Alexandre Reuben

List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/9340716/alexandre-reuben-publications-by-year.pdf

Version: 2024-04-10

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

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

87	8,160 citations	35	90
papers		h-index	g-index
109	11,894	12.9 avg, IF	5.51
ext. papers	ext. citations		L-index

#	Paper	IF	Citations
87	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	12.8	
86	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	17.4	1
85	Distinct tumor-infiltrating lymphocyte landscapes are associated with clinical outcomes in localized non-small-cell lung cancer. <i>Annals of Oncology</i> , 2021 ,	10.3	6
84	Short-term treatment with multi-drug regimens combining BRAF/MEK-targeted therapy and immunotherapy results in durable responses in -mutated melanoma. <i>OncoImmunology</i> , 2021 , 10, 19928	870 ²	2
83	Identification of bacteria-derived HLA-bound peptides in melanoma. <i>Nature</i> , 2021 , 592, 138-143	50.4	52
82	Global analysis of shared Titell specificities in human non-small cell lung cancer enables HLA inference and antigen discovery. <i>Immunity</i> , 2021 , 54, 586-602.e8	32.3	16
81	Response to Hypomethylating Agents in Myelodysplastic Syndrome Is Associated With Emergence of Novel TCR Clonotypes. <i>Frontiers in Immunology</i> , 2021 , 12, 659625	8.4	O
80	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	8.9	18
79	Intertumoral Genetic Heterogeneity Generates Distinct Tumor Microenvironments in a Novel Murine Synchronous Melanoma Model. <i>Cancers</i> , 2021 , 13,	6.6	1
78	Immune evolution from preneoplasia to invasive lung adenocarcinomas and underlying molecular features. <i>Nature Communications</i> , 2021 , 12, 2722	17.4	16
77	Gut microbiota signatures are associated with toxicity to combined CTLA-4 and PD-1 blockade. <i>Nature Medicine</i> , 2021 , 27, 1432-1441	50.5	57
76	Augmented Lipocalin-2 Is Associated with Chronic Obstructive Pulmonary Disease and Counteracts Lung Adenocarcinoma Development. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2021 , 203, 90-101	10.2	8
75	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	8.9	14
74	Evolution of DNA methylome from precancerous lesions to invasive lung adenocarcinomas. <i>Nature Communications</i> , 2021 , 12, 687	17.4	9
73	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	50.5	105
72	Oncogene-specific differences in tumor mutational burden, PD-L1 expression, and outcomes from immunotherapy in non-small cell lung cancer 2021 , 9,		13
71	Multiomics Analysis Reveals Distinct Immunogenomic Features of Lung Cancer with Ground-Glass Opacity. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2021 , 204, 1180-1192	10.2	1

(2020-2021)

