

Søren Møller

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

2,910
citations

516710

16
h-index

713466

21
g-index

25
all docs

25
docs citations

25
times ranked

4556
citing authors

#	ARTICLE	IF	CITATIONS
1	Single-Cell RNA Sequencing Reveals Stromal Evolution into LRRC15+ Myofibroblasts as a Determinant of Patient Response to Cancer Immunotherapy. <i>Cancer Discovery</i> , 2020, 10, 232-253.	9.4	466
2	Cross-tissue organization of the fibroblast lineage. <i>Nature</i> , 2021, 593, 575-579.	27.8	463
3	Single-cell profiling of human gliomas reveals macrophage ontogeny as a basis for regional differences in macrophage activation in the tumor microenvironment. <i>Genome Biology</i> , 2017, 18, 234.	8.8	448
4	Outer Radial Glia-like Cancer Stem Cells Contribute to Heterogeneity of Glioblastoma. <i>Cell Stem Cell</i> , 2020, 26, 48-63.e6.	11.1	222
5	Interpretation of T cell states from single-cell transcriptomics data using reference atlases. <i>Nature Communications</i> , 2021, 12, 2965.	12.8	210
6	The Phenotypes of Proliferating Glioblastoma Cells Reside on a Single Axis of Variation. <i>Cancer Discovery</i> , 2019, 9, 1708-1719.	9.4	205
7	Mesothelial cell-derived antigen-presenting cancer-associated fibroblasts induce expansion of regulatory T cells in pancreatic cancer. <i>Cancer Cell</i> , 2022, 40, 656-673.e7.	16.8	155
8	Single-cell analysis of human non-small cell lung cancer lesions refines tumor classification and patient stratification. <i>Cancer Cell</i> , 2021, 39, 1594-1609.e12.	16.8	151
9	Intratumoral plasma cells predict outcomes to PD-L1 blockade in non-small cell lung cancer. <i>Cancer Cell</i> , 2022, 40, 289-300.e4.	16.8	148
10	Single-cell sequencing maps gene expression to mutational phylogenies in PDGF- and EGF-driven gliomas. <i>Molecular Systems Biology</i> , 2016, 12, 889.	7.2	91
11	CONICS integrates scRNA-seq with DNA sequencing to map gene expression to tumor sub-clones. <i>Bioinformatics</i> , 2018, 34, 3217-3219.	4.1	87
12	Gremlin 1+ fibroblastic niche maintains dendritic cell homeostasis in lymphoid tissues. <i>Nature Immunology</i> , 2021, 22, 571-585.	14.5	44
13	Coexpression of Inhibitory Receptors Enriches for Activated and Functional CD8+ T Cells in Murine Syngeneic Tumor Models. <i>Cancer Immunology Research</i> , 2019, 7, 963-976.	3.4	36
14	Clonal ZEB1-Driven Mesenchymal Transition Promotes Targetable Oncologic Antiangiogenic Therapy Resistance. <i>Cancer Research</i> , 2020, 80, 1498-1511.	0.9	35
15	Single-Cell mRNA Sequencing in Cancer Research: Integrating the Genomic Fingerprint. <i>Frontiers in Genetics</i> , 2017, 8, 73.	2.3	27
16	A bird's eye view of fibroblast heterogeneity: A pan-disease, pan-cancer perspective. <i>Immunological Reviews</i> , 2021, 302, 299-320.	6.0	23
17	Peptide vaccine immunotherapy biomarkers and response patterns in pediatric gliomas. <i>JCI Insight</i> , 2018, 3, .	5.0	21
18	Single-cell Co-expression Subnetwork Analysis. <i>Scientific Reports</i> , 2017, 7, 15066.	3.3	19

#	ARTICLE	IF	CITATIONS
19	miRNA-independent function of long noncoding pri-miRNA loci. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	18
20	A membrane protein display platform for receptor interactome discovery. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	17
21	MHC class I H2-Kb negatively regulates neural progenitor cell proliferation by inhibiting FGFR signaling. PLoS Biology, 2021, 19, e3001311.	5.6	14