Eske M Derks

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5407283/publications.pdf

Version: 2024-02-01

146 papers 12,696 citations

43973 48 h-index 101 g-index

164 all docs

164 docs citations

times ranked

164

17596 citing authors

#	Article	IF	CITATIONS
1	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. Nature Genetics, 2018, 50, 668-681.	9.4	2,224
2	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	6.0	1,085
3	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. Cell, 2019, 179, 1469-1482.e11.	13.5	935
4	A tutorial on conducting genomeâ€wide association studies: Quality control and statistical analysis. International Journal of Methods in Psychiatric Research, 2018, 27, e1608.	1.1	465
5	GWAS of lifetime cannabis use reveals new risk loci, genetic overlap with psychiatric traits, and a causal effect of schizophrenia liability. Nature Neuroscience, 2018, 21, 1161-1170.	7.1	436
6	Revealing the complex genetic architecture of obsessive–compulsive disorder using meta-analysis. Molecular Psychiatry, 2018, 23, 1181-1188.	4.1	400
7	Using an atlas of gene regulation across 44 human tissues to inform complex disease- and trait-associated variation. Nature Genetics, 2018, 50, 956-967.	9.4	389
8	The Relationship of DNA Methylation with Age, Gender and Genotype in Twins and Healthy Controls. PLoS ONE, 2009, 4, e6767.	1.1	311
9	Partitioning the Heritability of Tourette Syndrome and Obsessive Compulsive Disorder Reveals Differences in Genetic Architecture. PLoS Genetics, 2013, 9, e1003864.	1.5	241
10	Netherlands Twin Register: A Focus on Longitudinal Research. Twin Research and Human Genetics, 2002, 5, 401-406.	1.5	195
11	Cohort profile: the Healthy Life in an Urban Setting (HELIUS) study in Amsterdam, The Netherlands. BMJ Open, 2017, 7, e017873.	0.8	163
12	Maternal Ratings of Attention Problems in ADHD: Evidence for the Existence of a Continuum. Journal of the American Academy of Child and Adolescent Psychiatry, 2009, 48, 1085-1093.	0.3	156
13	The Treatment of Hallucinations in Schizophrenia Spectrum Disorders. Schizophrenia Bulletin, 2012, 38, 704-714.	2.3	150
14	Across the continuum of attention skills: a twin study of the SWAN ADHD rating scale. Journal of Child Psychology and Psychiatry and Allied Disciplines, 2007, 48, 1080-1087.	3.1	148
15	Multisite prediction of 4-week and 52-week treatment outcomes in patients with first-episode psychosis: a machine learning approach. Lancet Psychiatry,the, 2016, 3, 935-946.	3.7	144
16	The Genetics of the Mood Disorder Spectrum: Genome-wide Association Analyses of More Than 185,000 Cases and 439,000 Controls. Biological Psychiatry, 2020, 88, 169-184.	0.7	137
17	Genome-wide association study of lifetime cannabis use based on a large meta-analytic sample of 32 330 subjects from the International Cannabis Consortium. Translational Psychiatry, 2016, 6, e769-e769.	2.4	136
18	Prevalence and Genetic Architecture of Child Behavior Checklist–Juvenile Bipolar Disorder. Biological Psychiatry, 2005, 58, 562-568.	0.7	133

