

# Priya Narasingarao

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12062315/publications.pdf>

Version: 2024-02-01

10

papers

777

citations

1040056

9

h-index

1372567

10

g-index

10

all docs

10

docs citations

10

times ranked

1130

citing authors

#	ARTICLE	IF	CITATIONS
1	High-quality draft genome sequence of <i>Sedimenticola selenatireducens</i> strain AK4OH1T, a gamma-proteobacterium isolated from estuarine sediment. <i>Standards in Genomic Sciences</i> , 2016, 11, 66.	1.5	5
2	Draft Genome Sequence of ‘ <i>Candidatus Halobonum tyrricense</i> ’ Strain G22, Isolated from the Hypersaline Waters of Lake Tyrrell, Australia. <i>Genome Announcements</i> , 2013, 1, .	0.8	13
3	Assembly-Driven Community Genomics of a Hypersaline Microbial Ecosystem. <i>PLoS ONE</i> , 2013, 8, e61692.	2.5	101
4	‘ <i>De novo</i> ’ metagenomic assembly reveals abundant novel major lineage of Archaea in hypersaline microbial communities. <i>ISME Journal</i> , 2012, 6, 81-93.	9.8	347
5	Complete genome sequence of <i>Desulfurispirillum indicum</i> strain S5T. <i>Standards in Genomic Sciences</i> , 2011, 5, 371-378.	1.5	11
6	<i>Desulfurispirillum indicum</i> sp. nov., a selenate- and selenite-respiring bacterium isolated from an estuarine canal. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 654-658.	1.7	47
7	Xenorhodopsins, an enigmatic new class of microbial rhodopsins horizontally transferred between archaea and bacteria. <i>Biology Direct</i> , 2011, 6, 52.	4.6	50
8	<i>Pelobacter seleniigenes</i> sp. nov., a selenate-respiring bacterium. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2007, 57, 1937-1942.	1.7	35
9	Identification of Anaerobic Selenate-Respiring Bacteria from Aquatic Sediments. <i>Applied and Environmental Microbiology</i> , 2007, 73, 3519-3527.	3.1	102
10	<i>Sedimenticola selenatireducens</i> , gen. nov., sp. nov., an anaerobic selenate-respiring bacterium isolated from estuarine sediment. <i>Systematic and Applied Microbiology</i> , 2006, 29, 382-388.	2.8	66