

# Namitha Mohandas

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11686793/publications.pdf>

Version: 2024-02-01

14

papers

326

citations

759233

12

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1058476

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docs citations

14

times ranked

517

citing authors

#	ARTICLE	IF	CITATIONS
1	Evidence for type-specific DNA methylation patterns in epilepsy: a discordant monozygotic twin approach. <i>Epigenomics</i> , 2019, 11, 951-968.	2.1	19
2	Deciphering the role of epigenetics in self-limited epilepsy with centrot temporal spikes. <i>Epilepsy Research</i> , 2019, 156, 106163.	1.6	5
3	Epigenome-wide analysis in newborn blood spots from monozygotic twins discordant for cerebral palsy reveals consistent regional differences in DNA methylation. <i>Clinical Epigenetics</i> , 2018, 10, 25.	4.1	47
4	Exploring molecular variation in <i>Schistosoma japonicum</i> in China. <i>Scientific Reports</i> , 2015, 5, 17345.	3.3	33
5	The mitochondrial genome of <i>Angiostrongylus mackerrasae</i> as a basis for molecular, epidemiological and population genetic studies. <i>Parasites and Vectors</i> , 2015, 8, 473.	2.5	16
6	The barber's pole worm CAP protein superfamily – A basis for fundamental discovery and biotechnology advances. <i>Biotechnology Advances</i> , 2015, 33, 1744-1754.	11.7	16
7	The mitochondrial genome of <i>Parascaris univalens</i> - implications for a "forgotten" parasite. <i>Parasites and Vectors</i> , 2014, 7, 428.	2.5	30
8	Mitochondrial genomes of <i>Anisakis simplex</i> and <i>Contra caecum osculatum</i> ( <i>sensu stricto</i> ) – Comparisons with selected nematodes. <i>Infection, Genetics and Evolution</i> , 2014, 21, 452-462.	2.3	25
9	Mitochondrial genomes of <i>Trichinella</i> species and genotypes – a basis for diagnosis, and systematic and epidemiological explorations. <i>International Journal for Parasitology</i> , 2014, 44, 1073-1080.	3.1	40
10	Characterisation of the mitochondrial genome of <i>Parafilaroides normani</i> (lungworm) of <i>Arctocephalus pusillus doriferus</i> (Australian fur seal). <i>Parasitology Research</i> , 2014, 113, 3049-3055.	1.6	9
11	The mitochondrial genome of <i>Aelurostrongylus abstrusus</i> – diagnostic, epidemiological and systematic implications. <i>Gene</i> , 2013, 516, 294-300.	2.2	17
12	The mitochondrial genome of <i>Protostrongylus rufescens</i> – implications for population and systematic studies. <i>Parasites and Vectors</i> , 2013, 6, 263.	2.5	18
13	Analyses of mitochondrial amino acid sequence datasets support the proposal that specimens of <i>Hypodontus macropi</i> from three species of macropodid hosts represent distinct species. <i>BMC Evolutionary Biology</i> , 2013, 13, 259.	3.2	17
14	Mitochondrial genome of <i>Angiostrongylus vasorum</i> : Comparison with congeners and implications for studying the population genetics and epidemiology of this parasite. <i>Infection, Genetics and Evolution</i> , 2012, 12, 1884-1891.	2.3	34