Donald J Macintyre

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/939371/publications.pdf

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55 papers 12,196 citations

201385 27 h-index 52 g-index

62 all docs

62 docs citations

times ranked

62

15581 citing authors

#	Article	IF	Citations
1	Language function following preterm birth: prediction using machine learning. Pediatric Research, 2022, 92, 480-489.	1.1	11
2	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
3	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. Nature Genetics, 2021, 53, 817-829.	9.4	629
4	Treatment resistant depression in the UK: sub-analysis of a European real-world evidence study. BJPsych Open, 2021, 7, S55-S55.	0.3	0
5	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. JAMA Psychiatry, 2021, 78, 1258.	6.0	88
6	A loss of mature microglial markers without immune activation in schizophrenia. Glia, 2021, 69, 1251-1267.	2.5	43
7	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. Biological Psychiatry, 2020, 87, 419-430.	0.7	27
8	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
9	Increased number of T-lymphocytes in post-mortem brain tissue of patients with schizophrenia Schizophrenia Research, 2020, 216, 526-528.	1.1	10
10	A validation of the diathesis-stress model for depression in Generation Scotland. Translational Psychiatry, 2019, 9, 25.	2.4	40
11	Genome-wide by environment interaction studies of depressive symptoms and psychosocial stress in UK Biobank and Generation Scotland. Translational Psychiatry, 2019, 9, 14.	2.4	87
12	Genome-wide association study identifies 30 loci associated with bipolar disorder. Nature Genetics, 2019, 51, 793-803.	9.4	1,191
13	Pharmaco-epidemiology of antidepressant exposure in a UK cohort record-linkage study. Journal of Psychopharmacology, 2019, 33, 482-493.	2.0	11
14	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
15	Cohort profile for the STratifying Resilience and Depression Longitudinally (STRADL) study: A depression-focused investigation of Generation Scotland, using detailed clinical, cognitive, and neuroimaging assessments. Wellcome Open Research, 2019, 4, 185.	0.9	27
16	Self-reported medication use validated through record linkage to national prescribing data. Journal of Clinical Epidemiology, 2018, 94, 132-142.	2.4	75
17	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
18	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

#	Article	IF	Citations
19	Digital Support Platform: a qualitative research study investigating the feasibility of an internet-based, postdiagnostic support platform for families living with dementia. BMJ Open, 2018, 8, e020281.	0.8	15
20	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	6.0	1,085
21	Digital Access in Working-Age and Older Adults and Their Caregivers Attending Psychiatry Outpatient Clinics: Quantitative Survey. JMIR Aging, 2018, 1, e4.	1.4	2
22	A Combined Pathway and Regional Heritability Analysis Indicates NETRIN1 Pathway Is Associated With Major Depressive Disorder. Biological Psychiatry, 2017, 81, 336-346.	0.7	32
23	Genome-wide Association for Major Depression Through Age at Onset Stratification: Major Depressive Disorder Working Group of the Psychiatric Genomics Consortium. Biological Psychiatry, 2017, 81, 325-335.	0.7	175
24	Evaluation of a brief art psychotherapy group for adults suffering from mild to moderate depression: Pilot pre, post and follow-up study. International Journal of Art Therapy: Inscape, 2017, 22, 106-117.	0.6	7
25	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. Biological Psychiatry, 2017, 82, 322-329.	0.7	84
26	Genome-wide Regional Heritability Mapping Identifies a Locus Within the TOX2 Gene Associated With Major Depressive Disorder. Biological Psychiatry, 2017, 82, 312-321.	0.7	26
27	Polygenic risk for alcohol dependence associates with alcohol consumption, cognitive function and social deprivation in a populationâ€based cohort. Addiction Biology, 2016, 21, 469-480.	1.4	27
28	Shared Genetics and Couple-Associated Environment Are Major Contributors to the Risk of Both Clinical and Self-Declared Depression. EBioMedicine, 2016, 14, 161-167.	2.7	32
29	Common polygenic risk for autism spectrum disorder (ASD) is associated with cognitive ability in the general population. Molecular Psychiatry, 2016, 21, 419-425.	4.1	145
30	Altered metabolic parameters in association with antipsychotic medication use in diabetes: A population based case-control study. Psychoneuroendocrinology, 2016, 66, 214-220.	1.3	10
31	Current Versus Lifetime Depression, APOE Variation, and Their Interaction on Cognitive Performance in Younger and Older Adults. Psychosomatic Medicine, 2015, 77, 480-492.	1.3	11
32	Epidemiology and Heritability of Major Depressive Disorder, Stratified by Age of Onset, Sex, and Illness Course in Generation Scotland: Scottish Family Health Study (GS:SFHS). PLoS ONE, 2015, 10, e0142197.	1.1	101
33	The association between lower educational attainment and depression owing to shared genetic effects? Results in ~25 000 subjects. Molecular Psychiatry, 2015, 20, 735-743.	4.1	59
34	Major depressive disorder and current psychological distress moderate the effect of polygenic risk for obesity on body mass index. Translational Psychiatry, 2015, 5, e592-e592.	2.4	24
35	Art psychotherapy practice with adults who suffer from depression in the UK: Qualitative findings from a depression-specific questionnaire. Arts in Psychotherapy, 2014, 41, 563-569.	0.6	11
36	Description of arts therapies practice with adults suffering from depression in the UK: Qualitative findings from the nationwide survey. Arts in Psychotherapy, 2014, 41, 535-544.	0.6	8

