## Timothy A Chan

# List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

163 64 29,990 173 h-index g-index papers citations 6.98 16.4 183 39,312 L-index avg, IF ext. citations ext. papers

#	Paper	IF	Citations
163	Aging-related cell type-specific pathophysiologic immune responses that exacerbate disease severity in aged COVID-19 patients <i>Aging Cell</i> , <b>2022</b> , e13544	9.9	2
162	Pre-treatment serum albumin and mutational burden as biomarkers of response to immune checkpoint blockade <i>Npj Precision Oncology</i> , <b>2022</b> , 6, 23	9.8	1
161	Phenotypic and molecular states of IDH1 mutation-induced CD24-positive glioma stem-like cells <i>Neoplasia</i> , <b>2022</b> , 28, 100790	6.4	O
160	Outcomes Among Patients With or Without Obesity and With Cancer Following Treatment With Immune Checkpoint Blockade <i>JAMA Network Open</i> , <b>2022</b> , 5, e220448	10.4	1
159	H3K9 methylation drives resistance to androgen receptor-antagonist therapy in prostate cancer <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2022</b> , 119, e211432411	9 <sup>11.5</sup>	2
158	The Genetic Evolution of Treatment-Resistant Cutaneous, Acral, and Uveal Melanomas. <i>Clinical Cancer Research</i> , <b>2021</b> , 27, 1516-1525	12.9	2
157	The Similarity of Class II HLA Genotypes Defines Patterns of Autoreactivity in Idiopathic Bone Marrow Failure Disorders. <i>Blood</i> , <b>2021</b> ,	2.2	2
156	Immune cytolytic activity is associated with reduced intra-tumoral genetic heterogeneity and with better clinical outcomes in triple negative breast cancer. <i>American Journal of Cancer Research</i> , <b>2021</b> , 11, 3628-3644	4.4	2
155	Molecular and phenotypic profiling of colorectal cancer patients in West Africa reveals biological insights. <i>Nature Communications</i> , <b>2021</b> , 12, 6821	17.4	2
154	Genomic and Transcriptomic Correlates of Thyroid Carcinoma Evolution after BRAF Inhibitor Therapy. <i>Molecular Cancer Research</i> , <b>2021</b> ,	6.6	2
153	Improved prediction of immune checkpoint blockade efficacy across multiple cancer types. <i>Nature Biotechnology</i> , <b>2021</b> ,	44.5	10
152	Mutations in and differentially affect the tumor microenvironment and response to checkpoint blockade immunotherapy. <i>Nature Cancer</i> , <b>2021</b> , 1, 1188-1203	15.4	43
151	Immunotherapy biomarkers: the long and winding road. <i>Nature Reviews Clinical Oncology</i> , <b>2021</b> , 18, 323	3-3 <u>3.4</u>	2
150	Qa-1 Modulates Resistance to Anti-PD-1 Immune Checkpoint Blockade in Tumors with Defects in Antigen Processing. <i>Molecular Cancer Research</i> , <b>2021</b> , 19, 1076-1084	6.6	3
149	Genomic Epidemiology of SARS-CoV-2 Infection During the Initial Pandemic Wave and Association With Disease Severity. <i>JAMA Network Open</i> , <b>2021</b> , 4, e217746	10.4	13
148	High Response Rate and Durability Driven by HLA Genetic Diversity in Patients with Kidney Cancer Treated with Lenvatinib and Pembrolizumab. <i>Molecular Cancer Research</i> , <b>2021</b> , 19, 1510-1521	6.6	2
147	Current Prospects for Treatment of Solid Tumors via Photodynamic, Photothermal, or Ionizing Radiation Therapies Combined with Immune Checkpoint Inhibition (A Review). <i>Pharmaceuticals</i> , <b>2021</b> , 14,	5.2	11

