

Ignacio I Wistuba

List of Publications by Year in descending order

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Version: 2024-02-01

312
papers

36,436
citations

6124

83
h-index

4622

176
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all docs

323
docs citations

323
times ranked

45717
citing authors

#	ARTICLE	IF	CITATIONS
1	The immunogenomic landscape of resected intrahepatic cholangiocarcinoma. <i>Hepatology</i> , 2022, 75, 297-308.	3.6	32
2	SRGN-Triggered Aggressive and Immunosuppressive Phenotype in a Subset of TTF-1â€“Negative Lung Adenocarcinomas. <i>Journal of the National Cancer Institute</i> , 2022, 114, 290-301.	3.0	18
3	Artificial intelligence strategy integrating morphologic and architectural biomarkers provides robust diagnostic accuracy for disease progression in chronic lymphocytic leukemia. <i>Journal of Pathology</i> , 2022, 256, 4-14.	2.1	18
4	Effector memory cytotoxic CD3+/CD8+/CD45RO+ T cells are predictive of good survival and a lower risk of recurrence in triple-negative breast cancer. <i>Modern Pathology</i> , 2022, 35, 601-608.	2.9	10
5	Surgical approach does not influence changes in circulating immune cell populations following lung cancer resection. <i>Lung Cancer</i> , 2022, 164, 69-75.	0.9	2
6	Expression of TRPS1 in phyllodes tumor and sarcoma of the breast. <i>Human Pathology</i> , 2022, 121, 73-80.	1.1	18
7	Combined IL-2, agonistic CD3 and 4-1BB stimulation preserve clonotype hierarchy in propagated non-small cell lung cancer tumor-infiltrating lymphocytes. , 2022, 10, e003082.		11
8	ISA101 and nivolumab for HPV-16⁺ cancer: updated clinical efficacy and immune correlates of response. , 2022, 10, e004232.		38
9	AXL targeting restores PD-1 blockade sensitivity of STK11/LKB1 mutant NSCLC through expansion of TCF1+ CD8 TAcells. <i>Cell Reports Medicine</i> , 2022, 3, 100554.	3.3	29
10	MTAP deficiency creates an exploitable target for antifolate therapy in 9p21-loss cancers. <i>Nature Communications</i> , 2022, 13, 1797.	5.8	23
11	Assessment of Clinical Response Following Atezolizumab and Bevacizumab Treatment in Patients With Neuroendocrine Tumors. <i>JAMA Oncology</i> , 2022, 8, 904.	3.4	13
12	Dynamic expression of Schlafen 11 (SLFN11) in circulating tumour cells as a liquid biomarker in small cell lung cancer. <i>British Journal of Cancer</i> , 2022, 127, 569-576.	2.9	8
13	Diminished Immune Surveillance during Histologic Progression of Intraductal Papillary Mucinous Neoplasms Offers a Therapeutic Opportunity for Cancer Interception. <i>Clinical Cancer Research</i> , 2022, 28, 1938-1947.	3.2	11
14	Immunogenomic intertumor heterogeneity across primary and metastatic sites in a patient with lung adenocarcinoma. <i>Journal of Experimental and Clinical Cancer Research</i> , 2022, 41, 172.	3.5	2
15	Chronic Lymphocytic Leukemia Progression Diagnosis with Intrinsic Cellular Patterns via Unsupervised Clustering. <i>Cancers</i> , 2022, 14, 2398.	1.7	6
16	Mutational Activation of the NRF2 Pathway Upregulates Kynureninase Resulting in Tumor Immunosuppression and Poor Outcome in Lung Adenocarcinoma. <i>Cancers</i> , 2022, 14, 2543.	1.7	16
17	The microRNA-183/96/182 cluster inhibits lung cancer progression and metastasis by inducing an interleukin-2-mediated antitumor CD8⁺ cytotoxic T-cell response. <i>Genes and Development</i> , 2022, 36, 582-600.	2.7	9
18	Naproxen chemoprevention promotes immune activation in Lynch syndrome colorectal mucosa. <i>Gut</i> , 2021, 70, 555-566.	6.1	37

