## Hadas Keren-Shaul

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3605869/publications.pdf

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#	Article	IF	CITATIONS
1	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
2	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
3	Transneuronal Dpr12/DIPâ€î interactions facilitate compartmentalized dopaminergic innervation of <i>Drosophila</i> mushroom body axons. EMBO Journal, 2021, 40, e105763.	7.8	15
4	Dissection of floral transition by single-meristem transcriptomes at high temporal resolution. Nature Plants, 2021, 7, 800-813.	9.3	26
5	Coupled scRNA-Seq and Intracellular Protein Activity Reveal an Immunosuppressive Role of TREM2 in Cancer. Cell, 2020, 182, 872-885.e19.	28.9	298
6	Lipid-Associated Macrophages Control Metabolic Homeostasis in a Trem2-Dependent Manner. Cell, 2019, 178, 686-698.e14.	28.9	718
7	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
8	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
9	Cross-Species Single-Cell Analysis Reveals Divergence of the Primate Microglia Program. Cell, 2019, 179, 1609-1622.e16.	28.9	292
10	Single cell dissection of plasma cell heterogeneity in symptomatic and asymptomatic myeloma. Nature Medicine, 2018, 24, 1867-1876.	30.7	179
11	Combining Developmental and Perturbation-Seq Uncovers Transcriptional Modules Orchestrating Neuronal Remodeling. Developmental Cell, 2018, 47, 38-52.e6.	7.0	56
12	Disease-Associated Microglia: A Universal Immune Sensor of Neurodegeneration. Cell, 2018, 173, 1073-1081.	28.9	765
13	The †TranSeq' 3′â€end sequencing method for highâ€ŧhroughput transcriptomics and gene space refine in plant genomes. Plant Journal, 2018, 96, 223-232.	ment 5.7	23
14	Single-cell transcriptome conservation in cryopreserved cells and tissues. Genome Biology, 2017, 18, 45.	8.8	134
15	Dicer Deficiency Differentially Impacts Microglia of the Developing and Adult Brain. Immunity, 2017, 46, 1030-1044.e8.	14.3	68
16	A Unique Microglia Type Associated with Restricting Development of Alzheimer's Disease. Cell, 2017, 169, 1276-1290.e17.	28.9	3,282
17	Dissecting Immune Circuits by Linking CRISPR-Pooled Screens with Single-Cell RNA-Seq. Cell, 2016, 167, 1883-1896.e15.	28.9	604
18	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

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19	The Spectrum and Regulatory Landscape of Intestinal Innate Lymphoid Cells Are Shaped by the Microbiome. Cell, 2016, 166, 1231-1246.e13.	28.9	465
20	Microglia development follows a stepwise program to regulate brain homeostasis. Science, 2016, 353, aad8670.	12.6	911
21	Each cell counts: Hematopoiesis and immunity research in the era of single cell genomics. Seminars in Immunology, 2015, 27, 67-71.	5.6	35
22	Transcriptional Heterogeneity and Lineage Commitment in Myeloid Progenitors. Cell, 2015, 163, 1663-1677.	28.9	875
23	Chronic exposure to <scp>TGF</scp> β1 regulates myeloid cell inflammatory response in an <scp>IRF</scp> 7â€dependent manner. EMBO Journal, 2014, 33, 2906-2921.	7.8	95
24	Tissue-Resident Macrophage Enhancer Landscapes Are Shaped by the Local Microenvironment. Cell, 2014, 159, 1312-1326.	28.9	1,705
25	Massively Parallel Single-Cell RNA-Seq for Marker-Free Decomposition of Tissues into Cell Types. Science, 2014, 343, 776-779.	12.6	1,563
26	Chromatin state dynamics during blood formation. Science, 2014, 345, 943-949.	12.6	699