### (2020-2021)

146	Response Rates to Anti-PD-1 Immunotherapy in Microsatellite-Stable Solid Tumors With 10 or More Mutations per Megabase. <i>JAMA Oncology</i> , <b>2021</b> , 7, 739-743	13.4	40
145	Single-cell sequencing links multiregional immune landscapes and tissue-resident Tīcells in ccRCC to tumor topology and therapy efficacy. <i>Cancer Cell</i> , <b>2021</b> , 39, 662-677.e6	24.3	28
144	Resource-efficient pooled sequencing expands translational impact in solid tumors. <i>Kidney Cancer Journal: Official Journal of the Kidney Cancer Association</i> , <b>2021</b> , 19, 18-23	0.1	О
143	Prevalence and Landscape of Actionable Genomic Alterations in Renal Cell Carcinoma. <i>Clinical Cancer Research</i> , <b>2021</b> , 27, 5595-5606	12.9	3
142	Multimodal single-cell omics analysis identifies epithelium-immune cell interactions and immune vulnerability associated with sex differences in COVID-19. <i>Signal Transduction and Targeted Therapy</i> , <b>2021</b> , 6, 292	21	3
141	Pathogenic ATM Mutations in Cancer and a Genetic Basis for Radiotherapeutic Efficacy. <i>Journal of the National Cancer Institute</i> , <b>2021</b> , 113, 266-273	9.7	15
140	Putative Drivers of Aggressiveness in TCEB1-mutant Renal Cell Carcinoma: An Emerging Entity with Variable Clinical Course. <i>European Urology Focus</i> , <b>2021</b> , 7, 381-389	5.1	7
139	The association between tumor mutational burden and prognosis is dependent on treatment context. <i>Nature Genetics</i> , <b>2021</b> , 53, 11-15	36.3	38
138	Precision Radiotherapy: Reduction in Radiation for Oropharyngeal Cancer in the 30 ROC Trial. <i>Journal of the National Cancer Institute</i> , <b>2021</b> , 113, 742-751	9.7	39
137	Precision regenerative medicine. Stem Cell Research and Therapy, <b>2021</b> , 12, 39	8.3	2
137	Precision regenerative medicine. Stem Cell Research and Therapy, 2021, 12, 39  Pretreatment neutrophil-to-lymphocyte ratio and mutational burden as biomarkers of tumor response to immune checkpoint inhibitors. Nature Communications, 2021, 12, 729	17.4	44
	Pretreatment neutrophil-to-lymphocyte ratio and mutational burden as biomarkers of tumor		
136	Pretreatment neutrophil-to-lymphocyte ratio and mutational burden as biomarkers of tumor response to immune checkpoint inhibitors. <i>Nature Communications</i> , <b>2021</b> , 12, 729  Inappropriate Use of the Same Cutoff by Different Sequencing Panels for Tumor Mutation Burden	17.4	44
136	Pretreatment neutrophil-to-lymphocyte ratio and mutational burden as biomarkers of tumor response to immune checkpoint inhibitors. <i>Nature Communications</i> , <b>2021</b> , 12, 729  Inappropriate Use of the Same Cutoff by Different Sequencing Panels for Tumor Mutation Burden as Immunotherapy Biomarker-Reply. <i>JAMA Oncology</i> , <b>2021</b> , 7, 1245-1246  Ectopic activation of the miR-200c-EpCAM axis enhances antitumor T cell responses in models of	17.4 13.4	1
136 135	Pretreatment neutrophil-to-lymphocyte ratio and mutational burden as biomarkers of tumor response to immune checkpoint inhibitors. <i>Nature Communications</i> , <b>2021</b> , 12, 729  Inappropriate Use of the Same Cutoff by Different Sequencing Panels for Tumor Mutation Burden as Immunotherapy Biomarker-Reply. <i>JAMA Oncology</i> , <b>2021</b> , 7, 1245-1246  Ectopic activation of the miR-200c-EpCAM axis enhances antitumor T cell responses in models of adoptive cell therapy. <i>Science Translational Medicine</i> , <b>2021</b> , 13, eabg4328  My personal mutanome: a computational genomic medicine platform for searching network	17.4 13.4 17.5	1
136 135 134	Pretreatment neutrophil-to-lymphocyte ratio and mutational burden as biomarkers of tumor response to immune checkpoint inhibitors. <i>Nature Communications</i> , <b>2021</b> , 12, 729  Inappropriate Use of the Same Cutoff by Different Sequencing Panels for Tumor Mutation Burden as Immunotherapy Biomarker-Reply. <i>JAMA Oncology</i> , <b>2021</b> , 7, 1245-1246  Ectopic activation of the miR-200c-EpCAM axis enhances antitumor T cell responses in models of adoptive cell therapy. <i>Science Translational Medicine</i> , <b>2021</b> , 13, eabg4328  My personal mutanome: a computational genomic medicine platform for searching network perturbing alleles linking genotype to phenotype. <i>Genome Biology</i> , <b>2021</b> , 22, 53  A Network Medicine Approach to Investigation and Population-based Validation of Disease	17.4 13.4 17.5	1 7
136 135 134 133	Pretreatment neutrophil-to-lymphocyte ratio and mutational burden as biomarkers of tumor response to immune checkpoint inhibitors. <i>Nature Communications</i> , <b>2021</b> , 12, 729  Inappropriate Use of the Same Cutoff by Different Sequencing Panels for Tumor Mutation Burden as Immunotherapy Biomarker-Reply. <i>JAMA Oncology</i> , <b>2021</b> , 7, 1245-1246  Ectopic activation of the miR-200c-EpCAM axis enhances antitumor T cell responses in models of adoptive cell therapy. <i>Science Translational Medicine</i> , <b>2021</b> , 13, eabg4328  My personal mutanome: a computational genomic medicine platform for searching network perturbing alleles linking genotype to phenotype. <i>Genome Biology</i> , <b>2021</b> , 22, 53  A Network Medicine Approach to Investigation and Population-based Validation of Disease Manifestations and Drug Repurposing for COVID-19. <i>ChemRxiv</i> , <b>2020</b> ,  Diverse Neoantigens and the Development of Cancer Therapies. <i>Seminars in Radiation Oncology</i> ,	17.4 13.4 17.5 18.3	1 7 5

