

Trever G Bivona

List of Publications by Year in descending order

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Version: 2024-02-01

79
papers

11,139
citations

81900

39
h-index

69250

77
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86
all docs

86
docs citations

86
times ranked

18010
citing authors

#	ARTICLE	IF	CITATIONS
1	Evolution of metastasis: new tools and insights. <i>Trends in Cancer</i> , 2022, 8, 98-109.	7.4	40
2	Inhibition of SHP2 as an approach to block RAS-driven cancers. <i>Advances in Cancer Research</i> , 2022, 153, 205-236.	5.0	7
3	Small-molecule targeted therapies induce dependence on DNA double-strand break repair in residual tumor cells. <i>Science Translational Medicine</i> , 2022, 14, eabc7480.	12.4	14
4	Lineage tracing reveals the phylodynamics, plasticity, and paths of tumor evolution. <i>Cell</i> , 2022, 185, 1905-1923.e25.	28.9	108
5	Deficiency of the splicing factor RBM10 limits EGFR inhibitor response in EGFR-mutant lung cancer. <i>Journal of Clinical Investigation</i> , 2022, 132, .	8.2	15
6	Single-cell lineages reveal the rates, routes, and drivers of metastasis in cancer xenografts. <i>Science</i> , 2021, 371, .	12.6	166
7	Kinase-mediated RAS signaling via membraneless cytoplasmic protein granules. <i>Cell</i> , 2021, 184, 2649-2664.e18.	28.9	102
8	Allosteric SHP2 inhibitors in cancer: Targeting the intersection of RAS, resistance, and the immune microenvironment. <i>Current Opinion in Chemical Biology</i> , 2021, 62, 1-12.	6.1	83
9	Stepwise evolution of therapy resistance in AML. <i>Cancer Cell</i> , 2021, 39, 904-906.	16.8	3
10	Targeting AXL in NSCLC. <i>Lung Cancer: Targets and Therapy</i> , 2021, Volume 12, 67-79.	2.7	8
11	Exocyst protein subnetworks integrate Hippo and mTOR signaling to promote virus detection and cancer. <i>Cell Reports</i> , 2021, 36, 109491.	6.4	11
12	Liquid Biopsy for Advanced NSCLC: A Consensus Statement From the International Association for the Study of Lung Cancer. <i>Journal of Thoracic Oncology</i> , 2021, 16, 1647-1662.	1.1	274
13	Profiling Sensitivity to Targeted Therapies in EGFR-Mutant NSCLC Patient-Derived Organoids. <i>Journal of Visualized Experiments</i> , 2021, , .	0.3	2
14	Co-occurring Alterations in the RAS-MAPK Pathway Limit Response to MET Inhibitor Treatment in MET Exon 14 Skipping Mutation-Positive Lung Cancer. <i>Clinical Cancer Research</i> , 2020, 26, 439-449.	7.0	64
15	Acquired Resistance in Lung Cancer. <i>Annual Review of Cancer Biology</i> , 2020, 4, 279-297.	4.5	13
16	Functional screening identifies aryl hydrocarbon receptor as suppressor of lung cancer metastasis. <i>Oncogenesis</i> , 2020, 9, 102.	4.9	24
17	Multi-faceted epigenetic dysregulation of gene expression promotes esophageal squamous cell carcinoma. <i>Nature Communications</i> , 2020, 11, 3675.	12.8	63
18	Long non-coding RNA ESCCAL-1 promotes esophageal squamous cell carcinoma by down regulating the negative regulator of APOBEC3G. <i>Cancer Letters</i> , 2020, 493, 217-227.	7.2	17

