

Malene Juul

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2261206/publications.pdf>

Version: 2024-02-01

9
papers

770
citations

1478505

6
h-index

1588992

8
g-index

14
all docs

14
docs citations

14
times ranked

2155
citing authors

#	ARTICLE	IF	CITATIONS
1	Analyses of non-coding somatic drivers in 2,658 cancer whole genomes. <i>Nature</i> , 2020, 578, 102-111.	27.8	424
2	Mutational Context and Diverse Clonal Development in Early and Late Bladder Cancer. <i>Cell Reports</i> , 2014, 7, 1649-1663.	6.4	128
3	Pan-cancer screen for mutations in non-coding elements with conservation and cancer specificity reveals correlations with expression and survival. <i>Npj Genomic Medicine</i> , 2018, 3, 1.	3.8	79
4	Non-coding cancer driver candidates identified with a sample- and position-specific model of the somatic mutation rate. <i>ELife</i> , 2017, 6, .	6.0	36
5	PINCAGE: probabilistic integration of cancer genomics data for perturbed gene identification and sample classification. <i>Bioinformatics</i> , 2016, 32, 1353-1365.	4.1	12
6	A site specific model and analysis of the neutral somatic mutation rate in whole-genome cancer data. <i>BMC Bioinformatics</i> , 2018, 19, 147.	2.6	11
7	The landscape and driver potential of site-specific hotspots across cancer genomes. <i>Npj Genomic Medicine</i> , 2021, 6, 33.	3.8	8
8	ncdDetect2: improved models of the site-specific mutation rate in cancer and driver detection with robust significance evaluation. <i>Bioinformatics</i> , 2019, 35, 189-199.	4.1	6
9	EBADIMEX: an empirical Bayes approach to detect joint differential expression and methylation and to classify samples. <i>Statistical Applications in Genetics and Molecular Biology</i> , 2019, 18, .	0.6	0