#	Article	IF	Citations
19	Individual differences in aggression: genetic analyses by age, gender, and informant in 3-, 7-, and 10-year-old Dutch twins. Behavior Genetics, 2003, 33, 575-589.	1.4	124
20	Childhood trauma and auditory verbal hallucinations. Psychological Medicine, 2012, 42, 2475-2484.	2.7	124
21	Evidence-based psychiatric genetics, AKA the false dichotomy between common and rare variant hypotheses. Molecular Psychiatry, 2012, 17, 474-485.	4.1	124
22	Netherlands Twin Register: A Focus on Longitudinal Research. , 2002, .		122
23	Genetic Schizophrenia Risk Variants Jointly Modulate Total Brain and White Matter Volume. Biological Psychiatry, 2013, 73, 525-531.	0.7	119
24	Social cognition and quality of life in schizophrenia. Schizophrenia Research, 2012, 137, 212-218.	1.1	118
25	Cross-Disorder Genome-Wide Analyses Suggest a Complex Genetic Relationship Between Tourette's Syndrome and OCD. American Journal of Psychiatry, 2015, 172, 82-93.	4.0	117
26	Genome-wide gene-environment analyses of major depressive disorder and reported lifetime traumatic experiences in UK Biobank. Molecular Psychiatry, 2020, 25, 1430-1446.	4.1	116
27	Young Netherlands Twin Register (Y-NTR): A Longitudinal Multiple Informant Study of Problem Behavior. Twin Research and Human Genetics, 2007, 10, 3-11.	0.3	113
28	Copy Number Variation in Obsessive-Compulsive Disorder and Tourette Syndrome: A Cross-Disorder Study. Journal of the American Academy of Child and Adolescent Psychiatry, 2014, 53, 910-919.	0.3	111
29	A Comparison of Ten Polygenic Score Methods for Psychiatric Disorders Applied Across Multiple Cohorts. Biological Psychiatry, 2021, 90, 611-620.	0.7	103
30	A Test of the Equal Environment Assumption (EEA) in Multivariate Twin Studies. Twin Research and Human Genetics, 2006, 9, 403-411.	0.3	102
31	Effects of Censoring on Parameter Estimates and Power in Genetic Modeling. Twin Research and Human Genetics, 2004, 7, 659-669.	1.5	96
32	The Biological Contributions to Gender Identity and Gender Diversity: Bringing Data to the Table. Behavior Genetics, 2018, 48, 95-108.	1.4	92
33	Attention Problems and Attention-Deficit/Hyperactivity Disorder in Discordant and Concordant Monozygotic Twins: Evidence of Environmental Mediators. Journal of the American Academy of Child and Adolescent Psychiatry, 2007, 46, 83-91.	0.3	89
34	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. JAMA Psychiatry, 2021, 78, 1258.	6.0	88
35	Does Childhood Trauma Moderate Polygenic Risk for Depression? A Meta-analysis of 5765 Subjects From the Psychiatric Genomics Consortium. Biological Psychiatry, 2018, 84, 138-147.	0.7	87
36	The Genetic and Environmental Contributions to Attention Deficit Hyperactivity Disorder as Measured by the Conners' Rating Scalesâ€"Revised. American Journal of Psychiatry, 2005, 162, 1614-1620.	4.0	82

#	Article	IF	Citations
37	Multi-tissue transcriptome analyses identify genetic mechanisms underlying neuropsychiatric traits. Nature Genetics, 2019, 51, 933-940.	9.4	77
38	A systems medicine research approach for studying alcohol addiction. Addiction Biology, 2013, 18, 883-896.	1.4	76
39	Longitudinal Stability of the CBCL-Juvenile Bipolar Disorder Phenotype: A Study in Dutch Twins. Biological Psychiatry, 2006, 60, 912-920.	0.7	75
40	A gene co-expression network-based analysis of multiple brain tissues reveals novel genes and molecular pathways underlying major depression. PLoS Genetics, 2019, 15, e1008245.	1.5	74
41	Evidence of causal effect of major depression on alcohol dependence: findings from the psychiatric genomics consortium. Psychological Medicine, 2019, 49, 1218-1226.	2.7	74
42	Why More Boys Than Girls With ADHD Receive Treatment: A Study of Dutch Twins. Twin Research and Human Genetics, 2007, 10, 765-770.	0.3	62
43	Investigation of the Genetic Association between Quantitative Measures of Psychosis and Schizophrenia: A Polygenic Risk Score Analysis. PLoS ONE, 2012, 7, e37852.	1.1	60
44	Measurement invariance testing of the PHQ-9 in a multi-ethnic population in Europe: the HELIUS study. BMC Psychiatry, 2017, 17, 349.	1.1	59
45	Antipsychotic Drug Treatment in First-Episode Psychosis. Journal of Clinical Psychopharmacology, 2010, 30, 176-180.	0.7	58
46	A structural MRI study in monozygotic twins concordant or discordant for attention/hyperactivity problems: Evidence for genetic and environmental heterogeneity in the developing brain. NeuroImage, 2007, 35, 1004-1020.	2.1	54
47	Genetic and Environmental Influences on the Relation Between Attention Problems and Attention Deficit Hyperactivity Disorder. Behavior Genetics, 2008, 38, 11-23.	1.4	53
48	Drug attitude as predictor for effectiveness in first-episode schizophrenia: Results of an open randomized trial (EUFEST). European Neuropsychopharmacology, 2010, 20, 310-316.	0.3	53
49	A Study of Genetic and Environmental Influences on Maternal and Paternal CBCL Syndrome Scores in a Large Sample of 3-Year-Old Dutch Twins. Behavior Genetics, 2004, 34, 571-583.	1.4	51
50	Genome-wide association study of monoamine metabolite levels in human cerebrospinal fluid. Molecular Psychiatry, 2014, 19, 228-234.	4.1	51
51	Underestimated Effect Sizes in GWAS: Fundamental Limitations of Single SNP Analysis for Dichotomous Phenotypes. PLoS ONE, 2011, 6, e27964.	1.1	48
52	Focal And Global Brain Measurements in Siblings of Patients With Schizophrenia. Schizophrenia Bulletin, 2012, 38, 814-825.	2.3	48
53	Comorbid substance abuse in first-episode schizophrenia: Effects on cognition and psychopathology in the EUFEST study. Schizophrenia Research, 2013, 147, 132-139.	1.1	48
54	Assessment and Etiology of Attention Deficit Hyperactivity Disorder and Oppositional Defiant Disorder in Boys and Girls. Behavior Genetics, 2007, 37, 559-566.	1.4	47

