell</i> , 2018, 173, 1705-1715.e16.	13.5	623
29	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.	1.1	17
30	GWAS meta-analysis reveals novel loci and genetic correlates for general cognitive function: a report from the COGENT consortium. <i>Molecular Psychiatry</i> , 2017, 22, 336-345.	4.1	194
31	Maob and Maob polymorphisms and personality traits in suicide attempters and healthy controls: a preliminary study. <i>Psychiatry Research</i> , 2017, 249, 212-217.	1.7	5
32	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.	2.8	28
33	Genome-wide association study of borderline personality disorder reveals genetic overlap with bipolar disorder, major depression and schizophrenia. <i>Translational Psychiatry</i> , 2017, 7, e1155-e1155.	2.4	150
34	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.	1.5	6
35	Large-Scale Cognitive GWAS Meta-Analysis Reveals Tissue-Specific Neural Expression and Potential Nootropic Drug Targets. <i>Cell Reports</i> , 2017, 21, 2597-2613.	2.9	103
36	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. <i>Nature Genetics</i> , 2017, 49, 27-35.	9.4	838

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37	Blood Levels of Monoamine Precursors and Smoking in Patients with Schizophrenia. <i>Frontiers in Public Health</i> , 2016, 4, 182.	1.3	5
38	Reciprocal moderation by <i>Toxoplasma gondii</i> seropositivity and blood phenylalanine : tyrosine ratio of their associations with trait aggression. <i>Pteridines</i> , 2016, 27, 77-85.	0.5	8
39	A sequence variant associating with educational attainment also affects childhood cognition. <i>Scientific Reports</i> , 2016, 6, 36189.	1.6	2
40	Kynurenine and Tryptophan Levels in Patients With Schizophrenia and Elevated Antigliadin Immunoglobulin G Antibodies. <i>Psychosomatic Medicine</i> , 2016, 78, 931-939.	1.3	24
41	Genome-wide autozygosity is associated with lower general cognitive ability. <i>Molecular Psychiatry</i> , 2016, 21, 837-843.	4.1	62
42	Meta-analysis of Genome-Wide Association Studies for Extraversion: Findings from the Genetics of Personality Consortium. <i>Behavior Genetics</i> , 2016, 46, 170-182.	1.4	178
43	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. <i>Behavior Genetics</i> , 2016, 46, 151-169.	1.4	98
44	Combined <i>Toxoplasma gondii</i> seropositivity and high blood kynurenine : Linked with nonfatal suicidal self-directed violence in patients with schizophrenia. <i>Journal of Psychiatric Research</i> , 2016, 72, 74-81.	1.5	29
45	No Reliable Association between Runs of Homozygosity and Schizophrenia in a Well-Powered Replication Study. <i>PLoS Genetics</i> , 2016, 12, e1006343.	1.5	24
46	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.	1.1	25
47	Meta-analysis of Genome-wide Association Studies for Neuroticism, and the Polygenic Association With Major Depressive Disorder. <i>JAMA Psychiatry</i> , 2015, 72, 642.	6.0	289
48	Joint Analysis of Psychiatric Disorders Increases Accuracy of Risk Prediction for Schizophrenia, Bipolar Disorder, and Major Depressive Disorder. <i>American Journal of Human Genetics</i> , 2015, 96, 283-294.	2.6	225
49	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. <i>American Journal of Human Genetics</i> , 2015, 97, 576-592.	2.6	1,098
50	Expression analysis in a rat psychosis model identifies novel candidate genes validated in a large case-control sample of schizophrenia. <i>Translational Psychiatry</i> , 2015, 5, e656-e656.	2.4	36
51	New data and an old puzzle: the negative association between schizophrenia and rheumatoid arthritis. <i>International Journal of Epidemiology</i> , 2015, 44, 1706-1721.	0.9	53
52	Latent 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.	1.5	92
53	Elevated Levels of Plasma Phenylalanine in Schizophrenia: A Guanosine Triphosphate Cyclohydrolase-1 Metabolic Pathway Abnormality?. <i>PLoS ONE</i> , 2014, 9, e85945.	1.1	19
54	Mild expression differences of MECP2 influencing aggressive social behavior. <i>EMBO Molecular Medicine</i> , 2014, 6, 662-684.	3.3	23

