Penelope A Lind

List of Publications by Year in descending order

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111 papers	18,686 citations	43973 48 h-index	24179 110 g-index
130 all docs	130 docs citations	130 times ranked	22423 citing authors

#	Article	IF	Citations
1	censusADHD Study: An Australian-wide medication-based recruitment study for Attention-Deficit/Hyperactivity Disorder. Australian and New Zealand Journal of Psychiatry, 2023, 57, 252-263.	1.3	0
2	Ten years of enhancing <scp>neuroâ€imaging</scp> genetics through <scp>metaâ€analysis</scp>: An overview from the <scp>ENIGMA Genetics Working Group</scp>. Human Brain Mapping, 2022, 43, 292-299.	1.9	19
3	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. Biological Psychiatry, 2022, 91, 102-117.	0.7	61
4	Identifying the Common Genetic Basis of Antidepressant Response. Biological Psychiatry Global Open Science, 2022, 2, 115-126.	1.0	31
5	Impact of CYP2C19 metaboliser status on SSRI response: a retrospective study of 9500 participants of the Australian Genetics of Depression Study. Pharmacogenomics Journal, 2022, 22, 130-135.	0.9	16
6	Australian Parkinsonâ€™s Genetics Study (APGS): pilot (n=1532). BMJ Open, 2022, 12, e052032.	0.8	1
7	Polygenic prediction of educational attainment within and between families from genome-wide association analyses in 3 million individuals. Nature Genetics, 2022, 54, 437-449.	9.4	215
8	Within-sibship genome-wide association analyses decrease bias in estimates of direct genetic effects. Nature Genetics, 2022, 54, 581-592.	9.4	142
9	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
10	Shared genetic risk between eating disorderâ€and substanceâ€useâ€related phenotypes: Evidence from genomeâ€wide association studies. Addiction Biology, 2021, 26, e12880.	1.4	28
11	The Evolutionary History of Common Genetic Variants Influencing Human Cortical Surface Area. Cerebral Cortex, 2021, 31, 1873-1887.	1.6	21
12	The psychosocial impact of nausea and vomiting during pregnancy as a predictor of postpartum depression. Journal of Health Psychology, 2021, 26, 1061-1072.	1.3	8
13	Bipolar multiplex families have an increased burden of common risk variants for psychiatric disorders. Molecular Psychiatry, 2021, 26, 1286-1298.	4.1	33
14	Model-based assessment of replicability for genome-wide association meta-analysis. Nature Communications, 2021, 12, 1964.	5.8	24
15	A Comparison of Ten Polygenic Score Methods for Psychiatric Disorders Applied Across Multiple Cohorts. Biological Psychiatry, 2021, 90, 611-620.	0.7	103
16	Identification of 371 genetic variants for age at first sex and birth linked to externalising behaviour. Nature Human Behaviour, 2021, 5, 1717-1730.	6.2	62
17	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. JAMA Psychiatry, 2021, 78, 1258.	6.0	88
18	Understanding genetic risk factors for common side effects of antidepressant medications. Communications Medicine, 2021, 1, .	1.9	15

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19	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. Biological Psychiatry, 2020, 87, 419-430.	0.7	27
20	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
21	Genetic correlations and genome-wide associations of cortical structure in general population samples of 22,824 adults. Nature Communications, 2020, 11, 4796.	5.8	61
22	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	3.7	200
23	The genetic architecture of sporadic and multiple consecutive miscarriage. Nature Communications, 2020, 11, 5980.	5.8	52
24	Rethinking Measurement of Parenting Stress in ADHD-Affected Families: A Principal Components Analysis of the Disruptive Behaviour Stress Inventory. Journal of Child and Family Studies, 2020, 29, 3253-3264.	0.7	0
25	Contributions of Nicholas Martin to Gambling Disorder Research. Twin Research and Human Genetics, 2020, 23, 127-128.	0.3	0
26	A phenome-wide association and Mendelian Randomisation study of polygenic risk for depression in UK Biobank. Nature Communications, 2020, 11, 2301.	5.8	81
27	The genetic architecture of the human cerebral cortex. Science, 2020, 367, .	6.0	450
28	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
29	New alcohol-related genes suggest shared genetic mechanisms with neuropsychiatric disorders. Nature Human Behaviour, 2019, 3, 950-961.	6.2	75
30	Genome-wide association and epidemiological analyses reveal common genetic origins between uterine leiomyomata and endometriosis. Nature Communications, 2019, 10, 4857.	5.8	90
31	Assessment of Bidirectional Relationships Between Physical Activity and Depression Among Adults. JAMA Psychiatry, 2019, 76, 399.	6.0	399
32	Integrated analysis of environmental and genetic influences on cord blood DNA methylation in new-borns. Nature Communications, 2019, 10, 2548.	5.8	94
33	Twenty-Five and Up (25Up) Study: A New Wave of the Brisbane Longitudinal Twin Study. Twin Research and Human Genetics, 2019, 22, 154-163.	0.3	19
34	Maternal and fetal genetic effects on birth weight and their relevance to cardio-metabolic risk factors. Nature Genetics, 2019, 51, 804-814.	9.4	402
35	Genome wide analysis for mouth ulcers identifies associations at immune regulatory loci. Nature Communications, 2019, 10, 1052.	5.8	50
36	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

