List of Publications by Year in descending order

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		16411	16127
232	17,641	64	124
papers	citations	h-index	g-index
234	234	234	21497
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	<i>STK11/LKB1</i> Mutations and PD-1 Inhibitor Resistance in <i>KRAS</i> -Mutant Lung Adenocarcinoma. Cancer Discovery, 2018, 8, 822-835.	7.7	1,108
2	PD-L1 as a biomarker of response to immune-checkpoint inhibitors. Nature Reviews Clinical Oncology, 2021, 18, 345-362.	12.5	646
3	Updated Molecular Testing Guideline for the Selection of Lung Cancer Patients for Treatment With Targeted Tyrosine Kinase Inhibitors: Guideline From the College of American Pathologists, the International Association for the Study of Lung Cancer, and the Association for Molecular Pathology, Archives of Pathology and Laboratory Medicine, 2018, 142, 321-346.	1.2	586
4	<i>MET</i> Exon 14 Mutations in Non–Small-Cell Lung Cancer Are Associated With Advanced Age and Stage-Dependent <i>MET</i> Genomic Amplification and c-Met Overexpression. Journal of Clinical Oncology, 2016, 34, 721-730.	0.8	549
5	Association of Polymerase e–Mutated and Microsatellite-Instable Endometrial Cancers With Neoantigen Load, Number of Tumor-Infiltrating Lymphocytes, and Expression of PD-1 and PD-L1. JAMA Oncology, 2015, 1, 1319.	3.4	523
6	Assessment of Resistance Mechanisms and Clinical Implications in Patients With <i>EGFR</i> T790M–Positive Lung Cancer and Acquired Resistance to Osimertinib. JAMA Oncology, 2018, 4, 1527.	3.4	522
7	Liquid Biopsy for Advanced Non-Small Cell LungÂCancer (NSCLC): A Statement Paper from theÂIASLC. Journal of Thoracic Oncology, 2018, 13, 1248-1268.	0.5	515
8	Acquired Resistance to KRAS ^{G12C} Inhibition in Cancer. New England Journal of Medicine, 2021, 384, 2382-2393.	13.9	482
9	Genomic correlates of response to immune checkpoint blockade in microsatellite-stable solid tumors. Nature Genetics, 2018, 50, 1271-1281.	9.4	438
10	Updated Molecular Testing Guideline for theÂSelection of Lung Cancer Patients for Treatment With Targeted Tyrosine Kinase Inhibitors. Journal of Thoracic Oncology, 2018, 13, 323-358.	0.5	408
11	PD-1 Inhibitor–Related Pneumonitis in Advanced Cancer Patients: Radiographic Patterns and Clinical Course. Clinical Cancer Research, 2016, 22, 6051-6060.	3.2	393
12	Multi-institutional Oncogenic Driver Mutation Analysis in Lung Adenocarcinoma: The Lung Cancer Mutation Consortium Experience. Journal of Thoracic Oncology, 2015, 10, 768-777.	0.5	357
13	Validation of OncoPanel: A Targeted Next-Generation Sequencing Assay for the Detection of Somatic Variants in Cancer. Archives of Pathology and Laboratory Medicine, 2017, 141, 751-758.	1.2	350
14	Institutional implementation of clinical tumor profiling on an unselected cancer population. JCI Insight, 2016, 1, e87062.	2.3	340
15	Clinicopathologic Features and Long-term Outcomes of NUT Midline Carcinoma. Clinical Cancer Research, 2012, 18, 5773-5779.	3.2	323
16	Suppression of STING Associated with LKB1 Loss in KRAS-Driven Lung Cancer. Cancer Discovery, 2019, 9, 34-45.	7.7	310
17	Clinical, Pathologic, and Biologic Features Associated with <i>BRAF</i> Mutations in Non–Small Cell Lung Cancer. Clinical Cancer Research, 2013, 19, 4532-4540.	3.2	307
18	The Reprogramming of Tumor Stroma by HSF1 Is a Potent Enabler of Malignancy. Cell, 2014, 158, 564-578.	13.5	298

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19	Implications of the tumor immune microenvironment for staging and therapeutics. Modern Pathology, 2018, 31, 214-234.	2.9	278
