

Samira Asgari

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

672
citations

758635

12
h-index

887659

17
g-index

26
all docs

26
docs citations

26
times ranked

1424
citing authors

#	ARTICLE	IF	CITATIONS
1	Severe viral respiratory infections in children with <i>IFIH1</i> loss-of-function mutations. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 8342-8347.	3.3	111
2	Differentiation and Transplantation of Human Induced Pluripotent Stem Cell-derived Hepatocyte-like Cells. Stem Cell Reviews and Reports, 2013, 9, 493-504.	5.6	91
3	Induced pluripotent stem cells: A new era for hepatology. Journal of Hepatology, 2010, 53, 738-751.	1.8	77
4	Single-cell eQTL models reveal dynamic T cell state dependence of disease loci. Nature, 2022, 606, 120-128.	13.7	75
5	Multimodally profiling memory T cells from a tuberculosis cohort identifies cell state associations with demographics, environment and disease. Nature Immunology, 2021, 22, 781-793.	7.0	52
6	Early progression to active tuberculosis is a highly heritable trait driven by 3q23 in Peruvians. Nature Communications, 2019, 10, 3765.	5.8	43
7	Disease-Corrected Hepatocyte-Like Cells from Familial Hypercholesterolemia-Induced Pluripotent Stem Cells. Molecular Biotechnology, 2013, 54, 863-873.	1.3	41
8	A positively selected <i>FBN1</i> missense variant reduces height in Peruvian individuals. Nature, 2020, 582, 234-239.	13.7	39
9	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
10	Exome Sequencing Reveals Primary Immunodeficiencies in Children with Community-Acquired <i>Pseudomonas aeruginosa</i> Sepsis. Frontiers in Immunology, 2016, 7, 357.	2.2	21
11	From Your Nose to Your Toes: A Review of Severe Acute Respiratory Syndrome Coronavirus 2 Pandemic-Associated Pernio. Journal of Investigative Dermatology, 2021, 141, 2791-2796.	0.3	21
12	Human genetic variants identified that affect COVID susceptibility and severity. Nature, 2021, 600, 390-391.	13.7	16
13	Whole-exome Sequencing for the Identification of Rare Variants in Primary Immunodeficiency Genes in Children With Sepsis: A Prospective, Population-based Cohort Study. Clinical Infectious Diseases, 2020, 71, e614-e623.	2.9	12
14	A sex-specific evolutionary interaction between <i>ADCY9</i> and <i>CETP</i> . ELife, 2021, 10, .	2.8	8
15	Higher native Peruvian genetic ancestry proportion is associated with tuberculosis progression risk. Cell Genomics, 2022, 2, 100151.	3.0	5
16	Human genomics of acute liver failure due to hepatitis B virus infection: An exome sequencing study in liver transplant recipients. Journal of Viral Hepatitis, 2019, 26, 271-277.	1.0	4