Amir Giladi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5210202/publications.pdf

Version: 2024-02-01

67 23,341 49 68 g-index

77 77 77 35513

times ranked

citing authors

docs citations

all docs

#	Article	IF	CITATIONS
1	Early antitumor activity of oral Langerhans cells is compromised by a carcinogen. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119 , .	7.1	15
2	Single-cell immunology: Past, present, and future. Immunity, 2022, 55, 393-404.	14.3	47
3	The interaction of CD4+ helper T cells with dendritic cells shapes the tumor microenvironment and immune checkpoint blockade response. Nature Cancer, 2022, 3, 303-317.	13.2	85
4	LGR5 expressing skin fibroblasts define a major cellular hub perturbed in scleroderma. Cell, 2022, 185, 1373-1388.e20.	28.9	50
5	Physically interacting beta-delta pairs in the regenerating pancreas revealed by single-cell sequencing. Molecular Metabolism, 2022, 60, 101467.	6.5	O
6	Alzheimer's disease modification mediated by bone marrow-derived macrophages via a TREM2-independent pathway in mouse model of amyloidosis. Nature Aging, 2022, 2, 60-73.	11.6	12
7	Digging for treasures in the tumour interactome. Nature Reviews Cancer, 2022, 22, 434-435.	28.4	1
8	Identification of the central intermediate in the extra-embryonic to embryonic endoderm transition through single-cell transcriptomics. Nature Cell Biology, 2022, 24, 833-844.	10.3	15
9	NF-κB activity during pancreas development regulates adult β-cell mass by modulating neonatal β-cell proliferation and apoptosis. Cell Death Discovery, 2021, 7, 2.	4.7	5
10	Identification of resistance pathways and therapeutic targets in relapsed multiple myeloma patients through single-cell sequencing. Nature Medicine, 2021, 27, 491-503.	30.7	118
11	XCR1+ type 1 conventional dendritic cells drive liver pathology in non-alcoholic steatohepatitis. Nature Medicine, 2021, 27, 1043-1054.	30.7	95
12	Clump sequencing exposes the spatial expression programs of intestinal secretory cells. Nature Communications, 2021, 12, 3074.	12.8	43
13	Single-cell analysis of regions of interest (SCARI) using a photosensitive tag. Nature Chemical Biology, 2021, 17, 1139-1147.	8.0	13
14	Meningeal lymphoid structures are activated under acute and chronic spinal cord pathologies. Life Science Alliance, 2021, 4, e202000907.	2.8	14
15	LifeTime and improving European healthcare through cell-based interceptive medicine. Nature, 2020, 587, 377-386.	27.8	108
16	Coupled scRNA-Seq and Intracellular Protein Activity Reveal an Immunosuppressive Role of TREM2 in Cancer. Cell, 2020, 182, 872-885.e19.	28.9	298
17	Host-Viral Infection Maps Reveal Signatures of Severe COVID-19 Patients. Cell, 2020, 181, 1475-1488.e12.	28.9	405
18	The Physiology, Pathology, and Potential Therapeutic Applications of the TREM2 Signaling Pathway. Cell, 2020, 181, 1207-1217.	28.9	279

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19	Dissecting cellular crosstalk by sequencing physically interacting cells. Nature Biotechnology, 2020, 38, 629-637.	17.5	187
20	C/EBPÎ 2 -Dependent Epigenetic Memory Induces Trained Immunity in Hematopoietic Stem Cells. Cell Stem Cell, 2020, 26, 657-674.e8.	11.1	180
21	Cancer-associated fibroblast compositions change with breast cancer progression linking the ratio of S100A4+ and PDPN+ CAFs to clinical outcome. Nature Cancer, 2020, 1, 692-708.	13.2	159
22	Spatiotemporal regulation of type I interferon expression determines the antiviral polarization of CD4+ T cells. Nature Immunology, 2020, 21, 321-330.	14.5	59
23	Cxcl10+ monocytes define a pathogenic subset in the central nervous system during autoimmune neuroinflammation. Nature Immunology, 2020, 21, 525-534.	14.5	74
24	A single cell atlas of the human liver tumor microenvironment. Molecular Systems Biology, 2020, 16, e9682.	7.2	99
25	Lipid-Associated Macrophages Control Metabolic Homeostasis in a Trem2-Dependent Manner. Cell, 2019, 178, 686-698.e14.	28.9	718
26	MetaCell: analysis of single-cell RNA-seq data using K-nn graph partitions. Genome Biology, 2019, 20, 206.	8.8	218
27	PD-1/PD-L1 checkpoint blockade harnesses monocyte-derived macrophages to combat cognitive impairment in a tauopathy mouse model. Nature Communications, 2019, 10, 465.	12.8	112
28	Plasmacytoid dendritic cells develop from Ly6D+ lymphoid progenitors distinct from the myeloid lineage. Nature Immunology, 2019, 20, 852-864.	14.5	162
29	MARS-seq2.0: an experimental and analytical pipeline for indexed sorting combined with single-cell RNA sequencing. Nature Protocols, 2019, 14, 1841-1862.	12.0	200
30	DC Respond to Cognate T Cell Interaction in the Antigen-Challenged Lymph Node. Frontiers in Immunology, 2019, 10, 863.	4.8	16
31	Single-Cell Analysis of Diverse Pathogen Responses Defines a Molecular Roadmap for Generating Antigen-Specific Immunity. Cell Systems, 2019, 8, 109-121.e6.	6.2	39
32	Cross-Species Single-Cell Analysis Reveals Divergence of the Primate Microglia Program. Cell, 2019, 179, 1609-1622.e16.	28.9	292
33	Dysfunctional CD8 T Cells Form a Proliferative, Dynamically Regulated Compartment within Human Melanoma. Cell, 2019, 176, 775-789.e18.	28.9	760
34	Embrace the fat when getting old. Aging, 2019, 11, 8730-8732.	3.1	3
35	Single-Cell Genomics: A Stepping Stone for Future Immunology Discoveries. Cell, 2018, 172, 14-21.	28.9	214
36	Impaired immune surveillance accelerates accumulation of senescent cells and aging. Nature Communications, 2018, 9, 5435.	12.8	325