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55	Genome-wide association study supports the role of the immunological system and of the neurodevelopmental processes in response to haloperidol treatment. <i>Pharmacogenetics and Genomics</i> , 2014, 24, 314-319.	0.7	28
56	Molecular genetic evidence for overlap between general cognitive ability and risk for schizophrenia: a report from the Cognitive Genomics consortium (COGENT). <i>Molecular Psychiatry</i> , 2014, 19, 168-174.	4.1	178
57	Partitioning Heritability of Regulatory and Cell-Type-Specific Variants across 11 Common Diseases. <i>American Journal of Human Genetics</i> , 2014, 95, 535-552.	2.6	569
58	Biological insights from 108 schizophrenia-associated genetic loci. <i>Nature</i> , 2014, 511, 421-427.	13.7	6,934
59	Influence of differentially expressed genes from suicide post-mortem study on personality traits as endophenotypes on healthy subjects and suicide attempters. <i>European Archives of Psychiatry and Clinical Neuroscience</i> , 2014, 264, 423-432.	1.8	11
60	Harmonization of Neuroticism and Extraversion phenotypes across inventories and cohorts in the Genetics of Personality Consortium: an application of Item Response Theory. <i>Behavior Genetics</i> , 2014, 44, 295-313.	1.4	103
61	Phadiatop Seropositivity in Schizophrenia Patients and Controls: A Preliminary Study. <i>AIMS Public Health</i> , 2014, 1, 43-50.	1.1	4
62	MAOA and MAOB polymorphisms and anger-related traits in suicidal participants and controls. <i>European Archives of Psychiatry and Clinical Neuroscience</i> , 2013, 263, 393-403.	1.8	32
63	In patients with schizophrenia, non-fatal suicidal self-directed violence is positively associated with present but not past smoking. <i>Schizophrenia Research</i> , 2013, 149, 194-195.	1.1	1
64	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. <i>Nature Genetics</i> , 2013, 45, 984-994.	9.4	2,067
65	Influence of ANKK1 and DRD2 polymorphisms in response to haloperidol. <i>European Archives of Psychiatry and Clinical Neuroscience</i> , 2013, 263, 65-74.	1.8	16
66	AKAP13, CACNA1, GRIK4 and GRIA1 genetic variations may be associated with haloperidol efficacy during acute treatment. <i>European Neuropsychopharmacology</i> , 2013, 23, 887-894.	0.3	27
67	Improved Detection of Common Variants Associated with Schizophrenia by Leveraging Pleiotropy with Cardiovascular-Disease Risk Factors. <i>American Journal of Human Genetics</i> , 2013, 92, 197-209.	2.6	422
68	CNTNAP2 polymorphisms and structural brain connectivity: A diffusion-tensor imaging study. <i>Journal of Psychiatric Research</i> , 2013, 47, 1349-1356.	1.5	37
69	Elevated gliadin antibody levels in individuals with schizophrenia. <i>World Journal of Biological Psychiatry</i> , 2013, 14, 509-515.	1.3	33
70	Improved Detection of Common Variants Associated with Schizophrenia and Bipolar Disorder Using Pleiotropy-Informed Conditional False Discovery Rate. <i>PLoS Genetics</i> , 2013, 9, e1003455.	1.5	298
71	Distinct Loci in the <i>CHRNA5</i> / <i>CHRNA3</i> / <i>CHRN4</i> Gene Cluster Are Associated With Onset of Regular Smoking. <i>Genetic Epidemiology</i> , 2013, 37, 846-859.	0.6	32
72	A Positive Association between <i>T. gondii</i> Seropositivity and Obesity. <i>Frontiers in Public Health</i> , 2013, 1, 73.	1.3	32

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73	Increased Genetic Vulnerability to Smoking at CHRNA5 in Early-Onset Smokers. Archives of General Psychiatry, 2012, 69, 854.	13.8	71
74	Lack of association of the 5-HT _{3A} receptor with schizophrenia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2012, 159B, 310-315.	1.1	8
75	Genome-wide association uncovers shared genetic effects among personality traits and mood states. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2012, 159B, 684-695.	1.1	112
76	Meta-analysis of genome-wide association studies for personality. Molecular Psychiatry, 2012, 17, 337-349.	4.1	340
77	Toxoplasma gondii antibody titers and history of suicide attempts in patients with schizophrenia. Schizophrenia Research, 2011, 133, 150-155.	1.1	108
78	Association of somatoform disorder symptoms with genetic variants potentially involved in the modulation of nociception. Psychiatric Genetics, 2011, 21, 50.	0.6	2