

# Nam-phuong Nguyen

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4278299/publications.pdf>

Version: 2024-02-01

19  
papers

4,033  
citations

706676

14  
h-index

843174

20  
g-index

22  
all docs

22  
docs citations

22  
times ranked

7067  
citing authors

#	ARTICLE	IF	CITATIONS
1	Long-distance dispersal of pigeons and doves generated new ecological opportunities for host-switching and adaptive radiation by their parasites. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2022, 289, 20220042.	1.2	13
2	Extrachromosomal DNA in HPV-Mediated Oropharyngeal Cancer Drives Diverse Oncogene Transcription. <i>Clinical Cancer Research</i> , 2021, 27, 6772-6786.	3.2	20
3	Extrachromosomal DNA is associated with oncogene amplification and poor outcome across multiple cancers. <i>Nature Genetics</i> , 2020, 52, 891-897.	9.4	273
4	Global phylogeography and ancient evolution of the widespread human gut virus crAssphage. <i>Nature Microbiology</i> , 2019, 4, 1727-1736.	5.9	184
5	Circular ecDNA promotes accessible chromatin and high oncogene expression. <i>Nature</i> , 2019, 575, 699-703.	13.7	343
6	One thousand plant transcriptomes and the phylogenomics of green plants. <i>Nature</i> , 2019, 574, 679-685.	13.7	1,162
7	Fully automated sequence alignment methods are comparable to, and much faster than, traditional methods in large data sets: an example with hepatitis B virus. <i>PeerJ</i> , 2019, 7, e6142.	0.9	3
8	ViFi: accurate detection of viral integration and mRNA fusion reveals indiscriminate and unregulated transcription in proximal genomic regions in cervical cancer. <i>Nucleic Acids Research</i> , 2018, 46, 3309-3325.	6.5	47
9	Simultaneous radiation of bird and mammal lice following the K-Pg boundary. <i>Biology Letters</i> , 2018, 14, 20180141.	1.0	33
10	Primates, Lice and Bacteria: Speciation and Genome Evolution in the Symbionts of Hominid Lice. <i>Molecular Biology and Evolution</i> , 2017, 34, 1743-1757.	3.5	51
11	Phylogenomics from Whole Genome Sequences Using aTRAM. <i>Systematic Biology</i> , 2017, 66, syw105.	2.7	47
12	Phylogenomics using Target-restricted Assembly Resolves Intra-generic Relationships of Parasitic Lice ( <i>Phthiraptera</i> : <i>Columbicola</i> ). <i>Systematic Biology</i> , 2017, 66, syx027.	2.7	22
13	HIPPI: highly accurate protein family classification with ensembles of HMMs. <i>BMC Genomics</i> , 2016, 17, 765.	1.2	17
14	High diversity of picornaviruses in rats from different continents revealed by deep sequencing. <i>Emerging Microbes and Infections</i> , 2016, 5, 1-8.	3.0	14
15	A perspective on 16S rRNA operational taxonomic unit clustering using sequence similarity. <i>Npj Biofilms and Microbiomes</i> , 2016, 2, 16004.	2.9	208
16	PASTA: Ultra-Large Multiple Sequence Alignment for Nucleotide and Amino-Acid Sequences. <i>Journal of Computational Biology</i> , 2015, 22, 377-386.	0.8	360
17	TIPP: taxonomic identification and phylogenetic profiling. <i>Bioinformatics</i> , 2014, 30, 3548-3555.	1.8	93
18	Phylotranscriptomic analysis of the origin and early diversification of land plants. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E4859-68.	3.3	1,123

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19	Design and analysis of a robust genetic Muller C-element. Journal of Theoretical Biology, 2010, 264, 174-187.	0.8	12