

Delphine Potier

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

1,901
citations

687363

13
h-index

642732

23
g-index

24
all docs

24
docs citations

24
times ranked

3965
citing authors

#	ARTICLE	IF	CITATIONS
1	Multiplexed single-cell RNA-sequencing of mouse thymic and splenic samples. STAR Protocols, 2022, 3, 101041.	1.2	2
2	Calcium Signaling Is Impaired in PTEN-Deficient T Cell Acute Lymphoblastic Leukemia. Frontiers in Immunology, 2022, 13, 797244.	4.8	4
3	Single-cell transcriptomics uncovers an instructive T α cell receptor role in adult β T α cell lineage commitment. EMBO Journal, 2022, 41, e110023.	7.8	7
4	Single-cell profiling identifies pre-existing CD19-negative subclones in a B-ALL patient with CD19-negative relapse after CAR-T therapy. Nature Communications, 2021, 12, 865.	12.8	81
5	MYC deficiency impairs the development of effector/memory T lymphocytes. iScience, 2021, 24, 102761.	4.1	10
6	GATA1 pathogenic variants disrupt MYH10 silencing during megakaryopoiesis. Journal of Thrombosis and Haemostasis, 2021, 19, 2287-2301.	3.8	6
7	Cortical Neurogenesis Requires Bcl6-Mediated Transcriptional Repression of Multiple Self-Renewal-Promoting Extrinsic Pathways. Neuron, 2019, 103, 1096-1108.e4.	8.1	38
8	Altering the Temporal Regulation of One Transcription Factor Drives Evolutionary Trade-Offs between Head Sensory Organs. Developmental Cell, 2019, 50, 780-792.e7.	7.0	34
9	Fit β T-cell receptor suppresses leukemogenesis of Pten-deficient thymocytes. Haematologica, 2018, 103, 999-1007.	3.5	6
10	The transcription factor Grainy head primes epithelial enhancers for spatiotemporal activation by displacing nucleosomes. Nature Genetics, 2018, 50, 1011-1020.	21.4	122
11	Desynchronization of the Germinal Center Dynamics and Remodeling of the Tumor Microenvironment Characterize KMT2D-Driven Lymphomagenesis. Blood, 2018, 132, 670-670.	1.4	8
12	Nuclear receptors connect progenitor transcription factors to cell cycle control. Scientific Reports, 2017, 7, 4845.	3.3	17
13	dachshund Potentiates Hedgehog Signaling during Drosophila Retinogenesis. PLoS Genetics, 2016, 12, e1006204.	3.5	15
14	An Ectopic Network of Transcription Factors Regulated by Hippo Signaling Drives Growth and Invasion of a Malignant Tumor Model. Current Biology, 2016, 26, 2101-2113.	3.9	87
15	Identification of Lineage-Specific Cis-Regulatory Modules Associated with Variation in Transcription Factor Binding and Chromatin Activity Using Ornstein-Uhlenbeck Models. Molecular Biology and Evolution, 2015, 32, 2441-2455.	8.9	11
16	Discovery of Transcription Factors and Regulatory Regions Driving In Vivo Tumor Development by ATAC-seq and FAIRE-seq Open Chromatin Profiling. PLoS Genetics, 2015, 11, e1004994.	3.5	155
17	i-cisTarget 2015 update: generalized cis-regulatory enrichment analysis in human, mouse and fly. Nucleic Acids Research, 2015, 43, W57-W64.	14.5	169
18	Mapping Gene Regulatory Networks in Drosophila Eye Development by Large-Scale Transcriptome Perturbations and Motif Inference. Cell Reports, 2014, 9, 2290-2303.	6.4	85

#	ARTICLE	IF	CITATIONS
19	iRegulon: From a Gene List to a Gene Regulatory Network Using Large Motif and Track Collections. PLoS Computational Biology, 2014, 10, e1003731.	3.2	787
20	Identification of cis-regulatory modules encoding temporal dynamics during development. BMC Genomics, 2014, 15, 534.	2.8	10
21	Comparative motif discovery combined with comparative transcriptomics yields accurate targetome and enhancer predictions. Genome Research, 2013, 23, 74-88.	5.5	54
22	i-cisTarget: an integrative genomics method for the prediction of regulatory features and cis-regulatory modules. Nucleic Acids Research, 2012, 40, e114-e114.	14.5	176
23	Using cisTargetX to Predict Transcriptional Targets and Networks in Drosophila. Methods in Molecular Biology, 2012, 786, 291-314.	0.9	17