

Hai Hu

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/165360/publications.pdf>

Version: 2024-02-01

31
papers

9,409
citations

471509

17
h-index

454955

30
g-index

32
all docs

32
docs citations

32
times ranked

16983
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Spatial Metrics of Interaction between CD163-Positive Macrophages and Cancer Cells and Progression-Free Survival in Chemo-Treated Breast Cancer. <i>Cancers</i> , 2022, 14, 308. | 3.7 | 8 |
| 2 | A Novel Blood-Based microRNA Diagnostic Model with High Accuracy for Multi-Cancer Early Detection. <i>Cancers</i> , 2022, 14, 1450. | 3.7 | 8 |
| 3 | Comparative analysis of differentially abundant proteins quantified by LC-MS/MS between flash frozen and laser microdissected OCT-embedded breast tumor samples. <i>Clinical Proteomics</i> , 2020, 17, 40. | 2.1 | 2 |
| 4 | Development and validation of prognostic gene signature for basal-like breast cancer and high-grade serous ovarian cancer. <i>Breast Cancer Research and Treatment</i> , 2020, 184, 689-698. | 2.5 | 4 |
| 5 | Malignant cell-specific pro-tumorigenic role of type I interferon receptor in breast cancers. <i>Cancer Biology and Therapy</i> , 2020, 21, 629-636. | 3.4 | 7 |
| 6 | PCA-PAM50 improves consistency between breast cancer intrinsic and clinical subtyping reclassifying a subset of luminal A tumors as luminal B. <i>Scientific Reports</i> , 2019, 9, 7956. | 3.3 | 37 |
| 7 | From Discovery to Practice and Survivorship: Building a National Real-World Data Learning Healthcare Framework for Military and Veteran Cancer Patients. <i>Clinical Pharmacology and Therapeutics</i> , 2019, 106, 52-57. | 4.7 | 18 |
| 8 | An Integrated TCGA Pan-Cancer Clinical Data Resource to Drive High-Quality Survival Outcome Analytics. <i>Cell</i> , 2018, 173, 400-416.e11. | 28.9 | 2,277 |
| 9 | The Immune Landscape of Cancer. <i>Immunity</i> , 2018, 48, 812-830.e14. | 14.3 | 3,706 |
| 10 | Genomic and Molecular Landscape of DNA Damage Repair Deficiency across The Cancer Genome Atlas. <i>Cell Reports</i> , 2018, 23, 239-254.e6. | 6.4 | 801 |
| 11 | Analysis of breast cancer in young women in the Department of Defense (DOD) database. <i>Breast Cancer Research and Treatment</i> , 2018, 168, 501-511. | 2.5 | 17 |
| 12 | Comparison of Breast Cancer Molecular Features and Survival by African and European Ancestry in The Cancer Genome Atlas. <i>JAMA Oncology</i> , 2017, 3, 1654. | 7.1 | 208 |
| 13 | Comparative Survival Analysis of Invasive Breast Cancer Patients Treated by a U.S. Military Medical Center and Matched Patients From the U.S. General Population. <i>Military Medicine</i> , 2017, 182, e1851-e1858. | 0.8 | 8 |
| 14 | Validation of tumor protein marker quantification by two independent automated immunofluorescence image analysis platforms. <i>Modern Pathology</i> , 2016, 29, 1143-1154. | 5.5 | 25 |
| 15 | Positive Association of Fibroadenomatoid Change with HER2-Negative Invasive Breast Cancer: A Co-Occurrence Study. <i>PLoS ONE</i> , 2015, 10, e0129500. | 2.5 | 7 |
| 16 | Comprehensive Molecular Portraits of Invasive Lobular Breast Cancer. <i>Cell</i> , 2015, 163, 506-519. | 28.9 | 1,485 |
| 17 | QAiT: A quality assurance issue tracking tool to facilitate the improvement of clinical data quality. <i>Computer Methods and Programs in Biomedicine</i> , 2013, 109, 86-91. | 4.7 | 5 |
| 18 | FH535 Inhibited Migration and Growth of Breast Cancer Cells. <i>PLoS ONE</i> , 2012, 7, e44418. | 2.5 | 27 |

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|----|---|-----|-----------|
| 19 | DW4TR: A Data Warehouse for Translational Research. <i>Journal of Biomedical Informatics</i> , 2011, 44, 1004-1019. | 4.3 | 48 |
| 20 | A Bayesian derived network of breast pathology co-occurrence. <i>Journal of Biomedical Informatics</i> , 2008, 41, 242-250. | 4.3 | 22 |
| 21 | MOF: An R Function to Detect Outlier Microarray. <i>Genomics, Proteomics and Bioinformatics</i> , 2008, 6, 186-189. | 6.9 | 2 |
| 22 | Clinical prediction of antidepressant response in mood disorders: Linear multivariate vs. neural network models. <i>Psychiatry Research</i> , 2007, 152, 223-231. | 3.3 | 24 |
| 23 | Detecting outlier microarray arrays by correlation and percentage of outliers spots. <i>Cancer Informatics</i> , 2007, 2, 351-60. | 1.9 | 5 |
| 24 | Detecting Outlier Microarray Arrays by Correlation and Percentage of Outliers Spots. <i>Cancer Informatics</i> , 2006, 2, 117693510600200. | 1.9 | 6 |
| 25 | Co-Occurrence Analysis for Discovery of Novel Breast Cancer Pathology Patterns. <i>IEEE Transactions on Information Technology in Biomedicine</i> , 2006, 10, 497-503. | 3.2 | 14 |
| 26 | Biomedical informatics: development of a comprehensive data warehouse for clinical and genomic breast cancer research. <i>Pharmacogenomics</i> , 2004, 5, 933-941. | 1.3 | 26 |
| 27 | Global search for chromosomal abnormalities in infiltrating ductal carcinoma of the breast using array-comparative genomic hybridization. <i>Cancer Genetics and Cytogenetics</i> , 2004, 155, 108-118. | 1.0 | 19 |
| 28 | A map of WW domain family interactions. <i>Proteomics</i> , 2004, 4, 643-655. | 2.2 | 122 |
| 29 | High-throughput proteomic analysis of human infiltrating ductal carcinoma of the breast. <i>Proteomics</i> , 2003, 3, 1863-1873. | 2.2 | 168 |
| 30 | Stretch-Activated Ion Channels in the Heart. <i>Journal of Molecular and Cellular Cardiology</i> , 1997, 29, 1511-1523. | 1.9 | 303 |
| 31 | A Novel Computational Analysis of Heterogeneity in Breast Tissue. , 0, , . | | 0 |