70	The histologic phenotype of lung cancers is associated with transcriptomic features rather than genomic characteristics. <i>Nature Communications</i> , 2021 , 12, 7081	17.4	1
69	Multiomics 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	18.3	13
68	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,		13
67	Neoantigen responses, immune correlates, and favorable outcomes after ipilimumab treatment of patients with prostate cancer. <i>Science Translational Medicine</i> , 2020 , 12,	17.5	54
66	Peripheral cytokines are not influenced by the type of surgical approach for non-small cell lung cancer by four weeks postoperatively. <i>Lung Cancer</i> , 2020 , 146, 303-309	5.9	2
65	Immuno-genomic landscape of osteosarcoma. <i>Nature Communications</i> , 2020 , 11, 1008	17.4	77
64	B cells and tertiary lymphoid structures promote immunotherapy response. <i>Nature</i> , 2020 , 577, 549-555	50.4	654
63	Discovery of a novel shared tumor antigen in human lung cancer <i>Journal of Clinical Oncology</i> , 2020 , 38, 3087-3087	2.2	
62	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	8.9	49
61	Comprehensive T cell repertoire characterization of non-small cell lung cancer. <i>Nature Communications</i> , 2020 , 11, 603	17.4	67
60	Immune and Circulating Tumor DNA Profiling After Radiation Treatment for Oligometastatic Non-Small Cell Lung Cancer: Translational Correlatives from a Mature Randomized Phase II Trial. <i>International Journal of Radiation Oncology Biology Physics</i> , 2020 , 106, 349-357	4	15
59	Multifactorial Deep Learning Reveals Pan-Cancer Genomic Tumor Clusters with Distinct Immunogenomic Landscape and Response to Immunotherapy. <i>Clinical Cancer Research</i> , 2020 , 26, 2908-3	2 92 8	12
58	Immune profiling of uveal melanoma identifies a potential signature associated with response to immunotherapy 2020 , 8,		14
57	Melanoma Evolves Complete Immunotherapy Resistance through the Acquisition of a Hypermetabolic Phenotype. <i>Cancer Immunology Research</i> , 2020 , 8, 1365-1380	12.5	13
56	Evolution of Genomic and T-cell Repertoire Heterogeneity of Malignant Pleural Mesothelioma Under Dasatinib Treatment. <i>Clinical Cancer Research</i> , 2020 , 26, 5477-5486	12.9	12
55	F-fluorodeoxyglucose positron emission tomography correlates with tumor immunometabolic phenotypes in resected lung cancer. <i>Cancer Immunology, Immunotherapy</i> , 2020 , 69, 1519-1534	7.4	11
54	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	4.3	9
53	Spatially resolved analyses link genomic and immune diversity and reveal unfavorable neutrophil activation in melanoma. <i>Nature Communications</i> , 2020 , 11, 1839	17.4	9

52	Correlative Analyses of the SARC028 Trial Reveal an Association Between Sarcoma-Associated Immune Infiltrate and Response to Pembrolizumab. <i>Clinical Cancer Research</i> , 2020 , 26, 1258-1266	12.9	55
51	Combination anti-CTLA-4 plus anti-PD-1 checkpoint blockade utilizes cellular mechanisms partially distinct from monotherapies. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 22699-22709	11.5	119
50	A Novel Mitochondrial Inhibitor Blocks MAPK Pathway and Overcomes MAPK Inhibitor Resistance in Melanoma. <i>Clinical Cancer Research</i> , 2019 , 25, 6429-6442	12.9	30
49	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	8.9	46
48	Poor Response to Neoadjuvant Chemotherapy Correlates with Mast Cell Infiltration in Inflammatory Breast Cancer. <i>Cancer Immunology Research</i> , 2019 , 7, 1025-1035	12.5	42
47	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	1.7	12
46	Multi-region exome sequencing reveals genomic evolution from preneoplasia to lung adenocarcinoma. <i>Nature Communications</i> , 2019 , 10, 2978	17.4	43
45	Suppressed immune microenvironment and repertoire in brain metastases from patients with resected non-small-cell lung cancer. <i>Annals of Oncology</i> , 2019 , 30, 1521-1530	10.3	47
44	Spatial and temporal heterogeneity of PD-L1 and its impact on benefit from immune checkpoint blockade in non-small cell lung cancer (NSCLC) <i>Journal of Clinical Oncology</i> , 2019 , 37, 9017-9017	2.2	7
43	Association of relative neutrophilia with a distinct immunoinhibitory milieu in non-small cell lung cancer <i>Journal of Clinical Oncology</i> , 2019 , 37, e14047-e14047	2.2	O
42	Characterization of Changes in the T-Cell Receptor Repertoire in Patients with Acute Myeloid Leukemia with Durable Remission Following Allogeneic Stem Cell Transplant. <i>Blood</i> , 2019 , 134, 5186-5	186 ²	
41	Molecular Profiling Reveals Unique Immune and Metabolic Features of Melanoma Brain Metastases. <i>Cancer Discovery</i> , 2019 , 9, 628-645	24.4	124
40	The Influence of the Gut Microbiome on Cancer, Immunity, and Cancer Immunotherapy. <i>Cancer Cell</i> , 2018 , 33, 570-580	24.3	527
39	A phase II study of combined therapy with a BRAF inhibitor (vemurafenib) and interleukin-2 (aldesleukin) in patients with metastatic melanoma. <i>OncoImmunology</i> , 2018 , 7, e1423172	7.2	20
38	Neoadjuvant plus adjuvant dabrafenib and trametinib versus standard of care in patients with high-risk, surgically resectable melanoma: a single-centre, open-label, randomised, phase 2 trial. <i>Lancet Oncology, The</i> , 2018 , 19, 181-193	21.7	168
37	Gut microbiome modulates response to anti-PD-1 immunotherapy in melanoma patients. <i>Science</i> , 2018 , 359, 97-103	33.3	1895
36	Concepts Collide: Genomic, Immune, and Microbial Influences on the Tumor Microenvironment and Response to Cancer Therapy. <i>Frontiers in Immunology</i> , 2018 , 9, 946	8.4	13
35	Association of the T-cell receptor landscape with survival in non-small cell lung cancer <i>Journal of Clinical Oncology</i> , 2018 , 36, 140-140	2.2	2

