

Abhilash Venugopalan

List of Publications by Year in descending order

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Version: 2024-02-01

14
papers

442
citations

933447

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1199594

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1126
citing authors

#	ARTICLE	IF	CITATIONS
1	APOBEC Mutagenesis and Copy-Number Alterations Are Drivers of Proteogenomic Tumor Evolution and Heterogeneity in Metastatic Thoracic Tumors. <i>Cell Reports</i> , 2019, 26, 2651-2666.e6.	6.4	92
2	Clonal Evolution and Heterogeneity of Osimertinib Acquired Resistance Mechanisms in EGFR Mutant Lung Cancer. <i>Cell Reports Medicine</i> , 2020, 1, 100007.	6.5	78
3	Loss of MIC6 Accelerates Initiation and Progression of Mutant Epidermal Growth Factor Receptor-Driven Lung Adenocarcinoma. <i>Cancer Discovery</i> , 2015, 5, 534-549.	9.4	57
4	A proteogenomic approach to map the proteome of an unsequenced pathogen <i>Leishmania donovani</i> . <i>Proteomics</i> , 2012, 12, 832-844.	2.2	42
5	Identifying novel targets of oncogenic EGF receptor signaling in lung cancer through global phosphoproteomics. <i>Proteomics</i> , 2015, 15, 340-355.	2.2	42
6	Quantitative Tyrosine Phosphoproteomics of Epidermal Growth Factor Receptor (EGFR) Tyrosine Kinase Inhibitor-treated Lung Adenocarcinoma Cells Reveals Potential Novel Biomarkers of Therapeutic Response. <i>Molecular and Cellular Proteomics</i> , 2017, 16, 891-910.	3.8	42
7	EGFR-targeted therapy results in dramatic early lung tumor regression accompanied by imaging response and immune infiltration in EGFR mutant transgenic mouse models. <i>Oncotarget</i> , 2016, 7, 54137-54156.	1.8	27
8	Genomic profiling of multiple sequentially acquired tumor metastatic sites from an exceptional responder lung adenocarcinoma patient reveals extensive genomic heterogeneity and novel somatic variants driving treatment response. <i>Journal of Physical Education and Sports Management</i> , 2016, 2, a001263.	1.2	18
9	Development of Highly Effective Anti-Mesothelin hYP218 Chimeric Antigen Receptor T Cells With Increased Tumor Infiltration and Persistence for Treating Solid Tumors. <i>Molecular Cancer Therapeutics</i> , 2022, 21, 1195-1206.	4.1	18
10	Quantitative Mass Spectrometry to Interrogate Proteomic Heterogeneity in Metastatic Lung Adenocarcinoma and Validate a Novel Somatic Mutation CDK12-G879V. <i>Molecular and Cellular Proteomics</i> , 2019, 18, 622-641.	3.8	15
11	SCAMP3 is a mutant EGFR phosphorylation target and a tumor suppressor in lung adenocarcinoma. <i>Oncogene</i> , 2021, 40, 3331-3346.	5.9	6
12	Alterations in HLA Class I-Presented Immunopeptidome and Class I-Interactome upon Osimertinib Resistance in EGFR Mutant Lung Adenocarcinoma. <i>Cancers</i> , 2021, 13, 4977.	3.7	5
13	Abstract 2927: Proteogenomic heterogeneity in metastatic lung adenocarcinoma revealed from rapid/warm autopsy. , 2017, , .		0
14	Abstract 4528: Quantitative mass spectrometry to interrogate proteomic heterogeneity in metastatic lung adenocarcinoma and validate a novel somatic mutation CDK12-G879V. , 2019, , .		0