## Fabio Parisi

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11933818/publications.pdf

Version: 2024-02-01

25 2,325 17
papers citations h-index

17 25
h-index g-index

26 26 all docs citations

26 times ranked 4905 citing authors

#	Article	IF	CITATIONS
1	Impact of healthcare worker shift scheduling on workforce preservation during the COVID-19 pandemic. Infection Control and Hospital Epidemiology, 2020, 41, 1443-1445.	1.8	34
2	MET Expression in Primary and Metastatic Clear Cell Renal Cell Carcinoma: Implications of Correlative Biomarker Assessment to MET Pathway Inhibitors. BioMed Research International, 2015, 2015, 1-7.	1.9	6
3	Regulation of Glutamine Carrier Proteins by RNF5 Determines Breast Cancer Response to ER Stress-Inducing Chemotherapies. Cancer Cell, 2015, 27, 354-369.	16.8	177
4	PLEKHA5 as a Biomarker and Potential Mediator of Melanoma Brain Metastasis. Clinical Cancer Research, 2015, 21, 2138-2147.	7.0	71
5	Preanalytical variables and phosphoepitope expression in FFPE tissue: quantitative epitope assessment after variable cold ischemic time. Laboratory Investigation, 2015, 95, 334-341.	3.7	52
6	Ranking and combining multiple predictors without labeled data. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 1253-1258.	7.1	74
7	A tissue quality index: an intrinsic control for measurement of effects of preanalytical variables on FFPE tissue. Laboratory Investigation, 2014, 94, 467-474.	3.7	48
8	Oct-1 Regulates IL-17 Expression by Directing Interchromosomal Associations in Conjunction with CTCF in T Cells. Molecular Cell, 2014, 54, 56-66.	9.7	44
9	NY-ESO-1 as a potential immunotherapeutic target in renal cell carcinoma. Oncotarget, 2014, 5, 5209-5217.	1.8	3
10	Interactions with RNA direct the Polycomb group protein SCML2 to chromatin where it represses target genes. ELife, 2014, 3, e02637.	6.0	46
11	Expression of drug targets in primary and matched metastatic renal cell carcinoma tumors. BMC Clinical Pathology, 2013, 13, 3.	1.8	15
12	Arpeggio: harmonic compression of ChIP-seq data reveals protein-chromatin interaction signatures. Nucleic Acids Research, 2013, 41, e161-e161.	14.5	7
13	TrAp: a tree approach for fingerprinting subclonal tumor composition. Nucleic Acids Research, 2013, 41, e165-e165.	14.5	113
14	Quantitative Assessment of Effect of Preanalytic Cold Ischemic Time on Protein Expression in Breast Cancer Tissues. Journal of the National Cancer Institute, 2012, 104, 1815-1824.	6.3	103
15	Picking ChIP-seq peak detectors for analyzing chromatin modification experiments. Nucleic Acids Research, 2012, 40, e70-e70.	14.5	64
16	PCGF Homologs, CBX Proteins, and RYBP Define Functionally Distinct PRC1 Family Complexes. Molecular Cell, 2012, 45, 344-356.	9.7	741
17	Integrated analysis of tumor samples sheds light on tumor heterogeneity. Yale Journal of Biology and Medicine, 2012, 85, 347-61.	0.2	14
18	Detecting copy number status and uncovering subclonal markers in heterogeneous tumor biopsies. BMC Genomics, 2011, 12, 230.	2.8	27

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19	Genome-wide remodeling of the epigenetic landscape during myogenic differentiation. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, E149-58.	7.1	254
20	VDA, a Method of Choosing a Better Algorithm with Fewer Validations. PLoS ONE, 2011, 6, e26074.	2.5	2
21	The Mammalian Sin3 Proteins Are Required for Muscle Development and Sarcomere Specification. Molecular and Cellular Biology, 2010, 30, 5686-5697.	2.3	59
22	Benefits of biomarker selection and clinico-pathological covariate inclusion in breast cancer prognostic models. Breast Cancer Research, 2010, 12, R66.	5.0	15
23	PLX4032, a selective BRAF <sup>V600E</sup> kinase inhibitor, activates the ERK pathway and enhances cell migration and proliferation of BRAF <sup>WT</sup> melanoma cells. Pigment Cell and Melanoma Research, 2010, 23, 190-200.	3.3	315
24	Relationship between estrogen receptor $\hat{l}_{\pm}$ location and gene induction reveals the importance of downstream sites and cofactors. BMC Genomics, 2009, 10, 381.	2.8	8
25	Identifying synergistic regulation involving c-Myc and sp1 in human tissues. Nucleic Acids Research, 2007, 35, 1098-1107.	14.5	33