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19	Therapy-Induced Evolution of Human Lung Cancer Revealed by Single-Cell RNA Sequencing. <i>Cell</i> , 2020, 182, 1232-1251.e22.	28.9	371
20	EGFR targeted therapy resistance: current status, challenges, and future outlook. <i>Journal of Thoracic Disease</i> , 2020, 12, 2849-2850.	1.4	2
21	Molecular Landscape of BRAF-Mutant NSCLC Reveals an Association Between Clonality and Driver Mutations and Identifies Targetable Non-V600 Driver Mutations. <i>Journal of Thoracic Oncology</i> , 2020, 15, 1611-1623.	1.1	43
22	Targeting Oncogenic BRAF: Past, Present, and Future. <i>Cancers</i> , 2019, 11, 1197.	3.7	143
23	Synthetic Essentiality of Metabolic Regulator PDHK1 in PTEN-Deficient Cells and Cancers. <i>Cell Reports</i> , 2019, 28, 2317-2330.e8.	6.4	12
24	BRAF Mutations Classes I, II, and III in NSCLC Patients Included in the SLLIP Trial: The Need for a New Pre-Clinical Treatment Rationale. <i>Cancers</i> , 2019, 11, 1381.	3.7	44
25	Principles of Resistance to Targeted Cancer Therapy: Lessons from Basic and Translational Cancer Biology. <i>Trends in Molecular Medicine</i> , 2019, 25, 185-197.	6.7	118
26	Use of nonsteroidal anti-inflammatory drugs predicts improved patient survival for <i>PIK3CA</i> -altered head and neck cancer. <i>Journal of Experimental Medicine</i> , 2019, 216, 419-427.	8.5	46
27	Engineering Multidimensional Evolutionary Forces to Combat Cancer. <i>Cancer Discovery</i> , 2019, 9, 587-604.	9.4	13
28	Dampening oncogenic RAS signaling. <i>Science</i> , 2019, 363, 1280-1281.	12.6	25
29	Polytherapy and Targeted Cancer Drug Resistance. <i>Trends in Cancer</i> , 2019, 5, 170-182.	7.4	183
30	Emerging Targeted Therapies for the Treatment of Non-small Cell Lung Cancer. <i>Current Oncology Reports</i> , 2019, 21, 21.	4.0	82
31	Pathologic Complete Response to Neoadjuvant Crizotinib in a Lung Adenocarcinoma Patient With a MET Exon 14 Skipping Mutation. <i>Clinical Lung Cancer</i> , 2019, 20, e137-e141.	2.6	8
32	Immunohistochemistry to Study YAP in Human Tissue Samples. <i>Methods in Molecular Biology</i> , 2019, 1893, 89-95.	0.9	6
33	Aurora kinase A drives the evolution of resistance to third-generation EGFR inhibitors in lung cancer. <i>Nature Medicine</i> , 2019, 25, 111-118.	30.7	196
34	Differential Subcellular Localization Regulates Oncogenic Signaling by ROS1 Kinase Fusion Proteins. <i>Cancer Research</i> , 2019, 79, 546-556.	0.9	59
35	Oncogenic Signaling Pathways in The Cancer Genome Atlas. <i>Cell</i> , 2018, 173, 321-337.e10.	28.9	2,111
36	<i>BRAF</i> in Lung Cancers: Analysis of Patient Cases Reveals Recurrent <i>BRAF</i> Mutations, Fusions, Kinase Duplications, and Concurrent Alterations. <i>JCO Precision Oncology</i> , 2018, 2, 1-15.	3.0	24

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37	Circulating tumor DNA analysis in patients with EGFR mutant lung cancer. <i>Journal of Thoracic Disease</i> , 2018, 10, S4061-S4064.	1.4	2
38	The evolving understanding of immunoediting and the clinical impact of immune escape. <i>Journal of Thoracic Disease</i> , 2018, 10, 1248-1252.	1.4	10
39	Liquid Biopsy for Advanced Non-Small Cell Lung Cancer (NSCLC): A Statement Paper from the IASLC. <i>Journal of Thoracic Oncology</i> , 2018, 13, 1248-1268.	1.1	515
40	RAS nucleotide cycling underlies the SHP2 phosphatase dependence of mutant BRAF-, NF1- and RAS-driven cancers. <i>Nature Cell Biology</i> , 2018, 20, 1064-1073.	10.3	276
41	TNF-driven adaptive response mediates resistance to EGFR inhibition in lung cancer. <i>Journal of Clinical Investigation</i> , 2018, 128, 2500-2518.	8.2	73
42	Emerging application of genomics-guided therapeutics in personalized lung cancer treatment. <i>Annals of Translational Medicine</i> , 2018, 6, 160-160.	1.7	22
43	Superior Efficacy and Selectivity of Novel Small-Molecule Kinase Inhibitors of T790M-Mutant EGFR in Preclinical Models of Lung Cancer. <i>Cancer Research</i> , 2017, 77, 1200-1211.	0.9	11
44	CD74 - ROS1 Fusion in NSCLC Detected by Hybrid Capture-Based Tissue Genomic Profiling and ctDNA Assays. <i>Journal of Thoracic Oncology</i> , 2017, 12, e19-e20.	1.1	6
45	Resistance is futile: overcoming resistance to targeted therapies in lung adenocarcinoma. <i>Npj Precision Oncology</i> , 2017, 1, .	5.4	70
46	A Case of Metastatic Atypical Neuroendocrine Tumor with <i>ALK</i> Translocation and Diffuse Brain Metastases. <i>Oncologist</i> , 2017, 22, 768-773.	3.7	33
47	Novel computational method for predicting polytherapy switching strategies to overcome tumor heterogeneity and evolution. <i>Scientific Reports</i> , 2017, 7, 44206.	3.3	28
48	Understanding and targeting resistance mechanisms in NSCLC. <i>Nature Reviews Cancer</i> , 2017, 17, 637-658.	28.4	679
49	Simultaneous evolutionary expansion and constraint of genomic heterogeneity in multifocal lung cancer. <i>Nature Communications</i> , 2017, 8, 823.	12.8	53
50	Co-activation of STAT3 and YES-Associated Protein 1 (YAP1) Pathway in EGFR-Mutant NSCLC. <i>Journal of the National Cancer Institute</i> , 2017, 109, .	6.3	128
51	Current Landscape of Targeted Therapy in Lung Cancer. <i>Clinical Pharmacology and Therapeutics</i> , 2017, 102, 757-764.	4.7	134
52	Evolution and clinical impact of co-occurring genetic alterations in advanced-stage EGFR-mutant lung cancers. <i>Nature Genetics</i> , 2017, 49, 1693-1704.	21.4	423
53	Inactivation of Capicua drives cancer metastasis. <i>Nature Genetics</i> , 2017, 49, 87-96.	21.4	130
54	Convergent Akt activation drives acquired EGFR inhibitor resistance in lung cancer. <i>Nature Communications</i> , 2017, 8, 410.	12.8	117

