

Elisa N Ferreira

List of Publications by Year in descending order

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Version: 2024-02-01

29
papers

636
citations

758635

12
h-index

580395

25
g-index

29
all docs

29
docs citations

29
times ranked

1530
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Recurrent somatic mutation in DROSHA induces microRNA profile changes in Wilms tumour. <i>Nature Communications</i> , 2014, 5, 4039. | 5.8 | 159 |
| 2 | Gene expression patterns through oral squamous cell carcinoma development: PD-L1 expression in primary tumor and circulating tumor cells. <i>Oncotarget</i> , 2015, 6, 20902-20920. | 0.8 | 96 |
| 3 | Reciprocal changes in gene expression profiles of cocultured breast epithelial cells and primary fibroblasts. <i>International Journal of Cancer</i> , 2009, 125, 2767-2777. | 2.3 | 52 |
| 4 | Genomic imbalances pinpoint potential oncogenes and tumor suppressors in Wilms tumors. <i>Molecular Cytogenetics</i> , 2016, 9, 20. | 0.4 | 36 |
| 5 | BRCA1 deficiency is a recurrent event in early-onset triple-negative breast cancer: a comprehensive analysis of germline mutations and somatic promoter methylation. <i>Breast Cancer Research and Treatment</i> , 2018, 167, 803-814. | 1.1 | 36 |
| 6 | Upregulated genes at 2q24 gains as candidate oncogenes in hepatoblastomas. <i>Future Oncology</i> , 2014, 10, 2449-2457. | 1.1 | 29 |
| 7 | A genomic case study of desmoplastic small round cell tumor: comprehensive analysis reveals insights into potential therapeutic targets and development of a monitoring tool for a rare and aggressive disease. <i>Human Genomics</i> , 2016, 10, 36. | 1.4 | 28 |
| 8 | A Transcript Finishing Initiative for Closing Gaps in the Human Transcriptome. <i>Genome Research</i> , 2004, 14, 1413-1423. | 2.4 | 22 |
| 9 | Complex Landscape of Germline Variants in Brazilian Patients With Hereditary and Early Onset Breast Cancer. <i>Frontiers in Genetics</i> , 2018, 9, 161. | 1.1 | 21 |
| 10 | Intratumoral heterogeneity of ADAM23 promotes tumor growth and metastasis through LGI4 and nitric oxide signals. <i>Oncogene</i> , 2015, 34, 1270-1279. | 2.6 | 20 |
| 11 | Poly (A)+ Transcriptome Assessment of ERBB2-Induced Alterations in Breast Cell Lines. <i>PLoS ONE</i> , 2011, 6, e21022. | 1.1 | 17 |
| 12 | TGIF1 splicing variant 8 is overexpressed in oral squamous cell carcinoma and is related to pathologic and clinical behavior. <i>Oral Surgery, Oral Medicine, Oral Pathology and Oral Radiology</i> , 2013, 116, 614-625. | 0.2 | 14 |
| 13 | Contribution of the GSTP1 c.313A>G variant to hearing loss risk in patients exposed to platin chemotherapy during childhood. <i>Clinical and Translational Oncology</i> , 2019, 21, 630-635. | 1.2 | 14 |
| 14 | Alternative splicing: a bioinformatics perspective. <i>Molecular BioSystems</i> , 2007, 3, 473. | 2.9 | 13 |
| 15 | Down-regulation of ANAPC13 and CLTCL1: Early Events in the Progression of Preinvasive Ductal Carcinoma of the Breast. <i>Translational Oncology</i> , 2012, 5, 113-IN8. | 1.7 | 13 |
| 16 | A transcriptome-based signature of pathological angiogenesis predicts breast cancer patient survival. <i>PLoS Genetics</i> , 2019, 15, e1008482. | 1.5 | 12 |
| 17 | Mutational Portrait of Lung Adenocarcinoma in Brazilian Patients: Past, Present, and Future of Molecular Profiling in the Clinic. <i>Frontiers in Oncology</i> , 2020, 10, 1068. | 1.3 | 11 |
| 18 | Alternative splicing enriched cDNA libraries identify breast cancer-associated transcripts. <i>BMC Genomics</i> , 2010, 11, S4. | 1.2 | 10 |

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|----|---|-----|-----------|
| 19 | Evaluation of Quantitative RT-PCR Using Nonamplified and Amplified RNA. Diagnostic Molecular Pathology, 2010, 19, 45-53. | 2.1 | 9 |
| 20 | Case report: Chondrosarcoma of the head and neck. Human Pathology: Case Reports, 2017, 7, 4-7. | 0.2 | 7 |
| 21 | Influence of BRCA1 Germline Mutations in the Somatic Mutational Burden of Triple-Negative Breast Cancer. Translational Oncology, 2019, 12, 1453-1460. | 1.7 | 6 |
| 22 | Epithelial cells captured from ductal carcinoma in situ reveal a gene expression signature associated with progression to invasive breast cancer. Oncotarget, 2016, 7, 75672-75684. | 0.8 | 5 |
| 23 | Linear mRNA amplification approach for RNAseq from limited amount of RNA. Gene, 2015, 564, 220-227. | 1.0 | 2 |
| 24 | Heteroduplex formation and S1 digestion for mapping alternative splicing sites. Genetics and Molecular Research, 2008, 7, 958-969. | 0.3 | 2 |
| 25 | Effects of tumor biobank storage on polysome stability. Applied Cancer Research, 2019, 39, . | 1.0 | 1 |
| 26 | A comparison between SOLiD 5500XL and Ion Torrent PGM-derived miRNA expression profiles in two breast cell lines. Genetics and Molecular Biology, 2020, 43, e20180351. | 0.6 | 1 |
| 27 | Abstract 452: Comprehensive gene expression analysis identifies molecular markers involved with the progression of ductal carcinoma in situ of the breast. , 2014, , . | | 0 |
| 28 | Abstract A32: Urine as a potential liquid biopsy for detecting tumor DNA in Wilms tumor patient: Detection of somatic mutations in urine opens perspectives of monitoring chemotherapy response in WT patients. , 2016, , . | | 0 |
| 29 | Identification and complete sequencing of novel human transcripts through the use of mouse orthologs and testis cDNA sequences. Genetics and Molecular Research, 2004, 3, 493-511. | 0.3 | 0 |