#	Article	IF	CITATIONS
55	Genetic heterogeneity in self-reported depressive symptoms identified through genetic analyses of the PHQ-9. Psychological Medicine, 2020, 50, 2385-2396.	2.7	46
56	Symptom-level modelling unravels the shared genetic architecture of anxiety and depression. Nature Human Behaviour, 2021, 5, 1432-1442.	6.2	45
57	Potential influence of socioeconomic status on genetic correlations between alcohol consumption measures and mental health. Psychological Medicine, 2020, 50, 484-498.	2.7	44
58	Association of polygenic score for major depression with response to lithium in patients with bipolar disorder. Molecular Psychiatry, 2021, 26, 2457-2470.	4.1	44
59	Efficacy of Antipsychotic Drugs Against Hostility in the European First-Episode Schizophrenia Trial (EUFEST). Journal of Clinical Psychiatry, 2011, 72, 955-961.	1.1	43
60	Sex differences in the genetic architecture of obsessive–compulsive disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 351-364.	1.1	41
61	Genetic correlates of socio-economic status influence the pattern of shared heritability across mental health traits. Nature Human Behaviour, 2021, 5, 1065-1073.	6.2	41
62	The relations between DISC-IV DSM diagnoses of ADHD and multi-informant CBCL-AP syndrome scores. Comprehensive Psychiatry, 2006, 47, 116-122.	1.5	40
63	The Latent Class Structure of ADHD Is Stable Across Informants. Twin Research and Human Genetics, 2006, 9, 507-522.	0.3	40
64	Kraepelin Was Right: A Latent Class Analysis of Symptom Dimensions in Patients and Controls. Schizophrenia Bulletin, 2012, 38, 495-505.	2.3	40
65	Schizophrenia genetic variants are not associated with intelligence. Psychological Medicine, 2013, 43, 2563-2570.	2.7	40
66	Genetic liability for schizophrenia predicts risk of immune disorders. Schizophrenia Research, 2014, 159, 347-352.	1.1	40
67	A test of the equal environment assumption (EEA) in multivariate twin studies. Twin Research and Human Genetics, 2006, 9, 403-11.	0.3	40
68	The Genetic and Environmental Contributions to Oppositional Defiant Behavior: A Multi-informant Twin Study. Journal of the American Academy of Child and Adolescent Psychiatry, 2005, 44, 907-914.	0.3	39
69	The influence of semantic top-down processing in auditory verbal hallucinations. Schizophrenia Research, 2012, 139, 82-86.	1.1	38
70	Using genetic drug-target networks to develop new drug hypotheses for major depressive disorder. Translational Psychiatry, 2019, 9, 117.	2.4	37
71	Genetic Analyses of Maternal and Teacher Ratings on Attention Problems in 7-year-old Dutch Twins. Behavior Genetics, 2006, 36, 833-844.	1.4	36
72	E-MAGMA: an eQTL-informed method to identify risk genes using genome-wide association study summary statistics. Bioinformatics, 2021, 37, 2245-2249.	1.8	34