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19	Female Gender Predicts Augmented Immune Infiltration in Lung Adenocarcinoma. <i>Clinical Lung Cancer</i> , 2021, 22, e415-e424.	1.1	10
20	TRPS1: a highly sensitive and specific marker for breast carcinoma, especially for triple-negative breast cancer. <i>Modern Pathology</i> , 2021, 34, 710-719.	2.9	90
21	Pathologic Assessment of Lung Squamous Cell Carcinoma After Neoadjuvant Immunotherapy. <i>Journal of Thoracic Oncology</i> , 2021, 16, e9-e10.	0.5	6
22	Altered Regulation of HIF-1 α in Naive- and Drug-Resistant EGFR-Mutant NSCLC: Implications for a Vascular Endothelial Growth Factor-Dependent Phenotype. <i>Journal of Thoracic Oncology</i> , 2021, 16, 439-451.	0.5	34
23	Expression of BCL2 alternative proteins and association with outcome in CLL patients treated with venetoclax. <i>Leukemia and Lymphoma</i> , 2021, 62, 1129-1135.	0.6	6
24	Neoadjuvant Chemotherapy Increases Cytotoxic T Cell, Tissue Resident Memory T Cell, and B Cell Infiltration in Resectable NSCLC. <i>Journal of Thoracic Oncology</i> , 2021, 16, 127-139.	0.5	48
25	Evolution of DNA methylome from precancerous lesions to invasive lung adenocarcinomas. <i>Nature Communications</i> , 2021, 12, 687.	5.8	30
26	Identification of distinct immune landscapes using an automated nine-color multiplex immunofluorescence staining panel and image analysis in paraffin tumor tissues. <i>Scientific Reports</i> , 2021, 11, 4530.	1.6	27
27	Neoadjuvant nivolumab or nivolumab plus ipilimumab in operable non-small cell lung cancer: the phase 2 randomized NEOSTAR trial. <i>Nature Medicine</i> , 2021, 27, 504-514.	15.2	357
28	Pathology and Classification of SCLC. <i>Cancers</i> , 2021, 13, 820.	1.7	68
29	PD-L1 as a biomarker of response to immune-checkpoint inhibitors. <i>Nature Reviews Clinical Oncology</i> , 2021, 18, 345-362.	12.5	646
30	Elevated NSD3 histone methylation activity drives squamous cell lung cancer. <i>Nature</i> , 2021, 590, 504-508.	13.7	79
31	Rare deleterious germline variants and risk of lung cancer. <i>Npj Precision Oncology</i> , 2021, 5, 12.	2.3	19
32	Single-Cell Expression Landscape of SARS-CoV-2 Receptor ACE2 and Host Proteases in Normal and Malignant Lung Tissues from Pulmonary Adenocarcinoma Patients. <i>Cancers</i> , 2021, 13, 1250.	1.7	7
33	Patterns of transcription factor programs and immune pathway activation define four major subtypes of SCLC with distinct therapeutic vulnerabilities. <i>Cancer Cell</i> , 2021, 39, 346-360.e7.	7.7	422
34	Cell-autonomous immune gene expression is repressed in pulmonary neuroendocrine cells and small cell lung cancer. <i>Communications Biology</i> , 2021, 4, 314.	2.0	44
35	Global analysis of shared T α cell specificities in human non-small cell lung cancer enables HLA inference and antigen discovery. <i>Immunity</i> , 2021, 54, 586-602.e8.	6.6	80
36	The Transcriptomic Landscape of Mismatch Repair-Deficient Intestinal Stem Cells. <i>Cancer Research</i> , 2021, 81, 2760-2773.	0.4	7

