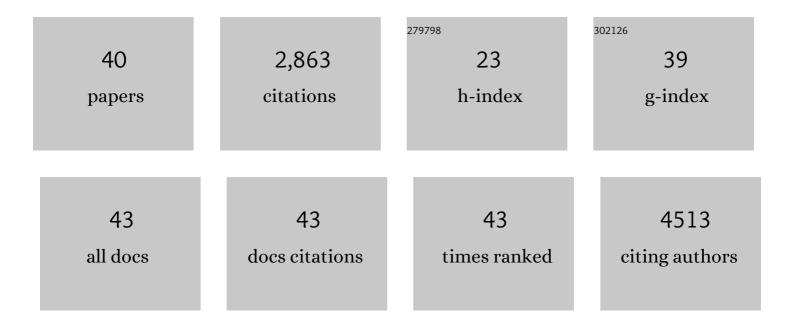
Jun R Yang

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Association of Genetic Ancestry With the Molecular Subtypes and Prognosis of Childhood Acute Lymphoblastic Leukemia. JAMA Oncology, 2022, 8, 354.	7.1	35
2	TERT Expression in Wilms Tumor Is Regulated by Promoter Mutation or Hypermethylation, WT1, and N-MYC. Cancers, 2022, 14, 1655.	3.7	3
3	Succinate dehydrogenase/complex II is critical for metabolic and epigenetic regulation of T cell proliferation and inflammation. Science Immunology, 2022, 7, eabm8161.	11.9	23
4	Targeting EP2 receptor with multifaceted mechanisms for high-risk neuroblastoma. Cell Reports, 2022, 39, 111000.	6.4	8
5	Recent Advances with KDM4 Inhibitors and Potential Applications. Journal of Medicinal Chemistry, 2022, 65, 9564-9579.	6.4	9
6	Targeting KDM4 for treating PAX3-FOXO1–driven alveolar rhabdomyosarcoma. Science Translational Medicine, 2022, 14, .	12.4	16
7	Association of <i>GATA3</i> Polymorphisms With Minimal Residual Disease and Relapse Risk in Childhood Acute Lymphoblastic Leukemia. Journal of the National Cancer Institute, 2021, 113, 408-417.	6.3	16
8	A protocol for high-throughput screening of histone lysine demethylase 4 inhibitors using TR-FRET assay. STAR Protocols, 2021, 2, 100702.	1.2	1
9	17-DMAG dually inhibits Hsp90 and histone lysine demethylases in alveolar rhabdomyosarcoma. IScience, 2021, 24, 101996.	4.1	7
10	Targeting the spliceosome through RBM39 degradation results in exceptional responses in high-risk neuroblastoma models. Science Advances, 2021, 7, eabj5405.	10.3	32
11	KDM6B promotes activation of the oncogenic CDK4/6-pRB-E2F pathway by maintaining enhancer activity in MYCN-amplified neuroblastoma. Nature Communications, 2021, 12, 7204.	12.8	22
12	MYCN drives glutaminolysis in neuroblastoma and confers sensitivity to an ROS augmenting agent. Cell Death and Disease, 2018, 9, 220.	6.3	46
13	KDM5A Regulates a Translational Program that Controls p53 Protein Expression. IScience, 2018, 9, 84-100.	4.1	25
14	Hypoxia and Hormone-Mediated Pathways Converge at the Histone Demethylase KDM4B in Cancer. International Journal of Molecular Sciences, 2018, 19, 240.	4.1	29
15	Systematic identification of hepatitis E virus ORF2 interactome reveals that TMEM134 engages in ORF2-mediated NF-κB pathway. Virus Research, 2017, 228, 102-108.	2.2	17
16	Targeting Histone Demethylases in MYC-Driven Neuroblastomas with Ciclopirox. Cancer Research, 2017, 77, 4626-4638.	0.9	42
17	RIG-I and IL-6 are negative-feedback regulators of STING induced by double-stranded DNA. PLoS ONE, 2017, 12, e0182961.	2.5	25
18	Histone demethylases and their roles in cancer epigenetics. , 2016, 1, 34-40.		47

Histone demethylases and their roles in cancer epigenetics. , 2016, 1, 34-40.

