

# Nuri A Temiz

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7858914/publications.pdf>

Version: 2024-02-01

24  
papers

2,852  
citations

471509

17  
h-index

610901

24  
g-index

28  
all docs

28  
docs citations

28  
times ranked

4168  
citing authors

#	ARTICLE	IF	CITATIONS
1	APOBEC3B is an enzymatic source of mutation in breast cancer. <i>Nature</i> , 2013, 494, 366-370.	27.8	758
2	Evidence for APOBEC3B mutagenesis in multiple human cancers. <i>Nature Genetics</i> , 2013, 45, 977-983.	21.4	660
3	A Sleeping Beauty forward genetic screen identifies new genes and pathways driving osteosarcoma development and metastasis. <i>Nature Genetics</i> , 2015, 47, 615-624.	21.4	207
4	The DNA cytosine deaminase APOBEC3B promotes tamoxifen resistance in ER-positive breast cancer. <i>Science Advances</i> , 2016, 2, e1601737.	10.3	175
5	Human Papillomavirus E6 Triggers Upregulation of the Antiviral and Cancer Genomic DNA Deaminase APOBEC3B. <i>MBio</i> , 2014, 5, .	4.1	172
6	APOBEC3B Upregulation and Genomic Mutation Patterns in Serous Ovarian Carcinoma. <i>Cancer Research</i> , 2013, 73, 7222-7231.	0.9	153
7	The DNA cytosine deaminase APOBEC3H haplotype I likely contributes to breast and lung cancer mutagenesis. <i>Nature Communications</i> , 2016, 7, 12918.	12.8	146
8	Comparative Transcriptome Analysis Quantifies Immune Cell Transcript Levels, Metastatic Progression, and Survival in Osteosarcoma. <i>Cancer Research</i> , 2018, 78, 326-337.	0.9	100
9	APOBEC3A catalyzes mutation and drives carcinogenesis in vivo. <i>Journal of Experimental Medicine</i> , 2020, 217, .	8.5	87
10	The Role of Methylation in the Intrinsic Dynamics of B- and Z-DNA. <i>PLoS ONE</i> , 2012, 7, e35558.	2.5	62
11	Mutation Signatures Including APOBEC in Cancer Cell Lines. <i>JNCI Cancer Spectrum</i> , 2018, 2, .	2.9	45
12	<i>Sleeping Beauty</i> Insertional Mutagenesis in Mice Identifies Drivers of Steatosis-Associated Hepatic Tumors. <i>Cancer Research</i> , 2017, 77, 6576-6588.	0.9	40
13	Guanine Holes Are Prominent Targets for Mutation in Cancer and Inherited Disease. <i>PLoS Genetics</i> , 2013, 9, e1003816.	3.5	34
14	Mutation Processes in 293-Based Clones Overexpressing the DNA Cytosine Deaminase APOBEC3B. <i>PLoS ONE</i> , 2016, 11, e0155391.	2.5	33
15	RNA sequencing of <i>Sleeping Beauty</i> transposon-induced tumors detects transposon-RNA fusions in forward genetic cancer screens. <i>Genome Research</i> , 2016, 26, 119-129.	5.5	28
16	<i>MYC</i> and <i>PVT1</i> synergize to regulate RSPO1 levels in breast cancer. <i>Cell Cycle</i> , 2016, 15, 881-885.	2.6	27
17	Cancer Stem Cell Phenotypes in ER+ Breast Cancer Models Are Promoted by PELP1/AIB1 Complexes. <i>Molecular Cancer Research</i> , 2018, 16, 707-719.	3.4	20
18	The somatic autosomal mutation matrix in cancer genomes. <i>Human Genetics</i> , 2015, 134, 851-864.	3.8	16

#	ARTICLE	IF	CITATIONS
19	PELP1/SRC-3-dependent regulation of metabolic PFKFB kinases drives therapy resistant ER+ breast cancer. <i>Oncogene</i> , 2021, 40, 4384-4397.	5.9	16
20	R&#oacute;spondin 2 Drives Liver Tumor Development in a Yes&#oacute;Associated Protein&#oacute;Dependent Manner. <i>Hepatology Communications</i> , 2019, 3, 1496-1509.	4.3	15
21	Insulin Receptor Substrate Suppression by the Tyrphostin NT157 Inhibits Responses to Insulin-Like Growth Factor-I and Insulin in Breast Cancer Cells. <i>Hormones and Cancer</i> , 2018, 9, 371-382.	4.9	14
22	Evaluating the landscape of gene cooperativity with receptor tyrosine kinases in liver tumorigenesis using transposon-mediated mutagenesis. <i>Journal of Hepatology</i> , 2019, 70, 470-482.	3.7	13
23	Functional and Structural Insights into a Vif/PPP2R5 Complex Elucidated Using Patient HIV-1 Isolates and Computational Modeling. <i>Journal of Virology</i> , 2020, 94, .	3.4	6
24	Novel modulation factor quantifies the role of water molecules in protein interactions. <i>Proteins: Structure, Function and Bioinformatics</i> , 2010, 78, 3226-3234.	2.6	5