

# Michele Berselli

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1714314/publications.pdf>

Version: 2024-02-01

7  
papers

435  
citations

1478505

6  
h-index

1720034

7  
g-index

8  
all docs

8  
docs citations

8  
times ranked

619  
citing authors

#	ARTICLE	IF	CITATIONS
1	The CAFA challenge reports improved protein function prediction and new functional annotations for hundreds of genes through experimental screens. <i>Genome Biology</i> , 2019, 20, 244.	8.8	261
2	G-quadruplex forming sequences in the genome of all known human viruses: A comprehensive guide. <i>PLoS Computational Biology</i> , 2018, 14, e1006675.	3.2	106
3	QPARSE: searching for long-looped or multimeric G-quadruplexes potentially distinctive and druggable. <i>Bioinformatics</i> , 2020, 36, 393-399.	4.1	16
4	Eliciting the Functional Taxonomy from protein annotations and taxa. <i>Scientific Reports</i> , 2016, 6, 31971.	3.3	14
5	Vimentin binds to G-quadruplex repeats found at telomeres and gene promoters. <i>Nucleic Acids Research</i> , 2022, 50, 1370-1381.	14.5	13
6	NeSSie: a tool for the identification of approximate DNA sequence symmetries. <i>Bioinformatics</i> , 2018, 34, 2503-2505.	4.1	11
7	BamSnap: a lightweight viewer for sequencing reads in BAM files. <i>Bioinformatics</i> , 2021, 37, 263-264.	4.1	5