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37	Identification of common genetic risk variants for autism spectrum disorder. Nature Genetics, 2019, 51, 431-444.	9.4	1,538
38	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
39	Association studies of up to 1.2 million individuals yield new insights into the genetic etiology of tobacco and alcohol use. Nature Genetics, 2019, 51, 237-244.	9.4	1,307
40	Genome-wide association analyses of risk tolerance and risky behaviors in over 1 million individuals identify hundreds of loci and shared genetic influences. Nature Genetics, 2019, 51, 245-257.	9.4	536
41	Genome-wide association study of offspring birth weight in 86â€‰577 women identifies five novel loci and highlights maternal genetic effects that are independent of fetal genetics. Human Molecular Genetics, 2018, 27, 742-756.	1.4	156
42	No evidence of association of oxytocin polymorphisms with breastfeeding in 2 independent samples. Genes, Brain and Behavior, 2018, 17, e12464.	1.1	4
43	Pathways to depression by age 16 years: Examining trajectories for self-reported psychological and somatic phenotypes across adolescence. Journal of Affective Disorders, 2018, 230, 1-6.	2.0	15
44	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
45	Nineteen and Up study (19Up): understanding pathways to mental health disorders in young Australian twins. BMJ Open, 2018, 8, e018959.	0.8	19
46	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
47	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. Nature Neuroscience, 2018, 21, 1656-1669.	7.1	490
48	Genome-wide interaction study of a proxy for stress-sensitivity and its prediction of major depressive disorder. PLoS ONE, 2018, 13, e0209160.	1.1	14
49	Applying polygenic risk scoring for psychiatric disorders to a large family with bipolar disorder and major depressive disorder. Communications Biology, 2018, 1, 163.	2.0	17
50	Gene discovery and polygenic prediction from a genome-wide association study of educational attainment in 1.1 million individuals. Nature Genetics, 2018, 50, 1112-1121.	9.4	1,835
51	Genomeâ€wide association metaâ€analysis of age at first cannabis use. Addiction, 2018, 113, 2073-2086.	1.7	24
52	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. Nature Genetics, 2017, 49, 834-841.	9.4	426
53	Genetic Association of Major Depression With Atypical Features and Obesity-Related Immunometabolic Dysregulations. JAMA Psychiatry, 2017, 74, 1214.	6.0	174
54	Personality Polygenes, Positive Affect, and Life Satisfaction. Twin Research and Human Genetics, 2016, 19, 407-417.	0.3	16

