

# Rahul S Desikan

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4383573/publications.pdf>

Version: 2024-02-01

63  
papers

5,918  
citations

109321

35  
h-index

133252

59  
g-index

73  
all docs

73  
docs citations

73  
times ranked

10849  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide meta-analysis identifies new loci and functional pathways influencing Alzheimer's disease risk. <i>Nature Genetics</i> , 2019, 51, 404-413.	21.4	1,625
2	Improved Detection of Common Variants Associated with Schizophrenia by Leveraging Pleiotropy with Cardiovascular-Disease Risk Factors. <i>American Journal of Human Genetics</i> , 2013, 92, 197-209.	6.2	422
3	Genetic assessment of age-associated Alzheimer disease risk: Development and validation of a polygenic hazard score. <i>PLoS Medicine</i> , 2017, 14, e1002258.	8.4	311
4	Improved Detection of Common Variants Associated with Schizophrenia and Bipolar Disorder Using Pleiotropy-Informed Conditional False Discovery Rate. <i>PLoS Genetics</i> , 2013, 9, e1003455.	3.5	298
5	Genome-wide Pleiotropy Between Parkinson Disease and Autoimmune Diseases. <i>JAMA Neurology</i> , 2017, 74, 780.	9.0	245
6	Association Between Genetic Traits for Immune-Mediated Diseases and Alzheimer Disease. <i>JAMA Neurology</i> , 2016, 73, 691.	9.0	151
7	Polygenic Overlap Between C-Reactive Protein, Plasma Lipids, and Alzheimer Disease. <i>Circulation</i> , 2015, 131, 2061-2069.	1.6	145
8	Magnetic resonance imaging in Alzheimer's Disease Neuroimaging Initiative 2. <i>Alzheimer's and Dementia</i> , 2015, 11, 740-756.	0.8	142
9	What Are Polygenic Scores and Why Are They Important?. <i>JAMA - Journal of the American Medical Association</i> , 2019, 321, 1820.	7.4	125
10	Amyloid- $\beta$ -Associated Clinical Decline Occurs Only in the Presence of Elevated P-tau. <i>Archives of Neurology</i> , 2012, 69, 709-13.	4.5	122
11	Beyond SNP heritability: Polygenicity and discoverability of phenotypes estimated with a univariate Gaussian mixture model. <i>PLoS Genetics</i> , 2020, 16, e1008612.	3.5	120
12	Immune-related genetic enrichment in frontotemporal dementia: An analysis of genome-wide association studies. <i>PLoS Medicine</i> , 2018, 15, e1002487.	8.4	111
13	Amyloid- $\beta$ associated volume loss occurs only in the presence of phospho-tau. <i>Annals of Neurology</i> , 2011, 70, 657-661.	5.3	109
14	Genetic architecture of sporadic frontotemporal dementia and overlap with Alzheimer's and Parkinson's diseases. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2017, 88, 152-164.	1.9	107
15	Dissecting the genetic relationship between cardiovascular risk factors and Alzheimer's disease. <i>Acta Neuropathologica</i> , 2019, 137, 209-226.	7.7	100
16	Malformations of cortical development. <i>Annals of Neurology</i> , 2016, 80, 797-810.	5.3	95
17	Genetic Markers of Human Evolution Are Enriched in Schizophrenia. <i>Biological Psychiatry</i> , 2016, 80, 284-292.	1.3	92
18	Shared genetic risk between corticobasal degeneration, progressive supranuclear palsy, and frontotemporal dementia. <i>Acta Neuropathologica</i> , 2017, 133, 825-837.	7.7	90

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19	Identifying Common Genetic Variants in Blood Pressure Due to Polygenic Pleiotropy With Associated Phenotypes. <i>Hypertension</i> , 2014, 63, 819-826.	2.7	83
20	Polygenic hazard score: an enrichment marker for Alzheimer's associated amyloid and tau deposition. <i>Acta Neuropathologica</i> , 2018, 135, 85-93.	7.7	80
21	Selective Genetic Overlap Between Amyotrophic Lateral Sclerosis and Diseases of the Frontotemporal Dementia Spectrum. <i>JAMA Neurology</i> , 2018, 75, 860.	9.0	79
22	Fine-mapping of the human leukocyte antigen locus as a risk factor for Alzheimer disease: A case-control study. <i>PLoS Medicine</i> , 2017, 14, e1002272.	8.4	67
23	The Role of Clusterin in Amyloid- $\beta$ -Associated Neurodegeneration. <i>JAMA Neurology</i> , 2014, 71, 180.	9.0	66
24	CXCR4 involvement in neurodegenerative diseases. <i>Translational Psychiatry</i> , 2018, 8, 73.	4.8	66
25	Polygenic hazard score, amyloid deposition and Alzheimer's neurodegeneration. <i>Brain</i> , 2019, 142, 460-470.	7.6	63
26	The relationship between complement factor C3, APOE $\epsilon$ 4, amyloid and tau in Alzheimer's disease. <i>Acta Neuropathologica Communications</i> , 2016, 4, 65.	5.2	60
27	Challenges in pediatric neuroimaging. <i>NeuroImage</i> , 2019, 185, 793-801.	4.2	54
28	Pediatric neuro MRI: tricks to minimize sedation. <i>Pediatric Radiology</i> , 2018, 48, 50-55.	2.0	53
29	Genome-wide Association Analysis of Parkinson's Disease and Schizophrenia Reveals Shared Genetic Architecture and Identifies Novel Risk Loci. <i>Biological Psychiatry</i> , 2021, 89, 227-235.	1.3	53
30	Identifying Novel Gene Variants in Coronary Artery Disease and Shared Genes With Several Cardiovascular Risk Factors. <i>Circulation Research</i> , 2016, 118, 83-94.	4.5	52
31	Heart fatty acid binding protein and $A\beta$ -associated Alzheimer's neurodegeneration. <i>Molecular Neurodegeneration</i> , 2013, 8, 39.	10.8	49
32	Polygenic hazard scores in preclinical Alzheimer disease. <i>Annals of Neurology</i> , 2017, 82, 484-488.	5.3	49
33	Meta-analysis of Alzheimer's disease on 9,751 samples from Norway and IGAP study identifies four risk loci. <i>Scientific Reports</i> , 2018, 8, 18088.	3.3	47
34	Age-dependent effects of APOE $\epsilon$ 4 in preclinical Alzheimer's disease. <i>Annals of Clinical and Translational Neurology</i> , 2016, 3, 668-677.	3.7	46
35	Sex-dependent autosomal effects on clinical progression of Alzheimer's disease. <i>Brain</i> , 2020, 143, 2272-2280.	7.6	46
36	Shared common variants in prostate cancer and blood lipids. <i>International Journal of Epidemiology</i> , 2014, 43, 1205-1214.	1.9	45

