

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 | The Molecular Taxonomy of Primary Prostate Cancer. <i>Cell</i> , 2015, 163, 1011-1025. | 28.9 | 2,435 |
| 2 | A large-scale evaluation of computational protein function prediction. <i>Nature Methods</i> , 2013, 10, 221-227. | 19.0 | 789 |
| 3 | Integrative Molecular Characterization of Malignant Pleural Mesothelioma. <i>Cancer Discovery</i> , 2018, 8, 1548-1565. | 9.4 | 422 |
| 4 | Inferring causal molecular networks: empirical assessment through a community-based effort. <i>Nature Methods</i> , 2016, 13, 310-318. | 19.0 | 209 |
| 5 | 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 |
| 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 | 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 |
| 8 | Modeling molecular development of breast cancer in canine mammary tumors. <i>Genome Research</i> , 2021, 31, 337-347. | 5.5 | 12 |
| 9 | Prophetic Granger Causality to infer gene regulatory networks. <i>PLoS ONE</i> , 2017, 12, e0170340. | 2.5 | 10 |
| 10 | Revealing cancer subtypes with higher-order correlations applied to imaging and omics data. <i>BMC Medical Genomics</i> , 2017, 10, 20. | 1.5 | 9 |
| 11 | PLATYPUS: A Multiple-View Learning Predictive Framework for Cancer Drug Sensitivity Prediction. , 2018, , . | | 9 |
| 12 | PLATYPUS: A Multiple-View Learning Predictive Framework for Cancer Drug Sensitivity Prediction. Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing, 2019, 24, 136-147. | 0.7 | 7 |
| 13 | Abstract 4177: Identification of pathways relevant for metastatic site prediction in prostate cancer. , 2014, , . | | 0 |
| 14 | Abstract PR02: Multiple Pathway Learning accurately predicts gene essentiality in the Cancer Cell Line Encyclopedia. , 2015, , . | | 0 |
| 15 | Abstract A2-64: A signature catalog to classify tumor mixtures: Application to recognition of metastatic disease in prostate cancer. , 2015, , . | | 0 |
| 16 | Abstract PR10: Multiple Pathway Learning accurately predicts gene essentiality in the Cancer Cell Line Encyclopedia. , 2015, , . | | 0 |