

Samira Asgari

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

17
papers

354
citations

8
h-index

18
g-index

26
ext. papers

516
ext. citations

18.5
avg, IF

3.05
L-index

#	Paper	IF	Citations
17	Differentiation and transplantation of human induced pluripotent stem cell-derived hepatocyte-like cells. <i>Stem Cell Reviews and Reports</i> , 2013 , 9, 493-504	6.4	79
16	Severe viral respiratory infections in children with loss-of-function mutations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017 , 114, 8342-8347	11.5	74
15	Induced pluripotent stem cells: a new era for hepatology. <i>Journal of Hepatology</i> , 2010 , 53, 738-51	13.4	69
14	Disease-corrected hepatocyte-like cells from familial hypercholesterolemia-induced pluripotent stem cells. <i>Molecular Biotechnology</i> , 2013 , 54, 863-73	3	34
13	Early progression to active tuberculosis is a highly heritable trait driven by 3q23 in Peruvians. <i>Nature Communications</i> , 2019 , 10, 3765	17.4	18
12	A positively selected FBN1 missense variant reduces height in Peruvian individuals. <i>Nature</i> , 2020 , 582, 234-239	50.4	15
11	Exome Sequencing Reveals Primary Immunodeficiencies in Children with Community-Acquired Sepsis. <i>Frontiers in Immunology</i> , 2016 , 7, 357	8.4	14
10	Multimodally profiling memory T cells from a tuberculosis cohort identifies cell state associations with demographics, environment and disease. <i>Nature Immunology</i> , 2021 , 22, 781-793	19.1	13
9	Multimodal memory T cell profiling identifies a reduction in a polyfunctional Th17 state associated with tuberculosis progression		7
8	From Your Nose to Your Toes: A Review of Severe Acute Respiratory Syndrome Coronavirus 2 Pandemic-Associated Pernio. <i>Journal of Investigative Dermatology</i> , 2021 , 141, 2791-2796	4.3	6
7	Whole-exome Sequencing for the Identification of Rare Variants in Primary Immunodeficiency Genes in Children With Sepsis: A Prospective, Population-based Cohort Study. <i>Clinical Infectious Diseases</i> , 2020 , 71, e614-e623	11.6	4
6	Single-cell eQTL models reveal dynamic T cell state dependence of disease loci. <i>Nature</i> , 2022 ,	50.4	4
5	A positively selected, common, missense variant in FBN1 confers a 2.2 centimeter reduction of height in the Peruvian population		3
4	Axes of inter-sample variability among transcriptional neighborhoods reveal disease associated cell states in single-cell data		2
3	Human genomics of acute liver failure due to hepatitis B virus infection: An exome sequencing study in liver transplant recipients. <i>Journal of Viral Hepatitis</i> , 2019 , 26, 271-277	3.4	2
2	Co-varying neighborhood analysis identifies cell populations associated with phenotypes of interest from single-cell transcriptomics. <i>Nature Biotechnology</i> , 2021 ,	44.5	1
1	Modeling memory T cell states at single-cell resolution identifies in vivo state-dependence of eQTLs influencing disease		1