20	Liquid Biopsy for Advanced NSCLC: A Consensus Statement From the International Association for the Study of Lung Cancer. Journal of Thoracic Oncology, 2021, 16, 1647-1662.	0.5	274
21	In situ detection of SARS-CoV-2 in lungs and airways of patients with COVID-19. Modern Pathology, 2020, 33, 2104-2114.	2.9	257
22	Updated Molecular Testing Guideline for the Selection of Lung Cancer Patients for Treatment With Targeted Tyrosine Kinase Inhibitors. Journal of Molecular Diagnostics, 2018, 20, 129-159.	1.2	241
23	A Grading System for Invasive Pulmonary Adenocarcinoma: A Proposal From the International Association for the Study of Lung Cancer Pathology Committee. Journal of Thoracic Oncology, 2020, 15, 1599-1610.	0.5	234
24	PD-L1 Testing for Lung Cancer in 2019: Perspective From the IASLC Pathology Committee. Journal of Thoracic Oncology, 2020, 15, 499-519.	0.5	203
25	Temporal and spatial heterogeneity of host response to SARS-CoV-2 pulmonary infection. Nature Communications, 2020, 11, 6319.	5.8	203
26	Succinate dehydrogenase-deficient renal cell carcinoma: detailed characterization of 11 tumors defining a unique subtype of renal cell carcinoma. Modern Pathology, 2015, 28, 80-94.	2.9	190
27	The Promises and Challenges of Tumor Mutation Burden as an Immunotherapy Biomarker: A Perspective from the International Association for the Study of Lung Cancer Pathology Committee. Journal of Thoracic Oncology, 2020, 15, 1409-1424.	0.5	182
28	ROS1 Immunohistochemistry for Detection of ROS1-Rearranged Lung Adenocarcinomas. American Journal of Surgical Pathology, 2013, 37, 1441-1449.	2.1	168
29	BreaKmer: detection of structural variation in targeted massively parallel sequencing data using kmers. Nucleic Acids Research, 2015, 43, e19-e19.	6.5	161
30	The Impact of Smoking and TP53 Mutations in Lung Adenocarcinoma Patients with Targetable Mutations—The Lung Cancer Mutation Consortium (LCMC2). Clinical Cancer Research, 2018, 24, 1038-1047.	3.2	154
31	The impact of tumor profiling approaches and genomic data strategies for cancer precision medicine. Genome Medicine, 2016, 8, 79.	3.6	151
32	Diminished Efficacy of Programmed Death-(Ligand)1 Inhibition in STK11- and KEAP1-Mutant Lung Adenocarcinoma Is Affected by KRAS Mutation Status. Journal of Thoracic Oncology, 2022, 17, 399-410.	0.5	151
33	Combined Use of ALK Immunohistochemistry and FISH for Optimal Detection of ALK-Rearranged Lung Adenocarcinomas. Journal of Thoracic Oncology, 2013, 8, 322-328.	0.5	145
34	Sox2 Protein Expression is an Independent Poor Prognostic Indicator in Stage I Lung Adenocarcinoma. American Journal of Surgical Pathology, 2010, 34, 1193-1198.	2.1	140
35	Clinical Sequencing Exploratory Research Consortium: Accelerating Evidence-Based Practice of Genomic Medicine. American Journal of Human Genetics, 2016, 98, 1051-1066.	2.6	137
36	Acquired <i>MET</i> D1228V Mutation and Resistance to MET Inhibition in Lung Cancer. Cancer Discovery, 2016, 6, 1334-1341.	7.7	133

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37	Association of High Tumor Mutation Burden in Non–Small Cell Lung Cancers With Increased Immune Infiltration and Improved Clinical Outcomes of PD-L1 Blockade Across PD-L1 Expression Levels. JAMA Oncology, 2022, 8, 1160.	3.4	117
38	Lung Adenocarcinoma with <i>EGFR</i> Amplification Has Distinct Clinicopathologic and Molecular Features in Never-Smokers. Cancer Research, 2009, 69, 8341-8348.	0.4	114
39	Targeted genomic profiling reveals recurrent KRAS mutations and gain of chromosome 1q in mesonephric carcinomas of the female genital tract. Modern Pathology, 2015, 28, 1504-1514.	2.9	111