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37	Pilot Clinical Trial of Perioperative Durvalumab and Tremelimumab in the Treatment of Resectable Colorectal Cancer Liver Metastases. <i>Clinical Cancer Research</i> , 2021, 27, 3039-3049.	3.2	28
38	CD8+ T cells inhibit metastasis and CXCL4 regulates its function. <i>British Journal of Cancer</i> , 2021, 125, 176-189.	2.9	21
39	Multiplex Immunofluorescence Tyramide Signal Amplification for Immune Cell Profiling of Paraffin-Embedded Tumor Tissues. <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 667067.	1.6	26
40	Contextual cues from cancer cells govern cancer-associated fibroblast heterogeneity. <i>Cell Reports</i> , 2021, 35, 109009.	2.9	18
41	Characterization of the Immune Landscape of EGFR-Mutant NSCLC Identifies CD73/Adenosine Pathway as a Potential Therapeutic Target. <i>Journal of Thoracic Oncology</i> , 2021, 16, 583-600.	0.5	62
42	Resolving the Spatial and Cellular Architecture of Lung Adenocarcinoma by Multiregion Single-Cell Sequencing. <i>Cancer Discovery</i> , 2021, 11, 2506-2523.	7.7	68
43	Immune evolution from preneoplasia to invasive lung adenocarcinomas and underlying molecular features. <i>Nature Communications</i> , 2021, 12, 2722.	5.8	74
44	Implementation of a Novel Web-Based Lesion Selection Tool to Improve Acquisition of Tumor Biopsy Specimens. <i>Journal of Immunotherapy and Precision Oncology</i> , 2021, 4, 45-52.	0.6	5
45	Efficacy, Safety, and Biomarker Analysis of Combined PD-L1 (Atezolizumab) and VEGF (Bevacizumab) Blockade in Advanced Malignant Peritoneal Mesothelioma. <i>Cancer Discovery</i> , 2021, 11, 2738-2747.	7.7	37
46	Blood biomarkers associated to complete pathological response on NSCLC patients treated with neoadjuvant chemoimmunotherapy included in NADIM clinical trial. <i>Clinical and Translational Medicine</i> , 2021, 11, e491.	1.7	26
47	Immune Phenotype and Response to Neoadjuvant Therapy in Triple-Negative Breast Cancer. <i>Clinical Cancer Research</i> , 2021, 27, 5365-5375.	3.2	29
48	A wake-up call for cancer DNA damage: the role of Schlafen 11 (SLFN11) across multiple cancers. <i>British Journal of Cancer</i> , 2021, 125, 1333-1340.	2.9	22
49	Multiplex Tissue Imaging Harmonization: A Multicenter Experience from CIMAC-CIDC Immuno-Oncology Biomarkers Network. <i>Clinical Cancer Research</i> , 2021, 27, 5072-5083.	3.2	10
50	Multi-institutional TSA-amplified Multiplexed Immunofluorescence Reproducibility Evaluation (MITRE) Study. , 2021, 9, e002197.		44
51	Immune Profiling Mass Cytometry Assay Harmonization: Multicenter Experience from CIMAC-CIDC. <i>Clinical Cancer Research</i> , 2021, 27, 5062-5071.	3.2	8
52	Estrogen Promotes Resistance to Bevacizumab in Murine Models of NSCLC. <i>Journal of Thoracic Oncology</i> , 2021, 16, 2051-2064.	0.5	6
53	The Combiome Hypothesis: Selecting Optimal Treatment for Cancer Patients. <i>Clinical Lung Cancer</i> , 2021, , .	1.1	4
54	Pretreatment Tissue TCR Repertoire Evenness Is Associated with Complete Pathologic Response in Patients with NSCLC Receiving Neoadjuvant Chemoimmunotherapy. <i>Clinical Cancer Research</i> , 2021, 27, 5878-5890.	3.2	30

