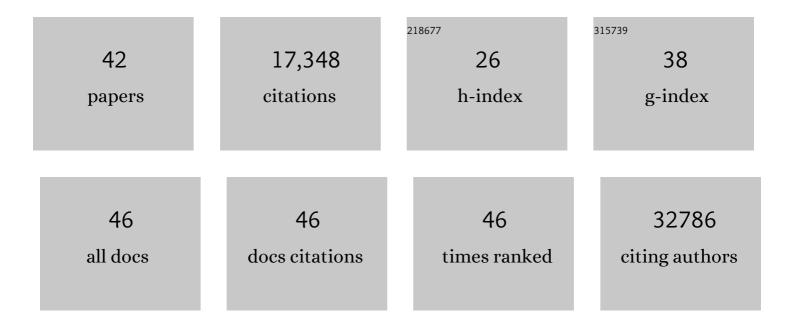
Mahmoud Mahdi Ghandi

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Sex-Biased <i>ZRSR2</i> Mutations in Myeloid Malignancies Impair Plasmacytoid Dendritic Cell Activation and Apoptosis. Cancer Discovery, 2022, 12, 522-541.	9.4	44
2	A tool for feature extraction from biological sequences. Briefings in Bioinformatics, 2022, 23, .	6.5	6
3	Integrated evaluation of telomerase activation and telomere maintenance across cancer cell lines. ELife, 2021, 10, .	6.0	8
4	A nonlinear acceleration method for iterative algorithms. Signal Processing, 2020, 168, 107346.	3.7	1
5	Distinct Classes of Complex Structural Variation Uncovered across Thousands of Cancer Genome Graphs. Cell, 2020, 183, 197-210.e32.	28.9	141
6	Multiplexed single-cell transcriptional response profiling to define cancer vulnerabilities and therapeutic mechanism of action. Nature Communications, 2020, 11, 4296.	12.8	98
7	Quantitative Proteomics of the Cancer Cell Line Encyclopedia. Cell, 2020, 180, 387-402.e16.	28.9	596
8	Small-Molecule and CRISPR Screening Converge to Reveal Receptor Tyrosine Kinase Dependencies in Pediatric Rhabdoid Tumors. Cell Reports, 2019, 28, 2331-2344.e8.	6.4	24
9	Next-generation characterization of the Cancer Cell Line Encyclopedia. Nature, 2019, 569, 503-508.	27.8	2,149
10	The landscape of cancer cell line metabolism. Nature Medicine, 2019, 25, 850-860.	30.7	350
11	WRN helicase is a synthetic lethal target in microsatellite unstable cancers. Nature, 2019, 568, 551-556.	27.8	253
12	DNA methyltransferase inhibition overcomes diphthamide pathway deficiencies underlying CD123-targeted treatment resistance. Journal of Clinical Investigation, 2019, 129, 5005-5019.	8.2	59
13	Targetable vulnerabilities in T- and NK-cell lymphomas identified through preclinical models. Nature Communications, 2018, 9, 2024.	12.8	80
14	Genetic and transcriptional evolution alters cancer cell line drug response. Nature, 2018, 560, 325-330.	27.8	662
15	Suppression of 19S proteasome subunits marks emergence of an altered cell state in diverse cancers. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 382-387.	7.1	47
16	Decomposing Oncogenic Transcriptional Signatures to Generate Maps of Divergent Cellular States. Cell Systems, 2017, 5, 105-118.e9.	6.2	40
17	Defining a Cancer Dependency Map. Cell, 2017, 170, 564-576.e16.	28.9	1,794
18	Allele-Specific DNA Methylation and Its Interplay with Repressive Histone Marks at Promoter-Mutant TERT Genes. Cell Reports, 2017, 21, 3700-3707.	6.4	68

