

Beatrice Bl Lloriod

List of Publications by Year in descending order

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38
papers

2,504
citations

279487

23
h-index

315357

38
g-index

38
all docs

38
docs citations

38
times ranked

4208
citing authors

#	ARTICLE	IF	CITATIONS
1	Effect of landiolol on sex-related transcriptomic changes in the myocardium during sepsis. <i>Intensive Care Medicine Experimental</i> , 2019, 7, 50.	0.9	5
2	Omics analysis of mouse brain models of human diseases. <i>Gene</i> , 2017, 600, 90-100.	1.0	10
3	Transcriptional Profiling of Midguts Prepared from Trypanosoma/T. congolense-Positive Glossina palpalis palpalis Collected from Two Distinct Cameroonian Foci: Coordinated Signatures of the Midguts' Remodeling As T. congolense-Supportive Niches. <i>Frontiers in Immunology</i> , 2017, 8, 876.	2.2	11
4	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.	1.0	17
5	CD8 + T Cells from Human Neonates Are Biased toward an Innate Immune Response. <i>Cell Reports</i> , 2016, 17, 2151-2160.	2.9	64
6	Vitamin D interacts with Esr1 and Igf1 to regulate molecular pathways relevant to Alzheimer's disease. <i>Molecular Neurodegeneration</i> , 2016, 11, 22.	4.4	58
7	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.	4.4	138
8	Identification of overexpressed genes in Sodalis glossinidius inhabiting trypanosome-infected self-cured tsetse flies. <i>Frontiers in Microbiology</i> , 2014, 5, 255.	1.5	6
9	The transcriptional signatures of Sodalis glossinidius in the Glossina palpalis gambiense flies negative for Trypanosoma brucei gambiense contrast with those of this symbiont in tsetse flies positive for the parasite: Possible involvement of a Sodalis-hosted prophage in fly Trypanosoma refractoriness?. <i>Infection, Genetics and Evolution</i> , 2014, 24, 41-56.	1.0	24
10	Genomic transcriptional profiling in LOU/C/Jall rats identifies genes for successful aging. <i>Brain Structure and Function</i> , 2013, 218, 1501-1512.	1.2	12
11	Divergent transcription is associated with promoters of transcriptional regulators. <i>BMC Genomics</i> , 2013, 14, 914.	1.2	95
12	Responder and nonresponder patients exhibit different peripheral transcriptional signatures during major depressive episode. <i>Translational Psychiatry</i> , 2012, 2, e185-e185.	2.4	192
13	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.	1.1	25
14	An Evaluation of the Role of Gene Expression in the Prediction and Diagnosis of Ventilator-associated Pneumonia. <i>Anesthesiology</i> , 2011, 115, 344-352.	1.3	13
15	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.	1.1	32
16	Population transcriptomics: insights from Drosophila simulans, Drosophila sechellia and their hybrids. <i>Genetica</i> , 2011, 139, 465-477.	0.5	13
17	Gene Expression Signature-Based Screening Identifies New Broadly Effective Influenza A Antivirals. <i>PLoS ONE</i> , 2010, 5, e13169.	1.1	59
18	Olfactory Stem Cells, a New Cellular Model for Studying Molecular Mechanisms Underlying Familial Dysautonomia. <i>PLoS ONE</i> , 2010, 5, e15590.	1.1	46

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19	Gene Expression Profiles Characterize Inflammation Stages in the Acute Lung Injury in Mice. PLoS ONE, 2010, 5, e11485.	1.1	23
20	Thrombin-induced endothelial microparticle generation: identification of a novel pathway involving ROCK-II activation by caspase-2. Blood, 2006, 108, 1868-1876.	0.6	194
21	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. Leukemia Research, 2006, 30, 665-676.	0.4	18
22	Gene profiling reveals specific oncogenic mechanisms and signaling pathways in oncocytic and papillary thyroid carcinoma. Oncogene, 2005, 24, 4155-4161.	2.6	40
23	Transcriptional Profiling Reveals Coordinated Up-Regulation of Oxidative Metabolism Genes in Thyroid Oncocytic Tumors. Journal of Clinical Endocrinology and Metabolism, 2004, 89, 994-1005.	1.8	80
24	Gene Expression Profiling for Molecular Characterization of Inflammatory Breast Cancer and Prediction of Response to Chemotherapy. Cancer Research, 2004, 64, 8558-8565.	0.4	177
25	A General Survey of Thymocyte Differentiation by Transcriptional Analysis of Knockout Mouse Models. Journal of Immunology, 2004, 173, 6109-6118.	0.4	27
26	Gene expression profiling of colon cancer by DNA microarrays and correlation with histoclinical parameters. Oncogene, 2004, 23, 1377-1391.	2.6	293
27	Gene expression profiling of 1p35-36 genes in neuroblastoma. Oncogene, 2004, 23, 5912-5922.	2.6	60
28	Feature extraction and signal processing for nylon DNA microarrays. BMC Genomics, 2004, 5, 38.	1.2	37
29	Large scale gene expression analysis of CBA/J mouse strain fetal thymus using cDNA-array hybridizations. Molecular and Cellular Biochemistry, 2004, 260, 65-68.	1.4	5
30	Small lymphocytic lymphoma, marginal zone B-cell lymphoma, and mantle cell lymphoma exhibit distinct gene-expression profiles allowing molecular diagnosis. Blood, 2004, 103, 2727-2737.	0.6	127
31	Gene expression profiling during thymus ontogeny and its association with TCRVbeta8.1-Dbeta2.1 rearrangements of inbred mouse strains. Molecular and Cellular Biochemistry, 2003, 252, 223-228.	1.4	10
32	Gene expression profiling of breast carcinomas using Nylon DNA arrays. Comptes Rendus - Biologies, 2003, 326, 1031-1039.	0.1	3
33	Gene expression profiling of multiple myeloma reveals molecular portraits in relation to the pathogenesis of the disease. Blood, 2003, 101, 4998-5006.	0.6	124
34	Gene expression profiles of poor-prognosis primary breast cancer correlate with survival. Human Molecular Genetics, 2002, 11, 863-872.	1.4	117
35	Gene expression profiling defines molecular subtypes of classical Hodgkin's disease. Oncogene, 2002, 21, 3095-3102.	2.6	91
36	Prognosis of Breast Cancer and Gene Expression Profiling Using DNA Arrays. Annals of the New York Academy of Sciences, 2002, 975, 217-231.	1.8	33

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37	Gene expression profiling of primary breast carcinomas using arrays of candidate genes. Human Molecular Genetics, 2000, 9, 2981-2991.	1.4	168
38	Expression scanning of an array of growth control genes in human tumor cell lines. Oncogene, 1999, 18, 3905-3912.	2.6	57