#	Article	IF	Citations
73	Categorical and dimensional psychopathology in Dutch and US offspring of parents with bipolar disorder: A preliminary cross-national comparison. Journal of Affective Disorders, 2016, 205, 95-102.	2.0	32
74	Symptom dimensions are associated with progressive brain volume changes in schizophrenia. Schizophrenia Research, 2012, 138, 171-176.	1.1	31
75	Identifying the Common Genetic Basis of Antidepressant Response. Biological Psychiatry Global Open Science, 2022, 2, 115-126.	1.0	31
76	Neuroimaging of response interference in twins concordant or discordant for inattention and hyperactivity symptoms. Neuroscience, 2009, 164, 16-29.	1.1	30
77	Measurement invariance of the SF-12 among different demographic groups: The HELIUS study. PLoS ONE, 2018, 13, e0203483.	1.1	30
78	Perceived ethnic discrimination in relation to smoking and alcohol consumption in ethnic minority groups in The Netherlands: the HELIUS study. International Journal of Public Health, 2017, 62, 879-887.	1.0	29
79	Cognitive biases and auditory verbal hallucinations in healthy and clinical individuals. Psychological Medicine, 2013, 43, 2339-2347.	2.7	28
80	Differential effects of antipsychotic drugs on insight in first episode schizophrenia: Data from the European First-Episode Schizophrenia Trial (EUFEST). European Neuropsychopharmacology, 2015, 25, 808-816.	0.3	28
81	Predictors of discontinuation of antipsychotic medication and subsequent outcomes in the European First Episode Schizophrenia Trial (EUFEST). Schizophrenia Research, 2016, 172, 145-151.	1.1	28
82	The Relative Contribution of Genes and Environment to Alcohol Use in Early Adolescents: Are Similar Factors Related to Initiation of Alcohol Use and Frequency of Drinking?. Alcoholism: Clinical and Experimental Research, 2008, 32, 975-982.	1.4	27
83	Does Assessment Type Matter? A Measurement Invariance Analysis of Online and Paper and Pencil Assessment of the Community Assessment of Psychic Experiences (CAPE). PLoS ONE, 2014, 9, e84011.	1.1	27
84	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. Biological Psychiatry, 2020, 87, 419-430.	0.7	27
85	Unmet needs in patients with first-episode schizophrenia: a longitudinal perspective. Psychological Medicine, 2012, 42, 1461-1473.	2.7	26
86	The prevalence of diabetes mellitus is increased in relatives of patients with a non-affective psychotic disorder. Schizophrenia Research, 2013, 143, 354-357.	1.1	26
87	Emotion processing in schizophrenia is state and trait dependent. Schizophrenia Research, 2015, 161, 392-398.	1.1	26
88	Genetic and environmental influences on the relationship between adult ADHD symptoms and self-reported problem drinking in 6024 Dutch twins. Psychological Medicine, 2014, 44, 2673-2683.	2.7	25
89	No evidence that common genetic risk variation is shared between schizophrenia and autism. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2013, 162, 55-60.	1.1	24
90	Impact of DSM-5 Changes on the Diagnosis and Acute Treatment of Schizophrenia. Schizophrenia Bulletin, 2015, 41, 637-643.	2.3	24