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37	Single cell dissection of plasma cell heterogeneity in symptomatic and asymptomatic myeloma. Nature Medicine, 2018, 24, 1867-1876.	30.7	179
38	Lung Single-Cell Signaling Interaction Map Reveals Basophil Role in Macrophage Imprinting. Cell, 2018, 175, 1031-1044.e18.	28.9	332
39	Paired-cell sequencing enables spatial gene expression mapping of liver endothelial cells. Nature Biotechnology, 2018, 36, 962-970.	17.5	262
40	Disease-Associated Microglia: A Universal Immune Sensor of Neurodegeneration. Cell, 2018, 173, 1073-1081.	28.9	765
41	Early metazoan cell type diversity and the evolution of multicellular gene regulation. Nature Ecology and Evolution, 2018, 2, 1176-1188.	7.8	226
42	From the Human Cell Atlas to dynamic immune maps in human disease. Nature Reviews Immunology, 2018, 18, 597-598.	22.7	23
43	Single-cell mapping of the thymic stroma identifies IL-25-producing tuft epithelial cells. Nature, 2018, 559, 622-626.	27.8	235
44	Differences in Cell Cycle Status Underlie Transcriptional Heterogeneity in the HSC Compartment. Cell Reports, 2018, 24, 766-780.	6.4	40
45	Single-cell characterization of haematopoietic progenitors and their trajectories in homeostasis and perturbed haematopoiesis. Nature Cell Biology, 2018, 20, 836-846.	10.3	267
46	Single-cell spatial reconstruction reveals global division of labour in the mammalian liver. Nature, 2017, 542, 352-356.	27.8	809
47	Single-cell transcriptome conservation in cryopreserved cells and tissues. Genome Biology, 2017, 18, 45.	8.8	134
48	Genomic Characterization of Murine Monocytes Reveals C/EBP \hat{l}^2 Transcription Factor Dependence of Ly6C \hat{a}^2 Cells. Immunity, 2017, 46, 849-862.e7.	14.3	233
49	Innate Immune Landscape in Early Lung Adenocarcinoma by Paired Single-Cell Analyses. Cell, 2017, 169, 750-765.e17.	28.9	937
50	A Unique Microglia Type Associated with Restricting Development of Alzheimer's Disease. Cell, 2017, 169, 1276-1290.e17.	28.9	3,282
51	Spatial reconstruction of immune niches by combining photoactivatable reporters and scRNA-seq. Science, 2017, 358, 1622-1626.	12.6	176
52	The Human Cell Atlas. ELife, 2017, 6, .	6.0	1,547
53	Immunology, one cell at a time. Nature, 2017, 547, 27-29.	27.8	33
54	Dissecting Immune Circuits by Linking CRISPR-Pooled Screens with Single-Cell RNA-Seq. Cell, 2016, 167, 1883-1896.e15.	28.9	604

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55	Extracellular Matrix Proteolysis by MT1-MMP Contributes to Influenza-Related Tissue Damage and Mortality. Cell Host and Microbe, 2016, 20, 458-470.	11.0	82
56	The Spectrum and Regulatory Landscape of Intestinal Innate Lymphoid Cells Are Shaped by the Microbiome. Cell, 2016, 166, 1231-1246.e13.	28.9	465
57	Microglia development follows a stepwise program to regulate brain homeostasis. Science, 2016, 353, aad8670.	12.6	911
58	PD-1 immune checkpoint blockade reduces pathology and improves memory in mouse models of Alzheimer's disease. Nature Medicine, 2016, 22, 135-137.	30.7	286
59	Sumoylation coordinates the repression of inflammatory and anti-viral gene-expression programs during innate sensing. Nature Immunology, 2016, 17, 140-149.	14.5	127
60	Transcription factor binding dynamics during human ES cell differentiation. Nature, 2015, 518, 344-349.	27.8	318
61	Transcriptional Heterogeneity and Lineage Commitment in Myeloid Progenitors. Cell, 2015, 163, 1663-1677.	28.9	875
62	Tissue-Resident Macrophage Enhancer Landscapes Are Shaped by the Local Microenvironment. Cell, 2014, 159, 1312-1326.	28.9	1,705
63	Massively Parallel Single-Cell RNA-Seq for Marker-Free Decomposition of Tissues into Cell Types. Science, 2014, 343, 776-779.	12.6	1,563
64	A Negative Feedback Loop of Transcription Factors Specifies Alternative Dendritic Cell Chromatin States. Molecular Cell, 2014, 56, 749-762.	9.7	58
65	Chromatin state dynamics during blood formation. Science, 2014, 345, 943-949.	12.6	699
66	Aging-induced type I interferon response at the choroid plexus negatively affects brain function. Science, 2014, 346, 89-93.	12.6	463
67	High-throughput chromatin immunoprecipitation for genome-wide mapping of in vivo protein-DNA interactions and epigenomic states. Nature Protocols, 2013, 8, 539-554.	12.0	246