128	DNA damage repair pathway alterations in metastatic clear cell renal cell carcinoma and implications on systemic therapy <b>2020</b> , 8,		20
127	Radiomic analysis identifies tumor subtypes associated with distinct molecular and microenvironmental factors in head and neck squamous cell carcinoma. <i>Oral Oncology</i> , <b>2020</b> , 110, 1048	7 <del>4</del> ·4	10
126	The Immune Microenvironment and Neoantigen Landscape of Aggressive Salivary Gland Carcinomas Differ by Subtype. <i>Clinical Cancer Research</i> , <b>2020</b> , 26, 2859-2870	12.9	31
125	Consensus guidelines for the definition, detection and interpretation of immunogenic cell death <b>2020</b> , 8,		233
124	Impact of HLA Evolutionary Divergence on Clinical Features of Patients with Aplastic Anemia and Paroxysmal Nocturnal Hemoglobinuria. <i>Blood</i> , <b>2020</b> , 136, 2-3	2.2	
123	Commensal bacteria stimulate antitumor responses via T cell cross-reactivity. <i>JCI Insight</i> , <b>2020</b> , 5,	9.9	30
122	Epigenetic driver mutations in ARID1A shape cancer immune phenotype and immunotherapy. <i>Journal of Clinical Investigation</i> , <b>2020</b> , 130, 2712-2726	15.9	45
121	A network medicine approach to investigation and population-based validation of disease manifestations and drug repurposing for COVID-19. <i>PLoS Biology</i> , <b>2020</b> , 18, e3000970	9.7	75
120	Transcriptomic signatures related to the obesity paradox in patients with clear cell renal cell carcinoma: a cohort study. <i>Lancet Oncology, The</i> , <b>2020</b> , 21, 283-293	21.7	55
119	TGF-Buppresses type 2 immunity to cancer. <i>Nature</i> , <b>2020</b> , 587, 115-120	50.4	63
118	Genomics-based immuno-oncology: bridging the gap between immunology and tumor biology. <i>Human Molecular Genetics</i> , <b>2020</b> , 29, R214-R225	5.6	1
117	Sarcomatoid renal cell carcinoma: biology, natural history and management. <i>Nature Reviews Urology</i> , <b>2020</b> , 17, 659-678	5.5	23
116	Genetic and environmental determinants of human TCR repertoire diversity. <i>Immunity and Ageing</i> , <b>2020</b> , 17, 26	9.7	17
115	HIF-1Iand HIF-2Idifferently regulate tumour development and inflammation of clear cell renal cell carcinoma in mice. <i>Nature Communications</i> , <b>2020</b> , 11, 4111	17.4	50
114	A pan-cancer analysis of PBAF complex mutations and their association with immunotherapy response. <i>Nature Communications</i> , <b>2020</b> , 11, 4168	17.4	15
113	Tumor mutational burden as a predictive biomarker for checkpoint inhibitor immunotherapy. <i>Human Vaccines and Immunotherapeutics</i> , <b>2020</b> , 16, 112-115	4.4	24
112	Comprehensive Genomic Analysis of Translocation Renal Cell Carcinoma Reveals Copy-Number Variations as Drivers of Disease Progression. <i>Clinical Cancer Research</i> , <b>2020</b> , 26, 3629-3640	12.9	16
111	A network medicine approach to investigation and population-based validation of disease manifestations and drug repurposing for COVID-19 <b>2020</b> , 18, e3000970		

