## Marine Jeanmougin

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/593116/publications.pdf

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933447 1125743 14 1,448 10 13 citations g-index h-index papers 14 14 14 3372 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	A comprehensive evaluation of normalization methods for Illumina high-throughput RNA sequencing data analysis. Briefings in Bioinformatics, 2013, 14, 671-683.	6.5	1,064
2	Should We Abandon the t-Test in the Analysis of Gene Expression Microarray Data: A Comparison of Variance Modeling Strategies. PLoS ONE, 2010, 5, e12336.	2.5	120
3	Human blood BDCA-1 dendritic cells differentiate into Langerhans-like cells with thymic stromal lymphopoietin and TGF-β. Blood, 2014, 124, 2411-2420.	1.4	72
4	Breast Cancer Cell–Derived GM-CSF Licenses Regulatory Th2 Induction by Plasmacytoid Predendritic Cells in Aggressive Disease Subtypes. Cancer Research, 2015, 75, 2775-2787.	0.9	49
5	CpG island methylator phenotype identifies high risk patients among microsatellite stable <i>BRAF</i> mutated colorectal cancers. International Journal of Cancer, 2017, 141, 967-976.	5.1	40
6	A robust internal control for high-precision DNA methylation analyses by droplet digital PCR. Clinical Epigenetics, 2018, 10, 24.	4.1	26
7	Multiple Testing in Large-Scale Genetic Studies. Methods in Molecular Biology, 2012, 888, 213-233.	0.9	16
8	Experimental factors affecting the robustness of DNA methylation analysis. Scientific Reports, 2016, 6, 33936.	3.3	15
9	Improved prognostication of glioblastoma beyond molecular subtyping by transcriptional profiling of the tumor microenvironment. Molecular Oncology, 2020, 14, 1016-1027.	4.6	15
10	Early and accurate detection of cholangiocarcinoma in patients with primary sclerosing cholangitis by methylation markers in bile. Hepatology, 2022, 75, 59-73.	7.3	15
11	Spatial analysis and CD25-expression identify regulatory T cells as predictors of a poor prognosis in colorectal cancer. Modern Pathology, 2022, 35, 1236-1246.	5.5	8
12	Prognostic relevance of an epigenetic biomarker panel in sentinel lymph nodes from colon cancer patients. Clinical Epigenetics, 2017, 9, 97.	4.1	4
13	Multiregional assessment of CIMP in primary colorectal cancers: Phenotype concordance but marker variability. International Journal of Cancer, 2021, 148, 1652-1657.	5.1	4
14	Network Inference in Breast Cancer with Gaussian Graphical Models and Extensions. , 2014, , 121-146.		0