

Anna Maria Cichonska

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7694752/publications.pdf>

Version: 2024-02-01

10
papers

649
citations

1039880

9
h-index

1372474

10
g-index

15
all docs

15
docs citations

15
times ranked

1090
citing authors

#	ARTICLE	IF	CITATIONS
1	metaCCA: summary statistics-based multivariate meta-analysis of genome-wide association studies using canonical correlation analysis. <i>Bioinformatics</i> , 2016, 32, 1981-1989.	1.8	138
2	Metabolic biomarker profiling for identification of susceptibility to severe pneumonia and COVID-19 in the general population. <i>ELife</i> , 2021, 10, .	2.8	112
3	Computational-experimental approach to drug-target interaction mapping: A case study on kinase inhibitors. <i>PLoS Computational Biology</i> , 2017, 13, e1005678.	1.5	84
4	Leveraging multi-way interactions for systematic prediction of pre-clinical drug combination effects. <i>Nature Communications</i> , 2020, 11, 6136.	5.8	63
5	Profiling persistent tubercule bacilli from patient sputa during therapy predicts early drug efficacy. <i>BMC Medicine</i> , 2016, 14, 68.	2.3	55
6	Identification of drug candidates and repurposing opportunities through compoundâ€target interaction networks. <i>Expert Opinion on Drug Discovery</i> , 2015, 10, 1333-1345.	2.5	54
7	Learning with multiple pairwise kernels for drug bioactivity prediction. <i>Bioinformatics</i> , 2018, 34, i509-i518.	1.8	51
8	Crowdsourced mapping of unexplored target space of kinase inhibitors. <i>Nature Communications</i> , 2021, 12, 3307.	5.8	41
9	An expanded analysis framework for multivariate GWAS connects inflammatory biomarkers to functional variants and disease. <i>European Journal of Human Genetics</i> , 2021, 29, 309-324.	1.4	19
10	Modeling drug combination effects via latent tensor reconstruction. <i>Bioinformatics</i> , 2021, 37, i93-i101.	1.8	9