Jeffrey E Mold

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8232031/publications.pdf

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		567281	839539
18	1,315	15	18
papers	citations	h-index	g-index
22	22	22	2001
23	23	23	2891
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	The Lifespan and Turnover of Microglia in the Human Brain. Cell Reports, 2017, 20, 779-784.	6.4	340
2	Antibody-secreting plasma cells persist for decades in human intestine. Journal of Experimental Medicine, 2017, 214, 309-317.	8.5	173
3	Analysis of allelic expression patterns in clonal somatic cells by single-cell RNA–seq. Nature Genetics, 2016, 48, 1430-1435.	21.4	142
4	Immunological Tolerance During Fetal Development. Advances in Immunology, 2012, 115, 73-111.	2.2	122
5	Unique transcriptional and protein-expression signature in human lung tissue-resident NK cells. Nature Communications, 2019, 10, 3841.	12.8	7 9
6	Transplanted Bone Marrow-Derived Cells Contribute to Human Adipogenesis. Cell Metabolism, 2015, 22, 408-417.	16.2	75
7	Comparison of whole genome amplification techniques for human single cell exome sequencing. PLoS ONE, 2017, 12, e0171566.	2.5	63
8	Activation of a neural stem cell transcriptional program in parenchymal astrocytes. ELife, 2020, 9, .	6.0	51
9	Cell generation dynamics underlying naive T-cell homeostasis in adult humans. PLoS Biology, 2019, 17, e3000383.	5.6	45
10	Expansions of adaptive-like NK cells with a tissue-resident phenotype in human lung and blood. Proceedings of the National Academy of Sciences of the United States of America, $2021,118,118$	7.1	43
11	At the crossroads between tolerance and aggression. Chimerism, 2011, 2, 35-41.	0.7	36
12	A biliary immune landscape map of primary sclerosing cholangitis reveals a dominant network of neutrophils and tissue-resident T cells. Science Translational Medicine, 2021, 13, .	12.4	31
13	Probabilistic count matrix factorization for single cell expression data analysis. Bioinformatics, 2019, 35, 4011-4019.	4.1	29
14	High dimensional classification with combined adaptive sparse PLS and logistic regression. Bioinformatics, 2018, 34, 485-493.	4.1	21
15	Conbase: a software for unsupervised discovery of clonal somatic mutations in single cells through read phasing. Genome Biology, 2019, 20, 68.	8.8	21
16	Divergent clonal differentiation trajectories establish CD8+ memory Tâcell heterogeneity during acute viral infections in humans. Cell Reports, 2021, 35, 109174.	6.4	9
17	A heterozygous germline CD100 mutation in a family with primary sclerosing cholangitis. Science Translational Medicine, 2021, 13, .	12.4	8
18	A discussion of immune tolerance and the layered immune system hypothesis. Chimerism, 2013, 4, 62-70.	0.7	5