Koei Chin

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1466840/publications.pdf

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26 3,759 13 24
papers citations h-index g-index

34 34 34 7585
all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	A collection of breast cancer cell lines for the study of functionally distinct cancer subtypes. Cancer Cell, 2006, 10, 515-527.	7.7	2,729
2	In situ analyses of genome instability in breast cancer. Nature Genetics, 2004, 36, 984-988.	9.4	337
3	Differentiation-state plasticity is a targetable resistance mechanism in basal-like breast cancer. Nature Communications, 2018, 9, 3815.	5 . 8	137
4	Protein Acetylation and Histone Deacetylase Expression Associated with Malignant Breast Cancer Progression. Clinical Cancer Research, 2009, 15, 3163-3171.	3.2	110
5	Microenvironment-Mediated Mechanisms of Resistance to HER2 Inhibitors Differ between HER2+ Breast Cancer Subtypes. Cell Systems, 2018, 6, 329-342.e6.	2.9	72
6	FOXP3-positive regulatory T lymphocytes and epithelial FOXP3 expression in synchronous normal, ductal carcinoma in situ, and invasive cancer of the breast. Breast Cancer Research and Treatment, 2013, 139, 381-390.	1.1	55
7	Copy Number Gain of hsa-miR-569 at 3q26.2 Leads to Loss of TP53INP1 and Aggressiveness of Epithelial Cancers. Cancer Cell, 2014, 26, 863-879.	7.7	46
8	Cyclic Multiplexed-Immunofluorescence (cmIF), a Highly Multiplexed Method for Single-Cell Analysis. Methods in Molecular Biology, 2020, 2055, 521-562.	0.4	33
9	Relevance of circulating hybrid cells as a non-invasive biomarker for myriad solid tumors. Scientific Reports, 2021, 11, 13630.	1.6	31
10	RESTORE: Robust intEnSiTy nORmalization mEthod for multiplexed imaging. Communications Biology, 2020, 3, 111.	2.0	28
11	Genomic Alterations during the <i>In Situ</i> to Invasive Ductal Breast Carcinoma Transition Shaped by the Immune System. Molecular Cancer Research, 2021, 19, 623-635.	1.5	24
12	Toward reproducible, scalable, and robust data analysis across multiplex tissue imaging platforms. Cell Reports Methods, 2021, 1, 100053.	1.4	22
13	An omic and multidimensional spatial atlas from serial biopsies of an evolving metastatic breast cancer. Cell Reports Medicine, 2022, 3, 100525.	3.3	22
14	The impact of tumor epithelial and microenvironmental heterogeneity on treatment responses in HER2-positive breast cancer. JCI Insight, 2021, 6, .	2.3	20
15	A framework for multiplex imaging optimization and reproducible analysis. Communications Biology, 2022, 5, 438.	2.0	17
16	Oligonucleotide conjugated antibodies permit highly multiplexed immunofluorescence for future use in clinical histopathology. Journal of Biomedical Optics, 2020, 25, 1.	1.4	16
17	Multiomics analysis of serial PARP inhibitor treated metastatic TNBC inform on rational combination therapies. Npj Precision Oncology, 2021, 5, 92.	2.3	11
18	Oligonucleotide conjugated antibody strategies for cyclic immunostaining. Scientific Reports, 2021, 11, 23844.	1.6	11

#	Article	IF	CITATIONS
19	GRB7 dependent proliferation of basalâ€like, HERâ€2 positive human breast cancer cell lines is mediated in part by HERâ€1 signaling. Molecular Carcinogenesis, 2019, 58, 699-707.	1.3	9
20	Sensitivity to targeted therapy differs between HER2-amplified breast cancer cells harboring kinase and helical domain mutations in PIK3CA. Breast Cancer Research, 2021, 23, 81.	2.2	7
21	Quantitative, in situ analysis of mRNAs and proteins with subcellular resolution. Scientific Reports, 2017, 7, 16459.	1.6	6
22	Proteomics advances for precision therapy in ovarian cancer. Expert Review of Proteomics, 2019, 16, 841-850.	1.3	5
23	cmIF: A Python Library for Scalable Multiplex Imaging Pipelines. Lecture Notes in Computer Science, 2019, , 37-43.	1.0	3
24	Antibody Conjugated Oligonucleotides as a Platform for Cyclic Immunofluorescent Staining. Microscopy and Microanalysis, 2019, 25, 1206-1207.	0.2	1
25	Signal removal methods for highly multiplexed immunofluorescent staining using antibody conjugated oligonucleotides., 2019, 10881, .		1
26	Simultaneous Detection of RNAs and Proteins with Subcellular Resolution. Methods in Molecular Biology, 2020, 2161, 59-73.	0.4	1