

Muralidharan Sargurupremraj

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

4,092
citations

430874

18
h-index

552781

26
g-index

33
all docs

33
docs citations

33
times ranked

8303
citing authors

#	ARTICLE	IF	CITATIONS
1	Multiancestry genome-wide association study of 520,000 subjects identifies 32 loci associated with stroke and stroke subtypes. <i>Nature Genetics</i> , 2018, 50, 524-537.	21.4	1,124
2	Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits. <i>Nature Genetics</i> , 2018, 50, 1412-1425.	21.4	924
3	Study of 300,486 individuals identifies 148 independent genetic loci influencing general cognitive function. <i>Nature Communications</i> , 2018, 9, 2098.	12.8	484
4	Trans-ethnic association study of blood pressure determinants in over 750,000 individuals. <i>Nature Genetics</i> , 2019, 51, 51-62.	21.4	328
5	Genetic architecture of subcortical brain structures in 38,851 individuals. <i>Nature Genetics</i> , 2019, 51, 1624-1636.	21.4	192
6	Genome-wide association study of intracranial aneurysms identifies 17 risk loci and genetic overlap with clinical risk factors. <i>Nature Genetics</i> , 2020, 52, 1303-1313.	21.4	163
7	Identification of additional risk loci for stroke and small vessel disease: a meta-analysis of genome-wide association studies. <i>Lancet Neurology</i> , The, 2016, 15, 695-707.	10.2	130
8	GWAS and colocalization analyses implicate carotid intima-media thickness and carotid plaque loci in cardiovascular outcomes. <i>Nature Communications</i> , 2018, 9, 5141.	12.8	119
9	Cerebral small vessel disease genomics and its implications across the lifespan. <i>Nature Communications</i> , 2020, 11, 6285.	12.8	89
10	Genome-wide meta-analysis identifies 3 novel loci associated with stroke. <i>Annals of Neurology</i> , 2018, 84, 934-939.	5.3	79
11	Genome-wide association study of cerebral small vessel disease reveals established and novel loci. <i>Brain</i> , 2019, 142, 3176-3189.	7.6	76
12	Common Genetic Variation Indicates Separate Causes for Periventricular and Deep White Matter Hyperintensities. <i>Stroke</i> , 2020, 51, 2111-2121.	2.0	71
13	Burden of Dilated Perivascular Spaces, an Emerging Marker of Cerebral Small Vessel Disease, Is Highly Heritable. <i>Stroke</i> , 2018, 49, 282-287.	2.0	62
14	Genome-wide association studies in asthma. <i>Current Opinion in Allergy and Clinical Immunology</i> , 2013, 13, 112-118.	2.3	39
15	A genome-wide association study identifies new loci for factor VII and implicates factor VII in ischemic stroke etiology. <i>Blood</i> , 2019, 133, 967-977.	1.4	34
16	Genetic and lifestyle risk factors for MRI-defined brain infarcts in a population-based setting. <i>Neurology</i> , 2019, 92, .	1.1	30
17	Genetics of common cerebral small vessel disease. <i>Nature Reviews Neurology</i> , 2022, 18, 84-101.	10.1	30
18	Migraine, Stroke, and Cervical Arterial Dissection. <i>Neurology: Genetics</i> , 2022, 8, 00.	1.9	18

#	ARTICLE	IF	CITATIONS
19	Circulating Metabolome and White Matter Hyperintensities in Women and Men. <i>Circulation</i> , 2022, 145, 1040-1052.	1.6	17
20	Multi-phenotype analyses of hemostatic traits with cardiovascular events reveal novel genetic associations. <i>Journal of Thrombosis and Haemostasis</i> , 2022, 20, 1331-1349.	3.8	12
21	Transposable elements and their potential role in complex lung disorder. <i>Respiratory Research</i> , 2013, 14, 99.	3.6	9
22	Intermediary quantitative traits—an alternative in the identification of disease genes in asthma?. <i>Genes and Immunity</i> , 2014, 15, 1-7.	4.1	6
23	Gene-mapping study of extremes of cerebral small vessel disease reveals TRIM47 as a strong candidate. <i>Brain</i> , 2022, 145, 1992-2007.	7.6	6
24	Genetics of patent foramen ovale—NKX2-5 and beyond. <i>Clinical Neurology and Neurosurgery</i> , 2010, 112, 457-458.	1.4	4
25	Genomic Studies Across the Lifespan Point to Early Mechanisms Determining Subcortical Volumes. <i>Biological Psychiatry: Cognitive Neuroscience and Neuroimaging</i> , 2022, 7, 616-628.	1.5	1
26	Comparing The Full Genome Sequence Derived Of Blood And Bronchial Brush Cells From COPD Patients. , 2012, , .		0
27	P1—Large-Scale Meta-Analysis of Genome-Wide Association Data on Delayed Recall Memory Performance: The Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium. <i>Alzheimer's and Dementia</i> , 2016, 12, P406.	0.8	0