Chuanfeng Wu

List of Publications by Year in descending order

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758635 794141 22 706 12 19 h-index citations g-index papers 24 24 24 1261 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Clonal tracking of haematopoietic cells: insights and clinical implications. British Journal of Haematology, 2021, 192, 819-831.	1.2	10
2	Comparative engraftment and clonality of macaque HSPCs expanded on human umbilical vein endothelial cells versus non-expanded cells. Molecular Therapy - Methods and Clinical Development, 2021, 20, 703-715.	1.8	1
3	Interrogation of clonal tracking data using barcodetrackR. Nature Computational Science, 2021, 1, 280-289.	3.8	13
4	Tissue Trafficking Kinetics of Rhesus Macaque Natural Killer Cells Measured by Serial Intravascular Staining. Frontiers in Immunology, 2021, 12, 772332.	2.2	2
5	Clonal tracking of erythropoiesis in rhesus macaques. Haematologica, 2020, 105, 1813-1824.	1.7	5
6	Aberrant Clonal Hematopoiesis following Lentiviral Vector Transduction of HSPCs in a Rhesus Macaque. Molecular Therapy, 2019, 27, 1074-1086.	3.7	34
7	Impact of CMV Infection on Natural Killer Cell Clonal Repertoire in CMV-NaÃ-ve Rhesus Macaques. Frontiers in Immunology, 2019, 10, 2381.	2.2	16
8	The impact of aging on primate hematopoiesis as interrogated by clonal tracking. Blood, 2018, 131, 1195-1205.	0.6	39
9	Geographic clonal tracking in macaques provides insights into HSPC migration and differentiation. Journal of Experimental Medicine, 2018, 215, 217-232.	4.2	32
10	Barcoding of Macaque Hematopoietic Stem and Progenitor Cells: A Robust Platform to Assess Vector Genotoxicity. Molecular Therapy - Methods and Clinical Development, 2018, 11, 143-154.	1.8	9
11	Clonal expansion and compartmentalized maintenance of rhesus macaque NK cell subsets. Science Immunology, 2018, 3, .	5.6	41
12	Quantitative stability of hematopoietic stem and progenitor cell clonal output in rhesus macaques receiving transplants. Blood, 2017, 129, 1448-1457.	0.6	53
13	Acquired somatic mutations in PNH reveal long-term maintenance of adaptive NK cells independent of HSPCs. Blood, 2017, 129, 1940-1946.	0.6	42
14	Stochastic Modeling of Hematopoietic Stem and Progenitor Cell Barcoding Data from Rhesus Macaques Challenges the Classic Model of Hematopoiesis. Blood, 2016, 128, 2643-2643.	0.6	0
15	Path to the Clinic: Assessment of iPSC-Based Cell Therapies InÂVivo in a Nonhuman Primate Model. Cell Reports, 2014, 7, 1298-1309.	2.9	84
16	Clonal Tracking of Rhesus Macaque Hematopoiesis Highlights a Distinct Lineage Origin for Natural Killer Cells. Cell Stem Cell, 2014, 14, 486-499.	5.2	149
17	Development of an inducible caspase-9 safety switch for pluripotent stem cell–based therapies. Molecular Therapy - Methods and Clinical Development, 2014, 1, 14053.	1.8	59
18	High Efficiency Restriction Enzyme–Free Linear Amplification-Mediated Polymerase Chain Reaction Approach for Tracking Lentiviral Integration Sites Does Not Abrogate Retrieval Bias. Human Gene Therapy, 2013, 24, 38-47.	1.4	24

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19	The Unique Ontogeny Of Natural Killer Cells As Revealed By Genetic Barcoding In The Nonhuman Primate Model. Blood, 2013, 122, 15-15.	0.6	2
20	Rhesus Macaque NK Cells Expanded Ex Vivo Undergo Similar Phenotypic and Functional Changes Observed With Expanded Human NK Cells Providing An Excellent Model To Optimize Adoptive NK Cell Transfer. Blood, 2013, 122, 2028-2028.	0.6	1
21	Stem cell gene therapy: the risks of insertional mutagenesis and approaches to minimize genotoxicity. Frontiers of Medicine, 2011, 5, 356-371.	1.5	90
22	Telomere Dynamics in Pluripotent Stem Cells Derived From Patients with Telomere Diseases. Blood, 2011, 118, 51-51.	0.6	0