

# Momeneh Foroutan

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3084755/publications.pdf>

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14  
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1040056

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1125743

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g-index

21  
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docs citations

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times ranked

1450  
citing authors

#	ARTICLE	IF	CITATIONS
1	Single sample scoring of molecular phenotypes. BMC Bioinformatics, 2018, 19, 404.	2.6	286
2	A Gene Signature Predicting Natural Killer Cell Infiltration and Improved Survival in Melanoma Patients. Cancer Immunology Research, 2019, 7, 1162-1174.	3.4	201
3	Combinatorial Targeting by MicroRNAs Co-ordinates Post-transcriptional Control of EMT. Cell Systems, 2018, 7, 77-91.e7.	6.2	92
4	A Transcriptional Program for Detecting TGF $\beta$ 2-Induced EMT in Cancer. Molecular Cancer Research, 2017, 15, 619-631.	3.4	63
5	Stimulus-dependent differences in signalling regulate epithelial-mesenchymal plasticity and change the effects of drugs in breast cancer cell lines. Cell Communication and Signaling, 2015, 13, 26.	6.5	47
6	The Ratio of Exhausted to Resident Infiltrating Lymphocytes Is Prognostic for Colorectal Cancer Patient Outcome. Cancer Immunology Research, 2021, 9, 1125-1140.	3.4	18
7	CSK-homologous kinase (CHK/MATK) is a potential colorectal cancer tumour suppressor gene epigenetically silenced by promoter methylation. Oncogene, 2021, 40, 3015-3029.	5.9	13
8	Using singscore to predict mutation status in acute myeloid leukemia from transcriptomic signatures. F1000Research, 2019, 8, 776.	1.6	12
9	Targeting WEE1/AKT Restores p53-Dependent Natural Killer Cell Activation to Induce Immune Checkpoint Blockade Responses in "Cold" Melanoma. Cancer Immunology Research, 2022, 10, 757-769.	3.4	11
10	TGF $\beta$ 2 and CIS Inhibition Overcomes NK-cell Suppression to Restore Antitumor Immunity. Cancer Immunology Research, 2022, 10, 1047-1054.	3.4	11
11	Using singscore to predict mutations in acute myeloid leukemia from transcriptomic signatures. F1000Research, 2019, 8, 776.	1.6	9
12	Networking and Bayesian analyses of genetic affinity in cotton germplasm. Nucleus (India), 2015, 58, 33-45.	2.2	5
13	Computational Screening of Anti-Cancer Drugs Identifies a New BRCA Independent Gene Expression Signature to Predict Breast Cancer Sensitivity to Cisplatin. Cancers, 2022, 14, 2404.	3.7	2
14	Using singscore to predict mutation status in acute myeloid leukemia from transcriptomic signatures. F1000Research, 0, 8, 776.	1.6	1