

Beatrice Bl Lloriod

List of Publications by Year in descending order

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Version: 2024-02-01

38
papers

2,504
citations

279798

23
h-index

315739

38
g-index

38
all docs

38
docs citations

38
times ranked

4208
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Gene expression profiling of colon cancer by DNA microarrays and correlation with histoclinical parameters. <i>Oncogene</i> , 2004, 23, 1377-1391. | 5.9 | 293 |
| 2 | Thrombin-induced endothelial microparticle generation: identification of a novel pathway involving ROCK-II activation by caspase-2. <i>Blood</i> , 2006, 108, 1868-1876. | 1.4 | 194 |
| 3 | Responder and nonresponder patients exhibit different peripheral transcriptional signatures during major depressive episode. <i>Translational Psychiatry</i> , 2012, 2, e185-e185. | 4.8 | 192 |
| 4 | Gene Expression Profiling for Molecular Characterization of Inflammatory Breast Cancer and Prediction of Response to Chemotherapy. <i>Cancer Research</i> , 2004, 64, 8558-8565. | 0.9 | 177 |
| 5 | Gene expression profiling of primary breast carcinomas using arrays of candidate genes. <i>Human Molecular Genetics</i> , 2000, 9, 2981-2991. | 2.9 | 168 |
| 6 | Temporal gene profiling of the 5XFAD transgenic mouse model highlights the importance of microglial activation in Alzheimer's disease. <i>Molecular Neurodegeneration</i> , 2014, 9, 33. | 10.8 | 138 |
| 7 | Small lymphocytic lymphoma, marginal zone B-cell lymphoma, and mantle cell lymphoma exhibit distinct gene-expression profiles allowing molecular diagnosis. <i>Blood</i> , 2004, 103, 2727-2737. | 1.4 | 127 |
| 8 | Gene expression profiling of multiple myeloma reveals molecular portraits in relation to the pathogenesis of the disease. <i>Blood</i> , 2003, 101, 4998-5006. | 1.4 | 124 |
| 9 | Gene expression profiles of poor-prognosis primary breast cancer correlate with survival. <i>Human Molecular Genetics</i> , 2002, 11, 863-872. | 2.9 | 117 |
| 10 | Divergent transcription is associated with promoters of transcriptional regulators. <i>BMC Genomics</i> , 2013, 14, 914. | 2.8 | 95 |
| 11 | Gene expression profiling defines molecular subtypes of classical Hodgkin's disease. <i>Oncogene</i> , 2002, 21, 3095-3102. | 5.9 | 91 |
| 12 | Transcriptional Profiling Reveals Coordinated Up-Regulation of Oxidative Metabolism Genes in Thyroid Oncocytic Tumors. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2004, 89, 994-1005. | 3.6 | 80 |
| 13 | CD8 + T Cells from Human Neonates Are Biased toward an Innate Immune Response. <i>Cell Reports</i> , 2016, 17, 2151-2160. | 6.4 | 64 |
| 14 | Gene expression profiling of 1p35-36 genes in neuroblastoma. <i>Oncogene</i> , 2004, 23, 5912-5922. | 5.9 | 60 |
| 15 | Gene Expression Signature-Based Screening Identifies New Broadly Effective Influenza A Antivirals. <i>PLoS ONE</i> , 2010, 5, e13169. | 2.5 | 59 |
| 16 | Vitamin D interacts with Esr1 and Igf1 to regulate molecular pathways relevant to Alzheimer's disease. <i>Molecular Neurodegeneration</i> , 2016, 11, 22. | 10.8 | 58 |
| 17 | Expression scanning of an array of growth control genes in human tumor cell lines. <i>Oncogene</i> , 1999, 18, 3905-3912. | 5.9 | 57 |
| 18 | Olfactory Stem Cells, a New Cellular Model for Studying Molecular Mechanisms Underlying Familial Dysautonomia. <i>PLoS ONE</i> , 2010, 5, e15590. | 2.5 | 46 |

