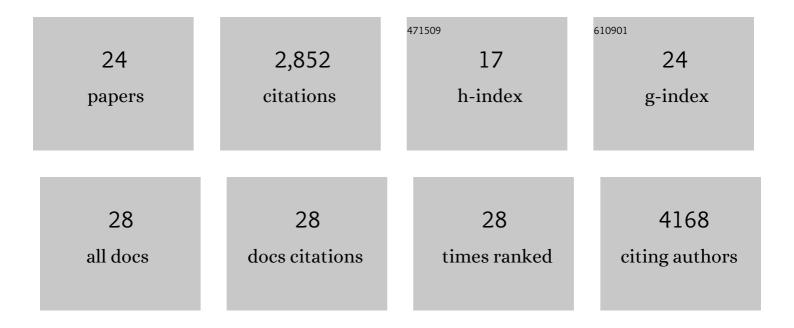
Nuri A Temiz

List of Publications by Year in descending order

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NIIDI A TEMIZ

#	Article	IF	CITATIONS
1	PELP1/SRC-3-dependent regulation of metabolic PFKFB kinases drives therapy resistant ER+ breast cancer. Oncogene, 2021, 40, 4384-4397.	5.9	16
2	APOBEC3A catalyzes mutation and drives carcinogenesis in vivo. Journal of Experimental Medicine, 2020, 217, .	8.5	87
3	Functional and Structural Insights into a Vif/PPP2R5 Complex Elucidated Using Patient HIV-1 Isolates and Computational Modeling. Journal of Virology, 2020, 94, .	3.4	6
4	Râ€spondin 2 Drives Liver Tumor Development in a Yesâ€Associated Proteinâ€Dependent Manner. Hepatology Communications, 2019, 3, 1496-1509.	4.3	15
5	Evaluating the landscape of gene cooperativity with receptor tyrosine kinases in liver tumorigenesis using transposon-mediated mutagenesis. Journal of Hepatology, 2019, 70, 470-482.	3.7	13
6	Cancer Stem Cell Phenotypes in ER+ Breast Cancer Models Are Promoted by PELP1/AIB1 Complexes. Molecular Cancer Research, 2018, 16, 707-719.	3.4	20
7	Comparative Transcriptome Analysis Quantifies Immune Cell Transcript Levels, Metastatic Progression, and Survival in Osteosarcoma. Cancer Research, 2018, 78, 326-337.	0.9	100
8	Insulin Receptor Substrate Suppression by the Tyrphostin NT157 Inhibits Responses to Insulin-Like Growth Factor-I and Insulin in Breast Cancer Cells. Hormones and Cancer, 2018, 9, 371-382.	4.9	14
9	Mutation Signatures Including APOBEC in Cancer Cell Lines. JNCI Cancer Spectrum, 2018, 2, .	2.9	45
10	<i>Sleeping Beauty</i> Insertional Mutagenesis in Mice Identifies Drivers of Steatosis-Associated Hepatic Tumors. Cancer Research, 2017, 77, 6576-6588.	0.9	40
11	<i>MYC</i> and <i>PVT1</i> synergize to regulate RSPO1 levels in breast cancer. Cell Cycle, 2016, 15, 881-885.	2.6	27
12	The DNA cytosine deaminase APOBEC3H haplotype I likely contributes to breast and lung cancer mutagenesis. Nature Communications, 2016, 7, 12918.	12.8	146
13	The DNA cytosine deaminase APOBEC3B promotes tamoxifen resistance in ER-positive breast cancer. Science Advances, 2016, 2, e1601737.	10.3	175
14	RNA sequencing of <i>Sleeping Beauty</i> transposon-induced tumors detects transposon-RNA fusions in forward genetic cancer screens. Genome Research, 2016, 26, 119-129.	5.5	28
15	Mutation Processes in 293-Based Clones Overexpressing the DNA Cytosine Deaminase APOBEC3B. PLoS ONE, 2016, 11, e0155391.	2.5	33
16	The somatic autosomal mutation matrix in cancer genomes. Human Genetics, 2015, 134, 851-864.	3.8	16
17	A Sleeping Beauty forward genetic screen identifies new genes and pathways driving osteosarcoma development and metastasis. Nature Genetics, 2015, 47, 615-624.	21.4	207
18	Human Papillomavirus E6 Triggers Upregulation of the Antiviral and Cancer Genomic DNA Deaminase APOBEC3B. MBio, 2014, 5, .	4.1	172

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#	Article	IF	CITATIONS
19	Evidence for APOBEC3B mutagenesis in multiple human cancers. Nature Genetics, 2013, 45, 977-983.	21.4	660
20	APOBEC3B is an enzymatic source of mutation in breast cancer. Nature, 2013, 494, 366-370.	27.8	758
21	APOBEC3B Upregulation and Genomic Mutation Patterns in Serous Ovarian Carcinoma. Cancer Research, 2013, 73, 7222-7231.	0.9	153
22	Guanine Holes Are Prominent Targets for Mutation in Cancer and Inherited Disease. PLoS Genetics, 2013, 9, e1003816.	3.5	34
23	The Role of Methylation in the Intrinsic Dynamics of B- and Z-DNA. PLoS ONE, 2012, 7, e35558.	2.5	62
24	Novel modulation factor quantifies the role of water molecules in protein interactions. Proteins: Structure, Function and Bioinformatics, 2010, 78, 3226-3234.	2.6	5