

Denise Harold

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

74
papers

13,527
citations

38
h-index

88
g-index

88
ext. papers

16,531
ext. citations

9.1
avg, IF

4.43
L-index

#	Paper	IF	Citations
74	Meta-analysis of 74,046 individuals identifies 11 new susceptibility loci for Alzheimer's disease. <i>Nature Genetics</i> , 2013 , 45, 1452-8	36.3	2714
73	Genome-wide association study identifies variants at CLU and PICALM associated with Alzheimer's disease. <i>Nature Genetics</i> , 2009 , 41, 1088-93	36.3	2018
72	Common variants at ABCA7, MS4A6A/MS4A4E, EPHA1, CD33 and CD2AP are associated with Alzheimer's disease. <i>Nature Genetics</i> , 2011 , 43, 429-35	36.3	1421
71	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates A β tau, immunity and lipid processing. <i>Nature Genetics</i> , 2019 , 51, 414-430	36.3	917
70	Genome-wide analysis of genetic loci associated with Alzheimer disease. <i>JAMA - Journal of the American Medical Association</i> , 2010 , 303, 1832-40	27.4	888
69	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , 2018 , 360,	33.3	666
68	Rare coding variants in PLCG2, ABI3, and TREM2 implicate microglial-mediated innate immunity in Alzheimer's disease. <i>Nature Genetics</i> , 2017 , 49, 1373-1384	36.3	508
67	Identification of Genetic Factors that Modify Clinical Onset of Huntington's Disease. <i>Cell</i> , 2015 , 162, 516-526	36.2	378
66	Genetic evidence implicates the immune system and cholesterol metabolism in the aetiology of Alzheimer's disease. <i>PLoS ONE</i> , 2010 , 5, e13950	3.7	276
65	GWAS of cerebrospinal fluid tau levels identifies risk variants for Alzheimer's disease. <i>Neuron</i> , 2013 , 78, 256-68	13.9	255
64	Increased expression of BIN1 mediates Alzheimer genetic risk by modulating tau pathology. <i>Molecular Psychiatry</i> , 2013 , 18, 1225-34	15.1	251
63	Strong evidence that KIAA0319 on chromosome 6p is a susceptibility gene for developmental dyslexia. <i>American Journal of Human Genetics</i> , 2005 , 76, 581-91	11	232
62	Common polygenic variation enhances risk prediction for Alzheimer's disease. <i>Brain</i> , 2015 , 138, 3673-84	11.2	227
61	Remapping the insulin gene/IDDM2 locus in type 1 diabetes. <i>Diabetes</i> , 2004 , 53, 1884-9	0.9	180
60	A novel Alzheimer disease locus located near the gene encoding tau protein. <i>Molecular Psychiatry</i> , 2016 , 21, 108-17	15.1	175
59	Estimation and partitioning of polygenic variation captured by common SNPs for Alzheimer's disease, multiple sclerosis and endometriosis. <i>Human Molecular Genetics</i> , 2013 , 22, 832-41	5.6	147
58	Convergent genetic and expression data implicate immunity in Alzheimer's disease. <i>Alzheimer's & Dementia</i> , 2015 , 11, 658-71	1.2	146

