

Daniel Nichol

List of Publications by Year in descending order

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Version: 2024-02-01

10
papers

677
citations

1040056

9
h-index

1372567

10
g-index

15
all docs

15
docs citations

15
times ranked

1193
citing authors

#	ARTICLE	IF	CITATIONS
1	Steering Evolution with Sequential Therapy to Prevent the Emergence of Bacterial Antibiotic Resistance. <i>PLoS Computational Biology</i> , 2015, 11, e1004493.	3.2	151
2	Antibiotic collateral sensitivity is contingent on the repeatability of evolution. <i>Nature Communications</i> , 2019, 10, 334.	12.8	135
3	Collateral sensitivity networks reveal evolutionary instability and novel treatment strategies in ALK mutated non-small cell lung cancer. <i>Scientific Reports</i> , 2017, 7, 1232.	3.3	79
4	Exploiting evolutionary steering to induce collateral drug sensitivity in cancer. <i>Nature Communications</i> , 2020, 11, 1923.	12.8	79
5	Subclonal reconstruction of tumors by using machine learning and population genetics. <i>Nature Genetics</i> , 2020, 52, 898-907.	21.4	77
6	Stochasticity in the Genotype-Phenotype Map: Implications for the Robustness and Persistence of Bet-Hedging. <i>Genetics</i> , 2016, 204, 1523-1539.	2.9	39
7	The Spatiotemporal Evolution of Lymph Node Spread in Early Breast Cancer. <i>Clinical Cancer Research</i> , 2018, 24, 4763-4770.	7.0	30
8	Model genotype–phenotype mappings and the algorithmic structure of evolution. <i>Journal of the Royal Society Interface</i> , 2019, 16, 20190332.	3.4	28
9	Mapping the breast cancer metastatic cascade onto ctDNA using genetic and epigenetic clonal tracking. <i>Nature Communications</i> , 2020, 11, 1446.	12.8	28
10	The role of memory in non-genetic inheritance and its impact on cancer treatment resistance. <i>PLoS Computational Biology</i> , 2021, 17, e1009348.	3.2	11