

# Yeguang Hu

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/995823/publications.pdf>

Version: 2024-02-01

19  
papers

1,340  
citations

759233

12  
h-index

839539

18  
g-index

21  
all docs

21  
docs citations

21  
times ranked

2364  
citing authors

#	ARTICLE	IF	CITATIONS
1	Early hematopoietic lineage restrictions directed by Ikaros. <i>Nature Immunology</i> , 2006, 7, 382-391.	14.5	272
2	Genome-wide Lineage-Specific Transcriptional Networks Underscore Ikaros-Dependent Lymphoid Priming in Hematopoietic Stem Cells. <i>Immunity</i> , 2009, 30, 493-507.	14.3	221
3	The role of the chromatin remodeler Mi-2 <sup>1</sup> in hematopoietic stem cell self-renewal and multilineage differentiation. <i>Genes and Development</i> , 2008, 22, 1174-1189.	5.9	168
4	Harnessing of the nucleosome-remodeling-deacetylase complex controls lymphocyte development and prevents leukemogenesis. <i>Nature Immunology</i> , 2012, 13, 86-94.	14.5	154
5	Loss of Ikaros DNA-binding function confers integrin-dependent survival on pre-B cells and progression to acute lymphoblastic leukemia. <i>Nature Immunology</i> , 2014, 15, 294-304.	14.5	136
6	Promoter Decommissioning by the NuRD Chromatin Remodeling Complex Triggers Synaptic Connectivity in the Mammalian Brain. <i>Neuron</i> , 2014, 83, 122-134.	8.1	92
7	Ikaros fingers on lymphocyte differentiation. <i>International Journal of Hematology</i> , 2014, 100, 220-229.	1.6	68
8	Superenhancer reprogramming drives a B-cell <sup>2</sup> epithelial transition and high-risk leukemia. <i>Genes and Development</i> , 2016, 30, 1971-1990.	5.9	59
9	Awakening lineage potential by Ikaros-mediated transcriptional priming. <i>Current Opinion in Immunology</i> , 2010, 22, 154-160.	5.5	44
10	Transcriptional regulation of the Ikaros locus. <i>Blood</i> , 2013, 122, 3149-3159.	1.4	30
11	Chromatin restriction by the nucleosome remodeler Mi-2 <sup>1</sup> and functional interplay with lineage-specific transcription regulators control B-cell differentiation. <i>Genes and Development</i> , 2019, 33, 763-781.	5.9	26
12	STAG2 regulates interferon signaling in melanoma via enhancer loop reprogramming. <i>Nature Communications</i> , 2022, 13, 1859.	12.8	21
13	Transcriptional circuits in B cell transformation. <i>Current Opinion in Hematology</i> , 2017, 24, 345-352.	2.5	19
14	Hypomorphic mutation of the mouse Huntington <sup>3</sup> disease gene orthologue. <i>PLoS Genetics</i> , 2019, 15, e1007765.	3.5	13
15	An Ikaros Promoter Element with Dual Epigenetic and Transcriptional Activities. <i>PLoS ONE</i> , 2015, 10, e0131568.	2.5	7
16	GATA-3 controls self-renewal in stressed HSCs. <i>Nature Immunology</i> , 2013, 14, 1032-1033.	14.5	5
17	Focal Adhesion Kinase Inhibitors Reverse the Stromal Adhesion Phenotype of Ikaros-Mutant B-ALL, Induce Apoptosis, and Synergize with ABL1 Tyrosine Kinase Inhibitors: A New Paradigm for Pathogenesis and Therapy of High-Risk B-ALL. <i>Blood</i> , 2014, 124, 285-285.	1.4	3
18	The Developmental Transcription Factor p63 Is Redeployed to Drive Allergic Skin Inflammation through Phosphorylation by p38 <sup>1</sup> . <i>Journal of Immunology</i> , 2022, 208, 2613-2621.	0.8	2

#	ARTICLE	IF	CITATIONS
19	Ikaros Mutation Confers Integrin-Dependent Survival Of Pre-B Cells and Progression To Acute Lymphoblastic Leukemia. Blood, 2013, 122, 1259-1259.	1.4	0