#	Article	IF	CITATIONS
19	Resistance to SL-401 in AML and BPDCN Is Associated with Loss of the Diphthamide Synthesis Pathway Enzyme DPH1 and Is Reversible By Azacitidine. Blood, 2017, 130, 797-797.	1.4	13
20	Characterizing genomic alterations in cancer by complementary functional associations. Nature Biotechnology, 2016, 34, 539-546.	17.5	78
21	gkmSVM: an R package for gapped-kmer SVM. Bioinformatics, 2016, 32, 2205-2207.	4.1	155
22	Pediatric-type nodal follicular lymphoma: a biologically distinct lymphoma with frequent MAPK pathway mutations. Blood, 2016, 128, 1093-1100.	1.4	126
23	Systematic Functional Characterization of Resistance to PI3K Inhibition in Breast Cancer. Cancer Discovery, 2016, 6, 1134-1147.	9.4	106
24	Blastic Plasmacytoid Dendritic Cell Neoplasm (BPDCN) Harbors Frequent Splicesosome Mutations That Cause Aberrant RNA Splicing Affecting Genes Critical in pDC Differentiation and Function. Blood, 2016, 128, 738-738.	1.4	5
25	The Molecular Signatures Database Hallmark Gene Set Collection. Cell Systems, 2015, 1, 417-425.	6.2	7,719
26	Pharmacogenomic agreement between two cancer cell line data sets. Nature, 2015, 528, 84-87.	27.8	358
27	Blastic Plasmacytoid Dendritic Cell Neoplasm (BPDCN) Patient-Derived Xenografts Are Faithful Genomic and Phenotypic Models of Primary Leukemia and Respond to the IL3 Receptor Targeting Agent SL-401 In Vivo. Blood, 2015, 126, 3797-3797.	1.4	1
28	Abstract PRO2: Identification of ERK1/2 mutations that confer resistance to MAPK pathway inhibitors. , 2015, , .		0
29	Enhanced Regulatory Sequence Prediction Using Gapped k-mer Features. PLoS Computational Biology, 2014, 10, e1003711.	3.2	426
30	Locally Disordered Methylation Forms the Basis of Intratumor Methylome Variation in Chronic Lymphocytic Leukemia. Cancer Cell, 2014, 26, 813-825.	16.8	323
31	ARID1B is a specific vulnerability in ARID1A-mutant cancers. Nature Medicine, 2014, 20, 251-254.	30.7	336
32	ERK Mutations Confer Resistance to Mitogen-Activated Protein Kinase Pathway Inhibitors. Cancer Research, 2014, 74, 7079-7089.	0.9	90
33	Robust \$\$k\$\$ k -mer frequency estimation using gapped \$\$k\$\$ k -mers. Journal of Mathematical Biology, 2014, 69, 469-500.	1.9	44
34	Abstract LB-126: ARID1B is a specific vulnerability in ARID1A-mutant cancers. Cancer Research, 2014, 74, LB-126-LB-126.	0.9	2
35	Punctuated Evolution of Prostate Cancer Genomes. Cell, 2013, 153, 666-677.	28.9	1,107
36	Increased Local Disorder of DNA Methylation Forms the Basis of High Intra-Leukemic Epigenetic Heterogeneity and Enhances CLL Evolution. Blood, 2013, 122, 596-596.	1.4	4

#	Article	IF	CITATIONS
37	Group Normalization for Genomic Data. PLoS ONE, 2012, 7, e38695.	2.5	4
38	A mechanistic study on the effect of dexamethasone in moderating cell death in Chinese Hamster Ovary cell cultures. Biotechnology Progress, 2012, 28, 490-496.	2.6	9
39	Some nonlinear/adaptive methods for fast recovery of the missing samples of signals. Signal Processing, 2008, 88, 624-638.	3.7	5
40	Multiple wavelet denoising for embolic signal enhancement. , 2007, , .		1
41	Bit-rate reduction of MPEG compressed video. , 0, , .		Ο
42	A novel context modeling scheme for motion vectors context-based arithmetic coding. , 0, , .		2