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55	The Genetic and Environmental Contributions to Internet Use and Associations With Psychopathology: A Twin Study. Twin Research and Human Genetics, 2016, 19, 1-9.	0.3	28
56	Genetic variants associated with subjective well-being, depressive symptoms, and neuroticism identified through genome-wide analyses. Nature Genetics, 2016, 48, 624-633.	9.4	870
57	Cohort Profile: Nausea and vomiting during pregnancy genetics consortium (NVP Genetics) Tj ETQq1 1 0.784314 rgBT /Overlock 10 T	0.9	11
58	Genome-wide association study identifies 74 loci associated with educational attainment. Nature, 2016, 533, 539-542.	13.7	1,204
59	Genome-wide analysis identifies 12 loci influencing human reproductive behavior. Nature Genetics, 2016, 48, 1462-1472.	9.4	284
60	Genetic variants linked to education predict longevity. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 13366-13371.	3.3	110
61	Genome-wide association study of pathological gambling. European Psychiatry, 2016, 36, 38-46.	0.1	82
62	Nausea and Vomiting During Pregnancy is Highly Heritable. Behavior Genetics, 2016, 46, 481-491.	1.4	24
63	International Genome-Wide Association Study Consortium Identifies Novel Loci Associated With Blood Pressure in Children and Adolescents. Circulation: Cardiovascular Genetics, 2016, 9, 266-278.	5.1	48
64	Connecting the dots, genome-wide association studies in substance use. Molecular Psychiatry, 2016, 21, 733-735.	4.1	31
65	Genetic Evidence for Causal Relationships Between Maternal Obesity-Related Traits and Birth Weight. JAMA - Journal of the American Medical Association, 2016, 315, 1129.	3.8	220
66	The association between lower educational attainment and depression owing to shared genetic effects? Results in ~25â€‰000 subjects. Molecular Psychiatry, 2015, 20, 735-743.	4.1	59
67	Directional dominance on stature and cognition inÂdiverse human populations. Nature, 2015, 523, 459-462.	13.7	173
68	A meta-analysis of genome-wide association studies identifies novel variants associated with osteoarthritis of the hip. Annals of the Rheumatic Diseases, 2014, 73, 2130-2136.	0.5	108
69	Polygenic scores associated with educational attainment in adults predict educational achievement and ADHD symptoms in children. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 510-520.	1.1	40
70	Genomeâ€wide association study of a quantitative disordered gambling trait. Addiction Biology, 2013, 18, 511-522.	1.4	94
71	Identification of heart rateâ€“associated loci and their effects on cardiac conduction and rhythm disorders. Nature Genetics, 2013, 45, 621-631.	9.4	282
72	GWAS of 126,559 Individuals Identifies Genetic Variants Associated with Educational Attainment. Science, 2013, 340, 1467-1471.	6.0	750

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73	Distinct Loci in the CHRNA5/CHRNA3/CHRNB4 Gene Cluster Are Associated With Onset of Regular Smoking. Genetic Epidemiology, 2013, 37, 846-859.	0.6	32
74	Case-Control Association Testing of Common Variants from Sequencing of DNA Pools. PLoS ONE, 2013, 8, e65410.	1.1	0
75	Coaction of Stress and Serotonin Transporter Genotype in Predicting Aggression at the Transition to Adulthood. Journal of Clinical Child and Adolescent Psychology, 2012, 41, 53-63.	2.2	34
76	Association Between In Vivo Alcohol Metabolism and Genetic Variation in Pathways that Metabolize the Carbon Skeleton of Ethanol and <scp>NADH</scp> Reoxidation in the Alcohol Challenge Twin Study. Alcoholism: Clinical and Experimental Research, 2012, 36, 2074-2085.	1.4	11
77	A Quantitative-Trait Genome-Wide Association Study of Alcoholism Risk in the Community: Findings and Implications. Biological Psychiatry, 2011, 70, 513-518.	0.7	184
78	Interactions between the COMT Val108/158Met polymorphism and maternal prenatal smoking predict aggressive behavior outcomes. Biological Psychology, 2011, 87, 99-105.	1.1	38
79	The Investigation into CYP2E1 in Relation to the Level of Response to Alcohol Through a Combination of Linkage and Association Analysis. Alcoholism: Clinical and Experimental Research, 2011, 35, 10-18.	1.4	33
80	GWAS of butyrylcholinesterase activity identifies four novel loci, independent effects within BCHE and secondary associations with metabolic risk factors. Human Molecular Genetics, 2011, 20, 4504-4514.	1.4	45
81	A Genomewide Association Study of Nicotine and Alcohol Dependence in Australian and Dutch Populations. Twin Research and Human Genetics, 2010, 13, 11-29.	0.3	3
82	Interaction of chronic stress with serotonin transporter and catechol-O -methyltransferase polymorphisms in predicting youth depression. Depression and Anxiety, 2010, 27, 737-745.	2.0	47
83	Linkage Analysis of Alcohol Dependence Symptoms in the Community. Alcoholism: Clinical and Experimental Research, 2010, 34, 158-163.	1.4	12
84	Dyslexia and DCDC2: normal variation in reading and spelling is associated with DCDC2 polymorphisms in an Australian population sample. European Journal of Human Genetics, 2010, 18, 668-673.	1.4	73
85	Dyslexia and DYX1C1: deficits in reading and spelling associated with a missense mutation. Molecular Psychiatry, 2010, 15, 1190-1196.	4.1	68
86	A Genomewide Association Study of Nicotine and Alcohol Dependence in Australian and Dutch Populations. Twin Research and Human Genetics, 2010, 13, 10-29.	0.3	98
87	Associations of ADH and ALDH2 gene variation with self report alcohol reactions, consumption and dependence: an integrated analysis. Human Molecular Genetics, 2009, 18, 580-593.	1.4	187
88	Genetic Variation in Female BMI Increases with Number of Children Born but Failure to Replicate Association betweenGNÎ²3Variants and Increased BMI in Parous Females. Twin Research and Human Genetics, 2009, 12, 276-285.	0.3	0
89	No Association Between Cholinergic Muscarinic Receptor 2 (CHRM2) Genetic Variation and Cognitive Abilities in Three Independent Samples. Behavior Genetics, 2009, 39, 513-523.	1.4	10
90	Variation in the dysbindin gene and normal cognitive function in three independent population samples. Genes, Brain and Behavior, 2009, 8, 218-227.	1.1	47

