

Giulia Franciosa

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/148022/publications.pdf>

Version: 2024-02-01

12
papers

270
citations

1307594

7
h-index

1199594

12
g-index

12
all docs

12
docs citations

12
times ranked

495
citing authors

#	ARTICLE	IF	CITATIONS
1	Oncogenic Mutations Rewire Signaling Pathways by Switching Protein Recruitment to Phosphotyrosine Sites. <i>Cell</i> , 2019, 179, 543-560.e26.	28.9	65
2	Prolyl-isomerase Pin1 controls Notch3 protein expression and regulates T-ALL progression. <i>Oncogene</i> , 2016, 35, 4741-4751.	5.9	45
3	NOTCH3 inactivation increases triple negative breast cancer sensitivity to gefitinib by promoting EGFR tyrosine dephosphorylation and its intracellular arrest. <i>Oncogenesis</i> , 2018, 7, 42.	4.9	39
4	ProAla is an Effective Alternative to Trypsin for Proteomics Applications and Disulfide Bond Mapping. <i>Molecular and Cellular Proteomics</i> , 2020, 19, 2139-2157.	3.8	27
5	Proteomics Reveals Global Regulation of Protein SUMOylation by ATM and ATR Kinases during Replication Stress. <i>Cell Reports</i> , 2017, 21, 546-558.	6.4	24
6	Proteomics of resistance to Notch1 inhibition in acute lymphoblastic leukemia reveals targetable kinase signatures. <i>Nature Communications</i> , 2021, 12, 2507.	12.8	22
7	Notch3 contributes to T-cell leukemia growth via regulation of the unfolded protein response. <i>Oncogenesis</i> , 2020, 9, 93.	4.9	13
8	Optimal analytical strategies for sensitive and quantitative phosphoproteomics using TMT-based multiplexing. <i>Proteomics</i> , 2022, 22, .	2.2	9
9	Phosphorylation of SHP2 at Tyr62 Enables Acquired Resistance to SHP2 Allosteric Inhibitors in FLT3-ITD-Driven AML. <i>Cancer Research</i> , 2022, 82, 2141-2155.	0.9	8
10	Loss of CBL E3-ligase activity in B-lineage childhood acute lymphoblastic leukaemia. <i>British Journal of Haematology</i> , 2012, 159, 115-119.	2.5	6
11	Numb-dependent integration of pre-TCR and p53 function in T-cell precursor development. <i>Cell Death and Disease</i> , 2014, 5, e1472-e1472.	6.3	6
12	Deciphering the human phosphoproteome. <i>Nature Biotechnology</i> , 2020, 38, 285-286.	17.5	6