Sameek Roychowdhury

List of Publications by Year in descending order

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65 papers

7,534 citations

35 h-index 63 g-index

65 all docs

65 docs citations

65 times ranked 13574 citing authors

#	Article	IF	CITATIONS
1	Mainstreaming germline genetic testing for patients with pancreatic cancer increases uptake. Familial Cancer, 2023, 22, 91-97.	1.9	6
2	Validation and Characterization of FGFR2 Rearrangements in Cholangiocarcinoma with Comprehensive Genomic Profiling. Journal of Molecular Diagnostics, 2022, 24, 351-364.	2.8	5
3	Characterization of Clonal Evolution in Microsatellite Unstable Metastatic Cancers through Multiregional Tumor Sequencing. Molecular Cancer Research, 2021, 19, 465-474.	3.4	2
4	Fibroblast growth factor receptors in cancer: genetic alterations, diagnostics, therapeutic targets and mechanisms of resistance. British Journal of Cancer, 2021, 124, 880-892.	6.4	150
5	Genomic and Transcriptomic Characterization of Relapsed SCLC Through Rapid Research Autopsy. JTO Clinical and Research Reports, 2021, 2, 100164.	1.1	6
6	Infigratinib (BCJ398) in previously treated patients with advanced or metastatic cholangiocarcinoma with FGFR2 fusions or rearrangements: mature results from a multicentre, open-label, single-arm, phase 2 study. The Lancet Gastroenterology and Hepatology, 2021, 6, 803-815.	8.1	205
7	Research Autopsy Demonstrates Polyclonal Acquired Resistance in a Patient With Metastatic GI Stromal Tumor. JCO Precision Oncology, 2020, 4, 131-138.	3.0	3
8	Infigratinib in patients with advanced cholangiocarcinoma with <i>FGFR2</i> gene fusions/translocations: the PROOF 301 trial. Future Oncology, 2020, 16, 2375-2384.	2.4	62
9	Co-occurrence of multiple endocrine neoplasia type 4 and spinal neurofibromatosis: a case report. Familial Cancer, 2020, 19, 189-192.	1.9	8
10	Phase I Trial of Trametinib with Neoadjuvant Chemoradiation in Patients with Locally Advanced Rectal Cancer. Clinical Cancer Research, 2020, 26, 3117-3125.	7.0	13
11	Efficacy of FGFR Inhibitors and Combination Therapies for Acquired Resistance in FGFR2-Fusion Cholangiocarcinoma. Molecular Cancer Therapeutics, 2020, 19, 847-857.	4.1	91
12	Detection of Microsatellite Instability Biomarkers via Next-Generation Sequencing. Methods in Molecular Biology, 2020, 2055, 119-132.	0.9	42
13	Pan-cancer analysis of FGFR1-3 genomic alterations to reveal a complex molecular landscape Journal of Clinical Oncology, 2020, 38, 3620-3620.	1.6	10
14	Implementing precision cancer medicine in the genomic era. Seminars in Cancer Biology, 2019, 55, 16-27.	9.6	24
15	Tumor heterogeneity and acquired drug resistance in FGFR2-fusion-positive cholangiocarcinoma through rapid research autopsy. Journal of Physical Education and Sports Management, 2019, 5, a004002.	1.2	60
16	Homologous recombination and DNA repair mutations in patients treated with carboplatin and nab-paclitaxel for metastatic non-small cell lung cancer. Lung Cancer, 2019, 134, 167-173.	2.0	9
17	Characterization of a KLK2-FGFR2 fusion gene in two cases of metastatic prostate cancer. Prostate Cancer and Prostatic Diseases, 2019, 22, 624-632.	3.9	5
18	Microsatellite Instability Occurs in a Subset of Follicular Thyroid Cancers. Thyroid, 2019, 29, 523-529.	4.5	31

