Vincent Chouraki

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Meta-analysis of 74,046 individuals identifies 11 new susceptibility loci for Alzheimer's disease. Nature Genetics, 2013, 45, 1452-1458.	21.4	3,741
2	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates Aβ, tau, immunity and lipid processing. Nature Genetics, 2019, 51, 414-430.	21.4	1,962
3	Common variants at ABCA7, MS4A6A/MS4A4E, EPHA1, CD33 and CD2AP are associated with Alzheimer's disease. Nature Genetics, 2011, 43, 429-435.	21.4	1,708
4	Incidence of Dementia over Three Decades in the Framingham Heart Study. New England Journal of Medicine, 2016, 374, 523-532.	27.0	788
5	Rare coding variants in PLCG2, ABI3, and TREM2 implicate microglial-mediated innate immunity in Alzheimer's disease. Nature Genetics, 2017, 49, 1373-1384.	21.4	783
6	Genetic associations at 53 loci highlight cell types and biological pathways relevant for kidney function. Nature Communications, 2016, 7, 10023.	12.8	412
7	Genetic contributions to variation in general cognitive function: a meta-analysis of genome-wide association studies in the CHARGE consortium (N=53 949). Molecular Psychiatry, 2015, 20, 183-192.	7.9	344
8	A common haplotype lowers PU.1 expression in myeloid cells and delays onset of Alzheimer's disease. Nature Neuroscience, 2017, 20, 1052-1061.	14.8	330
9	A novel Alzheimer disease locus located near the gene encoding tau protein. Molecular Psychiatry, 2016, 21, 108-117.	7.9	260
10	Novel genetic loci associated with hippocampal volume. Nature Communications, 2017, 8, 13624.	12.8	250
11	Novel genetic loci underlying human intracranial volume identified through genome-wide association. Nature Neuroscience, 2016, 19, 1569-1582.	14.8	213
12	Common variants at 12q14 and 12q24 are associated with hippocampal volume. Nature Genetics, 2012, 44, 545-551.	21.4	212
13	Association of branchedâ€chain amino acids and other circulating metabolites with risk of incident dementia and Alzheimer's disease: A prospective study in eight cohorts. Alzheimer's and Dementia, 2018, 14, 723-733.	0.8	182
14	Convergent genetic and expression data implicate immunity in Alzheimer's disease. Alzheimer's and Dementia, 2015, 11, 658-671.	0.8	173
15	Genome-Wide Association and Functional Follow-Up Reveals New Loci for Kidney Function. PLoS Genetics, 2012, 8, e1002584.	3.5	166
16	Multiethnic Genome-Wide Association Study of Cerebral White Matter Hyperintensities on MRI. Circulation: Cardiovascular Genetics, 2015, 8, 398-409.	5.1	162
17	Genetics of Alzheimer's Disease. Advances in Genetics, 2014, 87, 245-294.	1.8	159
18	Gene-Wide Analysis Detects Two New Susceptibility Genes for Alzheimer's Disease. PLoS ONE, 2014, 9, e94661	2.5	155

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19	Implication of the Immune System in Alzheimer's Disease: Evidence from Genome-Wide Pathway Analysis. Journal of Alzheimer's Disease, 2010, 20, 1107-1118.	2.6	152
20	Circulating metabolites and general cognitive ability and dementia: Evidence from 11 cohort studies. Alzheimer's and Dementia, 2018, 14, 707-722.	0.8	143
21	The changing pattern of Crohn's disease incidence in northern France: a continuing increase in the 10- to 19-year-old age bracket (1988-2007). Alimentary Pharmacology and Therapeutics, 2011, 33, 1133-1142.	3.7	138
22	GWAS for executive function and processing speed suggests involvement of the CADM2 gene. Molecular Psychiatry, 2016, 21, 189-197.	7.9	134
23	Genome-wide Association Studies Identify Genetic Loci Associated With Albuminuria in Diabetes. Diabetes, 2016, 65, 803-817.	0.6	131
24	Genome-wide association study of kidney function decline in individuals of European descent. Kidney International, 2015, 87, 1017-1029.	5.2	113
25	Genome-wide haplotype association study identifies the FRMD4A gene as a risk locus for Alzheimer's disease. Molecular Psychiatry, 2013, 18, 461-470.	7.9	103
26	Plasma amyloidâ€Î² and risk of Alzheimer's disease in the Framingham Heart Study. Alzheimer's and Dementia, 2015, 11, 249.	0.8	101
27	Follow-up of loci from the International Genomics of Alzheimer's Disease Project identifies TRIP4 as a novel susceptibility gene. Translational Psychiatry, 2014, 4, e358-e358.	4.8	98
28	1000 Genomes-based meta-analysis identifies 10 novel loci for kidney function. Scientific Reports, 2017, 7, 45040.	3.3	98
29	Association of amine biomarkers with incident dementia and Alzheimer's disease in the Framingham Study. Alzheimer's and Dementia, 2017, 13, 1327-1336.	0.8	93
30	Evidence of the association of BIN1 and PICALM with the AD risk in contrasting European populations. Neurobiology of Aging, 2011, 32, 756.e11-756.e15.	3.1	82
31	Evaluation of a Genetic Risk Score to Improve Risk Prediction for Alzheimer's Disease. Journal of Alzheimer's Disease, 2016, 53, 921-932.	2.6	77
32	Genome-wide Studies of Verbal Declarative Memory in Nondemented Older People: The Cohorts for Heart and Aging Research in Genomic Epidemiology Consortium. Biological Psychiatry, 2015, 77, 749-763.	1.3	67
33	Systematic Analysis of Candidate Genes for Alzheimer's Disease in a French, Genome-Wide Association Study. Journal of Alzheimer's Disease, 2010, 20, 1181-1188.	2.6	63
34	Shared genetic contribution to ischemic stroke and Alzheimer's disease. Annals of Neurology, 2016, 79, 739-747.	5.3	56
35	PLD3 variants in population studies. Nature, 2015, 520, E2-E3.	27.8	49
36	Rare Functional Variant in TM2D3 is Associated with Late-Onset Alzheimer's Disease. PLoS Genetics, 2016, 12, e1006327.	3.5	47

