

Ethan Cerami

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10608592/publications.pdf>

Version: 2024-02-01

19
papers

33,447
citations

623734

14
h-index

794594

19
g-index

21
all docs

21
docs citations

21
times ranked

45149
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | ThreadStates: State-based Visual Analysis of Disease Progression. IEEE Transactions on Visualization and Computer Graphics, 2022, 28, 238-247. | 4.4 | 10 |
| 2 | MITI minimum information guidelines for highly multiplexed tissue images. Nature Methods, 2022, 19, 262-267. | 19.0 | 37 |
| 3 | Genomic Characterization of <i>de novo</i> Metastatic Breast Cancer. Clinical Cancer Research, 2021, 27, 1105-1118. | 7.0 | 24 |
| 4 | OUP accepted manuscript. Bioinformatics, 2021, 37, i59-i66. | 4.1 | 1 |
| 5 | OncoTree: A Cancer Classification System for Precision Oncology. JCO Clinical Cancer Informatics, 2021, 5, 221-230. | 2.1 | 51 |
| 6 | Network for Biomarker Immunoprofiling for Cancer Immunotherapy: Cancer Immune Monitoring and Analysis Centers and Cancer Immunologic Data Commons (CIMAC-CIDC). Clinical Cancer Research, 2021, 27, 5038-5048. | 7.0 | 13 |
| 7 | Cross-Site Concordance Evaluation of Tumor DNA and RNA Sequencing Platforms for the CIMAC-CIDC Network. Clinical Cancer Research, 2021, 27, 5049-5061. | 7.0 | 6 |
| 8 | Pathway Commons 2019 Update: integration, analysis and exploration of pathway data. Nucleic Acids Research, 2020, 48, D489-D497. | 14.5 | 161 |
| 9 | The Human Tumor Atlas Network: Charting Tumor Transitions across Space and Time at Single-Cell Resolution. Cell, 2020, 181, 236-249. | 28.9 | 334 |
| 10 | Interactive or static reports to guide clinical interpretation of cancer genomics. Journal of the American Medical Informatics Association: JAMIA, 2018, 25, 458-464. | 4.4 | 14 |
| 11 | Institutional implementation of clinical tumor profiling on an unselected cancer population. JCI Insight, 2016, 1, e87062. | 5.0 | 340 |
| 12 | The Somatic Genomic Landscape of Glioblastoma. Cell, 2013, 155, 462-477. | 28.9 | 3,979 |
| 13 | Using MEMo to Discover Mutual Exclusivity Modules in Cancer. Current Protocols in Bioinformatics, 2013, 41, Unit 8.17. | 25.8 | 29 |
| 14 | Integrative Analysis of Complex Cancer Genomics and Clinical Profiles Using the cBioPortal. Science Signaling, 2013, 6, pl1. | 3.6 | 11,344 |
| 15 | Mutual exclusivity analysis identifies oncogenic network modules. Genome Research, 2012, 22, 398-406. | 5.5 | 597 |
| 16 | The cBio Cancer Genomics Portal: An Open Platform for Exploring Multidimensional Cancer Genomics Data. Cancer Discovery, 2012, 2, 401-404. | 9.4 | 12,801 |
| 17 | Integrative Genomic Profiling of Human Prostate Cancer. Cancer Cell, 2010, 18, 11-22. | 16.8 | 3,151 |
| 18 | Automated Network Analysis Identifies Core Pathways in Glioblastoma. PLoS ONE, 2010, 5, e8918. | 2.5 | 318 |

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 19 | Broadening the horizon “ level 2.5 of the HUPO-PSI format for molecular interactions. BMC Biology, 2007, 5, 44. | 3.8 | 237 |