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37	Combining Polygenic Hazard Score With Volumetric MRI and Cognitive Measures Improves Prediction of Progression From Mild Cognitive Impairment to Alzheimer's Disease. <i>Frontiers in Neuroscience</i> , 2018, 12, 260.	2.8	41
38	Abundant Genetic Overlap between Blood Lipids and Immune-Mediated Diseases Indicates Shared Molecular Genetic Mechanisms. <i>PLoS ONE</i> , 2015, 10, e0123057.	2.5	40
39	Identification of genetic heterogeneity of Alzheimer's disease across age. <i>Neurobiology of Aging</i> , 2019, 84, 243.e1-243.e9.	3.1	34
40	Identification of shared genetic variants between schizophrenia and lung cancer. <i>Scientific Reports</i> , 2018, 8, 674.	3.3	33
41	Large-scale genomics unveil polygenic architecture of human cortical surface area. <i>Nature Communications</i> , 2015, 6, 7549.	12.8	30
42	Entorhinal Cortex: Antemortem Cortical Thickness and Postmortem Neurofibrillary Tangles and Amyloid Pathology. <i>American Journal of Neuroradiology</i> , 2017, 38, 961-965.	2.4	30
43	Genetic overlap between multiple sclerosis and several cardiovascular disease risk factors. <i>Multiple Sclerosis Journal</i> , 2016, 22, 1783-1793.	3.0	25
44	Insulin-Like Growth Factor Binding Protein 2 Is Associated With Biomarkers of Alzheimer's Disease Pathology and Shows Differential Expression in Transgenic Mice. <i>Frontiers in Neuroscience</i> , 2018, 12, 476.	2.8	25
45	Genetic variation of oxidative phosphorylation genes in stroke and Alzheimer's disease. <i>Neurobiology of Aging</i> , 2014, 35, 1956.e1-1956.e8.	3.1	17
46	Microstructure of the Default Mode Network in Preterm Infants. <i>American Journal of Neuroradiology</i> , 2017, 38, 343-348.	2.4	17
47	Probing the Association between Early Evolutionary Markers and Schizophrenia. <i>PLoS ONE</i> , 2017, 12, e0169227.	2.5	17
48	Regionally specific TSC1 and TSC2 gene expression in tuberous sclerosis complex. <i>Scientific Reports</i> , 2018, 8, 13373.	3.3	13
49	Protein network analysis reveals selectively vulnerable regions and biological processes in FTD. <i>Neurology: Genetics</i> , 2018, 4, e266.	1.9	12
50	Abnormal Morphology of Select Cortical and Subcortical Regions in Neurofibromatosis Type 1. <i>Radiology</i> , 2018, 289, 499-508.	7.3	12
51	Linking tuberous sclerosis complex, excessive mTOR signaling, and age-related neurodegeneration: a new association between TSC1 mutation and frontotemporal dementia. <i>Acta Neuropathologica</i> , 2017, 134, 813-816.	7.7	11
52	The genetic architecture of human complex phenotypes is modulated by linkage disequilibrium and heterozygosity. <i>Genetics</i> , 2021, 217, .	2.9	10
53	Genetic variation across RNA metabolism and cell death gene networks is implicated in the semantic variant of primary progressive aphasia. <i>Scientific Reports</i> , 2019, 9, 10854.	3.3	9
54	Interpreting Alzheimer disease polygenic scores. <i>Annals of Neurology</i> , 2018, 83, 443-445.	5.3	6

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55	Hazards of Neurological Nomenclature. JAMA Neurology, 2017, 74, 1165.	9.0	5
56	Precision neuroradiology: mapping the nodes and networks that link genes to behaviour. British Journal of Radiology, 2019, 92, 20190093.	2.2	3
57	Association of Alzheimer Disease Susceptibility Variants and Gene Expression in the Human Brain—Reply. JAMA Neurology, 2016, 73, 1255.	9.0	1
58	Title is missing!. , 2020, 16, e1008612.		0
59	Title is missing!. , 2020, 16, e1008612.		0
60	Title is missing!. , 2020, 16, e1008612.		0
61	Title is missing!. , 2020, 16, e1008612.		0
62	Title is missing!. , 2020, 16, e1008612.		0
63	Title is missing!. , 2020, 16, e1008612.		0