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91	Replication of the association of common rs9939609 variant of FTO with increased BMI in an Australian adult twin population but no evidence for gene by environment (G Ã— E) interaction. International Journal of Obesity, 2009, 33, 75-79.	1.6	51
92	Can We Identify Genes For Alcohol Consumption In Samples Ascertained For Heterogeneous Purposes?. Alcoholism: Clinical and Experimental Research, 2009, 33, 729-739.	1.4	13
93	Association between harmful alcohol consumption behavior and dopamine transporter (DAT1) gene polymorphisms in a male Finnish population. Psychiatric Genetics, 2009, 19, 117-125.	0.6	25
94	Association between single nucleotide polymorphisms in the mu opioid receptor gene (OPRM1) and self-reported responses to alcohol in American Indians. BMC Medical Genetics, 2008, 9, 35.	2.1	47
95	Externalizing disorders in American Indians: Comorbidity and a genome wide linkage analysis. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2008, 147B, 690-698.	1.1	28
96	Association study of candidate variants of COMT with neuroticism, anxiety and depression. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2008, 147B, 1314-1318.	1.1	48
97	Testing replication of a 5-SNP set for general cognitive ability in six population samples. European Journal of Human Genetics, 2008, 16, 1388-1395.	1.4	8
98	The Role of GABRA2 in Alcohol Dependence, Smoking, and Illicit Drug Use in an Australian Population Sample. Alcoholism: Clinical and Experimental Research, 2008, 32, 1721-1731.	1.4	61
99	Recently-derived variants of brain-size genes ASPM, MCPH1, CDK5RAP and BRCA1 not associated with general cognition, reading or language. Intelligence, 2008, 36, 689-693.	1.6	18
100	Vitamin D Receptor Gene Polymorphisms Have Negligible Effect on Human Height. Twin Research and Human Genetics, 2008, 11, 488-494.	0.3	11
101	Long-Term Stability and Heritability of Telephone Interview Measures of Alcohol Consumption and Dependence. Twin Research and Human Genetics, 2008, 11, 287-305.	0.3	42
102	Effects ofGABRA2Variation on Physiological, Psychomotor and Subjective Responses in the Alcohol Challenge Twin Study. Twin Research and Human Genetics, 2008, 11, 174-182.	0.3	28
103	The role of aldehyde dehydrogenase-1 (ALDH1A1) polymorphisms in harmful alcohol consumption in a Finnish population. Human Genomics, 2008, 3, 24.	1.4	33
104	Association study of candidate variants from brain-derived neurotrophic factor and dystrobrevin-binding protein 1 with neuroticism, anxiety, and depression. Psychiatric Genetics, 2008, 18, 219-225.	0.6	21
105	The ongoing adaptive evolution of ASPM and Microcephalin is not explained by increased intelligence. Human Molecular Genetics, 2007, 16, 600-608.	1.4	93
106	Association Between Single Nucleotide Polymorphisms in the Cannabinoid Receptor Gene (CNR1) and Impulsivity in Southwest California Indians. Twin Research and Human Genetics, 2007, 10, 805-811.	0.3	66
107	A Haplotype Spanning KIAA0319 and TTRAP Is Associated with Normal Variation in Reading and Spelling Ability. Biological Psychiatry, 2007, 62, 811-817.	0.7	83
108	LRRTM1 on chromosome 2p12 is a maternally suppressed gene that is associated paternally with handedness and schizophrenia. Molecular Psychiatry, 2007, 12, 1129-1139.	4.1	300

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109	LRRTM1 protein is located in the endoplasmic reticulum (ER) in mammalian cells. Molecular Psychiatry, 2007, 12, 1057-1057.	4.1	8
110	Autosomal Linkage Analysis for the Level of Response to Alcohol. Alcoholism: Clinical and Experimental Research, 2005, 29, 1976-1982.	1.4	56
111	Esterase catalysis of substrate vapour: enzyme activity occurs at very low hydration. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2004, 1702, 103-110.	1.1	23