

Mark F Rogers

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1382488/publications.pdf>

Version: 2024-02-01

15
papers

1,112
citations

1040056

9
h-index

940533

16
g-index

16
all docs

16
docs citations

16
times ranked

2863
citing authors

#	ARTICLE	IF	CITATIONS
1	An integrative approach to predicting the functional effects of non-coding and coding sequence variation. <i>Bioinformatics</i> , 2015, 31, 1536-1543.	4.1	524
2	FATHMM-XF: accurate prediction of pathogenic point mutations via extended features. <i>Bioinformatics</i> , 2018, 34, 511-513.	4.1	296
3	Improving Illumina assemblies with Hi-C and long reads: An example with the North African dromedary. <i>Molecular Ecology Resources</i> , 2019, 19, 1015-1026.	4.8	67
4	CScape: a tool for predicting oncogenic single-point mutations in the cancer genome. <i>Scientific Reports</i> , 2017, 7, 11597.	3.3	52
5	Glucocorticoid Receptor-“Tethered Mineralocorticoid Receptors Increase Glucocorticoid-Induced Transcriptional Responses. <i>Endocrinology</i> , 2019, 160, 1044-1056.	2.8	35
6	RNA sequencing analysis of human podocytes reveals glucocorticoid regulated gene networks targeting non-immune pathways. <i>Scientific Reports</i> , 2016, 6, 35671.	3.3	25
7	iCLIP identifies novel roles for SAFB1 in regulating RNA processing and neuronal function. <i>BMC Biology</i> , 2015, 13, 111.	3.8	23
8	<i>CScape-somatic</i> : distinguishing driver and passenger point mutations in the cancer genome. <i>Bioinformatics</i> , 2020, 36, 3637-3644.	4.1	19
9	Nucleotide diversity of functionally different groups of immune response genes in Old World camels based on newly annotated and reference-guided assemblies. <i>BMC Genomics</i> , 2020, 21, 606.	2.8	15
10	Prediction of driver variants in the cancer genome via machine learning methodologies. <i>Briefings in Bioinformatics</i> , 2021, 22, .	6.5	13
11	Seasonal adaptations of the hypothalamo-neurohypophyseal system of the dromedary camel. <i>PLoS ONE</i> , 2019, 14, e0216679.	2.5	10
12	Corticosterone pattern-dependent glucocorticoid receptor binding and transcriptional regulation within the liver. <i>PLoS Genetics</i> , 2021, 17, e1009737.	3.5	10
13	High-resolution breakpoint junction mapping of proximally extended D4Z4 deletions in FSHD1 reveals evidence for a founder effect. <i>Human Molecular Genetics</i> , 2022, 31, 748-760.	2.9	8
14	Recognition memory-induced gene expression in the perirhinal cortex: A transcriptomic analysis. <i>Behavioural Brain Research</i> , 2017, 328, 1-12.	2.2	7
15	Estimating the Frequency of Single Point Driver Mutations across Common Solid Tumours. <i>Scientific Reports</i> , 2019, 9, 13452.	3.3	6