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55	Nodal immune flare mimics nodal disease progression following neoadjuvant immune checkpoint inhibitors in non-small cell lung cancer. <i>Nature Communications</i> , 2021, 12, 5045.	5.8	42
56	Overexpression of CD73 in pancreatic ductal adenocarcinoma is associated with immunosuppressive tumor microenvironment and poor survival. <i>Pancreatology</i> , 2021, 21, 942-949.	0.5	16
57	Evaluation of Pathologic Response in Lymph Nodes of Patients With Lung Cancer Receiving Neoadjuvant Chemotherapy. <i>Journal of Thoracic Oncology</i> , 2021, 16, 1289-1297.	0.5	27
58	BCL-W expression associates with poor outcome in patients with peripheral T-cell lymphoma not otherwise specified. <i>Blood Cancer Journal</i> , 2021, 11, 153.	2.8	1
59	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.	0.5	274
60	Network for Biomarker Immunoprofiling for Cancer Immunotherapy: Cancer Immune Monitoring and Analysis Centers and Cancer Immunologic Data Commons (CIMAC-CIDC). <i>Clinical Cancer Research</i> , 2021, 27, 5038-5048.	3.2	13
61	CD73 expression defines immune, molecular, and clinicopathological subgroups of lung adenocarcinoma. <i>Cancer Immunology, Immunotherapy</i> , 2021, 70, 1965-1976.	2.0	14
62	Transition From a Standard to a Hybrid On-Site and Remote Anatomic Pathology Training Model During the Coronavirus Disease 2019 (COVID-19) Pandemic. <i>Archives of Pathology and Laboratory Medicine</i> , 2021, 145, 22-31.	1.2	25
63	An analysis of research biopsy core variability from over 5000 prospectively collected core samples. <i>Npj Precision Oncology</i> , 2021, 5, 94.	2.3	4
64	NSD2 dimethylation at H3K36 promotes lung adenocarcinoma pathogenesis. <i>Molecular Cell</i> , 2021, 81, 4481-4492.e9.	4.5	42
65	Sirp α and CSF1R Positive Tissue Macrophages Are Increased in Patients with Follicular Lymphoma Who Relapse after Frontline Lenalidomide and Rituximab. <i>Blood</i> , 2021, 138, 1336-1336.	0.6	0
66	Cold and heterogeneous T cell repertoire is associated with copy number aberrations and loss of immune genes in small-cell lung cancer. <i>Nature Communications</i> , 2021, 12, 6655.	5.8	24
67	The histologic phenotype of lung cancers is associated with transcriptomic features rather than genomic characteristics. <i>Nature Communications</i> , 2021, 12, 7081.	5.8	16
68	The Prognostic and Therapeutic Role of Genomic Subtyping by Sequencing Tumor or Cell-Free DNA in Pulmonary Large-Cell Neuroendocrine Carcinoma. <i>Clinical Cancer Research</i> , 2020, 26, 892-901.	3.2	80
69	Lymphovascular Invasion Is Associated With Mutational Burden and PD-L1 in Resected Lung Cancer. <i>Annals of Thoracic Surgery</i> , 2020, 109, 358-366.	0.7	9
70	Therapeutic targeting of the PI4K2A/PKR lysosome network is critical for misfolded protein clearance and survival in cancer cells. <i>Oncogene</i> , 2020, 39, 801-813.	2.6	16
71	A proposal for pathologic processing of breast implant capsules in patients with suspected breast implant anaplastic large cell lymphoma. <i>Modern Pathology</i> , 2020, 33, 367-379.	2.9	29
72	RUVBL1/RUVBL2 ATPase Activity Drives PAQosome Maturation, DNA Replication and Radioresistance in Lung Cancer. <i>Cell Chemical Biology</i> , 2020, 27, 105-121.e14.	2.5	38