40	<i>EGFR</i> Mutation Is a Better Predictor of Response to Tyrosine Kinase Inhibitors in Non–Small Cell Lung Carcinoma Than FISH, CISH, and Immunohistochemistry. American Journal of Clinical Pathology, 2010, 133, 922-934.	0.4	110
41	Multiparametric profiling of non–small-cell lung cancers reveals distinct immunophenotypes. JCI Insight, 2016, 1, e89014.	2.3	110
42	Oncologists' and cancer patients' views on whole-exome sequencing and incidental findings: results from the CanSeq study. Genetics in Medicine, 2016, 18, 1011-1019.	1.1	108
43	Association of clonal hematopoiesis with chronic obstructive pulmonary disease. Blood, 2022, 139, 357-368.	0.6	106
44	Primary Pulmonary NUT Midline Carcinoma: Clinical, Radiographic, and Pathologic Characterizations. Journal of Thoracic Oncology, 2015, 10, 951-959.	0.5	100
45	Strategies for the successful implementation of plasma-based NSCLC genotyping in clinical practice. Nature Reviews Clinical Oncology, 2021, 18, 56-62.	12.5	99
46	Immunohistochemical Loss of LKB1 Is a Biomarker for More Aggressive Biology in <i>KRAS</i> -Mutant Lung Adenocarcinoma. Clinical Cancer Research, 2015, 21, 2851-2860.	3.2	96
47	Biomarker Testing in Lung Carcinoma Cytology Specimens: A Perspective From Members of the Pulmonary Pathology Society. Archives of Pathology and Laboratory Medicine, 2016, 140, 1267-1272.	1.2	95
48	Impact of DNA Damage Response and Repair (DDR) Gene Mutations on Efficacy of PD-(L)1 Immune Checkpoint Inhibition in Non–Small Cell Lung Cancer. Clinical Cancer Research, 2020, 26, 4135-4142.	3.2	95
49	Sox2 Expression in Pulmonary Non-small Cell and Neuroendocrine Carcinomas. Applied Immunohistochemistry and Molecular Morphology, 2010, 18, 55-61.	0.6	91
50	Overcoming MET-Dependent Resistance to Selective RET Inhibition in Patients with RET Fusion–Positive Lung Cancer by Combining Selpercatinib with Crizotinib. Clinical Cancer Research, 2021, 27, 34-42.	3.2	87
51	Intrinsic Immunogenicity of Small Cell Lung Carcinoma Revealed by Its Cellular Plasticity. Cancer Discovery, 2021, 11, 1952-1969.	7.7	87
52	Expression of ROS1 predicts ROS1 gene rearrangement in inflammatory myofibroblastic tumors. Modern Pathology, 2015, 28, 732-739.	2.9	85
53	Impact of MET inhibitors on survival among patients with non-small cell lung cancer harboring MET exon 14 mutations: a retrospective analysis. Lung Cancer, 2019, 133, 96-102.	0.9	85
54	Targeted genomic analysis of Müllerian adenosarcoma. Journal of Pathology, 2015, 235, 37-49.	2.1	84

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55	Clear cell ovarian cancers with microsatellite instability: A unique subset of ovarian cancers with increased tumor-infiltrating lymphocytes and PD-1/PD-L1 expression. Oncolmmunology, 2017, 6, e1277308.	2.1	84
56	Chemotherapy for locally advanced and metastatic pulmonary carcinoid tumors. Lung Cancer, 2014, 86, 241-246.	0.9	82
57	Lkb1 inactivation drives lung cancer lineage switching governed by Polycomb Repressive Complex 2. Nature Communications, 2017, 8, 14922.	5.8	80
58	Histopathology of Interstitial Lung Abnormalities in the Context of Lung Nodule Resections. American Journal of Respiratory and Critical Care Medicine, 2018, 197, 955-958.	2.5	78
59	Immunohistochemical Analysis of Langerin in Langerhans Cell Histiocytosis and Pulmonary Inflammatory and Infectious Diseases. American Journal of Surgical Pathology, 2007, 31, 947-952.	2.1	77
