Eske M Derks

List of Publications by Year
in descending order

146 papers	12,696 citations	43973 48 h-index	31759 101 g-index
164 all docs	164 docs citations	164 times ranked	17596 citing authors

#	Article	IF	Citations
1	Identifying the Common Genetic Basis of Antidepressant Response. Biological Psychiatry Global Open Science, 2022, 2, 115-126.	1.0	31
2	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
3	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
4	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
5	Evaluating the role of alcohol consumption in breast and ovarian cancer susceptibility using population‐based cohort studies and two‐sample Mendelian randomization analyses. International Journal of Cancer, 2021, 148, 1338-1350.	2.3	9
6	Associations between the <i>CADM2</i> gene, substance use, risky sexual behavior, and self‐control: A phenome‐wide association study. Addiction Biology, 2021, 26, e13015.	1.4	15
7	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
8	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
9	Symptom-level modelling unravels the shared genetic architecture of anxiety and depression. Nature Human Behaviour, 2021, 5, 1432-1442.	6.2	45
10	A Comparison of Ten Polygenic Score Methods for Psychiatric Disorders Applied Across Multiple Cohorts. Biological Psychiatry, 2021, 90, 611-620.	0.7	103
11	Risk and Protective Factors of Lifetime Cocaine-Associated Chest Pain. Frontiers in Psychiatry, 2021, 12, 704276.	1.3	3
12	Integrative Network-Based Analysis Reveals Gene Networks and Novel Drug Repositioning Candidates for Alzheimer Disease. Neurology: Genetics, 2021, 7, e622.	0.9	17
13	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. JAMA Psychiatry, 2021, 78, 1258.	6.0	88
14	An integrative systems‐based analysis of substance use: <scp>eQTL</scp>‐informed gene‐based tests, gene networks, and biological mechanisms. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2021, 186, 162-172.	1.1	5
15	Potential influence of socioeconomic status on genetic correlations between alcohol consumption measures and mental health. Psychological Medicine, 2020, 50, 484-498.	2.7	44
16	Genetic heterogeneity in self-reported depressive symptoms identified through genetic analyses of the PHQ-9. Psychological Medicine, 2020, 50, 2385-2396.	2.7	46
17	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. Biological Psychiatry, 2020, 87, 419-430.	0.7	27
18	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

#	Article	IF	Citations
19	Transcriptome‐wide association analysis offers novel opportunities for clinical translation of genetic discoveries on mental disorders. World Psychiatry, 2020, 19, 113-114.	4.8	4
20	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
21	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
22	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
23	Exploring Phenotypic and Genetic Overlap Between Cannabis Use and Schizotypy. Twin Research and Human Genetics, 2020, 23, 221-227.	0.3	7
24	Exploring Phenotypic and Genetic Overlap Between Cannabis Use and Schizotypy â€" Corrigendum. Twin Research and Human Genetics, 2020, 23, 306-306.	0.3	1
25	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
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	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
28	An analysis of genetically regulated gene expression across multiple tissues implicates novel gene candidates in Alzheimerâ€™s disease. Alzheimer's Research and Therapy, 2020, 12, 43.	3.0	20
29	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
30	High-potency cannabis and incident psychosis: correcting the causal assumption. Lancet Psychiatry,the, 2019, 6, 464.	3.7	6
31	Multi-tissue transcriptome analyses identify genetic mechanisms underlying neuropsychiatric traits. Nature Genetics, 2019, 51, 933-940.	9.4	77
32	Using genetic drug-target networks to develop new drug hypotheses for major depressive disorder. Translational Psychiatry, 2019, 9, 117.	2.4	37
33	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
34	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. Cell, 2019, 179, 1469-1482.e11.	13.5	935
35	Association of Whole-Genome and NETRIN1 Signaling Pathwayâ€"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
36	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

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37	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
38	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
39	A tutorial on conducting genome‐wide association studies: Quality control and statistical analysis. International Journal of Methods in Psychiatric Research, 2018, 27, e1608.	1.1	465
40	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
41	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.	1.4	7
42	The Biological Contributions to Gender Identity and Gender Diversity: Bringing Data to the Table. Behavior Genetics, 2018, 48, 95-108.	1.4	92
43	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
44	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
45	Revealing the complex genetic architecture of obsessive–compulsive disorder using meta-analysis. Molecular Psychiatry, 2018, 23, 1181-1188.	4.1	400
46	Measurement invariance of the SF-12 among different demographic groups: The HELIUS study. PLoS ONE, 2018, 13, e0203483.	1.1	30
47	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
48	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
49	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	6.0	1,085
50	Genome‐wide association meta‐analysis of age at first cannabis use. Addiction, 2018, 113, 2073-2086.	1.7	24
51	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
52	Evidence of selection on splicing-associated loci in human populations and relevance to disease loci mapping. Scientific Reports, 2017, 7, 5980.	1.6	10
53	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
54	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