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73	Multifactorial Deep Learning Reveals Pan-Cancer Genomic Tumor Clusters with Distinct Immunogenomic Landscape and Response to Immunotherapy. <i>Clinical Cancer Research</i> , 2020, 26, 2908-2920.	3.2	30
74	From clinical specimens to human cancer preclinical models—a journey the NCI cell line database 25 years later. <i>Journal of Cellular Biochemistry</i> , 2020, 121, 3986-3999.	1.2	6
75	Distinct co-acquired alterations and genomic evolution during TKI treatment in non-small-cell lung cancer patients with or without acquired T790M mutation. <i>Oncogene</i> , 2020, 39, 1846-1859.	2.6	29
76	Round Robin Evaluation of MET Protein Expression in Lung Adenocarcinomas Improves Interobserver Concordance. <i>Applied Immunohistochemistry and Molecular Morphology</i> , 2020, 28, 669-677.	0.6	5
77	Molecular differences across invasive lung adenocarcinoma morphological subgroups. <i>Translational Lung Cancer Research</i> , 2020, 9, 1029-1040.	1.3	3
78	Neoadjuvant chemotherapy and nivolumab in resectable non-small-cell lung cancer (NADIM): an open-label, multicentre, single-arm, phase 2 trial. <i>Lancet Oncology</i> , The, 2020, 21, 1413-1422.	5.1	475
79	Collagen promotes anti-PD-1/PD-L1 resistance in cancer through LAIR1-dependent CD8+ T cell exhaustion. <i>Nature Communications</i> , 2020, 11, 4520.	5.8	218
80	Evolution of Genomic and T-cell Repertoire Heterogeneity of Malignant Pleural Mesothelioma Under Dasatinib Treatment. <i>Clinical Cancer Research</i> , 2020, 26, 5477-5486.	3.2	15
81	Sarcomatoid Mesothelioma: A CDKN2A molecular analysis of 53 cases with immunohistochemical correlation with BAP1. <i>Pathology Research and Practice</i> , 2020, 216, 153267.	1.0	7
82	Multomics profiling of primary lung cancers and distant metastases reveals immunosuppression as a common characteristic of tumor cells with metastatic plasticity. <i>Genome Biology</i> , 2020, 21, 271.	3.8	36
83	The Society for Immunotherapy of Cancer statement on best practices for multiplex immunohistochemistry (IHC) and immunofluorescence (IF) staining and validation. , 2020, 8, e000155.		140
84	Neutrophil expansion defines an immunoinhibitory peripheral and intratumoral inflammatory milieu in resected non-small cell lung cancer: a descriptive analysis of a prospectively immunoprofiled cohort. , 2020, 8, e000405.		33
85	Hypertrophic lichenoid dermatitis immune-related adverse event during combined immune checkpoint and exportin inhibitor therapy: A diagnostic pitfall for superficially invasive squamous cell carcinoma. <i>Journal of Cutaneous Pathology</i> , 2020, 47, 954-959.	0.7	8
86	Plasma-Derived Extracellular Vesicles Convey Protein Signatures That Reflect Pathophysiology in Lung and Pancreatic Adenocarcinomas. <i>Cancers</i> , 2020, 12, 1147.	1.7	20
87	Clinical and Genomic Characteristics of Small Cell Lung Cancer in Never Smokers. <i>Chest</i> , 2020, 158, 1723-1733.	0.4	16
88	An in vivo functional genomics screen of nuclear receptors and their co-regulators identifies FOXA1 as an essential gene in lung tumorigenesis. <i>Neoplasia</i> , 2020, 22, 294-310.	2.3	21
89	Decrease in tumor content assessed in biopsies is associated with improved treatment outcome response to pembrolizumab in patients with rare tumors. , 2020, 8, e000665.		8
90	Interplay between estrogen and Stat3/NF- κ B-driven immunomodulation in lung cancer. <i>Carcinogenesis</i> , 2020, 41, 1529-1542.	1.3	9