(2016-2018)

34	P2.04-09 Driver Mutations are Associated with Distinct Patterns of Response to Immune Checkpoint Blockade in Non-Small Cell Lung Cancer. <i>Journal of Thoracic Oncology</i> , 2018 , 13, S733-S734	8.9	2
33	Neoadjuvant immune checkpoint blockade in high-risk resectable melanoma. <i>Nature Medicine</i> , 2018 , 24, 1649-1654	50.5	377
32	Defining T Cell States Associated with Response to Checkpoint Immunotherapy in Melanoma. <i>Cell</i> , 2018 , 175, 998-1013.e20	56.2	631
31	Combined Analysis of Antigen Presentation and T-cell Recognition Reveals Restricted Immune Responses in Melanoma. <i>Cancer Discovery</i> , 2018 , 8, 1366-1375	24.4	57
30	Integrated molecular analysis of tumor biopsies on sequential CTLA-4 and PD-1 blockade reveals markers of response and resistance. <i>Science Translational Medicine</i> , 2017 , 9,	17.5	409
29	An adaptive signaling network in melanoma inflammatory niches confers tolerance to MAPK signaling inhibition. <i>Journal of Experimental Medicine</i> , 2017 , 214, 1691-1710	16.6	51
28	Gene Targeting Meets Cell-Based Therapy: Raising the Tail, or Merely a Whimper?. <i>Clinical Cancer Research</i> , 2017 , 23, 327-329	12.9	1
27	Interaction of molecular alterations with immune response in melanoma. <i>Cancer</i> , 2017 , 123, 2130-2142	6.4	18
26	Genomic and immune heterogeneity are associated with differential responses to therapy in melanoma. <i>Npj Genomic Medicine</i> , 2017 , 2,	6.2	82
25	Potential role of intratumor bacteria in mediating tumor resistance to the chemotherapeutic drug gemcitabine. <i>Science</i> , 2017 , 357, 1156-1160	33-3	577
24	TCR Repertoire Intratumor Heterogeneity in Localized Lung Adenocarcinomas: An Association with Predicted Neoantigen Heterogeneity and Postsurgical Recurrence. <i>Cancer Discovery</i> , 2017 , 7, 1088-109	7 ^{24.4}	105
23	Toward a Molecular-Genetic Classification of Spitzoid Neoplasms. <i>Clinics in Laboratory Medicine</i> , 2017 , 37, 431-448	2.1	25
22	Comparative immunologic characterization of autoimmune giant cell myocarditis with ipilimumab. <i>OncoImmunology</i> , 2017 , 6, e1361097	7.2	31
21	Parallel profiling of immune infiltrate subsets in uveal melanoma versus cutaneous melanoma unveils similarities and differences: A pilot study. <i>OncoImmunology</i> , 2017 , 6, e1321187	7.2	32
20	The hemochromatosis protein HFE 20 years later: An emerging role in antigen presentation and in the immune system. <i>Immunity, Inflammation and Disease</i> , 2017 , 5, 218-232	2.4	24
19	Analysis of Immune Signatures in Longitudinal Tumor Samples Yields Insight into Biomarkers of Response and Mechanisms of Resistance to Immune Checkpoint Blockade. <i>Cancer Discovery</i> , 2016 , 6, 827-37	24.4	561
18	Distinct clinical patterns and immune infiltrates are observed at time of progression on targeted therapy versus immune checkpoint blockade for melanoma. <i>OncoImmunology</i> , 2016 , 5, e1136044	7.2	42
17	Targeted Therapies Combined With Immune Checkpoint Therapy. <i>Cancer Journal (Sudbury, Mass)</i> , 2016 , 22, 138-46	2.2	24

