

# Oleg Ursu

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7466574/publications.pdf>

Version: 2024-02-01

17  
papers

2,784  
citations

687363

13  
h-index

794594

19  
g-index

20  
all docs

20  
docs citations

20  
times ranked

5092  
citing authors

#	ARTICLE	IF	CITATIONS
1	A comprehensive map of molecular drug targets. <i>Nature Reviews Drug Discovery</i> , 2017, 16, 19-34.	46.4	1,608
2	Pharos: Collating protein information to shed light on the druggable genome. <i>Nucleic Acids Research</i> , 2017, 45, D995-D1002.	14.5	271
3	Unexplored therapeutic opportunities in the human genome. <i>Nature Reviews Drug Discovery</i> , 2018, 17, 317-332.	46.4	263
4	DrugCentral 2021 supports drug discovery and repositioning. <i>Nucleic Acids Research</i> , 2021, 49, D1160-D1169.	14.5	129
5	Badapple: promiscuity patterns from noisy evidence. <i>Journal of Cheminformatics</i> , 2016, 8, 29.	6.1	85
6	Drug target ontology to classify and integrate drug discovery data. <i>Journal of Biomedical Semantics</i> , 2017, 8, 50.	1.6	63
7	Discovery of a specific inhibitor of human GLUT5 by virtual screening and in vitro transport evaluation. <i>Scientific Reports</i> , 2016, 6, 24240.	3.3	45
8	Model-Free Drug-Likeness from Fragments. <i>Journal of Chemical Information and Modeling</i> , 2010, 50, 1387-1394.	5.4	28
9	TIN-X: target importance and novelty explorer. <i>Bioinformatics</i> , 2017, 33, 2601-2603.	4.1	27
10	Development and validation of a clinical prediction rule for candidemia in hospitalized patients with severe sepsis and septic shock. <i>Journal of Critical Care</i> , 2015, 30, 715-720.	2.2	26
11	Chronic obstructive pulmonary disease phenotypes using cluster analysis of electronic medical records. <i>Health Informatics Journal</i> , 2018, 24, 394-409.	2.1	22
12	Machine learning prediction and tau-based screening identifies potential Alzheimer's disease genes relevant to immunity. <i>Communications Biology</i> , 2022, 5, 125.	4.4	18
13	FRET detection of lymphocyte function-associated antigen-1 conformational extension. <i>Molecular Biology of the Cell</i> , 2015, 26, 43-54.	2.1	17
14	High-Throughput Flow Cytometry Screening of Multidrug Efflux Systems. <i>Methods in Molecular Biology</i> , 2018, 1700, 293-318.	0.9	12
15	Formalizing drug indications on the road to therapeutic intent. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2017, 24, 1169-1172.	4.4	8
16	CHEMICAL AND BIOLOGICAL DESCRIPTOR INTEGRATION IMPROVES COMPUTATIONAL MODELING OF RAT TOXICITY. <i>Revue Roumaine De Chimie</i> , 2015, 60, 219-226.	0.2	4
17	PepSeA: Peptide Sequence Alignment and Visualization Tools to Enable Lead Optimization. <i>Journal of Chemical Information and Modeling</i> , 2022, 62, 1259-1267.	5.4	3