

Bin Fang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6049780/publications.pdf>

Version: 2024-02-01

15
papers

689
citations

840776

11
h-index

1058476

14
g-index

15
all docs

15
docs citations

15
times ranked

1670
citing authors

#	ARTICLE	IF	CITATIONS
1	Unification of de novo and acquired ibrutinib resistance in mantle cell lymphoma. <i>Nature Communications</i> , 2017, 8, 14920.	12.8	122
2	An immunoproteomic approach to characterize the CAR interactome and signalosome. <i>Science Signaling</i> , 2019, 12, .	3.6	109
3	Proteogenomic landscape of squamous cell lung cancer. <i>Nature Communications</i> , 2019, 10, 3578.	12.8	84
4	Ligand-Independent EPHA2 Signaling Drives the Adoption of a Targeted Therapy-Mediated Metastatic Melanoma Phenotype. <i>Cancer Discovery</i> , 2015, 5, 264-273.	9.4	82
5	PLK1 stabilizes a MYC-dependent kinase network in aggressive B cell lymphomas. <i>Journal of Clinical Investigation</i> , 2018, 128, 5517-5530.	8.2	67
6	K27-linked ubiquitination of BRAF by ITCH engages cytokine response to maintain MEK-ERK signaling. <i>Nature Communications</i> , 2019, 10, 1870.	12.8	61
7	Combined BRAF and HSP90 Inhibition in Patients with Unresectable <i>BRAF</i> ^{V600E} -Mutant Melanoma. <i>Clinical Cancer Research</i> , 2018, 24, 5516-5524.	7.0	55
8	A Pilot Proteogenomic Study with Data Integration Identifies MCT1 and GLUT1 as Prognostic Markers in Lung Adenocarcinoma. <i>PLoS ONE</i> , 2015, 10, e0142162.	2.5	31
9	HDAC6 regulates DNA damage response via deacetylating MLH1. <i>Journal of Biological Chemistry</i> , 2019, 294, 5813-5826.	3.4	28
10	Relative protein quantification and accessible biology in lung tumor proteomes from four LC-MS/MS discovery platforms. <i>Proteomics</i> , 2017, 17, 1600300.	2.2	15
11	Chronologically modified androgen receptor in recurrent castration-resistant prostate cancer and its therapeutic targeting. <i>Science Translational Medicine</i> , 2022, 14, .	12.4	12
12	Targeted mass-spectrometry-based assays enable multiplex quantification of receptor tyrosine kinase, MAP kinase, and AKT signaling. <i>Cell Reports Methods</i> , 2021, 1, 100015.	2.9	10
13	Plasma cell dependence on histone/protein deacetylase 11 reveals a therapeutic target in multiple myeloma. <i>JCI Insight</i> , 2021, 6, .	5.0	8
14	Multi-Omics and Informatics Analysis of FFPE Tissues Derived from Melanoma Patients with Long/Short Responses to Anti-PD1 Therapy Reveals Pathways of Response. <i>Cancers</i> , 2020, 12, 3515.	3.7	5
15	Managing a Large-Scale Multiomics Project: A Team Science Case Study in Proteogenomics. <i>Methods in Molecular Biology</i> , 2021, 2194, 187-221.	0.9	0