#### (2019-2020)

110	A network medicine approach to investigation and population-based validation of disease manifestations and drug repurposing for COVID-19 <b>2020</b> , 18, e3000970		
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108	A network medicine approach to investigation and population-based validation of disease manifestations and drug repurposing for COVID-19 <b>2020</b> , 18, e3000970		
107	A network medicine approach to investigation and population-based validation of disease manifestations and drug repurposing for COVID-19 <b>2020</b> , 18, e3000970		
106	A network medicine approach to investigation and population-based validation of disease manifestations and drug repurposing for COVID-19 <b>2020</b> , 18, e3000970		
105	A network medicine approach to investigation and population-based validation of disease manifestations and drug repurposing for COVID-19 <b>2020</b> , 18, e3000970		
104	RIG-I activation is critical for responsiveness to checkpoint blockade. Science Immunology, 2019, 4,	28	47
103	Lung Cancer Evolution: Whatß Immunity Got to Do with It?. Cancer Cell, 2019, 35, 711-713	24.3	5
102	Immunogenic neoantigens derived from gene fusions stimulate T cell responses. <i>Nature Medicine</i> , <b>2019</b> , 25, 767-775	50.5	149
101	Genetic diversity of tumors with mismatch repair deficiency influences anti-PD-1 immunotherapy response. <i>Science</i> , <b>2019</b> , 364, 485-491	33.3	228
100	CD8 T cells regulate tumour ferroptosis during cancer immunotherapy. <i>Nature</i> , <b>2019</b> , 569, 270-274	50.4	632
99	High-dose radiation therapy is needed for intracranial control and long-term survival in patients with non-seminomatous germ cell tumor brain metastases. <i>Journal of Neuro-Oncology</i> , <b>2019</b> , 142, 523-5	5 <b>2</b> 18 <sup>8</sup>	3
98	The evolving landscape of biomarkers for checkpoint inhibitor immunotherapy. <i>Nature Reviews Cancer</i> , <b>2019</b> , 19, 133-150	31.3	996
97	G-quadruplex DNA drives genomic instability and represents a targetable molecular abnormality in ATRX-deficient malignant glioma. <i>Nature Communications</i> , <b>2019</b> , 10, 943	17.4	79
96	CD97 is a critical regulator of acute myeloid leukemia stem cell function. <i>Journal of Experimental Medicine</i> , <b>2019</b> , 216, 2362-2377	16.6	11
95	APOBEC mutagenesis is tightly linked to the immune landscape and immunotherapy biomarkers in head and neck squamous cell carcinoma. <i>Oral Oncology</i> , <b>2019</b> , 96, 140-147	4.4	21
94	The good, the bad, and the ugly: hyperprogression in cancer patients following immune checkpoint therapy. <i>Genome Medicine</i> , <b>2019</b> , 11, 43	14.4	13
93	Immunomodulatory and immunotherapeutic implications of tobacco smoking in squamous cell carcinomas and normal airway epithelium. <i>Oncotarget</i> , <b>2019</b> , 10, 3835-3839	3.3	5