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19	Significant and Durable Clinical Response to Sorafenib and Radiation Therapy for a Patient With Stage IV Hepatocellular Carcinoma and LRRK2 Mutation. JCO Precision Oncology, 2019, 3, 1-9.	3.0	1
20	Rapid Research Autopsy: Piecing the Puzzle of Tumor Heterogeneity. Trends in Cancer, 2019, 5, 1-5.	7.4	13
21	Precision Cancer Medicine and Clinical Trial Design. , 2019, , 49-63.		2
22	Genomic characterization of metastatic ultra-hypermutated interdigitating dendritic cell sarcoma through rapid research autopsy. Oncotarget, 2019, 10, 277-288.	1.8	6
23	Metaplastic breast cancer in a patient with neurofibromatosis type 1 and somatic loss of heterozygosity. Journal of Physical Education and Sports Management, 2018, 4, a002352.	1.2	11
24	Trametinib for the treatment of IGHV4-34, MAP2K1-mutant variant hairy cell leukemia. Leukemia and Lymphoma, 2018, 59, 1008-1011.	1.3	29
25	TargetingBRAFMutations in High-Grade Neuroendocrine Carcinoma of the Colon. Journal of the National Comprehensive Cancer Network: JNCCN, 2018, 16, 1035-1040.	4.9	24
26	Akt Activation Mediates Acquired Resistance to Fibroblast Growth Factor Receptor Inhibitor BGJ398. Molecular Cancer Therapeutics, 2017, 16, 614-624.	4.1	72
27	Validation of a Targeted RNA Sequencing Assay for Kinase Fusion Detection in Solid Tumors. Journal of Molecular Diagnostics, 2017, 19, 682-696.	2.8	56
28	Beyond Seed and Soil: Understanding and Targeting Metastatic Prostate Cancer; Report From the 2016 Coffey–Holden Prostate Cancer Academy Meeting. Prostate, 2017, 77, 123-144.	2.3	6
29	Performance evaluation for rapid detection of pan-cancer microsatellite instability with MANTIS. Oncotarget, 2017, 8, 7452-7463.	1.8	232
30	Landscape of Microsatellite Instability Across 39 Cancer Types. JCO Precision Oncology, 2017, 2017, 1-15.	3.0	796
31	Analytic validation and real-time clinical application of an amplicon-based targeted gene panel for advanced cancer. Oncotarget, 2017, 8, 75822-75833.	1.8	6
32	Inflammation-Induced Oxidative Stress Mediates Gene Fusion Formation in Prostate Cancer. Cell Reports, 2016, 17, 2620-2631.	6.4	68
33	The Bayesian basket design for genomic variant-driven phase II trials. Seminars in Oncology, 2016, 43, 13-18.	2.2	81
34	Translating cancer genomes and transcriptomes for precision oncology. Ca-A Cancer Journal for Clinicians, 2016, 66, 75-88.	329.8	133
35	Somatic cancer variant curation and harmonization through consensus minimum variant level data. Genome Medicine, 2016, 8, 117.	8.2	61
36	Targeted RNA Sequencing Assay to Characterize Gene Expression and Genomic Alterations. Journal of Visualized Experiments, 2016 , , .	0.3	10

