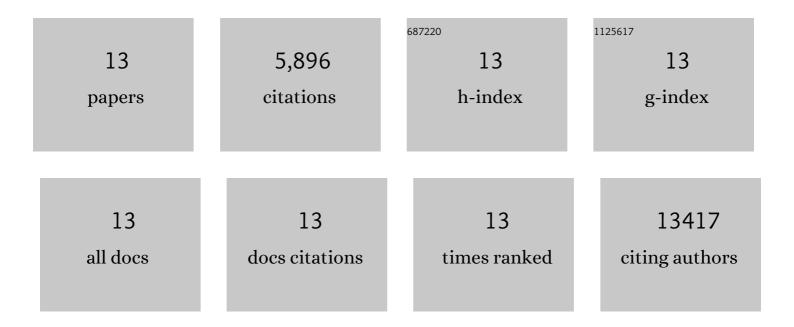
Xinli Hu

List of Publications by Year in descending order

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