57	Meta-analysis of the association between variants in SORL1 and Alzheimer disease. <i>Archives of Neurology</i> , 2011 , 68, 99-106		135
56	Further evidence that the KIAA0319 gene confers susceptibility to developmental dyslexia. <i>Molecular Psychiatry</i> , 2006 , 11, 1085-91, 1061	15.1	119
55	Genetic overlap between Alzheimer's disease and Parkinson's disease at the MAPT locus. <i>Molecular Psychiatry</i> , 2015 , 20, 1588-95	15.1	107
54	Characterization of a family with rare deletions in CNTNAP5 and DOCK4 suggests novel risk loci for autism and dyslexia. <i>Biological Psychiatry</i> , 2010 , 68, 320-8	7.9	103
53	Genome-wide association study of Alzheimer's disease with psychotic symptoms. <i>Molecular Psychiatry</i> , 2012 , 17, 1316-27	15.1	90
52	SNPs associated with cerebrospinal fluid phospho-tau levels influence rate of decline in Alzheimer's disease. <i>PLoS Genetics</i> , 2010 , 6, e1001101	6	90
51	Gene-wide analysis detects two new susceptibility genes for Alzheimer's disease. <i>PLoS ONE</i> , 2014 , 9, e94661	3.7	90
50	Missense variant in TREML2 protects against Alzheimer's disease. <i>Neurobiology of Aging</i> , 2014 , 35, 1510.e19-2684	3.7	90
49	Determining SNP allele frequencies in DNA pools. <i>BioTechniques</i> , 2000 , 28, 464-6, 468, 470	2.5	83
48	Genome-wide haplotype association study identifies the FRMD4A gene as a risk locus for Alzheimer's disease. <i>Molecular Psychiatry</i> , 2013 , 18, 461-70	15.1	77
47	Follow-up of loci from the International Genomics of Alzheimer's Disease Project identifies TRIP4 as a novel susceptibility gene. <i>Translational Psychiatry</i> , 2014 , 4, e358	8.6	70
46	Alzheimer's disease genetics: current knowledge and future challenges. <i>International Journal of Geriatric Psychiatry</i> , 2011 , 26, 793-802	3.9	69
45	A modifier of Huntington's disease onset at the MLH1 locus. <i>Human Molecular Genetics</i> , 2017 , 26, 3859-3867	3.67	59
44	Genetic evidence for the involvement of lipid metabolism in Alzheimer's disease. <i>Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids</i> , 2010 , 1801, 754-61	5	52
43	Genome-wide association interaction analysis for Alzheimer's disease. <i>Neurobiology of Aging</i> , 2014 , 35, 2436-2443	5.6	49
42	The role of variation at AβP, PSEN1, PSEN2, and MAPT in late onset Alzheimer's disease. <i>Journal of Alzheimer's Disease</i> , 2012 , 28, 377-87	4.3	47
41	No support for association between dyslexia susceptibility 1 candidate 1 and developmental dyslexia. <i>Molecular Psychiatry</i> , 2005 , 10, 237-8	15.1	47
40	Analysis of genome-wide association studies of Alzheimer disease and of Parkinson disease to determine if these 2 diseases share a common genetic risk. <i>JAMA Neurology</i> , 2013 , 70, 1268-76	17.2	46

39	Shared genetic contribution to Ischaemic Stroke and Alzheimer's Disease. <i>Annals of Neurology</i> , 2016 , 79, 739-747	9.4	42
38	Suggestive synergy between genetic variants in TF and HFE as risk factors for Alzheimer's disease. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2010 , 153B, 955-9	3.5	39
37	Alpha-T-catenin is expressed in human brain and interacts with the Wnt signaling pathway but is not responsible for linkage to chromosome 10 in Alzheimer's disease. <i>NeuroMolecular Medicine</i> , 2004 , 5, 133-46	4.6	38
36	No consistent evidence for association between mtDNA variants and Alzheimer disease. <i>Neurology</i> , 2012 , 78, 1038-42	6.5	37
35	Association analysis of 528 intra-genic SNPs in a region of chromosome 10 linked to late onset Alzheimer's disease. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008 , 147B, 727-31	3.5	35
34	A single nucleotide polymorphism in CHAT influences response to acetylcholinesterase inhibitors in Alzheimer's disease. <i>Pharmacogenetics and Genomics</i> , 2006 , 16, 75-7	1.9	31
33	Sequence variation in the CHAT locus shows no association with late-onset Alzheimer's disease. <i>Human Genetics</i> , 2003 , 113, 258-67	6.3	31
32	Polymorphisms in BACE2 may affect the age of onset Alzheimer's dementia in Down syndrome. <i>Neurobiology of Aging</i> , 2014 , 35, 1513.e1-5	5.6	30
31	Evidence that PICALM affects age at onset of Alzheimer's dementia in Down syndrome. <i>Neurobiology of Aging</i> , 2013 , 34, 2441.e1-5	5.6	30
30	Genetic variation at the CELF1 (CUGBP, elav-like family member 1 gene) locus is genome-wide associated with Alzheimer's disease and obesity. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2014 , 165B, 283-93	3.5	28
29	Can Studies of Neuroinflammation in a TSPO Genetic Subgroup (HAB or MAB) Be Applied to the Entire AD Cohort?. <i>Journal of Nuclear Medicine</i> , 2015 , 56, 707-13	8.9	26
28	A genome-wide study shows a limited contribution of rare copy number variants to Alzheimer's disease risk. <i>Human Molecular Genetics</i> , 2013 , 22, 816-24	5.6	26
27	Cognitive analysis of schizophrenia risk genes that function as epigenetic regulators of gene expression. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2016 , 171, 1170-1179	3.5	25
26	A review of pharmaceutical occurrence and pathways in the aquatic environment in the context of a changing climate and the COVID-19 pandemic. <i>Analytical Methods</i> , 2021 , 13, 575-594	3.2	25
25	The Genetic Modifiers of Motor OnsetAge[GeM MOA) Website: Genome-wide Association Analysis for Genetic Modifiers of Huntington's Disease. <i>Journal of Huntington's Disease</i> , 2015 , 4, 279-84	1.9	20
24	Alzheimer's disease and age-related macular degeneration have different genetic models for complement gene variation. <i>Neurobiology of Aging</i> , 2012 , 33, 1843.e9-17	5.6	19
23	MiR-137-derived polygenic risk: effects on cognitive performance in patients with schizophrenia and controls. <i>Translational Psychiatry</i> , 2017 , 7, e1012	8.6	18
22	Genetically predicted complement component 4A expression: effects on memory function and middle temporal lobe activation. <i>Psychological Medicine</i> , 2018 , 48, 1608-1615	6.9	18

