

Ritika Kundra

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

11,290
citations

279701

23
h-index

477173

29
g-index

30
all docs

30
docs citations

30
times ranked

20447
citing authors

#	ARTICLE	IF	CITATIONS
1	Mutational landscape of metastatic cancer revealed from prospective clinical sequencing of 10,000 patients. <i>Nature Medicine</i> , 2017, 23, 703-713.	15.2	2,473
2	Oncogenic Signaling Pathways in The Cancer Genome Atlas. <i>Cell</i> , 2018, 173, 321-337.e10.	13.5	2,111
3	OncoKB: A Precision Oncology Knowledge Base. <i>JCO Precision Oncology</i> , 2017, 2017, 1-16.	1.5	1,266
4	Comprehensive and Integrated Genomic Characterization of Adult Soft Tissue Sarcomas. <i>Cell</i> , 2017, 171, 950-965.e28.	13.5	738
5	The Genomic Landscape of Endocrine-Resistant Advanced Breast Cancers. <i>Cancer Cell</i> , 2018, 34, 427-438.e6.	7.7	633
6	The long tail of oncogenic drivers in prostate cancer. <i>Nature Genetics</i> , 2018, 50, 645-651.	9.4	601
7	Clinical Sequencing Defines the Genomic Landscape of Metastatic Colorectal Cancer. <i>Cancer Cell</i> , 2018, 33, 125-136.e3.	7.7	589
8	Prospective Comprehensive Molecular Characterization of Lung Adenocarcinomas for Efficient Patient Matching to Approved and Emerging Therapies. <i>Cancer Discovery</i> , 2017, 7, 596-609.	7.7	490
9	Prospective Genotyping of Hepatocellular Carcinoma: Clinical Implications of Next-Generation Sequencing for Matching Patients to Targeted and Immune Therapies. <i>Clinical Cancer Research</i> , 2019, 25, 2116-2126.	3.2	390
10	Prospective Genomic Profiling of Prostate Cancer Across Disease States Reveals Germline and Somatic Alterations That May Affect Clinical Decision Making. <i>JCO Precision Oncology</i> , 2017, 2017, 1-16.	1.5	286
11	Accelerating Discovery of Functional Mutant Alleles in Cancer. <i>Cancer Discovery</i> , 2018, 8, 174-183.	7.7	275
12	Genetic Predictors of Response to Systemic Therapy in Esophagogastric Cancer. <i>Cancer Discovery</i> , 2018, 8, 49-58.	7.7	275
13	The expanding landscape of α -oncohistone™ mutations in human cancers. <i>Nature</i> , 2019, 567, 473-478.	13.7	271
14	Genomic characterization of metastatic patterns from prospective clinical sequencing of 25,000 patients. <i>Cell</i> , 2022, 185, 563-575.e11.	13.5	223
15	TRK Fusions Are Enriched in Cancers with Uncommon Histologies and the Absence of Canonical Driver Mutations. <i>Clinical Cancer Research</i> , 2020, 26, 1624-1632.	3.2	103
16	Small-Cell Carcinomas of the Bladder and Lung Are Characterized by a Convergent but Distinct Pathogenesis. <i>Clinical Cancer Research</i> , 2018, 24, 1965-1973.	3.2	85
17	Therapeutic Implications of Germline Testing in Patients With Advanced Cancers. <i>Journal of Clinical Oncology</i> , 2021, 39, 2698-2709.	0.8	83
18	Clinical and Molecular Predictors of Response to Immune Checkpoint Inhibitors in Patients with Advanced Esophagogastric Cancer. <i>Clinical Cancer Research</i> , 2019, 25, 6160-6169.	3.2	73

#	ARTICLE	IF	CITATIONS
19	Genetic Determinants of Outcome in Intrahepatic Cholangiocarcinoma. <i>Hepatology</i> , 2021, 74, 1429-1444.	3.6	73
20	OncoTree: A Cancer Classification System for Precision Oncology. <i>JCO Clinical Cancer Informatics</i> , 2021, 5, 221-230.	1.0	51
21	Characteristics and Outcome of <i>AKT1</i> E17K-Mutant Breast Cancer Defined through AACR Project GENIE, a Clinicogenomic Registry. <i>Cancer Discovery</i> , 2020, 10, 526-535.	7.7	36
22	Regional differences in gallbladder cancer pathogenesis: Insights from a multi-institutional comparison of tumor mutations. <i>Cancer</i> , 2019, 125, 575-585.	2.0	34
23	Intrahepatic Cholangiocarcinoma with Lymph Node Metastasis: Treatment-Related Outcomes and the Role of Tumor Genomics in Patient Selection. <i>Clinical Cancer Research</i> , 2021, 27, 4101-4108.	3.2	24
24	FOLFICIS Treatment and Genomic Correlates of Response in Advanced Anal Squamous Cell Cancer. <i>Clinical Colorectal Cancer</i> , 2019, 18, e39-e52.	1.0	21
25	Correlation Between Surrogate End Points and Overall Survival in a Multi-institutional Clinicogenomic Cohort of Patients With Non-Small Cell Lung or Colorectal Cancer. <i>JAMA Network Open</i> , 2021, 4, e2117547.	2.8	20
26	Annotation of Somatic Genomic Variants in Hematologic Diseases Using OncoKB, a Precision Oncology Knowledgebase. <i>Blood</i> , 2019, 134, 2148-2148.	0.6	3
27	Linked Entity Attribute Pair (LEAP): A Harmonization Framework for Data Pooling. <i>JCO Clinical Cancer Informatics</i> , 2020, 4, 691-699.	1.0	2