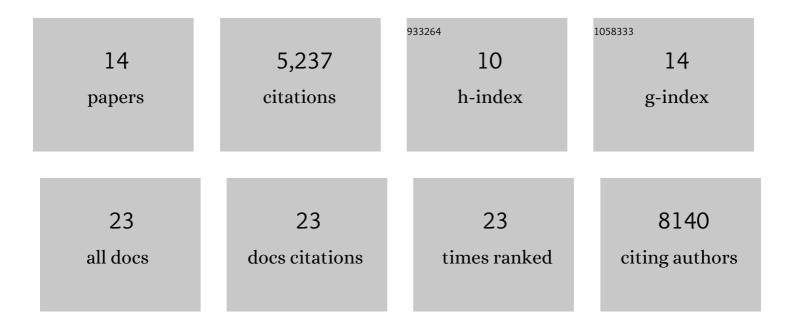
Ilya Korsunsky

List of Publications by Year in descending order

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LIVA KODSUNSKY

#	Article	IF	CITATIONS
1	Multimodal platform for assessing drug distribution and response in clinical trials. Neuro-Oncology, 2022, 24, 64-77.	0.6	4
2	Co-varying neighborhood analysis identifies cell populations associated with phenotypes of interest from single-cell transcriptomics. Nature Biotechnology, 2022, 40, 355-363.	9.4	30
3	Maternal oxytocin administration modulates gene expression in the brains of perinatal mice. Journal of Perinatal Medicine, 2022, 50, 207-218.	0.6	3
4	A framework for employing longitudinally collected multicenter electronic health records to stratify heterogeneous patient populations on disease history. Journal of the American Medical Informatics Association: JAMIA, 2022, 29, 761-769.	2.2	6
5	Cross-tissue, single-cell stromal atlas identifies shared pathological fibroblast phenotypes in four chronic inflammatory diseases. Med, 2022, 3, 481-518.e14.	2.2	51
6	IFN-γ and TNF-α drive a CXCL10+ CCL2+ macrophage phenotype expanded in severe COVID-19 lungs and inflammatory diseases with tissue inflammation. Genome Medicine, 2021, 13, 64.	3.6	128
7	IL-1-driven stromal–neutrophil interactions define a subset of patients with inflammatory bowel disease that does not respond to therapies. Nature Medicine, 2021, 27, 1970-1981.	15.2	117
8	Efficient and precise single-cell reference atlas mapping with Symphony. Nature Communications, 2021, 12, 5890.	5.8	100
9	Maximizing statistical power to detect differentially abundant cell states with scPOST. Cell Reports Methods, 2021, 1, 100120.	1.4	2
10	Notch signalling drives synovial fibroblast identity and arthritis pathology. Nature, 2020, 582, 259-264.	13.7	267
11	Distinct fibroblast subsets drive inflammation and damage in arthritis. Nature, 2019, 570, 246-251.	13.7	550
12	Lymphocyte innateness defined by transcriptional states reflects a balance between proliferation and effector functions. Nature Communications, 2019, 10, 687.	5.8	136
13	Fast, sensitive and accurate integration of single-cell data with Harmony. Nature Methods, 2019, 16, 1289-1296.	9.0	3,494
14	Mixed-effects association of single cells identifies an expanded effector CD4 ⁺ T cell subset in rheumatoid arthritis. Science Translational Medicine, 2018, 10, .	5.8	119