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19	A genome-wide association study of susceptibility to acute lymphoblastic leukemia in adolescents and young adults. Blood, 2015, 125, 680-686.	1.4	110
20	Estrogen receptor-α directly regulates the hypoxia-inducible factor 1 pathway associated with antiestrogen response in breast cancer. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 15172-15177.	7.1	110
21	Inherited coding variants at the CDKN2A locus influence susceptibility to acute lymphoblastic leukaemia in children. Nature Communications, 2015, 6, 7553.	12.8	72
22	The Role of Histone Demethylase KDM4B in Myc Signaling in Neuroblastoma. Journal of the National Cancer Institute, 2015, 107, djv080.	6.3	63
23	Inherited <i>NUDT15</i> Variant Is a Genetic Determinant of Mercaptopurine Intolerance in Children With Acute Lymphoblastic Leukemia. Journal of Clinical Oncology, 2015, 33, 1235-1242.	1.6	369
24	Germline genetic variation in ETV6 and risk of childhood acute lymphoblastic leukaemia: a systematic genetic study. Lancet Oncology, The, 2015, 16, 1659-1666.	10.7	161
25	Hepatitis E virus open reading frame 3 protein interacts with porcine liverâ€specific plasminogen and α2â€antiplasmin. Journal of Medical Virology, 2014, 86, 487-495.	5.0	3
26	Seven In Absentia Homolog 2 (SIAH2) downregulation is associated with tamoxifen resistance in MCF-7 breast cancer cells. Journal of Surgical Research, 2014, 190, 203-209.	1.6	12
27	Inherited GATA3 variants are associated with Ph-like childhood acute lymphoblastic leukemia and risk of relapse. Nature Genetics, 2013, 45, 1494-1498.	21.4	264
28	Novel Susceptibility Variants at 10p12.31-12.2 for Childhood Acute Lymphoblastic Leukemia in Ethnically Diverse Populations. Journal of the National Cancer Institute, 2013, 105, 733-742.	6.3	208
29	Virus Host Protein Interaction Network Analysis Reveals That the HEV ORF3 Protein May Interrupt the Blood Coagulation Process. PLoS ONE, 2013, 8, e56320.	2.5	27
30	<i>ARID5B</i> Genetic Polymorphisms Contribute to Racial Disparities in the Incidence and Treatment Outcome of Childhood Acute Lymphoblastic Leukemia. Journal of Clinical Oncology, 2012, 30, 751-757.	1.6	165
31	Genome-wide association study identifies germline polymorphisms associated with relapse of childhood acute lymphoblastic leukemia. Blood, 2012, 120, 4197-4204.	1.4	103
32	Human CHCHD4 mitochondrial proteins regulate cellular oxygen consumption rate and metabolism and provide a critical role in hypoxia signaling and tumor progression. Journal of Clinical Investigation, 2012, 122, 600-611.	8.2	82
33	The Histone Demethylase JMJD2B Is Regulated by Estrogen Receptor α and Hypoxia, and Is a Key Mediator of Estrogen Induced Growth. Cancer Research, 2010, 70, 6456-6466.	0.9	167
34	Small-Molecule Activation of p53 Blocks Hypoxia-Inducible Factor 1α and Vascular Endothelial Growth Factor Expression In Vivo and Leads to Tumor Cell Apoptosis in Normoxia and Hypoxia. Molecular and Cellular Biology, 2009, 29, 2243-2253.	2.3	89
35	Genome-wide Interrogation of Germline Genetic Variation Associated With Treatment Response in Childhood Acute Lymphoblastic Leukemia. JAMA - Journal of the American Medical Association, 2009, 301, 393.	7.4	193
36	Activation of a unique p53-dependent DNA damage response. Cell Cycle, 2009, 8, 1630-1632.	2.6	13

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37	Role of Hypoxiaâ€Inducible Factors in Epigenetic Regulation via Histone Demethylases. Annals of the New York Academy of Sciences, 2009, 1177, 185-197.	3.8	98
38	Crystal Structure of the Three Tandem FF Domains of the Transcription Elongation Regulator CA150. Journal of Molecular Biology, 2009, 393, 397-408.	4.2	11
39	Regulation of cell-cell interactions by phosphatidic acid phosphatase 2b/VCIP. EMBO Journal, 2003, 22, 1539-1554.	7.8	63
40	Analysis of VEGF-responsive genes involved in the activation of endothelial cells. Molecular Cancer, 2003, 2, 25.	19.2	76