

# Yang Li

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3357888/publications.pdf>

Version: 2024-02-01

15  
papers

709  
citations

933447

10  
h-index

1058476

14  
g-index

15  
all docs

15  
docs citations

15  
times ranked

1484  
citing authors

#	ARTICLE	IF	CITATIONS
1	UbiBrowser 2.0: a comprehensive resource for proteome-wide known and predicted ubiquitin ligase/deubiquitinase substrate interactions in eukaryotic species. <i>Nucleic Acids Research</i> , 2022, 50, D719-D728.	14.5	46
2	ALK1 signaling is required for the homeostasis of Kupffer cells and prevention of bacterial infection. <i>Journal of Clinical Investigation</i> , 2022, 132, .	8.2	21
3	The coupling of reduced type H vessels with unloading-induced bone loss and the protection role of <i>Panax quinquefolium</i> saponin in the male mice. <i>Bone</i> , 2021, 143, 115712.	2.9	12
4	<i>Panax quinquefolium</i> saponin attenuates cardiac remodeling induced by simulated microgravity. <i>Phytomedicine</i> , 2019, 56, 83-93.	5.3	12
5	UVGD 1.0: a gene-centric database bridging ultraviolet radiation and molecular biology effects in organisms. <i>International Journal of Radiation Biology</i> , 2019, 95, 1172-1177.	1.8	0
6	FibroAtlas: A Database for the Exploration of Fibrotic Diseases and Their Genes. <i>Cardiology Research and Practice</i> , 2019, 2019, 1-7.	1.1	10
7	OsteoporosAtlas: a human osteoporosis-related gene database. <i>PeerJ</i> , 2019, 7, e6778.	2.0	6
8	AllerGAtlas 1.0: a human allergy-related genes database. <i>Database: the Journal of Biological Databases and Curation</i> , 2018, 2018, .	3.0	11
9	Revealing the metabolic characteristics of human embryonic stem cells by genome-scale metabolic modeling. <i>FEBS Letters</i> , 2018, 592, 3670-3682.	2.8	9
10	Functional constraints on adaptive evolution of protein ubiquitination sites. <i>Scientific Reports</i> , 2017, 7, 39949.	3.3	6
11	An integrated bioinformatics platform for investigating the human E3 ubiquitin ligase-substrate interaction network. <i>Nature Communications</i> , 2017, 8, 347.	12.8	151
12	Similarity-based prediction for Anatomical Therapeutic Chemical classification of drugs by integrating multiple data sources. <i>Bioinformatics</i> , 2015, 31, 1788-1795.	4.1	59
13	The covalent modifier Nedd8 is critical for the activation of Smurf1 ubiquitin ligase in tumorigenesis. <i>Nature Communications</i> , 2014, 5, 3733.	12.8	157
14	CAPER 2.0: An Interactive, Configurable, and Extensible Workflow-Based Platform to Analyze Data Sets from the Chromosome-centric Human Proteome Project. <i>Journal of Proteome Research</i> , 2014, 13, 99-106.	3.7	10
15	Genome-wide association study identifies two new susceptibility loci for atopic dermatitis in the Chinese Han population. <i>Nature Genetics</i> , 2011, 43, 690-694.	21.4	199