#	Article	IF	CITATIONS
91	Genomeâ€wide association metaâ€analysis of age at first cannabis use. Addiction, 2018, 113, 2073-2086.	1.7	24
92	The relation between obesity and depressed mood in a multi-ethnic population. The HELIUS study. Social Psychiatry and Psychiatric Epidemiology, 2018, 53, 629-638.	1.6	20
93	An analysis of genetically regulated gene expression across multiple tissues implicates novel gene candidates in Alzheimer's disease. Alzheimer's Research and Therapy, 2020, 12, 43.	3.0	20
94	Generalizability of the Results of Efficacy Trials in First-Episode Schizophrenia. Journal of Clinical Psychiatry, 2010, 71, 58-65.	1.1	20
95	Post-GWAS analysis of six substance use traits improves the identification and functional interpretation of genetic risk loci. Drug and Alcohol Dependence, 2020, 206, 107703.	1.6	19
96	The latent class structure of ADHD is stable across informants. Twin Research and Human Genetics, 2006, 9, 507-22.	0.3	19
97	Using multidimensional modeling to combine self-report symptoms with clinical judgment of schizotypy. Psychiatry Research, 2013, 206, 75-80.	1.7	18
98	The one-carbon-cycle and methylenetetrahydrofolate reductase (MTHFR) C677T polymorphism in recurrent major depressive disorder; influence of antidepressant use and depressive state?. Journal of Affective Disorders, 2014, 166, 115-123.	2.0	17
99	Ethnic differences in current smoking and former smoking in the Netherlands and the contribution of socioeconomic factors: a cross-sectional analysis of the HELIUS study. BMJ Open, 2017, 7, e016041.	0.8	17
100	Integrative Network-Based Analysis Reveals Gene Networks and Novel Drug Repositioning Candidates for Alzheimer Disease. Neurology: Genetics, 2021, 7, e622.	0.9	17
101	A genome wide survey supports the involvement of large copy number variants in schizophrenia with and without intellectual disability. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2013, 162, 847-854.	1.1	16
102	Association of Whole-Genome and NETRIN1 Signaling Pathway–Derived Polygenic Risk Scores for Major Depressive Disorder and White Matter Microstructure in the UK Biobank. Biological Psychiatry: Cognitive Neuroscience and Neuroimaging, 2019, 4, 91-100.	1.1	16
103	The interrelation of needs and quality of life in first-episode schizophrenia. European Archives of Psychiatry and Clinical Neuroscience, 2012, 262, 207-216.	1.8	15
104	Associations between the <i>CADM2</i> gene, substance use, risky sexual behavior, and selfâ€control: A phenomeâ€wide association study. Addiction Biology, 2021, 26, e13015.	1.4	15
105	Insight and Hostility as Predictors and Correlates of Nonadherence in the European First Episode Schizophrenia Trial. Journal of Clinical Psychopharmacology, 2013, 33, 258-261.	0.7	14
106	The impact of second generation antipsychotics on insight in schizophrenia: Results from 14 randomized, placebo controlled trials. European Neuropsychopharmacology, 2017, 27, 82-86.	0.3	14
107	How antipsychotics impact the different dimensions of Schizophrenia: A test of competing hypotheses. European Neuropsychopharmacology, 2014, 24, 1279-1288.	0.3	13
108	A two-factor structure of first rank symptoms in patients with a psychotic disorder. Schizophrenia Research, 2013, 147, 269-274.	1.1	12

