Andrew J Woo

List of Publications by Year
in descending order

23 papers	1,470 citations	566801 15 h-index	752256 20 g-index
23 all docs	23 docs citations	23 times ranked	3487 citing authors

#	Article	IF	Citations
1	A Myc Network Accounts for Similarities between Embryonic Stem and Cancer Cell Transcription Programs. Cell, 2010, 143, 313-324.	13.5	606
2	Direct Recruitment of Polycomb Repressive Complex 1 to Chromatin by Core Binding Transcription Factors. Molecular Cell, 2012, 45, 330-343.	4.5	188
3	Distinct and Combinatorial Functions of Jmjd2b/Kdm4b and Jmjd2c/Kdm4c in Mouse Embryonic Stem Cell Identity. Molecular Cell, 2014, 53, 32-48.	4.5	112
4	Differentiation-Dependent Interactions between RUNX-1 and FLI-1 during Megakaryocyte Development. Molecular and Cellular Biology, 2009, 29, 4103-4115.	1.1	71
5	Identification of ZBP-89 as a Novel GATA-1-Associated Transcription Factor Involved in Megakaryocytic and Erythroid Development. Molecular and Cellular Biology, 2008, 28, 2675-2689.	1.1	62
6	Targeting RSPO3-LGR4 Signaling for Leukemia Stem Cell Eradication in Acute Myeloid Leukemia. Cancer Cell, 2020, 38, 263-278.e6.	7.7	59
7	Tumor penetrating peptides inhibiting MYC as a potent targeted therapeutic strategy for triple-negative breast cancers. Oncogene, 2019, 38, 140-150.	2.6	55
8	Surfactant Proteinâ€“C Chromatin-Bound Green Fluorescence Protein Reporter Mice Reveal Heterogeneity of Surfactant Protein Câ€“Expressing Lung Cells. American Journal of Respiratory Cell and Molecular Biology, 2013, 48, 288-298.	1.4	54
9	A Src family kinaseâ€“Shp2 axis controls RUNX1 activity in megakaryocyte and T-lymphocyte differentiation. Genes and Development, 2012, 26, 1587-1601.	2.7	52
10	Developmental differences in IFN signaling affect GATA1s-induced megakaryocyte hyperproliferation. Journal of Clinical Investigation, 2013, 123, 3292-3304.	3.9	37
11	A Proteomics Approach for the Identification of DNA Binding Activities Observed in the Electrophoretic Mobility Shift Assay. Molecular and Cellular Proteomics, 2002, 1, 472-478.	2.5	33
12	JMJD1C-mediated metabolic dysregulation contributes to HOXA9-dependent leukemogenesis. Leukemia, 2019, 33, 1400-1410.	3.3	31
13	Role of ZBP-89 in human globin gene regulation and erythroid differentiation. Blood, 2011, 118, 3684-3693.	0.6	26
14	CpG island-mediated global gene regulatory modes in mouse embryonic stem cells. Nature Communications, 2014, 5, 5490.	5.8	26
15	The oncogene AAMDC links PI3K-AKT-mTOR signaling with metabolic reprograming in estrogen receptor-positive breast cancer. Nature Communications, 2021, 12, 1920.	5.8	19
16	miR-101 suppresses the development of <i>MLL</i>-rearranged acute myeloid leukemia. Haematologica, 2019, 104, e296-e299.	1.7	14
17	The tumor suppressor miR-642a-5p targets Wilms Tumor 1 gene and cell-cycle progression in prostate cancer. Scientific Reports, 2021, 11, 18003.	1.6	10
18	Zfp281 (ZBP-99) plays a functionally redundant role with Zfp148 (ZBP-89) during erythroid development. Blood Advances, 2019, 3, 2499-2511.	2.5	7

#	Article	IF	Citations
19	Small nucleolar RNA networks are upâ€regulated during human anaphylaxis. Clinical and Experimental Allergy, 2021, 51, 1310-1321.	1.4	5
20	Identification of zfp148 (ZBP-89) as a Novel GATA-1 Associated Transcription Factor Involved in Megakaryopoiesis and Definitive Erythropoiesis.. Blood, 2005, 106, 828-828.	0.6	3
21	Abstract 2379: microRNA-7 replacement therapy: a promising approach for hepatocellular carcinoma. , 2021, , .		0
22	Role of the Kruì^ppel-Type Zinc Finger Transcription Factor ZBP-89 In Human Globin Gene Regulation and Erythroid Development. Blood, 2010, 116, 2067-2067.	0.6	0
23	Essential Role of the Transcription Factor ZBP-89 in Lymphopoiesis. Blood, 2012, 120, 277-277.	0.6	0