

# Xiaozhou Hu

## List of Publications by Year in descending order

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Version: 2024-02-01

11  
papers

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1478505

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times ranked

1167  
citing authors

#	ARTICLE	IF	CITATIONS
1	Activating mutations of STAT5B and STAT3 in lymphomas derived from $\hat{\beta}$ <sup>1</sup> -T or NK cells. Nature Communications, 2015, 6, 6025.	12.8	334
2	<i>PRDM1</i> is a tumor suppressor gene in natural killer cell malignancies. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 20119-20124.	7.1	103
3	Global Promoter Methylation Analysis Reveals Novel Candidate Tumor Suppressor Genes in Natural Killer Cell Lymphoma. Clinical Cancer Research, 2015, 21, 1699-1711.	7.0	78
4	HACE1 Is a Tumor Suppressor Gene Candidate in Natural Killer Cell Neoplasms. American Journal of Pathology, 2013, 182, 49-55.	3.8	52
5	Diagnostic and Biological Significance of KIR Expression Profile Determined by RNA-Seq in Natural Killer/T-Cell Lymphoma. American Journal of Pathology, 2016, 186, 1435-1441.	3.8	16
6	The relationship of REL proto-oncogene to pathobiology and chemoresistance in follicular and transformed follicular lymphoma. Leukemia Research, 2017, 54, 30-38.	0.8	6
7	PRDM1 decreases sensitivity of human NK cells to IL2-induced cell expansion by directly repressing CD25 (IL2RA). Journal of Leukocyte Biology, 2021, 109, 901-914.	3.3	5
8	Generation of a genetically engineered aggressive Nk-Cell leukemia cell line with stable IL2 expression. Acta Medica International, 2015, 2, 78.	0.2	1
9	Frequent Activating Mutations Of JAK-STAT Pathway Genes In Natural Killer Cell Lymphomas. Blood, 2013, 122, 812-812.	1.4	1
10	Lack of Evidence that HACE1 Is Not a Tumor Suppressor Gene in NKTCL. American Journal of Pathology, 2015, 185, 1167-1168.	3.8	0
11	Global Methylation Analysis Reveals Novel Candidate Tumor Suppressor Genes In Natural Killer Cell Lymphomas. Blood, 2013, 122, 1262-1262.	1.4	0