Alba Rodriguez-Meira

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2217629/publications.pdf

Version: 2024-02-01

14 papers

871 citations

1040056 9 h-index 1199594 12 g-index

20 all docs

20 docs citations

times ranked

20

1465 citing authors

#	Article	IF	Citations
1	Transitions in lineage specification and gene regulatory networks in hematopoietic stem/progenitor cells over human development. Cell Reports, 2021, 36, 109698.	6.4	38
2	Heterogeneous disease-propagating stem cells in juvenile myelomonocytic leukemia. Journal of Experimental Medicine, $2021, 218, \ldots$	8.5	25
3	Single-Cell Multi-Omics Reveals the Genetic, Cellular and Molecular Landscape of <i>TP53</i> Mutated Leukemic Transformation in MPN. Blood, 2021, 138, 3-3.	1.4	7
4	Ezh2 is essential for the generation of functional yolk sac derived erythro-myeloid progenitors. Nature Communications, 2021, 12, 7019.	12.8	8
5	TARGET-Seq: A Protocol for High-Sensitivity Single-Cell Mutational Analysis and Parallel RNA Sequencing. STAR Protocols, 2020, 1, 100125.	1.2	27
6	Rapid Emergence of Chronic Lymphocytic Leukemia During JAK2 Inhibitor Therapy in a Patient With Myelofibrosis. HemaSphere, 2020, 4, e356.	2.7	4
7	Single-Cell Analyses Reveal Megakaryocyte-Biased Hematopoiesis in Myelofibrosis and Identify Mutant Clone-Specific Targets. Molecular Cell, 2020, 78, 477-492.e8.	9.7	106
8	C/EBPα and GATA-2 Mutations Induce Bilineage Acute Erythroid Leukemia through Transformation of a Neomorphic Neutrophil-Erythroid Progenitor. Cancer Cell, 2020, 37, 690-704.e8.	16.8	16
9	Molecular Characterisation of Participants in the Phazar Trial Reveals Prognostic Impact of Mutations in Advanced-Phase-MPN. Blood, 2020, 136, 40-41.	1.4	O
10	Unravelling Intratumoral Heterogeneity through High-Sensitivity Single-Cell Mutational Analysis and Parallel RNA Sequencing. Molecular Cell, 2019, 73, 1292-1305.e8.	9.7	218
11	The Cellular Pathway of Leukemic Transformation in MDS: It's the Stem Cells, Stupid!. , 2019, 16, .		O
12	Single cell analysis of normal and leukemic hematopoiesis. Molecular Aspects of Medicine, 2018, 59, 85-94.	6.4	53
13	Single-cell transcriptomics uncovers distinct molecular signatures of stem cells in chronic myeloid leukemia. Nature Medicine, 2017, 23, 692-702.	30.7	336
14	Tumoral stem cell reprogramming as a driver of cancer: Theory, biological models, implications in cancer therapy. Seminars in Cancer Biology, 2015, 32, 3-9.	9.6	22