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91	A Phase II Trial of Alisertib (MLN8237) in Salvage Malignant Mesothelioma. <i>Oncologist</i> , 2020, 25, e1457-e1463.	1.9	7
92	A Grading System for Invasive Pulmonary Adenocarcinoma: A Proposal From the International Association for the Study of Lung Cancer Pathology Committee. <i>Journal of Thoracic Oncology</i> , 2020, 15, 1599-1610.	0.5	234
93	Genomic assessment distinguishes intrapulmonary metastases from synchronous primary lung cancers. <i>Journal of Thoracic Disease</i> , 2020, 12, 1952-1959.	0.6	6
94	IASLC Multidisciplinary Recommendations for Pathologic Assessment of Lung Cancer Resection Specimens After Neoadjuvant Therapy. <i>Journal of Thoracic Oncology</i> , 2020, 15, 709-740.	0.5	205
95	STING Pathway Expression Identifies NSCLC With an Immune-Responsive Phenotype. <i>Journal of Thoracic Oncology</i> , 2020, 15, 777-791.	0.5	94
96	Computational Staining of Pathology Images to Study the Tumor Microenvironment in Lung Cancer. <i>Cancer Research</i> , 2020, 80, 2056-2066.	0.4	88
97	PI4KIII ² is a therapeutic target in chromosome 1q amplified lung adenocarcinoma. <i>Science Translational Medicine</i> , 2020, 12, .	5.8	41
98	Procedural Requirements and Recommendations for Multiplex Immunofluorescence Tyramide Signal Amplification Assays to Support Translational Oncology Studies. <i>Cancers</i> , 2020, 12, 255.	1.7	58
99	A Phase I/II Study of Neoadjuvant Cisplatin, Docetaxel, and Nintedanib for Resectable Non-Small Cell Lung Cancer. <i>Clinical Cancer Research</i> , 2020, 26, 3525-3536.	3.2	22
100	Comprehensive Molecular Characterization Identifies Distinct Genomic and Immune Hallmarks of Renal Medullary Carcinoma. <i>Cancer Cell</i> , 2020, 37, 720-734.e13.	7.7	74
101	eIF5B drives integrated stress response-dependent translation of PD-L1 in lung cancer. <i>Nature Cancer</i> , 2020, 1, 533-545.	5.7	73
102	Agreement on Major Pathological Response in NSCLC Patients Receiving Neoadjuvant Chemotherapy. <i>Clinical Lung Cancer</i> , 2020, 21, 341-348.	1.1	70
103	¹⁸ F-fluorodeoxyglucose positron emission tomography correlates with tumor immunometabolic phenotypes in resected lung cancer. <i>Cancer Immunology, Immunotherapy</i> , 2020, 69, 1519-1534.	2.0	21
104	T-Cell Repertoire in Combination with T-Cell Density Predicts Clinical Outcomes in Patients with Merkel Cell Carcinoma. <i>Journal of Investigative Dermatology</i> , 2020, 140, 2146-2156.e4.	0.3	14
105	Multiplex Immunofluorescence Assays. <i>Methods in Molecular Biology</i> , 2020, 2055, 467-495.	0.4	44
106	LKB1/STK11 Expression in Lung Adenocarcinoma and Associations With Patterns of Recurrence. <i>Annals of Thoracic Surgery</i> , 2020, 110, 1131-1138.	0.7	8
107	Programmed Death-Ligand 1 Heterogeneity and Its Impact on Benefit From Immune Checkpoint Inhibitors in NSCLC. <i>Journal of Thoracic Oncology</i> , 2020, 15, 1449-1459.	0.5	109
108	The International Association for the Study of Lung Cancer Global Survey on Molecular Testing in Lung Cancer. <i>Journal of Thoracic Oncology</i> , 2020, 15, 1434-1448.	0.5	107

