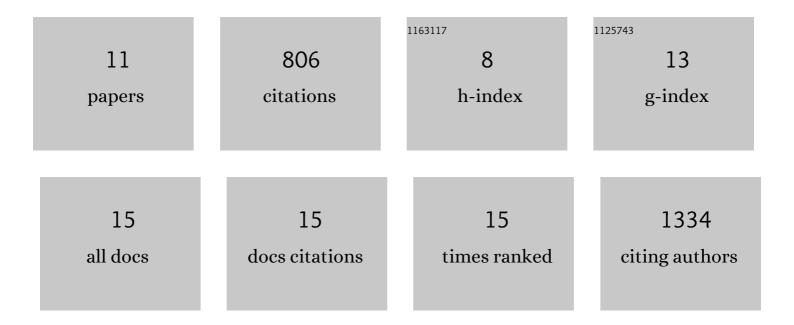
Yoji Nakamura

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8449824/publications.pdf

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#	Article	IF	CITATIONS
1	Biased biological functions of horizontally transferred genes in prokaryotic genomes. Nature Genetics, 2004, 36, 760-766.	21.4	459
2	Evolutionary changes of multiple visual pigment genes in the complete genome of Pacific bluefin tuna. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 11061-11066.	7.1	106
3	The genome stability in Corynebacterium species due to lack of the recombinational repair system. Gene, 2003, 317, 149-155.	2.2	66
4	A ddRAD-based genetic map and its integration with the genome assembly of Japanese eel (Anguilla) Tj ETQq0 0 C Genomics, 2014, 15, 233.) rgBT /Ove 2.8	erlock 10 Tf 63
5	Comparative genomics reveals that a fish pathogenic bacterium Edwardsiella tarda has acquired the locus of enterocyte effacement (LEE) through horizontal gene transfer. BMC Genomics, 2013, 14, 642.	2.8	40
6	Effects of plankton net characteristics on metagenetic community analysis of metazoan zooplankton in a coastal marine ecosystem. Journal of Experimental Marine Biology and Ecology, 2015, 469, 36-43.	1.5	28
7	Rhodopsin gene copies in Japanese eel originated in a teleost-specific genome duplication. Zoological Letters, 2017, 3, 18.	1.3	15
8	Targeted mutagenesis of the ryanodine receptor by Platinum TALENs causes slow swimming behaviour in Pacific bluefin tuna (Thunnus orientalis). Scientific Reports, 2019, 9, 13871.	3.3	14
9	Prediction of the Sex-Associated Genomic Region in Tunas (Thunnus Fishes). International Journal of Genomics, 2021, 2021, 1-14.	1.6	4
10	Prediction of Horizontally and Widely Transferred Genes in Prokaryotes. Evolutionary Bioinformatics, 2018, 14, 117693431881078.	1.2	3
11	Draft Whole-Genome Sequence of <i>Triparma laevis</i> f. <i>inornata</i> (Parmales, Bolidophyceae), Isolated from the Oyashio Region, Western North Pacific Ocean. Microbiology Resource Announcements, 2020, 9, .	0.6	3