

Laura Romero-PÃ©rez

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8711019/publications.pdf>

Version: 2024-02-01

28
papers

1,610
citations

361413

20
h-index

501196

28
g-index

38
all docs

38
docs citations

38
times ranked

3384
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Evidence for an alternative fatty acid desaturation pathway increasing cancer plasticity. <i>Nature</i> , 2019, 566, 403-406. | 27.8 | 326 |
| 2 | Sarcoma classification by DNA methylation profiling. <i>Nature Communications</i> , 2021, 12, 498. | 12.8 | 237 |
| 3 | MicroRNA signature of the epithelial-mesenchymal transition in endometrial carcinosarcoma. <i>Journal of Pathology</i> , 2011, 223, 72-80. | 4.5 | 194 |
| 4 | MicroRNA-200 Family Modulation in Distinct Breast Cancer Phenotypes. <i>PLoS ONE</i> , 2012, 7, e47709. | 2.5 | 85 |
| 5 | ZEB1 overexpression associated with E-cadherin and microRNA-200 downregulation is characteristic of undifferentiated endometrial carcinoma. <i>Modern Pathology</i> , 2013, 26, 1514-1524. | 5.5 | 68 |
| 6 | Nuclear TAZ expression associates with the triple-negative phenotype in breast cancer. <i>Endocrine-Related Cancer</i> , 2015, 22, 443-454. | 3.1 | 66 |
| 7 | VGLL1 expression is associated with a triple-negative basal-like phenotype in breast cancer. <i>Endocrine-Related Cancer</i> , 2014, 21, 587-599. | 3.1 | 53 |
| 8 | Improving the management of Inherited Retinal Dystrophies by targeted sequencing of a population-specific gene panel. <i>Scientific Reports</i> , 2016, 6, 23910. | 3.3 | 51 |
| 9 | Cooperation of cancer drivers with regulatory germline variants shapes clinical outcomes. <i>Nature Communications</i> , 2019, 10, 4128. | 12.8 | 51 |
| 10 | DNA methylation profiling distinguishes Ewing-like sarcoma with EWSR1-NFATc2 fusion from Ewing sarcoma. <i>Journal of Cancer Research and Clinical Oncology</i> , 2019, 145, 1273-1281. | 2.5 | 50 |
| 11 | Preclinical Efficacy of Endoglin-Targeting Antibody-Drug Conjugates for the Treatment of Ewing Sarcoma. <i>Clinical Cancer Research</i> , 2019, 25, 2228-2240. | 7.0 | 44 |
| 12 | STAG Mutations in Cancer. <i>Trends in Cancer</i> , 2019, 5, 506-520. | 7.4 | 38 |
| 13 | Oncogenic hijacking of a developmental transcription factor evokes vulnerability toward oxidative stress in Ewing sarcoma. <i>Nature Communications</i> , 2020, 11, 2423. | 12.8 | 35 |
| 14 | A comparative view on the expression patterns of PD-L1 and PD-1 in soft tissue sarcomas. <i>Cancer Immunology, Immunotherapy</i> , 2020, 69, 1353-1362. | 4.2 | 34 |
| 15 | Molecular events in endometrial carcinosarcomas and the role of high mobility group AT-hook 2 in endometrial carcinogenesis. <i>Human Pathology</i> , 2013, 44, 244-254. | 2.0 | 30 |
| 16 | DNA methylation-based profiling of uterine neoplasms: a novel tool to improve gynecologic cancer diagnostics. <i>Journal of Cancer Research and Clinical Oncology</i> , 2020, 146, 97-104. | 2.5 | 29 |
| 17 | Oncogene alterations in endometrial carcinosarcomas. <i>Human Pathology</i> , 2013, 44, 852-859. | 2.0 | 27 |
| 18 | Integrative gene network and functional analyses identify a prognostically relevant key regulator of metastasis in Ewing sarcoma. <i>Molecular Cancer</i> , 2022, 21, 1. | 19.2 | 25 |

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 19 | Translational evidence for RRM2 as a prognostic biomarker and therapeutic target in Ewing sarcoma. <i>Molecular Cancer</i> , 2021, 20, 97. | 19.2 | 24 |
| 20 | A role for the transducer of the Hippo pathway, TAZ, in the development of aggressive types of endometrial cancer. <i>Modern Pathology</i> , 2015, 28, 1492-1503. | 5.5 | 23 |
| 21 | Gene expression and immunohistochemical analyses identify SOX2 as major risk factor for overall survival and relapse in Ewing sarcoma patients. <i>EBioMedicine</i> , 2019, 47, 156-162. | 6.1 | 23 |
| 22 | Targeting the CALCB/RAMP1 axis inhibits growth of Ewing sarcoma. <i>Cell Death and Disease</i> , 2019, 10, 116. | 6.3 | 23 |
| 23 | Hippo pathway effectors YAP1/TAZ induce an <i>EWS-FLI1</i> opposing gene signature and associate with disease progression in Ewing sarcoma. <i>Journal of Pathology</i> , 2020, 250, 374-386. | 4.5 | 19 |
| 24 | High Specificity of BCL11B and GLG1 for EWSR1-FLI1 and EWSR1-ERG Positive Ewing Sarcoma. <i>Cancers</i> , 2020, 12, 644. | 3.7 | 16 |
| 25 | Integrative clinical transcriptome analysis reveals <i>TMPRSS2-ERG</i> dependency of prognostic biomarkers in prostate adenocarcinoma. <i>International Journal of Cancer</i> , 2020, 146, 2036-2046. | 5.1 | 13 |
| 26 | Therapeutic targeting of the PLK1-PRC1-axis triggers cell death in genomically silent childhood cancer. <i>Nature Communications</i> , 2021, 12, 5356. | 12.8 | 11 |
| 27 | Functional genomics identifies AMPD2 as a new prognostic marker for undifferentiated pleomorphic sarcoma. <i>International Journal of Cancer</i> , 2019, 144, 859-867. | 5.1 | 10 |
| 28 | Genetics of Endometrial Carcinoma. , 2013, , 349-390. | | 1 |