

# Beatrice Bl Lloriod

## List of Publications by Year in descending order

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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	Effect of landiolol on sex-related transcriptomic changes in the myocardium during sepsis. Intensive Care Medicine Experimental, 2019, 7, 50.	1.9	5
2	Omics analysis of mouse brain models of human diseases. Gene, 2017, 600, 90-100.	2.2	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's Remodeling As T. congolense-Supportive Niches. Frontiers in Immunology, 2017, 8, 876.	4.8	11
4	Deciphering the adaptation strategies of <i>Desulfovibrio piezophilus</i> to hydrostatic pressure through metabolic and transcriptional analyses. Environmental Microbiology Reports, 2016, 8, 520-526.	2.4	17
5	CD8 + T Cells from Human Neonates Are Biased toward an Innate Immune Response. Cell Reports, 2016, 17, 2151-2160.	6.4	64
6	Vitamin D interacts with Esr1 and Igf1 to regulate molecular pathways relevant to Alzheimer's disease. Molecular Neurodegeneration, 2016, 11, 22.	10.8	58
7	Temporal gene profiling of the 5XFAD transgenic mouse model highlights the importance of microglial activation in Alzheimer's disease. Molecular Neurodegeneration, 2014, 9, 33.	10.8	138
8	Identification of overexpressed genes in Sodalis glossinidius inhabiting trypanosome-infected self-cured tsetse flies. Frontiers in Microbiology, 2014, 5, 255.	3.5	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?. Infection, Genetics and Evolution, 2014, 24, 41-56.	2.3	24
10	Genomic transcriptional profiling in LOU/C/Jall rats identifies genes for successful aging. Brain Structure and Function, 2013, 218, 1501-1512.	2.3	12
11	Divergent transcription is associated with promoters of transcriptional regulators. BMC Genomics, 2013, 14, 914.	2.8	95
12	Responder and nonresponder patients exhibit different peripheral transcriptional signatures during major depressive episode. Translational Psychiatry, 2012, 2, e185-e185.	4.8	192
13	Genome-wide analysis of familial dysautonomia and kinetin target genes with patient olfactory ecto-mesenchymal stem cells. Human Mutation, 2012, 33, 530-540.	2.5	25
14	An Evaluation of the Role of Gene Expression in the Prediction and Diagnosis of Ventilator-associated Pneumonia. Anesthesiology, 2011, 115, 344-352.	2.5	13
15	Platelets Alter Gene Expression Profile in Human Brain Endothelial Cells in an In Vitro Model of Cerebral Malaria. PLoS ONE, 2011, 6, e19651.	2.5	32
16	Population transcriptomics: insights from Drosophila simulans, Drosophila sechellia and their hybrids. Genetica, 2011, 139, 465-477.	1.1	13
17	Gene Expression Signature-Based Screening Identifies New Broadly Effective Influenza A Antivirals. PLoS ONE, 2010, 5, e13169.	2.5	59
18	Olfactory Stem Cells, a New Cellular Model for Studying Molecular Mechanisms Underlying Familial Dysautonomia. PLoS ONE, 2010, 5, e15590.	2.5	46

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19	Gene Expression Profiles Characterize Inflammation Stages in the Acute Lung Injury in Mice. PLoS ONE, 2010, 5, e11485.	2.5	23
20	Thrombin-induced endothelial microparticle generation: identification of a novel pathway involving ROCK-II activation by caspase-2. Blood, 2006, 108, 1868-1876.	1.4	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.	0.8	18
22	Gene profiling reveals specific oncogenic mechanisms and signaling pathways in oncocytic and papillary thyroid carcinoma. Oncogene, 2005, 24, 4155-4161.	5.9	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.	3.6	80
24	Gene Expression Profiling for Molecular Characterization of Inflammatory Breast Cancer and Prediction of Response to Chemotherapy. Cancer Research, 2004, 64, 8558-8565.	0.9	177
25	A General Survey of Thymocyte Differentiation by Transcriptional Analysis of Knockout Mouse Models. Journal of Immunology, 2004, 173, 6109-6118.	0.8	27
26	Gene expression profiling of colon cancer by DNA microarrays and correlation with histoclinical parameters. Oncogene, 2004, 23, 1377-1391.	5.9	293
27	Gene expression profiling of 1p35-36 genes in neuroblastoma. Oncogene, 2004, 23, 5912-5922.	5.9	60
28	Feature extraction and signal processing for nylon DNA microarrays. BMC Genomics, 2004, 5, 38.	2.8	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.	3.1	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.	1.4	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.	3.1	10
32	Gene expression profiling of breast carcinomas using Nylon DNA arrays. Comptes Rendus - Biologies, 2003, 326, 1031-1039.	0.2	3
33	Gene expression profiling of multiple myeloma reveals molecular portraits in relation to the pathogenesis of the disease. Blood, 2003, 101, 4998-5006.	1.4	124
34	Gene expression profiles of poor-prognosis primary breast cancer correlate with survival. Human Molecular Genetics, 2002, 11, 863-872.	2.9	117
35	Gene expression profiling defines molecular subtypes of classical Hodgkin's disease. Oncogene, 2002, 21, 3095-3102.	5.9	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.	3.8	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.	2.9	168
38	Expression scanning of an array of growth control genes in human tumor cell lines. Oncogene, 1999, 18, 3905-3912.	5.9	57