92	Genomic Correlates of Disease Progression and Treatment Response in Prospectively Characterized Gliomas. <i>Clinical Cancer Research</i> , <b>2019</b> , 25, 5537-5547	12.9	48
91	Genetic hallmarks of recurrent/metastatic adenoid cystic carcinoma. <i>Journal of Clinical Investigation</i> , <b>2019</b> , 129, 4276-4289	15.9	64
90	RIG-I Activation Is Critical for Responsiveness to Checkpoint Blockade. <i>Blood</i> , <b>2019</b> , 134, 624-624	2.2	1
89	Evolutionary divergence of HLA class I genotype impacts efficacy of cancer immunotherapy. <i>Nature Medicine</i> , <b>2019</b> , 25, 1715-1720	50.5	89
88	Demethylation and epigenetic modification with 5-azacytidine reduces IDH1 mutant glioma growth in combination with temozolomide. <i>Neuro-Oncology</i> , <b>2019</b> , 21, 189-200	1	33
87	Transcriptomic Profiling of the Tumor Microenvironment Reveals Distinct Subgroups of Clear Cell Renal Cell Cancer: Data from a Randomized Phase III Trial. <i>Cancer Discovery</i> , <b>2019</b> , 9, 510-525	24.4	88
86	Tumor mutational load predicts survival after immunotherapy across multiple cancer types. <i>Nature Genetics</i> , <b>2019</b> , 51, 202-206	36.3	1435
85	Residual Tumor Volume, Cell Volume Fraction, and Tumor Cell Kill During Fractionated Chemoradiation Therapy of Human Glioblastoma using Quantitative Sodium MR Imaging. <i>Clinical Cancer Research</i> , <b>2019</b> , 25, 1226-1232	12.9	16
84	Tobacco Smoking-Associated Alterations in the Immune Microenvironment of Squamous Cell Carcinomas. <i>Journal of the National Cancer Institute</i> , <b>2018</b> , 110, 1386-1392	9.7	78
83	ImmunoMap: A Bioinformatics Tool for T-cell Repertoire Analysis. <i>Cancer Immunology Research</i> , <b>2018</b> , 6, 151-162	12.5	25
82	Merkel Cell Carcinoma Patients Presenting Without a Primary Lesion Have Elevated Markers of Immunity, Higher Tumor Mutation Burden, and Improved Survival. <i>Clinical Cancer Research</i> , <b>2018</b> , 24, 963-971	12.9	34
81	Genomics of NSCLC patients both affirm PD-L1 expression and predict their clinical responses to anti-PD-1 immunotherapy. <i>BMC Cancer</i> , <b>2018</b> , 18, 225	4.8	17
80	Atrx inactivation drives disease-defining phenotypes in glioma cells of origin through global epigenomic remodeling. <i>Nature Communications</i> , <b>2018</b> , 9, 1057	17.4	39
79	Phosphatidylinositol-3-Kinase Mutations Are Associated With Increased Local Failure in Brain Metastases Treated With Radiation. <i>International Journal of Radiation Oncology Biology Physics</i> , <b>2018</b> , 101, 833-844	4	1
78	Update on Tumor Neoantigens and Their Utility: Why It Is Good to Be Different. <i>Trends in Immunology</i> , <b>2018</b> , 39, 536-548	14.4	91
77	Integrated Genomic Analysis of Hithle Cell Cancer Reveals Oncogenic Drivers, Recurrent Mitochondrial Mutations, and Unique Chromosomal Landscapes. <i>Cancer Cell</i> , <b>2018</b> , 34, 256-270.e5	24.3	103
76	Mutant-IDH1-dependent chromatin state reprogramming, reversibility, and persistence. <i>Nature Genetics</i> , <b>2018</b> , 50, 62-72	36.3	86
75	Patient HLA class I genotype influences cancer response to checkpoint blockade immunotherapy. <i>Science</i> , <b>2018</b> , 359, 582-587	33.3	500