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109	Comprehensive T cell repertoire characterization of non-small cell lung cancer. <i>Nature Communications</i> , 2020, 11, 603.	5.8	140
110	Single-cell analyses reveal increased intratumoral heterogeneity after the onset of therapy resistance in small-cell lung cancer. <i>Nature Cancer</i> , 2020, 1, 423-436.	5.7	218
111	Removal of N-Linked Glycosylation Enhances PD-L1 Detection and Predicts Anti-PD-1/PD-L1 Therapeutic Efficacy. <i>Cancer Cell</i> , 2019, 36, 168-178.e4.	7.7	240
112	Tumor cellular proliferation is associated with enhanced immune checkpoint expression in stage I non-small cell lung cancer. <i>Journal of Thoracic and Cardiovascular Surgery</i> , 2019, 158, 911-919.e6.	0.4	21
113	Multi-region exome sequencing reveals genomic evolution from preneoplasia to lung adenocarcinoma. <i>Nature Communications</i> , 2019, 10, 2978.	5.8	91
114	Characterization and Comparison of GITR Expression in Solid Tumors. <i>Clinical Cancer Research</i> , 2019, 25, 6501-6510.	3.2	37
115	Tyrosine Threonine Kinase Inhibition Eliminates Lung Cancers by Augmenting Apoptosis and Polyploidy. <i>Molecular Cancer Therapeutics</i> , 2019, 18, 1775-1786.	1.9	21
116	Hybrid oncocytic/chromophobe renal tumors are molecularly distinct from oncocytoma and chromophobe renal cell carcinoma. <i>Modern Pathology</i> , 2019, 32, 1698-1707.	2.9	35
117	Development and Application of Duplex Sequencing Strategy for Cell-Free DNA-Based Longitudinal Monitoring of Stage IV Colorectal Cancer. <i>Journal of Molecular Diagnostics</i> , 2019, 21, 994-1009.	1.2	4
118	Non-invasive genotyping of metastatic colorectal cancer using circulating cell free DNA. <i>Cancer Genetics</i> , 2019, 237, 82-89.	0.2	7
119	Sphingosine Kinase 1 Signaling Promotes Metastasis of Triple-Negative Breast Cancer. <i>Cancer Research</i> , 2019, 79, 4211-4226.	0.4	48
120	PD-L1 Expression, Tumor Mutational Burden, and Cancer Gene Mutations Are Stronger Predictors of Benefit from Immune Checkpoint Blockade than HLA Class I Genotype in Non-small Cell Lung Cancer. <i>Journal of Thoracic Oncology</i> , 2019, 14, 1021-1031.	0.5	79
121	Genomic landscape of allelic imbalance in premalignant atypical adenomatous hyperplasias of the lung. <i>EBioMedicine</i> , 2019, 42, 296-303.	2.7	15
122	Gene expression profiling of lichenoid dermatitis immune-related adverse event from immune checkpoint inhibitors reveals increased CD14 ⁺ and CD16 ⁺ monocytes driving an innate immune response. <i>Journal of Cutaneous Pathology</i> , 2019, 46, 627-636.	0.7	27
123	ZEB1 suppression sensitizes KRAS mutant cancers to MEK inhibition by an IL17RD-dependent mechanism. <i>Science Translational Medicine</i> , 2019, 11, .	5.8	42
124	An Integrated Next-Generation Sequencing System for Analyzing DNA Mutations, Gene Fusions, and RNA Expression in Lung Cancer. <i>Translational Oncology</i> , 2019, 12, 836-845.	1.7	19
125	Rational Error Elimination Approach to Evaluating Molecular Barcoded Next-Generation Sequencing Data Identifies Low-Frequency Mutations in Hematologic Malignancies. <i>Journal of Molecular Diagnostics</i> , 2019, 21, 471-482.	1.2	1
126	Best Practices Recommendations for Diagnostic Immunohistochemistry in Lung Cancer. <i>Journal of Thoracic Oncology</i> , 2019, 14, 377-407.	0.5	212

