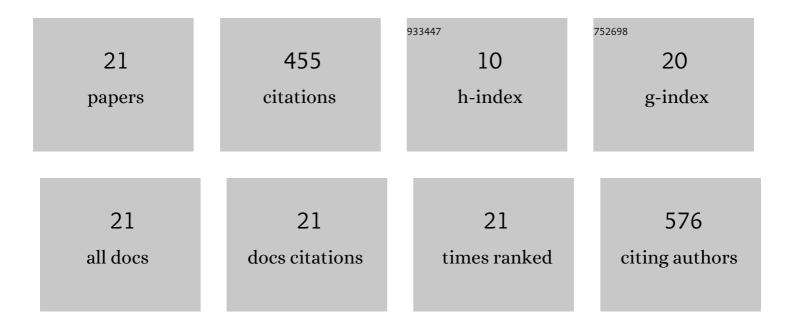
Rakhee Ds Khandeparker

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Phylogenetic diversity of culturable marine bacteria from sediments underlying the oxygen minimum zone of the Arabian Sea and their role in nitrate reduction. Marine Ecology, 2021, 42, e12646.	1.1	6
2	Xylanolytic enzyme consortia from Bacillus sp. NIORKP76 for improved biobleaching of kraft pulp. Bioprocess and Biosystems Engineering, 2021, 44, 2513-2524.	3.4	5
3	Denitrification rates of culturable bacteria from a coastal location turning temporally hypoxic. Journal of Marine Systems, 2020, 209, 103089.	2.1	8
4	High abundance of Vibrio in tarball-contaminated seawater from Vagator beach, Goa, India. Marine Pollution Bulletin, 2020, 150, 110773.	5.0	4
5	Seawater intrusion and resuspension of surface sediment control mercury (Hg) distribution and its bioavailability in water column of a monsoonal estuarine system. Science of the Total Environment, 2019, 660, 1441-1448.	8.0	33
6	Metagenomic analysis of tarball-associated bacteria from Goa, India. Marine Pollution Bulletin, 2019, 141, 398-403.	5.0	13
7	Bacterial Community Composition Markedly Altered by Coastal Hypoxia. Indian Journal of Microbiology, 2019, 59, 200-208.	2.7	7
8	Inter-annual variability of phytoplankton assemblage and Tetraspora gelatinosa bloom from anthropogenically affected harbour, Veraval, India. Environmental Monitoring and Assessment, 2019, 191, 87.	2.7	5
9	Nitrate Reductase Gene Expression in Idiomarina Strain cos21 Obtained from Oxygen Minimum Zone of Arabian Sea. Current Microbiology, 2019, 76, 63-69.	2.2	5
10	Quantitative analyses of denitrifying bacterial diversity from a seasonally hypoxic monsoon governed tropical coastal region. Deep-Sea Research Part II: Topical Studies in Oceanography, 2018, 156, 34-43.	1.4	11
11	Enzymatic saccharification of seaweeds into fermentable sugars by xylanase from marine Bacillus sp. strain BT21. 3 Biotech, 2017, 7, 296.	2.2	37
12	Recombinant Xylanase from Bacillus tequilensis BT21: Biochemical Characterization and Its Application in Production of Xylobiose from Agricultural Residues. Food Technology and Biotechnology, 2017, 55, 164-172.	2.1	31
13	Xylanolytic enzyme systems in <i>Arthrobacter</i> sp. MTCC 5214 and <i>Lactobacillus</i> sp Biotechnology and Applied Biochemistry, 2015, 62, 245-254.	3.1	8
14	Bacterial Diversity in Deep-Sea Sediments from Afanasy Nikitin Seamount, Equatorial Indian Ocean. Geomicrobiology Journal, 2014, 31, 942-949.	2.0	11
15	A study on bacteria associated with the intestinal tract of farmed yellow seahorse, Hippocampus kuda (Bleeker, 1852): characterization and extracellular enzymes. Aquaculture Research, 2012, 43, 386-394.	1.8	29
16	A novel halotolerant xylanase from marine isolate Bacillus subtilis cho40: gene cloning and sequencing. New Biotechnology, 2011, 28, 814-821.	4.4	61
17	Phylogenetic diversity of carbohydrate degrading culturable bacteria from Mandovi and Zuari estuaries, Goa, west coast of India. Estuarine, Coastal and Shelf Science, 2011, 95, 359-366.	2.1	19
18	Purification and characterization of α-l-arabinofuranosidase from Arthrobacter sp. MTCC 5214 in solid-state fermentation. Process Biochemistry, 2008, 43, 707-712.	3.7	9

#	Article	IF	CITATIONS
19	Bifunctional xylanases and their potential use in biotechnology. Journal of Industrial Microbiology and Biotechnology, 2008, 35, 635-644.	3.0	118
20	Extracellular polymeric substances of the marine fouling diatom <i>amphora rostrata</i> Wm.Sm. Biofouling, 2001, 17, 117-127.	2.2	35
21	Spatio-temporal variations in culturable bacterial community associated with denitrification in the Arabian Sea oxygen minimum zone. Marine Biology Research, 0, , 1-14.	0.7	Ο