#	Article	IF	CITATIONS
109	Investigating the genetic and causal relationship between initiation or use of alcohol, caffeine, cannabis and nicotine. Drug and Alcohol Dependence, 2020, 210, 107966.	1.6	12
110	Quantitative and qualitative symptomatic differences in individuals at Ultra-High Risk for psychosis and healthy controls. Psychiatry Research, 2013, 210, 432-437.	1.7	11
111	Assumptions and Properties of Limiting Pathway Models for Analysis of Epistasis in Complex Traits. PLoS ONE, 2013, 8, e68913.	1.1	11
112	Self-reported cue-induced physical symptoms of craving as an indicator of cocaine dependence. American Journal on Addictions, 2015, 24, 740-743.	1.3	11
113	Genetics of ADHD, Hyperactivity, and Attention Problems. , 2009, , 361-378.		11
114	Evidence of selection on splicing-associated loci in human populations and relevance to disease loci mapping. Scientific Reports, 2017, 7, 5980.	1.6	10
115	Exploring the role of low-frequency and rare exonic variants in alcohol and tobacco use. Drug and Alcohol Dependence, 2018, 188, 94-101.	1.6	10
116	Use of the Fagerström test to assess differences in the degree of nicotine dependence in smokers from five ethnic groups: The HELIUS study. Drug and Alcohol Dependence, 2019, 194, 197-204.	1.6	10
117	Ethnic and sex differences in the association of child maltreatment and depressed mood. The HELIUS study. Child Abuse and Neglect, 2020, 99, 104239.	1.3	10
118	Effects of Censoring on Parameter Estimates and Power in Genetic Modeling. , 0, .		10
118	Effects of Censoring on Parameter Estimates and Power in Genetic Modeling., 0, . Evaluating the role of alcohol consumption in breast and ovarian cancer susceptibility using populationâ€based cohort studies and twoâ€sample Mendelian randomization analyses. International Journal of Cancer, 2021, 148, 1338-1350.	2.3	10
	Evaluating the role of alcohol consumption in breast and ovarian cancer susceptibility using populationâ€based cohort studies and twoâ€sample Mendelian randomization analyses. International	2.3	
119	Evaluating the role of alcohol consumption in breast and ovarian cancer susceptibility using populationâ€based cohort studies and twoâ€sample Mendelian randomization analyses. International Journal of Cancer, 2021, 148, 1338-1350. The Influence of Informant Characteristics on the Reliability of Family History Interviews. Twin		9
119	Evaluating the role of alcohol consumption in breast and ovarian cancer susceptibility using populationâ€based cohort studies and twoâ€sample Mendelian randomization analyses. International Journal of Cancer, 2021, 148, 1338-1350. The Influence of Informant Characteristics on the Reliability of Family History Interviews. Twin Research and Human Genetics, 2011, 14, 217-220. Segment-Wise Genome-Wide Association Analysis Identifies a Candidate Region Associated with	0.3	9
119 120 121	Evaluating the role of alcohol consumption in breast and ovarian cancer susceptibility using populationâ€based cohort studies and twoâ€sample Mendelian randomization analyses. International Journal of Cancer, 2021, 148, 1338-1350. The Influence of Informant Characteristics on the Reliability of Family History Interviews. Twin Research and Human Genetics, 2011, 14, 217-220. Segment-Wise Genome-Wide Association Analysis Identifies a Candidate Region Associated with Schizophrenia in Three Independent Samples. PLoS ONE, 2012, 7, e38828. Parental Smoking and Adult Offspring's Smoking Behaviors in Ethnic Minority Groups: An	0.3	9 7 7
119 120 121 122	Evaluating the role of alcohol consumption in breast and ovarian cancer susceptibility using populationâ€based cohort studies and twoâ€sample Mendelian randomization analyses. International Journal of Cancer, 2021, 148, 1338-1350. The Influence of Informant Characteristics on the Reliability of Family History Interviews. Twin Research and Human Genetics, 2011, 14, 217-220. Segment-Wise Genome-Wide Association Analysis Identifies a Candidate Region Associated with Schizophrenia in Three Independent Samples. PLoS ONE, 2012, 7, e38828. Parental Smoking and Adult Offspring's Smoking Behaviors in Ethnic Minority Groups: An Intergenerational Analysis in the HELIUS Study. Nicotine and Tobacco Research, 2018, 20, 766-774. Exploring Phenotypic and Genetic Overlap Between Cannabis Use and Schizotypy. Twin Research and	0.3	9 7 7
119 120 121 122	Evaluating the role of alcohol consumption in breast and ovarian cancer susceptibility using populationâ€based cohort studies and twoâ€sample Mendelian randomization analyses. International journal of Cancer, 2021, 148, 1338-1350. The Influence of Informant Characteristics on the Reliability of Family History Interviews. Twin Research and Human Genetics, 2011, 14, 217-220. Segment-Wise Genome-Wide Association Analysis Identifies a Candidate Region Associated with Schizophrenia in Three Independent Samples. PLoS ONE, 2012, 7, e38828. Parental Smoking and Adult Offspring's Smoking Behaviors in Ethnic Minority Groups: An Intergenerational Analysis in the HELIUS Study. Nicotine and Tobacco Research, 2018, 20, 766-774. Exploring Phenotypic and Genetic Overlap Between Cannabis Use and Schizotypy. Twin Research and Human Genetics, 2020, 23, 221-227. Statistical Power to Detect Genetic and Environmental Influences in the Presence of Data Missing at	0.3 1.1 1.4	9 7 7 7