60	Neurotrophin receptor TrkB promotes lung adenocarcinoma metastasis. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 10299-10304.	3.3	77
61	ldentification of Existing Drugs That Effectively Target <i>NTRK1</i> and <i>ROS1</i> Rearrangements in Lung Cancer. Clinical Cancer Research, 2017, 23, 204-213.	3.2	73
62	Targeted genomic sequencing of follicular dendritic cell sarcoma reveals recurrent alterations in NF-IºB regulatory genes. Modern Pathology, 2016, 29, 67-74.	2.9	71
63	Clinical and Molecular Characteristics of <i>NF1</i> -Mutant Lung Cancer. Clinical Cancer Research, 2016, 22, 3148-3156.	3.2	71
64	Prospective Enterprise-Level Molecular Genotyping of a Cohort of Cancer Patients. Journal of Molecular Diagnostics, 2014, 16, 660-672.	1.2	70
65	Cytologicâ€histologic correlation of programmed deathâ€ligand 1 immunohistochemistry in lung carcinomas. Cancer Cytopathology, 2018, 126, 253-263.	1.4	70
66	Identification of diverse activating mutations of the RAS-MAPK pathway in histiocytic sarcoma. Modern Pathology, 2019, 32, 830-843.	2.9	68
67	A Pilot Study Linking Endothelial Injury in Lungs and Kidneys in Chronic Obstructive Pulmonary Disease. American Journal of Respiratory and Critical Care Medicine, 2017, 195, 1464-1476.	2.5	67
68	Sensitivity of next-generation sequencing assays detecting oncogenic fusions in plasma cell-free DNA. Lung Cancer, 2019, 134, 96-99.	0.9	67
69	Biomarkers in Lung Adenocarcinoma: A Decade of Progress. Archives of Pathology and Laboratory Medicine, 2015, 139, 469-480.	1.2	66
70	KRAS and NKX2-1 Mutations in Invasive Mucinous Adenocarcinoma of the Lung. Journal of Thoracic Oncology, 2016, 11, 496-503.	0.5	65
71	Radiationâ€associated neoplasia: clinical, pathological and genomic correlates. Histopathology, 2017, 70, 70-80.	1.6	65
72	Amplification of Wild-type <i>KRAS</i> Imparts Resistance to Crizotinib in <i>MET</i> Exon 14 Mutant Non–Small Cell Lung Cancer. Clinical Cancer Research, 2018, 24, 5963-5976.	3.2	63

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73	Cytotoxic T Cells in PD-L1–Positive Malignant Pleural Mesotheliomas Are Counterbalanced by Distinct Immunosuppressive Factors. Cancer Immunology Research, 2016, 4, 1038-1048.	1.6	62
74	Genomic Analysis of Plasma Cell-Free DNA in Patients With Cancer. JAMA Oncology, 2017, 3, 740.	3.4	61
75	Long-term Benefit of PD-L1 Blockade in Lung Cancer Associated with <i>JAK3</i> Activation. Cancer Immunology Research, 2015, 3, 855-863.	1.6	60
76	Use of targeted next generation sequencing to characterize tumor mutational burden and efficacy of immune checkpoint inhibition in small cell lung cancer. , 2019, 7, 87.		60
77	Sarcoid-Like Granulomatosis of the Lung Related to Immune-Checkpoint Inhibitors: Distinct Clinical and Imaging Features of a Unique Immune-Related Adverse Event. Cancer Immunology Research, 2018, 6, 630-635.	1.6	59
78	Consistency and reproducibility of nextâ€generation sequencing and other multigene mutational assays: A worldwide ring trial study on quantitative cytological molecular reference specimens. Cancer Cytopathology, 2017, 125, 615-626.	1.4	58
79	Harmonization of Tumor Mutational Burden Quantification and Association With Response to Immune Checkpoint Blockade in Non–Small-Cell Lung Cancer. JCO Precision Oncology, 2019, 3, 1-12.	1.5	58
80	Thoracic Complications of Precision Cancer Therapies: A Practical Guide for Radiologists in the New Era of Cancer Care. Radiographics, 2017, 37, 1371-1387.	1.4	56
81	Clinical Implications of Variant ALK FISH Rearrangement Patterns. Journal of Thoracic Oncology, 2015, 10, 1648-1652.	0.5	52
