

# Mathangi Thiagarajan

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12185497/publications.pdf>

Version: 2024-02-01

11

papers

2,665

citations

840776

11

h-index

1281871

11

g-index

11

all docs

11

docs citations

11

times ranked

5152

citing authors

#	ARTICLE	IF	CITATIONS
1	Metabolic Reconstruction for Metagenomic Data and Its Application to the Human Microbiome. PLoS Computational Biology, 2012, 8, e1002358.	3.2	939
2	Macronuclear Genome Sequence of the Ciliate <i>Tetrahymena thermophila</i> , a Model Eukaryote. PLoS Biology, 2006, 4, e286.	5.6	657
3	Targeted metagenomics and ecology of globally important uncultured eukaryotic phytoplankton. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 14679-14684.	7.1	257
4	Functional Tradeoffs Underpin Salinity-Driven Divergence in Microbial Community Composition. PLoS ONE, 2014, 9, e89549.	2.5	184
5	A Metagenomic Framework for the Study of Airborne Microbial Communities. PLoS ONE, 2013, 8, e81862.	2.5	127
6	Influence of nutrients and currents on the genomic composition of microbes across an upwelling mosaic. ISME Journal, 2012, 6, 1403-1414.	9.8	120
7	METAREP: JCVI metagenomics reports—“an open source tool for high-performance comparative metagenomics. Bioinformatics, 2010, 26, 2631-2632.	4.1	118
8	Metagenomic Exploration of Viruses throughout the Indian Ocean. PLoS ONE, 2012, 7, e42047.	2.5	113
9	Refined annotation and assembly of the <i>Tetrahymena thermophila</i> genome sequence through EST analysis, comparative genomic hybridization, and targeted gap closure. BMC Genomics, 2008, 9, 562.	2.8	84
10	The JCVI standard operating procedure for annotating prokaryotic metagenomic shotgun sequencing data. Standards in Genomic Sciences, 2010, 2, 229-237.	1.5	55
11	A Case Study for Large-Scale Human Microbiome Analysis Using JCVI’s Metagenomics Reports (METAREP). PLoS ONE, 2012, 7, e29044.	2.5	11