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55	Cohort profile: the Healthy Life in an Urban Setting (HELIUS) study in Amsterdam, The Netherlands. BMJ Open, 2017, 7, e017873.	0.8	163
56	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
57	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
58	Body integrity identity disorder crosses culture: case reports in the Japanese and Chinese literature. Neuropsychiatric Disease and Treatment, 2016, 12, 1419.	1.0	6
59	Genome-wide association study of lifetime cannabis use based on a large meta-analytic sample of 32â€‰330 subjects from the International Cannabis Consortium. Translational Psychiatry, 2016, 6, e769-e769.	2.4	136
60	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
61	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
62	What Cure Models Can Teach us About Genome-Wide Survival Analysis. Behavior Genetics, 2016, 46, 269-280.	1.4	5
63	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
64	A guide on gene prioritization in studies of psychiatric disorders. International Journal of Methods in Psychiatric Research, 2015, 24, 245-256.	1.1	6
65	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
66	Emotion processing in schizophrenia is state and trait dependent. Schizophrenia Research, 2015, 161, 392-398.	1.1	26
67	Cross-Disorder Genome-Wide Analyses Suggest a Complex Genetic Relationship Between Touretteâ€™s Syndrome and OCD. American Journal of Psychiatry, 2015, 172, 82-93.	4.0	117
68	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
69	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
70	Impact of DSM-5 Changes on the Diagnosis and Acute Treatment of Schizophrenia. Schizophrenia Bulletin, 2015, 41, 637-643.	2.3	24
71	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
72	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

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73	How antipsychotics impact the different dimensions of Schizophrenia: A test of competing hypotheses. European Neuropsychopharmacology, 2014, 24, 1279-1288.	0.3	13
74	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
75	Genetic liability for schizophrenia predicts risk of immune disorders. Schizophrenia Research, 2014, 159, 347-352.	1.1	40
76	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
77	Genome-wide association study of monoamine metabolite levels in human cerebrospinal fluid. Molecular Psychiatry, 2014, 19, 228-234.	4.1	51
78	Using multidimensional modeling to combine self-report symptoms with clinical judgment of schizotypy. Psychiatry Research, 2013, 206, 75-80.	1.7	18
79	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
80	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
81	A systems medicine research approach for studying alcohol addiction. Addiction Biology, 2013, 18, 883-896.	1.4	76
82	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
83	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
84	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
85	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
86	Genetic Schizophrenia Risk Variants Jointly Modulate Total Brain and White Matter Volume. Biological Psychiatry, 2013, 73, 525-531.	0.7	119
87	A two-factor structure of first rank symptoms in patients with a psychotic disorder. Schizophrenia Research, 2013, 147, 269-274.	1.1	12
88	Partitioning the Heritability of Tourette Syndrome and Obsessive Compulsive Disorder Reveals Differences in Genetic Architecture. PLoS Genetics, 2013, 9, e1003864.	1.5	241
89	Cognitive biases and auditory verbal hallucinations in healthy and clinical individuals. Psychological Medicine, 2013, 43, 2339-2347.	2.7	28
90	Schizophrenia genetic variants are not associated with intelligence. Psychological Medicine, 2013, 43, 2563-2570.	2.7	40

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91	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
92	Assumptions and Properties of Limiting Pathway Models for Analysis of Epistasis in Complex Traits. PLoS ONE, 2013, 8, e68913.	1.1	11
93	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
94	Kraepelin Was Right: A Latent Class Analysis of Symptom Dimensions in Patients and Controls. Schizophrenia Bulletin, 2012, 38, 495-505.	2.3	40
95	Childhood trauma and auditory verbal hallucinations. Psychological Medicine, 2012, 42, 2475-2484.	2.7	124
96	Focal And Global Brain Measurements in Siblings of Patients With Schizophrenia. Schizophrenia Bulletin, 2012, 38, 814-825.	2.3	48
97	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
98	Unmet needs in patients with first-episode schizophrenia: a longitudinal perspective. Psychological Medicine, 2012, 42, 1461-1473.	2.7	26
99	Evidence-based psychiatric genetics, AKA the false dichotomy between common and rare variant hypotheses. Molecular Psychiatry, 2012, 17, 474-485.	4.1	124
100	Social cognition and quality of life in schizophrenia. Schizophrenia Research, 2012, 137, 212-218.	1.1	118
101	Interpreting treatment trials in schizophrenia patients: Lessons learned from EUFEST. Schizophrenia Research, 2012, 138, 39-40.	1.1	3
102	Symptom dimensions are associated with progressive brain volume changes in schizophrenia. Schizophrenia Research, 2012, 138, 171-176.	1.1	31
103	The influence of semantic top-down processing in auditory verbal hallucinations. Schizophrenia Research, 2012, 139, 82-86.	1.1	38
104	The identification of family subtype based on the assessment of subclinical levels of psychosis in relatives. BMC Psychiatry, 2012, 12, 71.	1.1	4
105	Segment-Wise Genome-Wide Association Analysis Identifies a Candidate Region Associated with Schizophrenia in Three Independent Samples. PLoS ONE, 2012, 7, e38828.	1.1	7
106	The Treatment of Hallucinations in Schizophrenia Spectrum Disorders. Schizophrenia Bulletin, 2012, 38, 704-714.	2.3	150
107	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
108	Underestimated Effect Sizes in GWAS: Fundamental Limitations of Single SNP Analysis for Dichotomous Phenotypes. PLoS ONE, 2011, 6, e27964.	1.1	48