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37	SUCLG2 identified as both a determinator of CSF Aβ1–42 levels and an attenuator of cognitive decline in Alzheimer's disease. Human Molecular Genetics, 2014, 23, 6644-6658.	2.9	45
38	PLCG2 protective variant p.P522R modulates tau pathology and disease progression in patients with mild cognitive impairment. Acta Neuropathologica, 2020, 139, 1025-1044.	7.7	40
39	A genome-wide association meta-analysis of plasma Aβ peptides concentrations in the elderly. Molecular Psychiatry, 2014, 19, 1326-1335.	7.9	36
40	Plasma clusterin levels and risk of dementia, Alzheimer's disease, and stroke. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2016, 3, 103-109.	2.4	32
41	Genetic and lifestyle risk factors for MRI-defined brain infarcts in a population-based setting. Neurology, 2019, 92, .	1.1	30
42	Plasma amyloid β levels are driven by genetic variants near <i>APOE, BACE1, APP, PSEN2</i> : A genomeâ€wide association study in over 12,000 nonâ€demented participants. Alzheimer's and Dementia, 2021, 17, 1663-1674.	0.8	20
43	Whole exome sequence-based association analyses of plasma amyloid-Î ² in African and European Americans; the Atherosclerosis Risk in Communities-Neurocognitive Study. PLoS ONE, 2017, 12, e0180046.	2.5	18
44	Smoking habits, waist circumference and coronary artery disease risk relationship: the PRIME study. European Journal of Cardiovascular Prevention and Rehabilitation, 2008, 15, 625-630.	2.8	15
45	Meta-analysis of genome-wide association studies identifies ancestry-specific associations underlying circulating total tau levels. Communications Biology, 2022, 5, 336.	4.4	6
46	Genetically elevated highâ€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.	2.4	2
47	O4-05-03: Whole exome sequence analysis of white matter hyperintensities on cranial MRI. , 2015, 11, P278-P279.		1
48	DT-02-02: Low-frequency variant imputation identifies rare variant candidate loci in a gwas of late-onset Alzheimer's disease in the igap consortium. , 2015, 11, P333-P334.		1
49	Genomeâ€wide metaâ€analysis of lateâ€onset Alzheimer's disease using rare variant imputation in 65,602 subjects identifies risk loci with roles in memory, neurodevelopment, and cardiometabolic traits: The international genomics of Alzheimer's project (IGAP). Alzheimer's and Dementia, 2020, 16, e044193.	0.8	1
50	The changing epidemiology of paediatric inflammatory bowel disease: authors' reply. Alimentary Pharmacology and Therapeutics, 2011, 33, 1381-1382.	3.7	0
51	O4-02-01: Plasma clusterin levels and risk of dementia and Alzheimer's disease: The Framingham Heart Study. , 2013, 9, P681-P681.		0
52	O4-05-02: Genome-wide association study of lobar brain volumes. , 2015, 11, P278-P278.		0
53	O1-04-06: Association of plasma biomarkers with risk of incident dementia in the framingham heart study: A metabolomics approach. , 2015, 11, P134-P135.		0
54	O2â€10â€06: A Common Allele in <i>SPI1</i> Lowers Risk and Delays Age at Onset for Alzheimer's Disease. Alzheimer's and Dementia, 2016, 12, P253.	0.8	0

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55	Identification of hippocampal volume as a mediator of the association between APOE4 and dementia. Alzheimer's and Dementia, 2020, 16, e047425.	0.8	0