

# Olga I Zolotareva

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

19  
papers

322  
citations

9  
h-index

17  
g-index

24  
ext. papers

573  
ext. citations

11.2  
avg, IF

3.51  
L-index

#	Paper	IF	Citations
19	Flimma: a federated and privacy-aware tool for differential gene expression analysis.. <i>Genome Biology</i> , <b>2021</b> , 22, 338	18.3	2
18	Out-of-distribution generalization from labelled and unlabelled gene expression data for drug response prediction. <i>Nature Machine Intelligence</i> , <b>2021</b> , 3, 962-972	22.5	1
17	The AIme registry for artificial intelligence in biomedical research. <i>Nature Methods</i> , <b>2021</b> , 18, 1128-1131	21.6	11
16	Interpretable Drug Response Prediction using a Knowledge-based Neural Network <b>2021</b> ,		2
15	AITL: Adversarial Inductive Transfer Learning with input and output space adaptation for pharmacogenomics. <i>Bioinformatics</i> , <b>2020</b> , 36, i380-i388	7.2	10
14	A Survey of Gene Prioritization Tools for Mendelian and Complex Human Diseases. <i>Journal of Integrative Bioinformatics</i> , <b>2019</b> , 16,	3.8	7
13	MOLI: multi-omics late integration with deep neural networks for drug response prediction. <i>Bioinformatics</i> , <b>2019</b> , 35, i501-i509	7.2	85
12	Comorbidity of asthma and hypertension may be mediated by shared genetic dysregulation and drug side effects. <i>Scientific Reports</i> , <b>2019</b> , 9, 16302	4.9	9
11	Novel candidate genes important for asthma and hypertension comorbidity revealed from associative gene networks. <i>BMC Medical Genomics</i> , <b>2018</b> , 11, 15	3.7	19
10	Search for New Candidate Genes Involved in the Comorbidity of Asthma and Hypertension Based on Automatic Analysis of Scientific Literature. <i>Journal of Integrative Bioinformatics</i> , <b>2018</b> , 15,	3.8	4
9	GenCoNet - A Graph Database for the Analysis of Comorbidities by Gene Networks. <i>Journal of Integrative Bioinformatics</i> , <b>2018</b> , 15,	3.8	6
8	Molecular Relationships between Bronchial Asthma and Hypertension as Comorbid Diseases. <i>Journal of Integrative Bioinformatics</i> , <b>2018</b> , 15,	3.8	9
7	Antigen receptor repertoire profiling from RNA-seq data. <i>Nature Biotechnology</i> , <b>2017</b> , 35, 908-911	44.5	125
6	Robotic QM/MM-driven maturation of antibody combining sites. <i>Science Advances</i> , <b>2016</b> , 2, e1501695	14.3	10
5	Whole exome sequencing links dental tumor to an autosomal-dominant mutation in ANO5 gene associated with gnathodiaphyseal dysplasia and muscle dystrophies. <i>Scientific Reports</i> , <b>2016</b> , 6, 26440	4.9	15
4	O3-13-01: Whole genome sequencing of late-onset Alzheimer's disease patients from genetic isolate <b>2015</b> , 11, P250-P251		
3	MOLI: Multi-Omics Late Integration with deep neural networks for drug response prediction		2

2 AITL: Adversarial Inductive Transfer Learning with input and output space adaptation for pharmacogenomics 3

1 BDKANN - Biological Domain Knowledge-based Artificial Neural Network for drug response prediction 1