#### (2017-2018)

74	Multicenter Phase IB Trial of Carboxyamidotriazole Orotate and Temozolomide for Recurrent and Newly Diagnosed Glioblastoma and Other Anaplastic Gliomas. <i>Journal of Clinical Oncology</i> , <b>2018</b> , 36, 1702-1709	2.2	27
73	Genomically annotated risk model for advanced renal-cell carcinoma: a retrospective cohort study. Lancet Oncology, The, <b>2018</b> , 19, 1688-1698	21.7	70
72	ATRX, DAXX or MEN1 mutant pancreatic neuroendocrine tumors are a distinct alpha-cell signature subgroup. <i>Nature Communications</i> , <b>2018</b> , 9, 4158	17.4	80
71	The hallmarks of successful anticancer immunotherapy. Science Translational Medicine, 2018, 10,	17.5	<b>26</b> 0
70	Transcriptional Mechanisms of Resistance to Anti-PD-1 Therapy. Clinical Cancer Research, 2017, 23, 3168	s <u>13</u> 1680	51
69	Genomic Alterations in Fatal Forms of Non-Anaplastic Thyroid Cancer: Identification of and as Novel Thyroid Cancer Genes Associated with Tumor Virulence. <i>Clinical Cancer Research</i> , <b>2017</b> , 23, 5970-5	<del>198</del> 8	64
68	Dissecting microsatellite instability in colorectal cancer: one size does not fit all. <i>Genome Medicine</i> , <b>2017</b> , 9, 45	14.4	3
67	Clinical outcomes of patients with limited brain metastases treated with hypofractionated (5BGy) conformal radiotherapy. <i>Radiotherapy and Oncology</i> , <b>2017</b> , 123, 203-208	5.3	12
66	Stratification of Pancreatic Ductal Adenocarcinoma: Combinatorial Genetic, Stromal, and Immunologic Markers. <i>Clinical Cancer Research</i> , <b>2017</b> , 23, 4429-4440	12.9	108
65	Pan-Cancer Analysis Links PARK2 to BCL-XL-Dependent Control of Apoptosis. <i>Neoplasia</i> , <b>2017</b> , 19, 75-83	6.4	23
64	Malignant Astrocytic Tumor Progression Potentiated by JAK-mediated Recruitment of Myeloid Cells. <i>Clinical Cancer Research</i> , <b>2017</b> , 23, 3109-3119	12.9	13
63	Multi-dimensional genomic analysis of myoepithelial carcinoma identifies prevalent oncogenic gene fusions. <i>Nature Communications</i> , <b>2017</b> , 8, 1197	17.4	46
62	Tumor and Microenvironment Evolution during Immunotherapy with Nivolumab. <i>Cell</i> , <b>2017</b> , 171, 934-94	<b>3.6.1</b> 6	831
61	Melanoma brain metastases treated with stereotactic radiosurgery and concurrent pembrolizumab display marked regression; efficacy and safety of combined treatment <b>2017</b> , 5, 76		78
60	NF- <b>B</b> c-Rel Is Crucial for the Regulatory T Cell Immune Checkpoint in Cancer. <i>Cell</i> , <b>2017</b> , 170, 1096-1108.	<b>65163</b> 2	151
59	An Integrated Systems Biology Approach Identifies TRIM25 as a Key Determinant of Breast Cancer Metastasis. <i>Cell Reports</i> , <b>2017</b> , 20, 1623-1640	10.6	59
58	Identification of unique neoantigen qualities in long-term survivors of pancreatic cancer. <i>Nature</i> , <b>2017</b> , 551, 512-516	50.4	533
57	A neoantigen fitness model predicts tumour response to checkpoint blockade immunotherapy.  Nature, <b>2017</b> , 551, 517-520	50.4	336

