

Kiley Graim

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4583277/publications.pdf>

Version: 2024-02-01

16
papers

4,148
citations

1040056

9
h-index

1281871

11
g-index

16
all docs

16
docs citations

16
times ranked

9338
citing authors

#	ARTICLE	IF	CITATIONS
1	Modeling molecular development of breast cancer in canine mammary tumors. <i>Genome Research</i> , 2021, 31, 337-347.	5.5	12
2	PLATYPUS: A Multiple-View Learning Predictive Framework for Cancer Drug Sensitivity Prediction. <i>Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing</i> , 2019, 24, 136-147.	0.7	7
3	PLATYPUS: A Multiple-View Learning Predictive Framework for Cancer Drug Sensitivity Prediction. , 2018, , .		9
4	Integrative Molecular Characterization of Malignant Pleural Mesothelioma. <i>Cancer Discovery</i> , 2018, 8, 1548-1565.	9.4	422
5	A Community Challenge for Inferring Genetic Predictors of Gene Essentialities through Analysis of a Functional Screen of Cancer Cell Lines. <i>Cell Systems</i> , 2017, 5, 485-497.e3.	6.2	19
6	TumorMap: Exploring the Molecular Similarities of Cancer Samples in an Interactive Portal. <i>Cancer Research</i> , 2017, 77, e111-e114.	0.9	59
7	Revealing cancer subtypes with higher-order correlations applied to imaging and omics data. <i>BMC Medical Genomics</i> , 2017, 10, 20.	1.5	9
8	Prophetic Granger Causality to infer gene regulatory networks. <i>PLoS ONE</i> , 2017, 12, e0170340.	2.5	10
9	Inferring causal molecular networks: empirical assessment through a community-based effort. <i>Nature Methods</i> , 2016, 13, 310-318.	19.0	209
10	The Molecular Taxonomy of Primary Prostate Cancer. <i>Cell</i> , 2015, 163, 1011-1025.	28.9	2,435
11	A basal stem cell signature identifies aggressive prostate cancer phenotypes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E6544-52.	7.1	168
12	Abstract PR02: Multiple Pathway Learning accurately predicts gene essentiality in the Cancer Cell Line Encyclopedia. , 2015, , .		0
13	Abstract A2-64: A signature catalog to classify tumor mixtures: Application to recognition of metastatic disease in prostate cancer. , 2015, , .		0
14	Abstract PR10: Multiple Pathway Learning accurately predicts gene essentiality in the Cancer Cell Line Encyclopedia. , 2015, , .		0
15	Abstract 4177: Identification of pathways relevant for metastatic site prediction in prostate cancer. , 2014, , .		0
16	A large-scale evaluation of computational protein function prediction. <i>Nature Methods</i> , 2013, 10, 221-227.	19.0	789