21	Interaction between the ADAM12 and SH3MD1 genes may confer susceptibility to late-onset Alzheimer's disease. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2007 , 144B, 448-52	3.5	18
20	Deep sequencing of the Nicastrin gene in pooled DNA, the identification of genetic variants that affect risk of Alzheimer's disease. <i>PLoS ONE</i> , 2011 , 6, e17298	3.7	18
19	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	8.7	16
18	No evidence that extended tracts of homozygosity are associated with Alzheimer's disease. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2011 , 156B, 764-71	3.5	15
17	Haplotype-based stratification of Huntington's disease. <i>European Journal of Human Genetics</i> , 2017 , 25, 1202-1209	5.3	14
16	Meta-analysis of genetic association with diagnosed Alzheimer's disease identifies novel risk loci and implicates Abeta, Tau, immunity and lipid processing		9
15	Beyond C4: Analysis of the complement gene pathway shows enrichment for IQ in patients with psychotic disorders and healthy controls. <i>Genes, Brain and Behavior</i> , 2019 , 18, e12602	3.6	8
14	Characterisation and validation of insertions and deletions in 173 patient exomes. <i>PLoS ONE</i> , 2012 , 7, e51292	3.7	8
13	Linkage disequilibrium structure of KIAA0319 and DCDC2, two candidate susceptibility genes for developmental dyslexia. <i>Molecular Psychiatry</i> , 2006 , 11, 1061-1061	15.1	8
12	Identifying schizophrenia patients who carry pathogenic genetic copy number variants using standard clinical assessment: retrospective cohort study. <i>British Journal of Psychiatry</i> , 2020 , 216, 275-279	5.4	7
11	Effects of MiR-137 genetic risk score on brain volume and cortical measures in patients with schizophrenia and controls. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018 , 177, 369-376	3.5	6
10	Rare Copy Number Variants Are Associated With Poorer Cognition in Schizophrenia. <i>Biological Psychiatry</i> , 2021 , 90, 28-34	7.9	5
9	From molecule to clinic and community for neurodegeneration: research to bridge translational gaps. <i>Journal of Alzheimer's Disease</i> , 2013 , 33 Suppl 1, S385-96	4.3	4
8	Population-based identity-by-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	3.5	2
7	Biomarkers for Psychosis: the Molecular Genetics of Psychosis. <i>Current Behavioral Neuroscience Reports</i> , 2015 , 2, 112-118	1.7	1
6	A03 Genetic Modifiers Affecting The Age At Motor Onset In Huntington's Disease. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2014 , 85, A1-A2	5.5	1
5	Monitoring of emerging contaminants of concern in the aquatic environment: a review of studies showing the application of effect-based measures. <i>Analytical Methods</i> , 2021 , 13, 5120-5143	3.2	1
4	Effects of complement gene-set polygenic risk score on brain volume and cortical measures in patients with psychotic disorders and healthy controls. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2020 , 183, 445-453	3.5	1

- 3 The Differential Influence of Immune, Endocytotic, and Lipid Metabolism Genes on Amyloid Deposition and Neurodegeneration in Subjects at Risk of Alzheimer's Disease. *Journal of Alzheimer's Disease*, **2021**, 79, 127-139 4.3 1
- 2 RNA-seq analysis of murine peyer's patches at 6 and 18 h post infection with *Fasciola hepatica* metacecariae.. *Veterinary Parasitology*, **2021**, 302, 109643 2.8
- 1 Genetically elevated high-density lipoprotein cholesterol through the cholesteryl ester transfer protein gene does not associate with risk of Alzheimer's disease. *Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring*, **2018**, 10, 595-598 5.2