## Ranit Aharonov

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11760959/publications.pdf

Version: 2024-02-01

26 papers

4,726 citations

471509 17 h-index 18 g-index

26 all docs

26 docs citations

26 times ranked 6207 citing authors

#	Article	IF	CITATIONS
1	Identification of hundreds of conserved and nonconserved human microRNAs. Nature Genetics, 2005, 37, 766-770.	21.4	1,720
2	MicroRNAs accurately identify cancer tissue origin. Nature Biotechnology, 2008, 26, 462-469.	17.5	909
3	MicroRNA expression detected by oligonucleotide microarrays: System establishment and expression profiling in human tissues. Genome Research, 2004, 14, 2486-2494.	5.5	495
4	Diagnostic Assay Based on hsa-miR-205 Expression Distinguishes Squamous From Nonsquamous Non–Small-Cell Lung Carcinoma. Journal of Clinical Oncology, 2009, 27, 2030-2037.	1.6	381
5	A Second-Generation MicroRNA-Based Assay for Diagnosing Tumor Tissue Origin. Oncologist, 2012, 17, 801-812.	3.7	144
6	hsa-miR-29c* Is Linked to the Prognosis of Malignant Pleural Mesothelioma. Cancer Research, 2010, 70, 1916-1924.	0.9	140
7	Validation of a microRNA-based qRT-PCR test for accurate identification of tumor tissue origin. Modern Pathology, 2010, 23, 814-823.	5.5	129
8	An autonomous debating system. Nature, 2021, 591, 379-384.	27.8	127
9	A Diagnostic Assay Based on MicroRNA Expression Accurately Identifies Malignant Pleural Mesothelioma. Journal of Molecular Diagnostics, 2010, 12, 771-779.	2.8	111
10	Prospective Gene Signature Study Using microRNA to Identify the Tissue of Origin in Patients with Carcinoma of Unknown Primary. Clinical Cancer Research, 2011, 17, 4063-4070.	7.0	110
11	Classification of the Four Main Types of Lung Cancer Using a MicroRNA-Based Diagnostic Assay. Journal of Molecular Diagnostics, 2012, 14, 510-517.	2.8	107
12	Tumor microRNA-29a expression and the risk of recurrence in stage II colon cancer. International Journal of Oncology, 2012, 40, 2097-103.	3.3	48
13	Novel microRNA-based assay demonstrates 92% agreement with diagnosis based on clinicopathologic and management data in a cohort of patients with carcinoma of unknown primary. Molecular Cancer, 2013, 12, 57.	19.2	43
14	Predicting progression of bladder urothelial carcinoma using micro <scp>RNA</scp> expression. BJU International, 2013, 112, 1027-1034.	2.5	38
15	A Large-Scale Dataset for Argument Quality Ranking: Construction and Analysis. Proceedings of the AAAI Conference on Artificial Intelligence, 2020, 34, 7805-7813.	4.9	34
16	Accurate Classification of Metastatic Brain Tumors Using a Novel MicroRNAâ€Based Test. Oncologist, 2011, 16, 165-174.	3.7	33
17	Will it Blend? Blending Weak and Strong Labeled Data in a Neural Network for Argumentation Mining. , 2018, , .		28
18	Are You Convinced? Choosing the More Convincing Evidence with a Siamese Network. , 2019, , .		24

#	Article	IF	CITATIONS
19	Unsupervised corpus–wide claim detection. , 2017, , .		24
20	Global microRNA profiling in favorable prognosis subgroups of cancer of unknown primary (CUP) demonstrates no significant expression differences with metastases of matched known primary tumors. Clinical and Experimental Metastasis, 2013, 30, 431-439.	3 <b>.</b> 3	23
21	Learning Thematic Similarity Metric from Article Sections Using Triplet Networks. , 2018, , .		17
22	Corpus Wide Argument Mining—A Working Solution. Proceedings of the AAAI Conference on Artificial Intelligence, 2020, 34, 7683-7691.	4.9	16
23	Listening Comprehension over Argumentative Content. , 2018, , .		7
24	A Dataset of General-Purpose Rebuttal., 2019,,.		7
25	Towards Effective Rebuttal: Listening Comprehension Using Corpus-Wide Claim Mining. , 2019, , .		6
26	From Surrogacy to Adoption; From Bitcoin to Cryptocurrency: Debate Topic Expansion. , 2019, , .		5