56	Multicenter, Phase 1, Dose Escalation Study of Hypofractionated Stereotactic Radiation Therapy With Bevacizumab for Recurrent Glioblastoma and Anaplastic Astrocytoma. <i>International Journal of Radiation Oncology Biology Physics</i> , <b>2017</b> , 99, 797-804	4	27
55	The Molecular Landscape of Recurrent and Metastatic Head and Neck Cancers: Insights From a Precision Oncology Sequencing Platform. <i>JAMA Oncology</i> , <b>2017</b> , 3, 244-255	13.4	141
54	Genomic analysis of exceptional responders to radiotherapy reveals somatic mutations in ATM. <i>Oncotarget</i> , <b>2017</b> , 8, 10312-10323	3.3	23
53	PLK1 inhibition enhances temozolomide efficacy in IDH1 mutant gliomas. <i>Oncotarget</i> , <b>2017</b> , 8, 15827-1	5837	13
52	Tumor immune microenvironment characterization in clear cell renal cell carcinoma identifies prognostic and immunotherapeutically relevant messenger RNA signatures. <i>Genome Biology</i> , <b>2016</b> , 17, 231	18.3	391
51	FAT1 mutations cause a glomerulotubular nephropathy. <i>Nature Communications</i> , <b>2016</b> , 7, 10822	17.4	69
50	Genomic profile, smoking, and response to anti-PD-1 therapy in non-small cell lung carcinoma. <i>Molecular and Cellular Oncology</i> , <b>2016</b> , 3, e1048929	1.2	24
49	Regularized quantile regression under heterogeneous sparsity with application to quantitative genetic traits. <i>Computational Statistics and Data Analysis</i> , <b>2016</b> , 95, 222-239	1.6	15
48	Immunotherapy and Oncogenic Pathways: The PTEN Connection. Cancer Discovery, 2016, 6, 128-9	24.4	16
47	Clonal neoantigens elicit T cell immunoreactivity and sensitivity to immune checkpoint blockade. <i>Science</i> , <b>2016</b> , 351, 1463-9	33.3	1758
46	Integrated genomic characterization of IDH1-mutant glioma malignant progression. <i>Nature Genetics</i> , <b>2016</b> , 48, 59-66	36.3	185
45			
15	Cancer Neoantigens and Applications for Immunotherapy. Clinical Cancer Research, <b>2016</b> , 22, 807-12	12.9	146
44	Cancer Neoantigens and Applications for Immunotherapy. <i>Clinical Cancer Research</i> , <b>2016</b> , 22, 807-12  CD97 Is a Critical Regulator of Acute Myeloid Leukemia Stem Cell Function. <i>Blood</i> , <b>2016</b> , 128, 1077-107		146
44	CD97 Is a Critical Regulator of Acute Myeloid Leukemia Stem Cell Function. <i>Blood</i> , <b>2016</b> , 128, 1077-107  The head and neck cancer immune landscape and its immunotherapeutic implications. <i>JCI Insight</i> ,	72.2	2
44	CD97 Is a Critical Regulator of Acute Myeloid Leukemia Stem Cell Function. <i>Blood</i> , <b>2016</b> , 128, 1077-107  The head and neck cancer immune landscape and its immunotherapeutic implications. <i>JCI Insight</i> , <b>2016</b> , 1, e89829  Mutational landscape of MCPyV-positive and MCPyV-negative Merkel cell carcinomas with	9.9	356
44 43 42	CD97 Is a Critical Regulator of Acute Myeloid Leukemia Stem Cell Function. <i>Blood</i> , <b>2016</b> , 128, 1077-107. The head and neck cancer immune landscape and its immunotherapeutic implications. <i>JCI Insight</i> , <b>2016</b> , 1, e89829.  Mutational landscape of MCPyV-positive and MCPyV-negative Merkel cell carcinomas with implications for immunotherapy. <i>Oncotarget</i> , <b>2016</b> , 7, 3403-15.  Pan-cancer analysis of intratumor heterogeneity as a prognostic determinant of survival.	9.9 3.3	2 356 229

## (2014-2016)

38	Molecular and Clinical Effects of Notch Inhibition in Glioma Patients: A Phase 0/I Trial. <i>Clinical Cancer Research</i> , <b>2016</b> , 22, 4786-4796	12.9	69
37	Integrated Genomics for Pinpointing Survival Loci within Arm-Level Somatic Copy Number Alterations. <i>Cancer Cell</i> , <b>2016</b> , 29, 737-750	24.3	31
36	Personalized Oncology Meets Immunology: The Path toward Precision Immunotherapy. <i>Cancer Discovery</i> , <b>2016</b> , 6, 703-13	24.4	64
35	Recurrent SERPINB3 and SERPINB4 mutations in patients who respond to anti-CTLA4 immunotherapy. <i>Nature Genetics</i> , <b>2016</b> , 48, 1327-1329	36.3	84
34	Spatial Proximity to Fibroblasts Impacts Molecular Features and Therapeutic Sensitivity of Breast Cancer Cells Influencing Clinical Outcomes. <i>Cancer Research</i> , <b>2016</b> , 76, 6495-6506	10.1	68
33	Comprehensive, Integrative Genomic Analysis of Diffuse Lower-Grade Gliomas. <i>New England Journal of Medicine</i> , <b>2015</b> , 372, 2481-98	59.2	1828
32	Cancer immunology. Mutational landscape determines sensitivity to PD-1 blockade in non-small cell lung cancer. <i>Science</i> , <b>2015</b> , 348, 124-8	33.3	5003
31	Genetics and immunology: reinvigorated. <i>OncoImmunology</i> , <b>2015</b> , 4, e1029705	7.2	7
30	Genomic landscape of adenoid cystic carcinoma of the breast. <i>Journal of Pathology</i> , <b>2015</b> , 237, 179-89	9.4	101
29	Long-term risk of radionecrosis and imaging changes after stereotactic radiosurgery for brain metastases. <i>Journal of Neuro-Oncology</i> , <b>2015</b> , 125, 149-56	4.8	165
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