#	Article	IF	CITATIONS
127	High-potency cannabis and incident psychosis: correcting the causal assumption. Lancet Psychiatry,the, 2019, 6, 464.	3.7	6
128	A Local Genetic Correlation Analysis Provides Biological Insights Into the Shared Genetic Architecture of Psychiatric and Substance Use Phenotypes. Biological Psychiatry, 2022, 92, 583-591.	0.7	6
129	The calculation of familial loading in schizophrenia. Schizophrenia Research, 2009, 111, 198-199.	1.1	5
130	The Association between Intelligence Scores and Family History of Psychiatric Disorder in Schizophrenia Patients, Their Siblings and Healthy Controls. PLoS ONE, 2013, 8, e77215.	1.1	5
131	What Cure Models Can Teach us About Genome-Wide Survival Analysis. Behavior Genetics, 2016, 46, 269-280.	1.4	5
132	An integrative systemsâ€based analysis of substance use: <scp>eQTL</scp> â€informed geneâ€based tests, gene networks, and biological mechanisms. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2021, 186, 162-172.	1.1	5
133	The identification of family subtype based on the assessment of subclinical levels of psychosis in relatives. BMC Psychiatry, 2012, 12, 71.	1,1	4
134	Transcriptomeâ€wide association analysis offers novel opportunities for clinical translation of genetic discoveries on mental disorders. World Psychiatry, 2020, 19, 113-114.	4.8	4
135	A Comparison of Excessive Drinking, Binge Drinking and Alcohol Dependence in Ethnic Minority Groups in the Netherlands: The HELIUS Study. European Addiction Research, 2020, 26, 66-76.	1.3	4
136	Contribution of Alcohol and Nicotine Dependence to the Prevalence of Depressed Mood in Different Ethnic Groups in The Netherlands: The HELIUS Study. Journal of Dual Diagnosis, 2020, 16, 271-284.	0.7	4
137	Interpreting treatment trials in schizophrenia patients: Lessons learned from EUFEST. Schizophrenia Research, 2012, 138, 39-40.	1.1	3
138	Risk and Protective Factors of Lifetime Cocaine-Associated Chest Pain. Frontiers in Psychiatry, 2021, 12, 704276.	1.3	3
139	An analysis of genetically regulated gene expression and the role of co-expression networks across 16 psychiatric and substance use phenotypes. European Journal of Human Genetics, 2022, 30, 560-566.	1.4	3
140	"Forward Genetics―as a Method to Maximize Power and Cost-Efficiency in Studies of Human Complex Traits. Behavior Genetics, 2010, 40, 564-571.	1.4	2
141	Qualitative and quantitative aspects of information processing in first psychosis: Latent class analyses in patients, atâ€risk subjects, and controls. Psychophysiology, 2015, 52, 585-593.	1.2	2
142	Differential item functioning analysis of the CUDIT and relations with alcohol and tobacco use among men across five ethnic groups: The HELIUS study Psychology of Addictive Behaviors, 2019, 33, 697-709.	1.4	2
143	Replication and refinement of the role of rs548181 in schizophrenia: Results from a family based study. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2013, 162, 75-77.	1.1	1
144	Exploring Phenotypic and Genetic Overlap Between Cannabis Use and Schizotypy — Corrigendum. Twin Research and Human Genetics, 2020, 23, 306-306.	0.3	1

#	Article	IF	CITATIONS
145	SEGMENT-WISE GENOME-WIDE ASSOCIATION ANALYSIS IDENTIFIES A LIMITED NUMBER OF REPLICABLE CANDIDATE REGIONS ASSOCIATED WITH SCHIZOPHRENIA. Schizophrenia Research, 2010, 117, 219.	1.1	O
146	CORTICAL THICKNESS IN PATIENTS WITH SCHIZOPHRENIA AND THEIR SIBLINGS. Schizophrenia Research, 2010, 117, 223.	1.1	0