16	Monitoring immune responses in the tumor microenvironment. <i>Current Opinion in Immunology</i> , 2016 , 41, 23-31	7.8	76
15	Influences of BRAF Inhibitors on the Immune Microenvironment and the Rationale for Combined Molecular and Immune Targeted Therapy. <i>Current Oncology Reports</i> , 2016 , 18, 42	6.3	36
14	Clinical, Molecular, and Immune Analysis of Dabrafenib-Trametinib Combination Treatment for BRAF Inhibitor-Refractory Metastatic Melanoma: A Phase 2 Clinical Trial. <i>JAMA Oncology</i> , 2016 , 2, 1056	-64·4	34
13	Density, Distribution, and Composition of Immune Infiltrates Correlate with Survival in Merkel Cell Carcinoma. <i>Clinical Cancer Research</i> , 2016 , 22, 5553-5563	12.9	70
12	Immune Effects of Chemotherapy, Radiation, and Targeted Therapy and Opportunities for Combination With Immunotherapy. <i>Seminars in Oncology</i> , 2015 , 42, 601-16	5.5	83
11	Update on use of aldesleukin for treatment of high-risk metastatic melanoma. <i>ImmunoTargets and Therapy</i> , 2015 , 4, 79-89	9	14
10	Does It MEK a Difference? Understanding Immune Effects of Targeted Therapy. <i>Clinical Cancer Research</i> , 2015 , 21, 3102-4	12.9	22
9	T lymphocyte-derived TNF and IFN-Trepress HFE expression in cancer cells. <i>Molecular Immunology</i> , 2015 , 65, 259-66	4.3	5
8	Raising the bar: optimizing combinations of targeted therapy and immunotherapy. <i>Annals of Translational Medicine</i> , 2015 , 3, 272	3.2	
7	The WT hemochromatosis protein HFE inhibits CD8+ T-lymphocyte activation. <i>European Journal of Immunology</i> , 2014 , 44, 1604-14	6.1	15
6	Evidence of synergy with combined BRAF-targeted therapy and immune checkpoint blockade for metastatic melanoma. <i>Oncolmmunology</i> , 2014 , 3, e954956	7.2	17
5	Combining targeted therapy and immune checkpoint inhibitors in the treatment of metastatic melanoma. <i>Cancer Biology and Medicine</i> , 2014 , 11, 237-46	5.2	57
4	Polyfunctionality of a DKK1 self-antigen-specific CD8(+) T lymphocyte clone in lung cancer. <i>Cancer Immunology, Immunotherapy</i> , 2011 , 60, 1119-25	7.4	1
3	Antifungal susceptibility of 44 clinical isolates of Fusarium species determined by using a broth microdilution method. <i>Antimicrobial Agents and Chemotherapy</i> , 1989 , 33, 1647-9	5.9	75
2	Immune evolution from preneoplasia to invasive lung adenocarcinomas and underlying molecular featu	ıres	2
1	Deep learning-based prediction of the T cell receptor Intigen binding specificity. <i>Nature Machine Intelligence</i> ,	22.5	9