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37	Impact of genomic sequencing on precision medicine for clinical oncology. Expert Review of Precision Medicine and Drug Development, 2016, 1, 255-265.	0.7	O
38	Germline Findings in Tumor-Only Sequencing: Points to Consider for Clinicians and Laboratories: Table 1 Journal of the National Cancer Institute, 2016, 108, djv351.	6.3	86
39	Evaluation of Hybridization Capture Versus Ampliconâ€Based Methods for Wholeâ€Exome Sequencing. Human Mutation, 2015, 36, 903-914.	2.5	206
40	Clinical Tumor Sequencing: Opportunities and Challenges for Precision Cancer Medicine. American Society of Clinical Oncology Educational Book / ASCO American Society of Clinical Oncology Meeting, 2015, , e175-e182.	3.8	47
41	Cancer Driver Log (CanDL). Journal of Molecular Diagnostics, 2015, 17, 554-559.	2.8	56
42	Integrative Clinical Sequencing in the Management of Refractory or Relapsed Cancer in Youth. JAMA - Journal of the American Medical Association, 2015, 314, 913.	7.4	333
43	Comparison of Custom Capture for Targeted Next-Generation DNA Sequencing. Journal of Molecular Diagnostics, 2015, 17, 64-75.	2.8	65
44	Clinical Tumor Sequencing: An Incidental Casualty of the American College of Medical Genetics and Genomics Recommendations for Reporting of Incidental Findings. Journal of Clinical Oncology, 2014, 32, 2203-2205.	1.6	36
45	Return of Genomic Results to Research Participants: The Floor, the Ceiling, and the Choices In Between. American Journal of Human Genetics, 2014, 94, 818-826.	6.2	342
46	Translating Genomics for Precision Cancer Medicine. Annual Review of Genomics and Human Genetics, 2014, 15, 395-415.	6.2	63
47	Identification of Targetable FGFR Gene Fusions in Diverse Cancers. Cancer Discovery, 2013, 3, 636-647.	9.4	614
48	Identification of recurrent NAB2-STAT6 gene fusions in solitary fibrous tumor by integrative sequencing. Nature Genetics, 2013, 45, 180-185.	21.4	662
49	Activating ESR1 mutations in hormone-resistant metastatic breast cancer. Nature Genetics, 2013, 45, 1446-1451.	21.4	925
50	Advancing Precision Medicine for Prostate Cancer Through Genomics. Journal of Clinical Oncology, 2013, 31, 1866-1873.	1.6	84
51	Implementing personalized cancer genomics in clinical trials. Nature Reviews Drug Discovery, 2013, 12, 358-369.	46.4	267
52	Cancer genomics meets clinical trials: the challenge ahead. Personalized Medicine, 2012, 9, 459-461.	1.5	3
53	Managing resistance in chronic myeloid leukemia. Blood Reviews, 2011, 25, 279-290.	5.7	56
54	Personalized Oncology Through Integrative High-Throughput Sequencing: A Pilot Study. Science Translational Medicine, 2011, 3, 111ra121.	12.4	531

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55	IFN- \hat{l}^3 gene polymorphisms associate with development of EBV+ lymphoproliferative disease in hu PBL-SCID mice. Blood, 2005, 105, 1558-1565.	1.4	35
56	Donor-derived IL-15 is critical for acute allogeneic graft-versus-host disease. Blood, 2005, 105, 894-901.	1.4	75
57	Anti–human CTLA-4 monoclonal antibody promotes T-cell expansion and immunity in a hu-PBL-SCID model: a new method for preclinical screening of costimulatory monoclonal antibodies. Blood, 2005, 105, 1114-1120.	1.4	27
58	IL-15 but not IL-2 rapidly induces lethal xenogeneic graft-versus-host disease. Blood, 2005, 106, 2433-2435.	1.4	45
59	A Human CD34(+) Subset Resides in Lymph Nodes and Differentiates into CD56brightNatural Killer Cells. Immunity, 2005, 22, 295-304.	14.3	331
60	Combination Immunotherapy of B-Cell Non-Hodgkin's Lymphoma with Rituximab and Interleukin-2. Clinical Cancer Research, 2004, 10, 6101-6110.	7.0	74
61	Failed Adoptive Immunotherapy with Tumor-Specific T Cells. Cancer Research, 2004, 64, 8062-8067.	0.9	66
62	Selective Efficacy of Depsipeptide in a Xenograft Model of Epstein-Barr Virus-Positive Lymphoproliferative Disorder. Journal of the National Cancer Institute, 2004, 96, 1447-1457.	6.3	29
63	A Novel Human CD34(+) Subset That Constitutively Expresses the High Affinity Interleukin-2 Receptor Traffics to Lymph Nodes and Differentiates into CD56Bright Natural Killer Cells Blood, 2004, 104, 314-314.	1.4	13
64	Experimental treatment of Epstein-Barr virus-associated primary central nervous system lymphoma. Cancer Research, 2003, 63, 965-71.	0.9	70
65	Successful treatment of posttransplantation lymphoproliferative disorder (PTLD) following renal allografting is associated with sustained CD8+ T-cell restoration. Blood, 2002, 100, 2341-2348.	1.4	54