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109	The Influence of Informant Characteristics on the Reliability of Family History Interviews. Twin Research and Human Genetics, 2011, 14, 217-220.	0.3	7
110	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
111	Antipsychotic Drug Treatment in First-Episode Psychosis. Journal of Clinical Psychopharmacology, 2010, 30, 176-180.	0.7	58
112	â€œForward Geneticsâ€•as a Method to Maximize Power and Cost-Efficiency in Studies of Human Complex Traits. Behavior Genetics, 2010, 40, 564-571.	1.4	2
113	SEGMENT-WISE GENOME-WIDE ASSOCIATION ANALYSIS IDENTIFIES A LIMITED NUMBER OF REPLICABLE CANDIDATE REGIONS ASSOCIATED WITH SCHIZOPHRENIA. Schizophrenia Research, 2010, 117, 219.	1.1	0
114	CORTICAL THICKNESS IN PATIENTS WITH SCHIZOPHRENIA AND THEIR SIBLINGS. Schizophrenia Research, 2010, 117, 223.	1.1	0
115	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
116	Generalizability of the Results of Efficacy Trials in First-Episode Schizophrenia. Journal of Clinical Psychiatry, 2010, 71, 58-65.	1.1	20
117	The Relationship of DNA Methylation with Age, Gender and Genotype in Twins and Healthy Controls. PLoS ONE, 2009, 4, e6767.	1.1	311
118	Neuroimaging of response interference in twins concordant or discordant for inattention and hyperactivity symptoms. Neuroscience, 2009, 164, 16-29.	1.1	30
119	The calculation of familial loading in schizophrenia. Schizophrenia Research, 2009, 111, 198-199.	1.1	5
120	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
121	Genetics of ADHD, Hyperactivity, and Attention Problems. , 2009, , 361-378.		11
122	Genetic and Environmental Influences on the Relation Between Attention Problems and Attention Deficit Hyperactivity Disorder. Behavior Genetics, 2008, 38, 11-23.	1.4	53
123	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
124	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
125	Statistical Power to Detect Genetic and Environmental Influences in the Presence of Data Missing at Random. Twin Research and Human Genetics, 2007, 10, 159-167.	0.3	6
126	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

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127	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
128	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
129	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
130	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
131	Longitudinal Stability of the CBCL-Juvenile Bipolar Disorder Phenotype: A Study in Dutch Twins. Biological Psychiatry, 2006, 60, 912-920.	0.7	75
132	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
133	The Latent Class Structure of ADHD Is Stable Across Informants. Twin Research and Human Genetics, 2006, 9, 507-522.	0.3	40
134	A Test of the Equal Environment Assumption (EEA) in Multivariate Twin Studies. Twin Research and Human Genetics, 2006, 9, 403-411.	0.3	102
135	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
136	A test of the equal environment assumption (EEA) in multivariate twin studies. Twin Research and Human Genetics, 2006, 9, 403-11.	0.3	40
137	The latent class structure of ADHD is stable across informants. Twin Research and Human Genetics, 2006, 9, 507-22.	0.3	19
138	The Genetic and Environmental Contributions to Attention Deficit Hyperactivity Disorder as Measured by the Conners' Rating Scales—Revised. American Journal of Psychiatry, 2005, 162, 1614-1620.	4.0	82
139	Prevalence and Genetic Architecture of Child Behavior Checklist—Juvenile Bipolar Disorder. Biological Psychiatry, 2005, 58, 562-568.	0.7	133
140	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
141	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
142	Effects of Censoring on Parameter Estimates and Power in Genetic Modeling. Twin Research and Human Genetics, 2004, 7, 659-669.	1.5	96
143	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
144	Netherlands Twin Register: A Focus on Longitudinal Research. Twin Research and Human Genetics, 2002, 5, 401-406.	1.5	195

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145	Netherlands Twin Register: A Focus on Longitudinal Research. , 2002, .		122
146	Effects of Censoring on Parameter Estimates and Power in Genetic Modeling. , 0, .		10