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37	708 Common and 2010 rare DISC1 locus variants identified in 1542 subjects: analysis for association with psychiatric disorder and cognitive traits. Molecular Psychiatry, 2014, 19, 668-675.	4.1	59
38	Description of arts therapies practice with adults suffering from depression in the UK: Quantitative results from the nationwide survey. Arts in Psychotherapy, 2013, 40, 458-464.	0.6	17
39	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. Nature Genetics, 2013, 45, 984-994.	9.4	2,067
40	Association of DISC1 variants with age of onset in a population-based sample of recurrent major depression. Molecular Psychiatry, 2013, 18, 745-747.	4.1	14
41	Cohort Profile: Generation Scotland: Scottish Family Health Study (GS:SFHS). The study, its participants and their potential for genetic research on health and illness. International Journal of Epidemiology, 2013, 42, 689-700.	0.9	353
42	Alzheimer's disease risk factor complement receptor 1 is associated with depression. Neuroscience Letters, 2012, 510, 6-9.	1.0	18
43	Genetic variation in Hyperpolarization-activated cyclic nucleotide-gated channels and its relationship with neuroticism, cognition and risk of depression. Frontiers in Genetics, 2012, 3, 116.	1.1	12
44	Genome-wide association study of major depressive disorder: new results, meta-analysis, and lessons learned. Molecular Psychiatry, 2012, 17, 36-48.	4.1	405
45	Facial emotion recognition in Scottish prisoners. International Journal of Law and Psychiatry, 2012, 35, 57-61.	0.5	21
46	Association of GPR50, an X-linked orphan G protein-coupled receptor, and affective disorder in an independent sample of the Scottish population. Neuroscience Letters, 2010, 475, 169-173.	1.0	18
47	Genome-wide association for major depressive disorder: a possible role for the presynaptic protein piccolo. Molecular Psychiatry, 2009, 14, 359-375.	4.1	354
48	A Cytogenetic Abnormality and Rare Coding Variants Identify ABCA13 as a Candidate Gene in Schizophrenia, Bipolar Disorder, and Depression. American Journal of Human Genetics, 2009, 85, 833-846.	2.6	102
49	Collaborative genome-wide association analysis supports a role for ANK3 and CACNA1C in bipolar disorder. Nature Genetics, 2008, 40, 1056-1058.	9.4	1,102
50	Whole-genome association study of bipolar disorder. Molecular Psychiatry, 2008, 13, 558-569.	4.1	642
51	Attitudes of patients and family members towards implantable psychiatric medication. Schizophrenia Research, 2008, 105, 279-286.	1.1	9
52	Wake-up call for British psychiatry. British Journal of Psychiatry, 2008, 193, 6-9.	1.7	183
53	Chromosomal abnormalities and mental illness. Molecular Psychiatry, 2003, 8, 275-287.	4.1	111
54	A GP guide to schizophrenia. Practitioner, 2003, 247, 692-4, 696, 698 passim.	0.3	0

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55	Cohort profile for the STratifying Resilience and Depression Longitudinally (STRADL) study: A depression-focused investigation of Generation Scotland, using detailed clinical, cognitive, and neuroimaging assessments. Wellcome Open Research, 0, 4, 185.	0.9	12