82	Quantitative assessment of PD-L1 as an analyte in immunohistochemistry diagnostic assays using a standardized cell line tissue microarray. Laboratory Investigation, 2020, 100, 4-15.	1.7	52
83	Validation of chromogenic in situ hybridization for detection of EGFR copy number amplification in nonsmall cell lung carcinoma. Modern Pathology, 2007, 20, 1028-1035.	2.9	51
84	Treatment-Related Toxicities in a Phase II Trial of Dasatinib in Patients with Squamous Cell Carcinoma of the Lung. Journal of Thoracic Oncology, 2013, 8, 1434-1437.	0.5	51
85	Nextâ€generation sequencing of cytologic preparations: An analysis of quality metrics. Cancer Cytopathology, 2017, 125, 786-794.	1.4	51
86	OncoTree: A Cancer Classification System for Precision Oncology. JCO Clinical Cancer Informatics, 2021, 5, 221-230.	1.0	51
87	Concurrent TP53 Mutations Facilitate Resistance Evolution in EGFR-Mutant Lung Adenocarcinoma. Journal of Thoracic Oncology, 2022, 17, 779-792.	0.5	50
88	Targeted next-generation sequencing reveals high frequency of mutations in epigenetic regulators across treatment-naÃ ⁻ ve patient melanomas. Clinical Epigenetics, 2015, 7, 59.	1.8	49
89	Validation of a targeted next-generation sequencing approach to detect mismatch repair deficiency in colorectal adenocarcinoma. Modern Pathology, 2018, 31, 1882-1890.	2.9	49
90	Characteristics of mismatch repair deficiency in sarcomas. Modern Pathology, 2019, 32, 977-987.	2.9	49

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91	SMARCA4 and Other SWItch/Sucrose NonFermentable Family Genomic Alterations in NSCLC: Clinicopathologic Characteristics and Outcomes to Immune Checkpoint Inhibition. Journal of Thoracic Oncology, 2021, 16, 1176-1187.	0.5	49
92	Assigning clinical meaning to somatic and germ-line whole-exome sequencing data in a prospective cancer precision medicine study. Genetics in Medicine, 2017, 19, 787-795.	1.1	46
93	Liquid biopsy of fine-needle aspiration supernatant for lung cancer genotyping. Lung Cancer, 2018, 122, 72-75.	0.9	46
94	Metaplastic thymoma: a distinctive thymic neoplasm characterized by YAP1-MAML2 gene fusions. Modern Pathology, 2020, 33, 560-565.	2.9	46
95	Imaging of Precision Therapy for Lung Cancer: Current State of the Art. Radiology, 2019, 293, 15-29.	3.6	45
96	Molecular diagnostics of lung cancer in the clinic. Translational Lung Cancer Research, 2017, 6, 560-569.	1.3	43
97	Generation of Genetically Engineered Mouse Lung Organoid Models for Squamous Cell Lung Cancers Allows for the Study of Combinatorial Immunotherapy. Clinical Cancer Research, 2020, 26, 3431-3442.	3.2	41
98	Pulmonary Large Cell Carcinoma Lacking Squamous Differentiation Is Clinicopathologically Indistinguishable From Solid-Subtype Adenocarcinoma. Archives of Pathology and Laboratory Medicine, 2014, 138, 626-635.	1.2	39
99	Detection of activating <i>MAP2K1</i> mutations in atypical hairy cell leukemia and hairy cell leukemia variant. Leukemia and Lymphoma, 2017, 58, 233-236.	0.6	39
100	Consistency and reproducibility of nextâ€generation sequencing in cytopathology: A second worldwide ring trial study on improved cytological molecular reference specimens. Cancer Cytopathology, 2019, 127, 285-296.	1.4	39
101	Abnormal p53 and p16 staining patterns distinguish uterine leiomyosarcoma from inflammatory myofibroblastic tumour. Histopathology, 2017, 70, 1138-1146.	1.6	38
102	Detection of ERBB2 Amplification by Next-Generation Sequencing Predicts HER2 Expression in Colorectal Carcinoma. American Journal of Clinical Pathology, 2019, 152, 97-108.	0.4	36
