## Giulia Franciosa

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Phosphorylation of SHP2 at Tyr62 Enables Acquired Resistance to SHP2 Allosteric Inhibitors in FLT3-ITDâ€"Driven AML. Cancer Research, 2022, 82, 2141-2155.	0.9	8
2	Optimal analytical strategies for sensitive and quantitative phosphoproteomics using TMTâ€based multiplexing. Proteomics, 2022, 22, .	2.2	9
3	Proteomics of resistance to Notch1 inhibition in acute lymphoblastic leukemia reveals targetable kinase signatures. Nature Communications, 2021, 12, 2507.	12.8	22
4	Notch3 contributes to T-cell leukemia growth via regulation of the unfolded protein response. Oncogenesis, 2020, 9, 93.	4.9	13
5	ProAlanase is an Effective Alternative to Trypsin for Proteomics Applications and Disulfide Bond Mapping. Molecular and Cellular Proteomics, 2020, 19, 2139-2157.	3 <b>.</b> 8	27
6	Deciphering the human phosphoproteome. Nature Biotechnology, 2020, 38, 285-286.	17.5	6
7	Oncogenic Mutations Rewire Signaling Pathways by Switching Protein Recruitment to Phosphotyrosine Sites. Cell, 2019, 179, 543-560.e26.	28.9	65
8	NOTCH3 inactivation increases triple negative breast cancer sensitivity to gefitinib by promoting EGFR tyrosine dephosphorylation and its intracellular arrest. Oncogenesis, 2018, 7, 42.	4.9	39
9	Proteomics Reveals Global Regulation of Protein SUMOylation by ATM and ATR Kinases during Replication Stress. Cell Reports, 2017, 21, 546-558.	6.4	24
10	Prolyl-isomerase Pin1 controls Notch3 protein expression and regulates T-ALL progression. Oncogene, 2016, 35, 4741-4751.	5.9	45
11	Numb-dependent integration of pre-TCR and p53 function in T-cell precursor development. Cell Death and Disease, 2014, 5, e1472-e1472.	6.3	6
12	Loss of <scp>CBL</scp> E3â€ligase activity in Bâ€lineage childhood acute lymphoblastic leukaemia. British Journal of Haematology, 2012, 159, 115-119.	2.5	6