Francisco MartÃ-nez-Jiménez

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1343003/publications.pdf

Version: 2024-02-01

12 papers 1,149 citations

933447 10 h-index 1199594 12 g-index

14 all docs

14 docs citations

times ranked

14

2455 citing authors

#	Article	IF	CITATIONS
1	A compendium of mutational cancer driver genes. Nature Reviews Cancer, 2020, 20, 555-572.	28.4	605
2	Genomic epidemiology of superspreading events in Austria reveals mutational dynamics and transmission properties of SARS-CoV-2. Science Translational Medicine, 2020, 12, .	12.4	203
3	Genomic and evolutionary classification of lung cancer in never smokers. Nature Genetics, 2021, 53, 1348-1359.	21.4	81
4	In silico saturation mutagenesis of cancer genes. Nature, 2021, 596, 428-432.	27.8	61
5	Target Prediction for an Open Access Set of Compounds Active against Mycobacterium tuberculosis. PLoS Computational Biology, 2013, 9, e1003253.	3.2	51
6	Release of 50 new, drug-like compounds and their computational target predictions for open source anti-tubercular drug discovery. PLoS ONE, 2015, 10, e0142293.	2.5	38
7	A Novel Family of Soluble Minimal Scaffolds Provides Structural Insight into the Catalytic Domains of Integral Membrane Metallopeptidases. Journal of Biological Chemistry, 2013, 288, 21279-21294.	3.4	35
8	Systematic analysis of alterations in the ubiquitin proteolysis system reveals its contribution to driver mutations in cancer. Nature Cancer, 2020, 1, 122-135.	13.2	30
9	Ligand-Target Prediction by Structural Network Biology Using nAnnoLyze. PLoS Computational Biology, 2015, 11, e1004157.	3.2	16
10	Rational design of non-resistant targeted cancer therapies. Scientific Reports, 2017, 7, 46632.	3.3	11
11	Detection of oncogenic and clinically actionable mutations in cancer genomes critically depends on variant calling tools. Bioinformatics, 2022, 38, 3181-3191.	4.1	9
12	Should network biology be used for drug discovery?. Expert Opinion on Drug Discovery, 2016, 11, 1135-1137.	5.0	4