103	Loss of SMAD4 protein expression in gastrointestinal and extraâ€gastrointestinal carcinomas. Histopathology, 2019, 75, 546-551.	1.6	35
104	Morphologic correlates of molecular alterations in extrauterine Müllerian carcinomas. Modern Pathology, 2016, 29, 893-903.	2.9	33
105	GNAS mutations in primary mucinous and non-mucinous lung adenocarcinomas. Modern Pathology, 2017, 30, 1720-1727.	2.9	33
106	Biomarkers of response to checkpoint inhibitors beyond PD-L1 in lung cancer. Modern Pathology, 2022, 35, 66-74.	2.9	33
107	Kinase Domain Activation of FGFR2 Yields High-Grade Lung Adenocarcinoma Sensitive to a Pan-FGFR Inhibitor in a Mouse Model of NSCLC. Cancer Research, 2014, 74, 4676-4684.	0.4	31
108	Identification of Oncogenic and Drug-Sensitizing Mutations in the Extracellular Domain of FGFR2. Cancer Research, 2015, 75, 3139-3146.	0.4	30

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109	Cabozantinib in Patients with Advanced Merkel Cell Carcinoma. Oncologist, 2018, 23, 814-821.	1.9	30
110	An Acquired NRAS Q61K Mutation in BRAF V600E-Mutant Lung Adenocarcinoma Resistant toÂDabrafenib Plus Trametinib. Journal of Thoracic Oncology, 2018, 13, e131-e133.	0.5	30
111	Molecular Characterization of Neuroendocrine Carcinomas of the Endometrium. American Journal of Surgical Pathology, 2020, 44, 1541-1548.	2.1	26
112	Impact of MET inhibitors on survival among patients (pts) with <i>MET</i> exon 14 mutant (<i>MET</i> del14) non-small cell lung cancer (NSCLC) Journal of Clinical Oncology, 2017, 35, 8511-8511.	0.8	26
113	ARID1A mutations and expression loss in non-small cell lung carcinomas: clinicopathologic and molecular analysis. Modern Pathology, 2020, 33, 2256-2268.	2.9	25
114	Phase II Clinical Trial of Everolimus in a Pan-Cancer Cohort of Patients with mTOR Pathway Alterations. Clinical Cancer Research, 2021, 27, 3845-3853.	3.2	25
115	The Molecular Pathology of Lung Cancer. Surgical Pathology Clinics, 2016, 9, 353-378.	0.7	23
116	Pulmonary Clinicopathological Correlation after Allogeneic Hematopoietic Stem Cell Transplantation: An Autopsy Series. Biology of Blood and Marrow Transplantation, 2017, 23, 1767-1772.	2.0	23
117	Next-generation sequencing informs diagnosis and identifies unexpected therapeutic targets in lung squamous cell carcinomas. Lung Cancer, 2020, 140, 35-41.	0.9	22
118	Variation in targetable genomic alterations in non-small cell lung cancer by genetic ancestry, sex, smoking history, and histology. Genome Medicine, 2022, 14, 39.	3.6	22
119	Successful Management of a Patient with Malignant Thyroid Teratoma. Thyroid, 2017, 27, 125-128.	2.4	21
120	Lung Adenocarcinoma Syndecan-2 Potentiates Cell Invasiveness. American Journal of Respiratory Cell and Molecular Biology, 2019, 60, 659-666.	1.4	20
121	Biallelic PTCH1 Inactivation Is a Dominant Genomic Change in Sporadic Keratocystic Odontogenic Tumors. American Journal of Surgical Pathology, 2020, 44, 553-560.	2.1	20
122	Malignant tumours of the uterus and ovaries with Mullerian and germ cell or trophoblastic components have a somatic origin and are characterised by genomic instability ^Â . Histopathology, 2020, 77, 788-797.	1.6	20
123	Correlation of methylthioadenosine phosphorylase (MTAP) protein expression with <i>MTAP</i> and <i>CDKN2A</i> copy number in malignant pleural mesothelioma. Histopathology, 2021, 78, 1032-1042.	1.6	20
124	Effect of expanded genomic testing in lung adenocarcinoma (LUCA) on survival benefit: The Lung Cancer Mutation Consortium II (LCMC II) experience Journal of Clinical Oncology, 2016, 34, 11510-11510.	0.8	20
