# Juan DÃ-az MartÃ-n

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/4303954/publications.pdf

Version: 2024-02-01

| 33 | 1,333 | 393982 | 433756 |
|---|---|---|---|
| papers | citations | 19 h-index | 31 g-index |

| 34 | 34 | 34 | 2680 |
|---|---|---|---|
| all docs | docs citations | times ranked | citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | Characterizing the Invasive Tumor Front of Aggressive Uterine Adenocarcinoma and Leiomyosarcoma. Frontiers in Cell and Developmental Biology, 2021, 9, 670185. | 1.8 | 5 |
| 2 | A Novel NFIX-STAT6 Gene Fusion in Solitary Fibrous Tumor: A Case Report. International Journal of Molecular Sciences, 2021, 22, 7514. | 1.8 | 4 |
| 3 | Sarcoma classification by DNA methylation profiling. Nature Communications, 2021, 12, 498. | 5.8 | 237 |
| 4 | Evaluation of NAB2-STAT6 Fusion Variants and Other Molecular Alterations as Prognostic Biomarkers in a Case Series of 83 Solitary Fibrous Tumors. Cancers, 2021, 13, 5237. | 1.7 | 9 |
| 5 | Molecular Approaches to Diagnosis in Ewing Sarcoma: Targeted RNA Sequencing. Methods in Molecular Biology, 2021, 2226, 105-116. | 0.4 | 1 |
| 6 | Hippo pathway effectors YAP1/TAZ induce an EWSâ€“FLI1â€opposing gene signature and associate with disease progression in Ewing sarcoma. Journal of Pathology, 2020, 250, 374-386. | 2.1 | 19 |
| 7 | Pazopanib for treatment of typical solitary fibrous tumours: a multicentre, single-arm, phase 2 trial. Lancet Oncology, The, 2020, 21, 456-466. | 5.1 | 51 |
| 8 | Breakthrough Technologies Reshape the Ewing Sarcoma Molecular Landscape. Cells, 2020, 9, 804. | 1.8 | 8 |
| 9 | Abstract 6158: The relevance of endoglin and MMP14 to the metastatic potential of Ewing sarcoma cells. , 2020, , . | | 0 |
| 10 | DNA methylation profiling distinguishes Ewing-like sarcoma with EWSR1â€“NFATc2 fusion from Ewing sarcoma. Journal of Cancer Research and Clinical Oncology, 2019, 145, 1273-1281. | 1.2 | 50 |
| 11 | Pazopanib for treatment of advanced malignant and dedifferentiated solitary fibrous tumour: a multicentre, single-arm, phase 2 trial. Lancet Oncology, The, 2019, 20, 134-144. | 5.1 | 97 |
| 12 | What's in a name? Molecular subclassification of sarcomas creates fresh challenges. Journal of Pathology, 2019, 247, 409-412. | 2.1 | 2 |
| 13 | Preclinical Efficacy of Endoglin-Targeting Antibodyâ€“Drug Conjugates for the Treatment of Ewing Sarcoma. Clinical Cancer Research, 2019, 25, 2228-2240. | 3.2 | 44 |
| 14 | Nuclear TAZ expression associates with the triple-negative phenotype in breast cancer. Endocrine-Related Cancer, 2015, 22, 443-454. | 1.6 | 66 |
| 15 | A role for the transducer of the Hippo pathway, TAZ, in the development of aggressive types of endometrial cancer. Modern Pathology, 2015, 28, 1492-1503. | 2.9 | 23 |
| 16 | Zeb1 and <scp>S</scp>nail1 engage mi<scp>R</scp>â€200f transcriptional and epigenetic regulation during <scp>EMT</scp>. International Journal of Cancer, 2015, 136, E62-73. | 2.3 | 52 |
| 17 | A core microRNA signature associated with inducers of the epithelial-to-mesenchymal transition. Journal of Pathology, 2014, 232, 319-329. | 2.1 | 66 |
| 18 | VGLL1 expression is associated with a triple-negative basal-like phenotype in breast cancer. Endocrine-Related Cancer, 2014, 21, 587-599. | 1.6 | 53 |

| # | Article | IF | Citations |
|---|---|---|---|
| 19 | Molecular events in endometrial carcinosarcomas and the role of high mobility group AT-hook 2 in endometrial carcinogenesis. Human Pathology, 2013, 44, 244-254. | 1.1 | 30 |
| 20 | Oncogene alterations in endometrial carcinosarcomas. Human Pathology, 2013, 44, 852-859. | 1.1 | 27 |
| 21 | ZEB1 overexpression associated with E-cadherin and microRNA-200 downregulation is characteristic of undifferentiated endometrial carcinoma. Modern Pathology, 2013, 26, 1514-1524. | 2.9 | 68 |
| 22 | Age-Mediated Transcriptomic Changes in Adult Mouse Substantia Nigra. PLoS ONE, 2013, 8, e62456. | 1.1 | 15 |
| 23 | Genetics of Endometrial Carcinoma. , 2013, , 349-390. | | 1 |
| 24 | MicroRNA-200 Family Modulation in Distinct Breast Cancer Phenotypes. PLoS ONE, 2012, 7, e47709. | 1.1 | 85 |
| 25 | Intermediate alleles at the FRAXA and FRAXE loci in Parkinsonâ€™s disease. Parkinsonism and Related Disorders, 2011, 17, 281-284. | 1.1 | 16 |
| 26 | Heat shock protein 70 kDa over-expression and 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine-induced nigrostriatal degeneration in mice. Neuroscience, 2011, 193, 323-329. | 1.1 | 4 |
| 27 | Mesencephalic and striatal protein profiles in mice over-expressing glucose-6-phosphate dehydrogenase in dopaminergic neurons. Journal of Proteomics, 2010, 73, 1747-1757. | 1.2 | 5 |
| 28 | Brain-derived neurotrophic factor G196A polymorphism and clinical features in Parkinsonâ€™s disease. Acta Neurologica Scandinavica, 2010, 122, 41-45. | 1.0 | 37 |
| 29 | The HaDREB2 transcription factor enhances basal thermotolerance and longevity of seeds through functional interaction with HaHSFA9. BMC Plant Biology, 2009, 9, 75. | 1.6 | 57 |
| 30 | Prevalence and clinical features ofLRRK2mutations in patients with Parkinsonâ€™s disease in southern Spain. European Journal of Neurology, 2009, 16, 957-960. | 1.7 | 32 |
| 31 | Glucose-6-phosphate dehydrogenase activity in Parkinsonâ€™s disease. Journal of Neurology, 2008, 255, 1850-1851. | 1.8 | 8 |
| 32 | Functional Interaction between Two Transcription Factors Involved in the Developmental Regulation of a Small Heat Stress Protein Gene Promoter. Plant Physiology, 2005, 139, 1483-1494. | 2.3 | 80 |
| 33 | A Seed-specific Heat-shock Transcription Factor Involved in Developmental Regulation during Embryogenesis in Sunflower. Journal of Biological Chemistry, 2002, 277, 43866-43872. | 1.6 | 81 |