

Xinli Hu

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11431548/publications.pdf>

Version: 2024-02-01

13
papers

5,896
citations

687220

13
h-index

1125617

13
g-index

13
all docs

13
docs citations

13
times ranked

13417
citing authors

#	ARTICLE	IF	CITATIONS
1	Host-microbe interactions have shaped the genetic architecture of inflammatory bowel disease. <i>Nature</i> , 2012, 491, 119-124.	13.7	4,038
2	High-density genetic mapping identifies new susceptibility loci for rheumatoid arthritis. <i>Nature Genetics</i> , 2012, 44, 1336-1340.	9.4	558
3	Automated high-dimensional flow cytometric data analysis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 8519-8524.	3.3	355
4	Additive and interaction effects at three amino acid positions in HLA-DQ and HLA-DR molecules drive type 1 diabetes risk. <i>Nature Genetics</i> , 2015, 47, 898-905.	9.4	235
5	Widespread non-additive and interaction effects within HLA loci modulate the risk of autoimmune diseases. <i>Nature Genetics</i> , 2015, 47, 1085-1090.	9.4	164
6	Integrating Autoimmune Risk Loci with Gene-Expression Data Identifies Specific Pathogenic Immune Cell Subsets. <i>American Journal of Human Genetics</i> , 2011, 89, 496-506.	2.6	159
7	Disentangling the Effects of Colocalizing Genomic Annotations to Functionally Prioritize Non-coding Variants within Complex-Trait Loci. <i>American Journal of Human Genetics</i> , 2015, 97, 139-152.	2.6	122
8	A method to decipher pleiotropy by detecting underlying heterogeneity driven by hidden subgroups applied to autoimmune and neuropsychiatric diseases. <i>Nature Genetics</i> , 2016, 48, 803-810.	9.4	62
9	SNPsea: an algorithm to identify cell types, tissues and pathways affected by risk loci. <i>Bioinformatics</i> , 2014, 30, 2496-2497.	1.8	60
10	New data and an old puzzle: the negative association between schizophrenia and rheumatoid arthritis. <i>International Journal of Epidemiology</i> , 2015, 44, 1706-1721.	0.9	53
11	Regulation of Gene Expression in Autoimmune Disease Loci and the Genetic Basis of Proliferation in CD4+ Effector Memory T Cells. <i>PLoS Genetics</i> , 2014, 10, e1004404.	1.5	46
12	What have we learned from six years of GWAS in autoimmune diseases, and what is next?. <i>Current Opinion in Immunology</i> , 2012, 24, 571-575.	2.4	28
13	Application of user-guided automated cytometric data analysis to large-scale immunoprofiling of invariant natural killer T cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 19030-19035.	3.3	16