125	Incidental nonneoplastic parenchymal findings in patients undergoing lung resection for mass lesions. Human Pathology, 2019, 86, 93-101.	1.1	19
126	Targeted Cancer Next-Generation Sequencing as a Primary Screening Tool for Microsatellite Instability and Lynch Syndrome in Upper Gastrointestinal Tract Cancers. Cancer Epidemiology Biomarkers and Prevention, 2019, 28, 1246-1251.	1.1	18

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127	Phase IB Study of Osimertinib in Combination with Navitoclax in <i>EGFR</i> -mutant NSCLC Following Resistance to Initial <i>EGFR</i> Therapy (ETCTN 9903). Clinical Cancer Research, 2021, 27, 1604-1611.	3.2	18
128	Myc protein expression correlates with <i><scp>MYC</scp></i> amplification in small ell lung carcinoma. Histopathology, 2015, 67, 81-89.	1.6	17
129	Medical Oncologists' Experiences in Using Genomic Testing for Lung and Colorectal Cancer Care. Journal of Oncology Practice, 2017, 13, e185-e196.	2.5	17
130	Pulmonary Pathology Society Perspective on the 2018 American Thoracic Society, European Respiratory Society, Japanese Respiratory Society, and Latin American Thoracic Society Idiopathic Pulmonary Fibrosis Clinical Practice Guidelines. Annals of the American Thoracic Society, 2020, 17, 550-554.	1.5	17
131	Epitope mapping of spontaneous autoantibodies to anaplastic lymphoma kinase (ALK) in non-small cell lung cancer. Oncotarget, 2017, 8, 92265-92274.	0.8	17
132	Clinical and molecular validation of BAP1, MTAP, P53, and Merlin immunohistochemistry in diagnosis of pleural mesothelioma. Modern Pathology, 2022, 35, 1383-1397.	2.9	17
133	Ewing sarcoma mimicking atypical carcinoid tumor: detection of unexpected genomic alterations demonstrates the use of next generation sequencing as a diagnostic tool. Cancer Genetics, 2014, 207, 335-339.	0.2	16
134	Pseudoendocrine Sarcoma. American Journal of Surgical Pathology, 2022, 46, 33-43.	2.1	16
135	The fuzzy world of precision medicine: deliberations of a precision medicine tumor board. Personalized Medicine, 2017, 14, 37-50.	0.8	15
136	Diagnostic and Predictive Immunohistochemistry for Non–Small Cell Lung Carcinomas. Advances in Anatomic Pathology, 2018, 25, 374-386.	2.4	15
137	Biobanking and cryopreservation of human lung explants for omic analysis. European Respiratory Journal, 2020, 55, 1801635.	3.1	15
138	Genomic and pathological heterogeneity in clinically diagnosed small cell lung cancer in never/light smokers identifies therapeutically targetable alterations. Molecular Oncology, 2021, 15, 27-42.	2.1	15
139	Comparative molecular analysis of testicular Leydig cell tumors demonstrates distinct subsets of neoplasms with aggressive histopathologic features. Modern Pathology, 2021, 34, 1935-1946.	2.9	15
140	Thymic Measurements in Pathologically Proven Normal Thymus and Thymic Hyperplasia. Academic Radiology, 2014, 21, 733-742.	1.3	14
141	Interactive or static reports to guide clinical interpretation of cancer genomics. Journal of the American Medical Informatics Association: JAMIA, 2018, 25, 458-464.	2.2	14
142	Histopathologic Assessment of Suspected Idiopathic Pulmonary Fibrosis: Where We Are and Where We Need to Go. Archives of Pathology and Laboratory Medicine, 2020, 144, 1477-1489.	1.2	14
143	Osimertinib (Osi) plus necitumumab (Neci) in EGFR-mutant NSCLC: An ETCTN California cancer consortium phase I study Journal of Clinical Oncology, 2019, 37, 9057-9057.	0.8	14
144	Phase I Trial of a Tablet Formulation of Pilaralisib, a Pan lass I PI3K Inhibitor, in Patients with Advanced Solid Tumors. Oncologist, 2018, 23, 401.	1.9	13

