## **Christophe Battail**

List of Publications by Year in descending order

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CHRISTORHE RATTAIL

#	Article	IF	CITATIONS
1	Different cardiovascular and pulmonary phenotypes for single- and double-knock-out mice deficient in BMP9 and BMP10. Cardiovascular Research, 2022, 118, 1805-1820.	3.8	26
2	RNA sequencing of chronic GVHD skin lesions defines shared and unique inflammatory pathways characterizing lichen planus and morphea. Blood Advances, 2022, 6, 2805-2811.	5.2	6
3	Temporal Gene Expression Profiles Reflect the Dynamics of Lymphoid Differentiation. International Journal of Molecular Sciences, 2022, 23, 1115.	4.1	0
4	Cooperative Blockade of CK2 and ATM Kinases Drives Apoptosis in VHL-Deficient Renal Carcinoma Cells through ROS Overproduction. Cancers, 2021, 13, 576.	3.7	19
5	Region-specific expression of young small-scale duplications in the human central nervous system. Bmc Ecology and Evolution, 2021, 21, 59.	1.6	4
6	NLRP7 Promotes Choriocarcinoma Growth and Progression through the Establishment of an Immunosuppressive Microenvironment. Cancers, 2021, 13, 2999.	3.7	16
7	Transcriptomic Characterization of Postmolar Gestational Choriocarcinoma. Biomedicines, 2021, 9, 1474.	3.2	4
8	Battle of the Sex Chromosomes: Competition between X and Y Chromosome-Encoded Proteins for Partner Interaction and Chromatin Occupancy Drives Multicopy Gene Expression and Evolution in Muroid Rodents. Molecular Biology and Evolution, 2020, 37, 3453-3468.	8.9	25
9	Gene expression and response prediction to amisulpride in the OPTiMiSE first episode psychoses. Neuropsychopharmacology, 2020, 45, 1637-1644.	5.4	5
10	Multi-omic analysis of gametogenesis reveals a novel signature at the promoters and distal enhancers of active genes. Nucleic Acids Research, 2020, 48, 4115-4138.	14.5	24
11	Systematic analysis of TruSeq, SMARTer and SMARTer Ultra-Low RNA-seq kits for standard, low and ultra-low quantity samples. Scientific Reports, 2019, 9, 7550.	3.3	22
12	Disentangling the Causes for Faster-X Evolution in Aphids. Genome Biology and Evolution, 2018, 10, 507-520.	2.5	42
13	Gene flow contributes to diversification of the major fungal pathogen Candida albicans. Nature Communications, 2018, 9, 2253.	12.8	131
14	Immortalized human myotonic dystrophy muscle cell lines to assess therapeutic compounds. DMM Disease Models and Mechanisms, 2017, 10, 487-497.	2.4	65
15	Bdf1 Bromodomains Are Essential for Meiosis and the Expression of Meiotic-Specific Genes. PLoS Genetics, 2017, 13, e1006541.	3.5	13
16	The Arabidopsis hnRNP-Q Protein LIF2 and the PRC1 Subunit LHP1 Function in Concert to Regulate the Transcription of Stress-Responsive Genes. Plant Cell, 2016, 28, 2197-2211.	6.6	37
17	In Vitro and In Vivo Modulation of Alternative Splicing by the Biguanide Metformin. Molecular Therapy - Nucleic Acids, 2015, 4, e262.	5.1	63
18	The Spatiotemporal Program of DNA Replication Is Associated with Specific Combinations of Chromatin Marks in Human Cells. PLoS Genetics, 2014, 10, e1004282.	3.5	123

#	Article	IF	CITATIONS
19	Early allopolyploid evolution in the post-Neolithic <i>Brassica napus</i> oilseed genome. Science, 2014, 345, 950-953.	12.6	2,089
20	The Innate Immune Database (IIDB). BMC Immunology, 2008, 9, 7.	2.2	45