Fernanda Bernardi Bertonha

List of Publications by Year
in descending order

17	161	1163117 8	1199594 12
papers	citations	h-index	g-index

18	18	18	255
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Methods for Gene Co-expression Network Visualization and Analysis. , 2022, , 143-163.		2
2	Hippocampal CA3 transcriptional modules associated with granule cell alterations and cognitive impairment in refractory mesial temporal lobe epilepsy patients. Scientific Reports, 2021, 11, 10257.	3.3	7
3	Prevalence of Inflammatory Pathways Over Immuno-Tolerance in Peripheral Blood Mononuclear Cells of Recent-Onset Type 1 Diabetes. Frontiers in Immunology, 2021, 12, 765264.	4.8	8
4	Age-related transcriptional modules and TF-miRNA-mRNA interactions in neonatal and infant human thymus. PLoS ONE, 2020, 15, e0227547.	2.5	8
5	Dynamic Gene Network Analysis of Caco-2 Cell Response to Shiga Toxin-Producing Escherichia coli-Associated Hemolyticâ€“Uremic Syndrome. Microorganisms, 2019, 7, 195.	3.6	12
6	Functional Genomics of the Infant Human Thymus: AIRE and Minipuberty. , 2019, , 235-245.		0
7	1696-P: PBMC of Recent-Onset Patients with Type 1 Diabetes Present a Differential Gene Expression Profile. Diabetes, 2019, 68, .	0.6	0
8	Minipuberty and Sexual Dimorphism in the Infant Human Thymus. Scientific Reports, 2018, 8, 13169.	3.3	17
9	Temporal analysis of hippocampal CA3 gene co-expression networks in a rat model of febrile seizures. DMM Disease Models and Mechanisms, 2017, 11, .	2.4	11
10	Modular transcriptional repertoire and MicroRNA target analyses characterize genomic dysregulation in the thymus of Down syndrome infants. Oncotarget, 2016, 7, 7497-7533.	1.8	19
11	Redes de interaÃ§Ã£o gÃªnica e controle epigenÃ©tico na transiÃ§Ã£o saÃºde-doenÃ§a. , 2015, 94, 223.	0.1	2
12	Community Structure Analysis of Transcriptional Networks Reveals Distinct Molecular Pathways for Early- and Late-Onset Temporal Lobe Epilepsy with Childhood Febrile Seizures. PLoS ONE, 2015, 10, e0128174.	2.5	14
13	PHF21B as a candidate tumor suppressor gene in head and neck squamous cell carcinomas. Molecular Oncology, 2015, 9, 450-462.	4.6	18
14	Methods for Gene Coexpression Network Visualization and Analysis. , 2014, , 79-94.		1
15	Thymus Gene Coexpression Networks: A Comparative Study in Children with and Without Down Syndrome. , 2014, , 123-136.		0
16	Recurrent copy number gains of ACVR1 and corresponding transcript overexpression are associated with survival in head and neck squamous cell carcinomas. Histopathology, 2011, 59, 81-89.	2.9	6
17	Evaluation of estrogen receptor Î± and Î² and progesterone receptor expression and correlation with clinicopathologic factors and proliferative marker Ki-67 in breast cancers. Human Pathology, 2008, 39, 720-730.	2.0	36