

Kirby D Johnson

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

25
papers

913
citations

16
h-index

30
g-index

30
ext. papers

1,100
ext. citations

8.8
avg, IF

3.67
L-index

#	Paper	IF	Citations
25	Spacing Constraints of Neighboring Zinc Finger Modules within GATA2. <i>Blood</i> , 2021 , 138, 3306-3306	2.2	
24	Gata2 -77 enhancer regulates adult hematopoietic stem cell survival. <i>Leukemia</i> , 2021 , 35, 901-905	10.7	0
23	Conditionally pathogenic genetic variants of a hematopoietic disease-suppressing enhancer. <i>Science Advances</i> , 2021 , 7, eabk3521	14.3	0
22	GATA2 Enhancer Modules Governing Hematopoietic Regeneration. <i>Blood</i> , 2020 , 136, 15-16	2.2	
21	Constructing and deconstructing GATA2-regulated cell fate programs to establish developmental trajectories. <i>Journal of Experimental Medicine</i> , 2020 , 217,	16.6	8
20	Single-nucleotide human disease mutation inactivates a blood-regenerative GATA2 enhancer. <i>Journal of Clinical Investigation</i> , 2019 , 129, 1180-1192	15.9	19
19	Blood disease-causing and -suppressing transcriptional enhancers: general principles and mechanisms. <i>Blood Advances</i> , 2019 , 3, 2045-2056	7.8	10
18	Mechanisms of erythrocyte development and regeneration: implications for regenerative medicine and beyond. <i>Development (Cambridge)</i> , 2018 , 145,	6.6	36
17	Human leukemia mutations corrupt but do not abrogate GATA-2 function. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018 , 115, E10109-E10118	11.5	21
16	p53 ^{-/-} synergizes with enhanced NrasG12D signaling to transform megakaryocyte-erythroid progenitors in acute myeloid leukemia. <i>Blood</i> , 2017 , 129, 358-370	2.2	17
15	Integrating Enhancer Mechanisms to Establish a Hierarchical Blood Development Program. <i>Cell Reports</i> , 2017 , 20, 2966-2979	10.6	29
14	GATA Factor-G-Protein-Coupled Receptor Circuit Suppresses Hematopoiesis. <i>Stem Cell Reports</i> , 2016 , 6, 368-82	8	8
13	Hematopoietic Signaling Mechanism Revealed from a Stem/Progenitor Cell Cistrome. <i>Molecular Cell</i> , 2015 , 59, 62-74	17.6	29
12	Cis-regulatory mechanisms governing stem and progenitor cell transitions. <i>Science Advances</i> , 2015 , 1, e1500503	14.3	34
11	Mechanism governing a stem cell-generating cis-regulatory element. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, E1091-100	11.5	34
10	Hematopoietic transcriptional mechanisms: from locus-specific to genome-wide vantage points. <i>Experimental Hematology</i> , 2014 , 42, 618-29	3.1	19
9	Gata2 cis-element is required for hematopoietic stem cell generation in the mammalian embryo. <i>Journal of Experimental Medicine</i> , 2013 , 210, 2833-42	16.6	101

8	GATA2 haploinsufficiency caused by mutations in a conserved intronic element leads to MonoMAC syndrome. <i>Blood</i> , 2013 , 121, 3830-7, S1-7	2.2	169
7	Cis-element mutated in GATA2-dependent immunodeficiency governs hematopoiesis and vascular integrity. <i>Journal of Clinical Investigation</i> , 2012 , 122, 3692-704	15.9	127
6	Context-dependent function of "GATA switch" sites in vivo. <i>Blood</i> , 2011 , 117, 4769-72	2.2	45
5	A Master Regulatory Cis-element Governs the Hematopoietic Stem/Progenitor Cell Compartment, Vascular Integrity, and Cardiovascular Development. <i>Blood</i> , 2011 , 118, 1304-1304	2.2	
4	Controlling Hematopoiesis through Sumoylation-Dependent Regulation of a GATA Factor.. <i>Blood</i> , 2009 , 114, 1467-1467	2.2	
3	Friend of GATA-1-independent transcriptional repression: a novel mode of GATA-1 function. <i>Blood</i> , 2007 , 109, 5230-3	2.2	54
2	Differential sensitivities of transcription factor target genes underlie cell type-specific gene expression profiles. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006 , 103, 15939-44	11.5	43
1	Chromatin domain activation via GATA-1 utilization of a small subset of dispersed GATA motifs within a broad chromosomal region. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 17065-70	11.5	109