

Peter MÄ;jek

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

1,416
citations

840119

11
h-index

1199166

12
g-index

12
all docs

12
docs citations

12
times ranked

3572
citing authors

#	ARTICLE	IF	CITATIONS
1	24-Norursodeoxycholic acid reshapes immunometabolism in CD8+ T cells and alleviates hepatic inflammation. <i>Journal of Hepatology</i> , 2021, 75, 1164-1176.	1.8	20
2	Systematic characterization of BAF mutations provides insights into intracomplex synthetic lethalties in human cancers. <i>Nature Genetics</i> , 2019, 51, 1399-1410.	9.4	92
3	Type I Interferon Signaling Disrupts the Hepatic Urea Cycle and Alters Systemic Metabolism to Suppress T Cell Function. <i>Immunity</i> , 2019, 51, 1074-1087.e9.	6.6	72
4	Characterization of host proteins interacting with the lymphocytic choriomeningitis virus L protein. <i>PLoS Pathogens</i> , 2017, 13, e1006758.	2.1	19
5	FASIL-MS: An Integrated Proteomic and Bioinformatic Workflow To Universally Quantitate In Vivo-Acetylated Positional Isomers. <i>Journal of Proteome Research</i> , 2016, 15, 2579-2594.	1.8	6
6	Gene essentiality and synthetic lethality in haploid human cells. <i>Science</i> , 2015, 350, 1092-1096.	6.0	773
7	Early-onset inflammatory bowel disease and common variable immunodeficiency-like disease caused by IL-21 deficiency. <i>Journal of Allergy and Clinical Immunology</i> , 2014, 133, 1651-1659.e12.	1.5	124
8	Biallelic loss-of-function mutation in NIK causes a primary immunodeficiency with multifaceted aberrant lymphoid immunity. <i>Nature Communications</i> , 2014, 5, 5360.	5.8	116
9	Experiments and Comprehensive Simulations of the Formation of a Helical Turn. <i>Journal of Physical Chemistry B</i> , 2012, 116, 6598-6610.	1.2	22
10	Milestoning without a Reaction Coordinate. <i>Journal of Chemical Theory and Computation</i> , 2010, 6, 1805-1817.	2.3	100
11	A coarse-grained potential for fold recognition and molecular dynamics simulations of proteins. <i>Proteins: Structure, Function and Bioinformatics</i> , 2009, 76, 822-836.	1.5	47
12	Building and assessing atomic models of proteins from structural templates: Learning and benchmarks. <i>Proteins: Structure, Function and Bioinformatics</i> , 2009, 76, 930-945.	1.5	25