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List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	MegaPath: sensitive and rapid pathogen detection using metagenomic NGS data. BMC Genomics, 2020, 21, 500.	2.8	6
2	Potential utility of metagenomic sequencing for improving etiologic diagnosis of infective endocarditis. Future Cardiology, 2019, 15, 411-424.	1.2	15
3	AC-DIAMOND v1: accelerating large-scale DNA–protein alignment. Bioinformatics, 2018, 34, 3744-3746.	4.1	8
4	MegaGTA: a sensitive and accurate metagenomic gene-targeted assembler using iterative de Bruijn graphs. BMC Bioinformatics, 2017, 18, 408.	2.6	11
5	BASE: a practical de novo assembler for large genomes using long NGS reads. BMC Genomics, 2016, 17, 499.	2.8	6
6	MEGAHIT v1.0: A fast and scalable metagenome assembler driven by advanced methodologies and community practices. Methods, 2016, 102, 3-11.	3.8	1,174
7	MEGAHIT: an ultra-fast single-node solution for large and complex metagenomics assembly via succinct <i>de Bruijn</i> graph. Bioinformatics, 2015, 31, 1674-1676.	4.1	4,864