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145	Identification of a RAS-activating <i>TMEM87A–RASGRF1</i> Fusion in an Exceptional Responder to Sunitinib with Non–Small Cell Lung Cancer. Clinical Cancer Research, 2020, 26, 4072-4079.	3.2	13
146	IER5, a DNA damage response gene, is required for Notch-mediated induction of squamous cell differentiation. ELife, 2020, 9, .	2.8	13
147	Mast cells in lung damage of COVIDâ€19 autopsies: A descriptive study. Allergy: European Journal of Allergy and Clinical Immunology, 2022, 77, 2237-2239.	2.7	13
148	Clinical Pan-Cancer Assessment of Mismatch Repair Deficiency Using Tumor-Only, Targeted Next-Generation Sequencing. JCO Precision Oncology, 2020, 4, 1084-1097.	1.5	11
149	Mural nodules in mucinous ovarian tumors represent a morphologic spectrum of clonal neoplasms: a morphologic, immunohistochemical, and molecular analysis of 13 cases. Modern Pathology, 2021, 34, 613-626.	2.9	11
150	A Novel Protective Role for Matrix Metalloproteinase-8 in the Pulmonary Vasculature. American Journal of Respiratory and Critical Care Medicine, 2021, 204, 1433-1451.	2.5	11
151	Quantitative computed tomography assessment of bronchiolitis obliterans syndrome after lung transplantation. Clinical Transplantation, 2017, 31, e12943.	0.8	10
152	Engaging Patients in Precision Oncology: Development and Usability of a Web-Based Patient-Facing Genomic Sequencing Report. JCO Precision Oncology, 2020, 4, 307-318.	1.5	10
153	Plasma cfDNA Genotyping in Hospitalized Patients With Suspected Metastatic NSCLC. JCO Precision Oncology, 2021, 5, 726-732.	1.5	10
154	Oncogenic switch and single-agent MET inhibitor sensitivity in a subset of <i>EGFR</i> -mutant lung cancer. Science Translational Medicine, 2021, 13, eabb3738.	5.8	10
155	Association Between Immune-Related Adverse Events and Clinical Outcomes to Programmed Cell Death Protein 1/Programmed Death-Ligand 1 Blockade in SCLC. JTO Clinical and Research Reports, 2020, 1, 100074.	0.6	10
156	Emerging Evidence for MicroRNAs as Regulators of Cancer Stem Cells. Cancers, 2011, 3, 3957-3971.	1.7	9
157	Protein correlates of molecular alterations in lung adenocarcinoma: Immunohistochemistry as a surrogate for molecular analysis. Seminars in Diagnostic Pathology, 2015, 32, 325-333.	1.0	9
158	Meningioma transcription factors link cell lineage with systemic metabolic cues. Neuro-Oncology, 2018, 20, 1331-1343.	0.6	9
159	Intestinal metaplasia of the urinary tract harbors potentially oncogenic genetic variants. Modern Pathology, 2021, 34, 457-468.	2.9	9
160	Molecular assessment of testicular adult granulosa cell tumor demonstrates significant differences when compared to ovarian counterparts. Modern Pathology, 2022, 35, 697-704.	2.9	9
161	Micropapillary adenocarcinoma of lung: Morphological criteria and diagnostic reproducibility among pulmonary pathologists. Annals of Diagnostic Pathology, 2019, 41, 43-50.	0.6	8
162	Re-evaluating tumors of purported specialized prostatic stromal origin reveals molecular heterogeneity, including non-recurring gene fusions characteristic of uterine and soft tissue sarcoma subtypes. Modern Pathology, 2021, 34, 1763-1779.	2.9	8

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163	Abstract LB002: Mechanisms of acquired resistance to KRAS G12C inhibition in cancer. , 2021, , .		8
164	Traditional Diagnostics versus Disruptive Technology: The Role of the Pathologist in the Era of Liquid Biopsy. Cancer Research, 2020, 80, 3197-3199.	0.4	7
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