

Tiphaine C Martin

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

2,669
citations

430874

18
h-index

552781

26
g-index

33
all docs

33
docs citations

33
times ranked

5979
citing authors

#	ARTICLE	IF	CITATIONS
1	Machine Learning Approaches for Early Prostate Cancer Prediction Based on Healthcare Utilization Patterns. <i>Studies in Health Technology and Informatics</i> , 2022, 289, 65-68.	0.3	1
2	Altered BAF occupancy and transcription factor dynamics in PBAF-deficient melanoma. <i>Cell Reports</i> , 2022, 39, 110637.	6.4	12
3	NOTCH and EZH2 collaborate to repress PTEN expression in breast cancer. <i>Communications Biology</i> , 2021, 4, 312.	4.4	16
4	AKT Degradation Selectively Inhibits the Growth of PI3K/PTEN Pathway Mutant Cancers with Wild-Type KRAS and BRAF by Destabilizing Aurora Kinase B. <i>Cancer Discovery</i> , 2021, 11, 3064-3089.	9.4	32
5	Altered N-glycan profile of IgG-depleted serum proteins in Hashimoto's thyroiditis. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2020, 1864, 129464.	2.4	8
6	Discovery of a first-in-class EZH2 selective degrader. <i>Nature Chemical Biology</i> , 2020, 16, 214-222.	8.0	148
7	Cooperation Between Distinct Cancer Driver Genes Underlies Intertumor Heterogeneity in Hepatocellular Carcinoma. <i>Gastroenterology</i> , 2020, 159, 2203-2220.e14.	1.3	47
8	Dysregulated Antibody, Natural Killer Cell and Immune Mediator Profiles in Autoimmune Thyroid Diseases. <i>Cells</i> , 2020, 9, 665.	4.1	18
9	Decreased Immunoglobulin G Core Fucosylation, A Player in Antibody-dependent Cell-mediated Cytotoxicity, is Associated with Autoimmune Thyroid Diseases. <i>Molecular and Cellular Proteomics</i> , 2020, 19, 774-792.	3.8	32
10	Limited Mitochondrial Activity Coupled With Strong Expression of CD34, CD90 and EPCR Determines the Functional Fitness of ex vivo Expanded Human Hematopoietic Stem Cells. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 592348.	3.7	8
11	Interplay between the human gut microbiome and host metabolism. <i>Nature Communications</i> , 2019, 10, 4505.	12.8	450
12	An integrative cross-omics analysis of DNA methylation sites of glucose and insulin homeostasis. <i>Nature Communications</i> , 2019, 10, 2581.	12.8	62
13	Smoking induces coordinated DNA methylation and gene expression changes in adipose tissue with consequences for metabolic health. <i>Clinical Epigenetics</i> , 2018, 10, 126.	4.1	110
14	Conducting metagenomic studies in microbiology and clinical research. <i>Applied Microbiology and Biotechnology</i> , 2018, 102, 8629-8646.	3.6	26
15	Gut microbiota associations with common diseases and prescription medications in a population-based cohort. <i>Nature Communications</i> , 2018, 9, 2655.	12.8	411
16	YAMP: a containerized workflow enabling reproducibility in metagenomics research. <i>GigaScience</i> , 2018, 7, .	6.4	22
17	Hippurate as a metabolomic marker of gut microbiome diversity: Modulation by diet and relationship to metabolic syndrome. <i>Scientific Reports</i> , 2017, 7, 13670.	3.3	193
18	Metabolites of milk intake: a metabolomic approach in UK twins with findings replicated in two European cohorts. <i>European Journal of Nutrition</i> , 2017, 56, 2379-2391.	3.9	24

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19	coMET: visualisation of regional epigenome-wide association scan results and DNA co-methylation patterns. BMC Bioinformatics, 2015, 16, 131.	2.6	104
20	Twin Studies and Epigenetics. , 2015, , 683-702.		3
21	Transcriptional diversity during lineage commitment of human blood progenitors. Science, 2014, 345, 1251033.	12.6	253
22	The complete genome of Blastobotrys (Arxula) adenivorans LS3 - a yeast of biotechnological interest. Biotechnology for Biofuels, 2014, 7, 66.	6.2	57
23	Comparative genomics of emerging pathogens in the Candida glabrata clade. BMC Genomics, 2013, 14, 623.	2.8	174
24	<i>Pichia sorbitophila</i> , an Interspecies Yeast Hybrid, Reveals Early Steps of Genome Resolution After Polyploidization. G3: Genes, Genomes, Genetics, 2012, 2, 299-311.	1.8	113
25	The GÃ©nolevures database. Comptes Rendus - Biologies, 2011, 334, 585-589.	0.2	9
26	GÃ©nolevures: protein families and synteny among complete hemiascomycetous yeast proteomes and genomes. Nucleic Acids Research, 2009, 37, D550-D554.	14.5	117
27	Comparative genomics of protoploid <i>Saccharomycetaceae</i> . Genome Research, 2009, 19, 1696-1709.	5.5	207