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|----|---|-----|-----------|
| 19 | Gene profiling reveals specific oncogenic mechanisms and signaling pathways in oncocytic and papillary thyroid carcinoma. <i>Oncogene</i> , 2005, 24, 4155-4161. | 5.9 | 40 |
| 20 | Feature extraction and signal processing for nylon DNA microarrays. <i>BMC Genomics</i> , 2004, 5, 38. | 2.8 | 37 |
| 21 | Prognosis of Breast Cancer and Gene Expression Profiling Using DNA Arrays. <i>Annals of the New York Academy of Sciences</i> , 2002, 975, 217-231. | 3.8 | 33 |
| 22 | Platelets Alter Gene Expression Profile in Human Brain Endothelial Cells in an In Vitro Model of Cerebral Malaria. <i>PLoS ONE</i> , 2011, 6, e19651. | 2.5 | 32 |
| 23 | A General Survey of Thymocyte Differentiation by Transcriptional Analysis of Knockout Mouse Models. <i>Journal of Immunology</i> , 2004, 173, 6109-6118. | 0.8 | 27 |
| 24 | Genome-wide analysis of familial dysautonomia and kinetin target genes with patient olfactory ecto-mesenchymal stem cells. <i>Human Mutation</i> , 2012, 33, 530-540. | 2.5 | 25 |
| 25 | The transcriptional signatures of <i>Sodalis glossinidius</i> in the <i>Glossina palpalis gambiense</i> flies negative for <i>Trypanosoma brucei gambiense</i> contrast with those of this symbiont in tsetse flies positive for the parasite: Possible involvement of a <i>Sodalis</i> -hosted prophage in fly <i>Trypanosoma</i> refractoriness?. <i>Infection, Genetics and Evolution</i> , 2014, 24, 41-56. | 2.3 | 24 |
| 26 | Gene Expression Profiles Characterize Inflammation Stages in the Acute Lung Injury in Mice. <i>PLoS ONE</i> , 2010, 5, e11485. | 2.5 | 23 |
| 27 | The chemosensitivity to therapy of childhood early B acute lymphoblastic leukemia could be determined by the combined expression of CD34, SPI-B and BCR genes. <i>Leukemia Research</i> , 2006, 30, 665-676. | 0.8 | 18 |
| 28 | Deciphering the adaptation strategies of <i>Desulfovibrio piezophilus</i> to hydrostatic pressure through metabolic and transcriptional analyses. <i>Environmental Microbiology Reports</i> , 2016, 8, 520-526. | 2.4 | 17 |
| 29 | An Evaluation of the Role of Gene Expression in the Prediction and Diagnosis of Ventilator-associated Pneumonia. <i>Anesthesiology</i> , 2011, 115, 344-352. | 2.5 | 13 |
| 30 | Population transcriptomics: insights from <i>Drosophila simulans</i> , <i>Drosophila sechellia</i> and their hybrids. <i>Genetica</i> , 2011, 139, 465-477. | 1.1 | 13 |
| 31 | Genomic transcriptional profiling in LOU/C/Jall rats identifies genes for successful aging. <i>Brain Structure and Function</i> , 2013, 218, 1501-1512. | 2.3 | 12 |
| 32 | Transcriptional Profiling of Midguts Prepared from <i>Trypanosoma/T. congolense</i> -Positive <i>Glossina palpalis palpalis</i> Collected from Two Distinct Cameroonian Foci: Coordinated Signatures of the Midguts' Remodeling As <i>T. congolense</i> -Supportive Niches. <i>Frontiers in Immunology</i> , 2017, 8, 876. | 4.8 | 11 |
| 33 | Gene expression profiling during thymus ontogeny and its association with TCRVbeta8.1-Dbeta2.1 rearrangements of inbred mouse strains. <i>Molecular and Cellular Biochemistry</i> , 2003, 252, 223-228. | 3.1 | 10 |
| 34 | Omics analysis of mouse brain models of human diseases. <i>Gene</i> , 2017, 600, 90-100. | 2.2 | 10 |
| 35 | Identification of overexpressed genes in <i>Sodalis glossinidius</i> inhabiting trypanosome-infected self-cured tsetse flies. <i>Frontiers in Microbiology</i> , 2014, 5, 255. | 3.5 | 6 |
| 36 | Large scale gene expression analysis of CBA/J mouse strain fetal thymus using cDNA-array hybridizations. <i>Molecular and Cellular Biochemistry</i> , 2004, 260, 65-68. | 3.1 | 5 |

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|----|---|-----|-----------|
| 37 | Effect of landiolol on sex-related transcriptomic changes in the myocardium during sepsis. Intensive Care Medicine Experimental, 2019, 7, 50. | 1.9 | 5 |
| 38 | Gene expression profiling of breast carcinomas using Nylon DNA arrays. Comptes Rendus - Biologies, 2003, 326, 1031-1039. | 0.2 | 3 |