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127	B7-H3 Expression in Merkel Cell Carcinoma Associated Endothelial Cells Correlates with Locally Aggressive Primary Tumor Features and Increased Vascular Density. <i>Clinical Cancer Research</i> , 2019, 25, 3455-3467.	3.2	24
128	Targeting DNA Damage Response Promotes Antitumor Immunity through STING-Mediated T-cell Activation in Small Cell Lung Cancer. <i>Cancer Discovery</i> , 2019, 9, 646-661.	7.7	555
129	High OX-40 expression in the tumor immune infiltrate is a favorable prognostic factor of overall survival in non-small cell lung cancer. , 2019, 7, 351.		39
130	ConvPath: A software tool for lung adenocarcinoma digital pathological image analysis aided by a convolutional neural network. <i>EBioMedicine</i> , 2019, 50, 103-110.	2.7	66
131	Immunotherapy response evaluation with magnetic resonance elastography (MRE) in advanced HCC. , 2019, 7, 329.		33
132	TMPRSS4: A Novel Tumor Prognostic Indicator for the Stratification of Stage IA Tumors and a Liquid Biopsy Biomarker for NSCLC Patients. <i>Journal of Clinical Medicine</i> , 2019, 8, 2134.	1.0	17
133	Inhibitor of Differentiation-1 Sustains Mutant <i>KRAS</i> -Driven Progression, Maintenance, and Metastasis of Lung Adenocarcinoma via Regulation of a FOSL1 Network. <i>Cancer Research</i> , 2019, 79, 625-638.	0.4	19
134	LCE: an open web portal to explore gene expression and clinical associations in lung cancer. <i>Oncogene</i> , 2019, 38, 2551-2564.	2.6	78
135	5 protein-based signature for resectable lung squamous cell carcinoma improves the prognostic performance of the TNM staging. <i>Thorax</i> , 2019, 74, 371-379.	2.7	9
136	Genetic characterization of medullary thyroid cancer in childhood survivors of the Chernobyl accident. <i>Surgery</i> , 2019, 165, 58-63.	1.0	5
137	Centrifuged supernatants from FNA provide a liquid biopsy option for clinical next-generation sequencing of thyroid nodules. <i>Cancer Cytopathology</i> , 2019, 127, 146-160.	1.4	37
138	Combining Immune Checkpoint Blockade and Tumor-Specific Vaccine for Patients With Incurable Human Papillomavirus Related Cancer. <i>JAMA Oncology</i> , 2019, 5, 67.	3.4	344
139	Validation of the 12-gene Predictive Signature for Adjuvant Chemotherapy Response in Lung Cancer. <i>Clinical Cancer Research</i> , 2019, 25, 150-157.	3.2	13
140	Primary Mediastinal Yolk Sac Tumors: An Immunohistochemical Analysis of 14 Cases. <i>Applied Immunohistochemistry and Molecular Morphology</i> , 2019, 27, 125-133.	0.6	12
141	RCRnorm: An integrated system of random-coefficient hierarchical regression models for normalizing NanoString nCounter data. <i>Annals of Applied Statistics</i> , 2019, 13, 1617-1647.	0.5	6
142	Salvaging the supernatant: next generation cytopathology for solid tumor mutation profiling. <i>Modern Pathology</i> , 2018, 31, 1036-1045.	2.9	60
143	Osteoblast-Secreted Factors Mediate Dormancy of Metastatic Prostate Cancer in the Bone via Activation of the TGF β 2/RIII-p38MAPK-pS249/T252RB Pathway. <i>Cancer Research</i> , 2018, 78, 2911-2924.	0.4	117
144	Major pathologic response and RAD51 predict survival in lung cancer patients receiving neoadjuvant chemotherapy. <i>Cancer Medicine</i> , 2018, 7, 2405-2414.	1.3	22

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145	Chemistry-First Approach for Nomination of Personalized Treatment in Lung Cancer. <i>Cell</i> , 2018, 173, 864-878.e29.	13.5	102
146	Increased Tumor Glycolysis Characterizes Immune Resistance to Adoptive T Cell Therapy. <i>Cell Metabolism</i> , 2018, 27, 977-987.e4.	7.2	398
147	Spatial interaction of tumor cells and regulatory T cells correlates with survival in non-small cell lung cancer. <i>Lung Cancer</i> , 2018, 117, 73-79.	0.9	135
148	Multiregion gene expression profiling reveals heterogeneity in molecular subtypes and immunotherapy response signatures in lung cancer. <i>Modern Pathology</i> , 2018, 31, 947-955.	2.9	56
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