

Ilan Shomorony

List of Publications by Year in descending order

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Version: 2024-02-01

18
papers

424
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1163117

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1125743

13
g-index

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all docs

18
docs citations

18
times ranked

383
citing authors

#	ARTICLE	IF	CITATIONS
1	HINGE: long-read assembly achieves optimal repeat resolution. <i>Genome Research</i> , 2017, 27, 747-756.	5.5	88
2	Fundamental limits of DNA storage systems. , 2017, , .		73
3	Degrees of Freedom of Two-Hop Wireless Networks: Everyone Gets the Entire Cake. <i>IEEE Transactions on Information Theory</i> , 2014, 60, 2417-2431.	2.4	61
4	Worst-Case Additive Noise in Wireless Networks. <i>IEEE Transactions on Information Theory</i> , 2013, 59, 3833-3847.	2.4	44
5	Capacity Results for the Noisy Shuffling Channel. , 2019, , .		38
6	DNA-Based Storage: Models and Fundamental Limits. <i>IEEE Transactions on Information Theory</i> , 2021, 67, 3675-3689.	2.4	32
7	Information-optimal genome assembly via sparse read-overlap graphs. <i>Bioinformatics</i> , 2016, 32, i494-i502.	4.1	23
8	Computing Half-Duplex Schedules in Gaussian Relay Networks via Min-Cut Approximations. <i>IEEE Transactions on Information Theory</i> , 2014, 60, 7204-7220.	2.4	15
9	Information-Theoretic Foundations of DNA Data Storage. <i>Foundations and Trends in Communications and Information Theory</i> , 2022, 19, 1-106.	3.1	11
10	Fundamental Limits of Genome Assembly Under an Adversarial Erasure Model. <i>IEEE Transactions on Molecular, Biological, and Multi-Scale Communications</i> , 2016, 2, 199-208.	2.1	10
11	Spectral Jaccard Similarity: A New Approach to Estimating Pairwise Sequence Alignments. <i>Patterns</i> , 2020, 1, 100081.	5.9	9
12	Torn-Paper Coding. <i>IEEE Transactions on Information Theory</i> , 2021, 67, 7904-7913.	2.4	5
13	Communicating over the Torn-Paper Channel. , 2020, , .		5
14	Coded Shotgun Sequencing. <i>IEEE Journal on Selected Areas in Information Theory</i> , 2022, 3, 147-159.	2.5	4
15	Network Compression: Worst Case Analysis. <i>IEEE Transactions on Information Theory</i> , 2015, 61, 3980-3995.	2.4	3
16	JIND: joint integration and discrimination for automated single-cell annotation. <i>Bioinformatics</i> , 2022, 38, 2488-2495.	4.1	3
17	Sketching and Sequence Alignment: A Rate-Distortion Perspective. , 2021, , .		0
18	Achieving the Capacity of a DNA Storage Channel with Linear Coding Schemes. , 2022, , .		0