

Andrew J Woo

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

1,470
citations

567281

15
h-index

752698

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. <i>Cell</i> , 2010, 143, 313-324.	28.9	606
2	Direct Recruitment of Polycomb Repressive Complex 1 to Chromatin by Core Binding Transcription Factors. <i>Molecular Cell</i> , 2012, 45, 330-343.	9.7	188
3	Distinct and Combinatorial Functions of Jmjd2b/Kdm4b and Jmjd2c/Kdm4c in Mouse Embryonic Stem Cell Identity. <i>Molecular Cell</i> , 2014, 53, 32-48.	9.7	112
4	Differentiation-Dependent Interactions between RUNX-1 and FLI-1 during Megakaryocyte Development. <i>Molecular and Cellular Biology</i> , 2009, 29, 4103-4115.	2.3	71
5	Identification of ZBP-89 as a Novel GATA-1-Associated Transcription Factor Involved in Megakaryocytic and Erythroid Development. <i>Molecular and Cellular Biology</i> , 2008, 28, 2675-2689.	2.3	62
6	Targeting RSPO3-LGR4 Signaling for Leukemia Stem Cell Eradication in Acute Myeloid Leukemia. <i>Cancer Cell</i> , 2020, 38, 263-278.e6.	16.8	59
7	Tumor penetrating peptides inhibiting MYC as a potent targeted therapeutic strategy for triple-negative breast cancers. <i>Oncogene</i> , 2019, 38, 140-150.	5.9	55
8	Surfactant Protein C Chromatin-Bound Green Fluorescence Protein Reporter Mice Reveal Heterogeneity of Surfactant Protein C-Expressing Lung Cells. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2013, 48, 288-298.	2.9	54
9	A Src family kinase-Shp2 axis controls RUNX1 activity in megakaryocyte and T-lymphocyte differentiation. <i>Genes and Development</i> , 2012, 26, 1587-1601.	5.9	52
10	Developmental differences in IFN signaling affect GATA1s-induced megakaryocyte hyperproliferation. <i>Journal of Clinical Investigation</i> , 2013, 123, 3292-3304.	8.2	37
11	A Proteomics Approach for the Identification of DNA Binding Activities Observed in the Electrophoretic Mobility Shift Assay. <i>Molecular and Cellular Proteomics</i> , 2002, 1, 472-478.	3.8	33
12	JMJD1C-mediated metabolic dysregulation contributes to HOXA9-dependent leukemogenesis. <i>Leukemia</i> , 2019, 33, 1400-1410.	7.2	31
13	Role of ZBP-89 in human globin gene regulation and erythroid differentiation. <i>Blood</i> , 2011, 118, 3684-3693.	1.4	26
14	CpG island-mediated global gene regulatory modes in mouse embryonic stem cells. <i>Nature Communications</i> , 2014, 5, 5490.	12.8	26
15	The oncogene AAMDC links PI3K-AKT-mTOR signaling with metabolic reprogramming in estrogen receptor-positive breast cancer. <i>Nature Communications</i> , 2021, 12, 1920.	12.8	19
16	miR-101 suppresses the development of <i>MLL</i> -rearranged acute myeloid leukemia. <i>Haematologica</i> , 2019, 104, e296-e299.	3.5	14
17	The tumor suppressor miR-642a-5p targets Wilms Tumor 1 gene and cell-cycle progression in prostate cancer. <i>Scientific Reports</i> , 2021, 11, 18003.	3.3	10
18	Zfp281 (ZBP-99) plays a functionally redundant role with Zfp148 (ZBP-89) during erythroid development. <i>Blood Advances</i> , 2019, 3, 2499-2511.	5.2	7

#	ARTICLE	IF	CITATIONS
19	Small nucleolar RNA networks are upregulated during human anaphylaxis. <i>Clinical and Experimental Allergy</i> , 2021, 51, 1310-1321.	2.9	5
20	Identification of zfp148 (ZBP-89) as a Novel GATA-1 Associated Transcription Factor Involved in Megakaryopoiesis and Definitive Erythropoiesis.. <i>Blood</i> , 2005, 106, 828-828.	1.4	3
21	Abstract 2379: microRNA-7 replacement therapy: a promising approach for hepatocellular carcinoma. , 2021, , .		0
22	Role of the Krüppel-Type Zinc Finger Transcription Factor ZBP-89 In Human Globin Gene Regulation and Erythroid Development. <i>Blood</i> , 2010, 116, 2067-2067.	1.4	0
23	Essential Role of the Transcription Factor ZBP-89 in Lymphopoiesis. <i>Blood</i> , 2012, 120, 277-277.	1.4	0