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55	Dividing and conquering the variation among variants in EML4-ALK lung cancer. <i>Translational Cancer Research</i> , 2017, 6, S369-S370.	1.0	0
56	Neoadjuvant Oncogene-Targeted Therapy in Early Stage Non-Small-Cell Lung Cancer as a Strategy to Improve Clinical Outcome and Identify Early Mechanisms of Resistance. <i>Clinical Lung Cancer</i> , 2016, 17, 466-469.	2.6	16
57	RAS-MAPK signaling influences the efficacy of ALK-targeting agents in lung cancer. <i>Molecular and Cellular Oncology</i> , 2016, 3, e1091061.	0.7	7
58	A framework for understanding and targeting residual disease in oncogene-driven solid cancers. <i>Nature Medicine</i> , 2016, 22, 472-478.	30.7	145
59	Combined chemical-genetic approach identifies cytosolic HSP70 dependence in rhabdomyosarcoma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 9015-9020.	7.1	33
60	Preclinical efficacy of a RAF inhibitor that evades paradoxical MAPK pathway activation in protein kinase <i>BRAF</i> -mutant lung cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 13456-13461.	7.1	65
61	An improved CTC isolation scheme for pairing with downstream genomics: Demonstrating clinical utility in metastatic prostate, lung and pancreatic cancer. <i>Cancer Letters</i> , 2016, 380, 144-152.	7.2	26
62	The Hippo effector YAP regulates the response of cancer cells to MAPK pathway inhibitors. <i>Molecular and Cellular Oncology</i> , 2016, 3, e1021441.	0.7	29
63	RAS signaling in ALK fusion lung cancer. <i>Small GTPases</i> , 2016, 7, 32-33.	1.6	20
64	Tracking Down Response and Resistance to TRK Inhibitors. <i>Cancer Discovery</i> , 2016, 6, 14-16.	9.4	14
65	Transcriptomic-metabolomic reprogramming in EGFR-mutant NSCLC early adaptive drug escape linking TGF β 2-bioenergetics-mitochondrial priming. <i>Oncotarget</i> , 2016, 7, 82013-82027.	1.8	23
66	AXL receptor tyrosine kinase as a therapeutic target in NSCLC. <i>Lung Cancer: Targets and Therapy</i> , 2015, 6, 27.	2.7	18
67	AUY922 Effectively Overcomes MET- and AXL-Mediated Resistance to EGFR-TKI in Lung Cancer Cells. <i>PLoS ONE</i> , 2015, 10, e0119832.	2.5	28
68	RAS-MAPK in ALK targeted therapy resistance. <i>Cell Cycle</i> , 2015, 14, 3661-3662.	2.6	12
69	The Hippo effector YAP promotes resistance to RAF- and MEK-targeted cancer therapies. <i>Nature Genetics</i> , 2015, 47, 250-256.	21.4	434
70	Cell-Free DNA Next-Generation Sequencing in Pancreatobiliary Carcinomas. <i>Cancer Discovery</i> , 2015, 5, 1040-1048.	9.4	226
71	NF- κ B-Activating Complex Engaged in Response to EGFR Oncogene Inhibition Drives Tumor Cell Survival and Residual Disease in Lung Cancer. <i>Cell Reports</i> , 2015, 11, 98-110.	6.4	178
72	AXL Mediates Resistance to PI3K Inhibition by Activating the EGFR/PKC/mTOR Axis in Head and Neck and Esophageal Squamous Cell Carcinomas. <i>Cancer Cell</i> , 2015, 27, 533-546.	16.8	263

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73	RAS-MAPK dependence underlies a rational polytherapy strategy in EML4-ALK ⁺ positive lung cancer. <i>Nature Medicine</i> , 2015, 21, 1038-1047.	30.7	245
74	Mapping the molecular determinants of BRAF oncogene dependence in human lung cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E748-57.	7.1	90
75	Plasma mRNA expression levels of BRCA1 and TS as potential predictive biomarkers for chemotherapy in gastric cancer. <i>Journal of Translational Medicine</i> , 2014, 12, 355.	4.4	18
76	Recent advances in personalized lung cancer medicine. <i>Personalized Medicine</i> , 2014, 11, 309-321.	1.5	22
77	Activation of the AXL kinase causes resistance to EGFR-targeted therapy in lung cancer. <i>Nature Genetics</i> , 2012, 44, 852-860.	21.4	1,049
78	FAS and NF- κ B signalling modulate dependence of lung cancers on mutant EGFR. <i>Nature</i> , 2011, 471, 523-526.	27.8	374
79	Ras signalling on the endoplasmic reticulum and the Golgi. <i>Nature Cell Biology</i> , 2002, 4, 343-350.	10.3	582