

Yan Jaszczyszyn

List of Publications by Year in descending order

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Version: 2024-02-01

13
papers

1,485
citations

933447

10
h-index

1058476

14
g-index

18
all docs

18
docs citations

18
times ranked

2787
citing authors

#	ARTICLE	IF	CITATIONS
1	Pattern and causes of the establishment of the invasive bacterial potato pathogen <i>Dickeya solani</i> and of the maintenance of the resident pathogen <i>D. adianthicola</i> . <i>Molecular Ecology</i> , 2021, 30, 608-624.	3.9	13
2	DNA molecular combing-based replication fork directionality profiling. <i>Nucleic Acids Research</i> , 2021, 49, e69-e69.	14.5	7
3	Data from crosslinking and analysis of cDNAs (CRAC) of Nab3 in yeast cells expressing a circular ncRNA decoy. <i>Data in Brief</i> , 2021, 35, 106951.	1.0	1
4	The antibody response induced FMDV vaccines in sheep correlates with early transcriptomic responses in blood. <i>Npj Vaccines</i> , 2020, 5, 1.	6.0	101
5	Transcription-mediated organization of the replication initiation program across large genes sets common fragile sites genome-wide. <i>Nature Communications</i> , 2019, 10, 5693.	12.8	73
6	Developmental and cancer-associated plasticity of DNA replication preferentially targets GC-poor, lowly expressed and late-replicating regions. <i>Nucleic Acids Research</i> , 2018, 46, 10157-10172.	14.5	30
7	The evolution of the temporal program of genome replication. <i>Nature Communications</i> , 2018, 9, 2199.	12.8	19
8	The Third Revolution in Sequencing Technology. <i>Trends in Genetics</i> , 2018, 34, 666-681.	6.7	759
9	Dual Roles of Poly(dA:dT) Tracts in Replication Initiation and Fork Collapse. <i>Cell</i> , 2018, 174, 1127-1142.e19.	28.9	167
10	Finding sRNA-associated phenotypes by competition assays: An example with <i>Staphylococcus aureus</i> . <i>Methods</i> , 2017, 117, 21-27.	3.8	19
11	Holistic and Affordable Analyses of MicroRNA Expression Profiles Using Tagged cDNA Libraries and a Multiplex Sequencing Strategy. <i>Methods in Molecular Biology</i> , 2017, 1654, 179-196.	0.9	4
12	Replication landscape of the human genome. <i>Nature Communications</i> , 2016, 7, 10208.	12.8	259
13	DNA Physical Properties and Nucleosome Positions Are Major Determinants of HIV-1 Integrase Selectivity. <i>PLoS ONE</i> , 2015, 10, e0129427.	2.5	21