

# Andrea Bagnacani

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7025487/publications.pdf>

Version: 2024-02-01

8  
papers

245  
citations

1478505

6  
h-index

1588992

8  
g-index

9  
all docs

9  
docs citations

9  
times ranked

446  
citing authors

#	ARTICLE	IF	CITATIONS
1	The RNA workbench 2.0: next generation RNA data analysis. <i>Nucleic Acids Research</i> , 2019, 47, W511-W515.	14.5	13
2	Workflow Development for the Functional Characterization of ncRNAs. <i>Methods in Molecular Biology</i> , 2019, 1912, 111-132.	0.9	7
3	Tools for Understanding miRNA-mRNA Interactions for Reproducible RNA Analysis. <i>Methods in Molecular Biology</i> , 2019, 1912, 199-214.	0.9	3
4	Evolution of computational models in BioModels Database and the Physiome Model Repository. <i>BMC Systems Biology</i> , 2018, 12, 53.	3.0	3
5	Community-Driven Data Analysis Training for Biology. <i>Cell Systems</i> , 2018, 6, 752-758.e1.	6.2	141
6	The RNA workbench: best practices for RNA and high-throughput sequencing bioinformatics in Galaxy. <i>Nucleic Acids Research</i> , 2017, 45, W560-W566.	14.5	38
7	Robust Cross-Platform Workflows: How Technical and Scientific Communities Collaborate to Develop, Test and Share Best Practices for Data Analysis. <i>Data Science and Engineering</i> , 2017, 2, 232-244.	6.4	19
8	Customized workflow development and data modularization concepts for RNA-Sequencing and metatranscriptome experiments. <i>Journal of Biotechnology</i> , 2017, 261, 85-96.	3.8	16