## Eduardo Moraes Rego Reis

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genes regulated by DNA methylation are involved in distinct phenotypes during melanoma progression and are prognostic factors for patients. Molecular Oncology, 2022, 16, 1913-1930.	4.6	1
2	Prognostic value of integrin αV expression and localization pattern in invasive breast carcinomas. Neoplasia, 2022, 30, 100803.	5.3	3
3	Annotation and functional characterization of long noncoding RNAs deregulated in pancreatic adenocarcinoma. Cellular Oncology (Dordrecht), 2022, 45, 479-504.	4.4	4
4	Genome-wide promoter methylation profiling in a cellular model of melanoma progression reveals markers of malignancy and metastasis that predict melanoma survival. Clinical Epigenetics, 2022, 14, .	4.1	3
5	Transcriptome of iPSC-derived neuronal cells reveals a module of co-expressed genes consistently associated with autism spectrum disorder. Molecular Psychiatry, 2021, 26, 1589-1605.	7.9	44
6	Noncoding Gene Families of the Human Genome. , 2021, , 139-180.		1
7	Transcriptional signatures underlying dynamic phenotypic switching and novel disease biomarkers in a linear cellular model of melanoma progression. Neoplasia, 2021, 23, 439-455.	5.3	5
8	Gene co-expression and histone modification signatures are associated with melanoma progression, epithelial-to-mesenchymal transition, and metastasis. Clinical Epigenetics, 2020, 12, 127.	4.1	9
9	Aurora A kinase and its activator TPX2 are potential therapeutic targets in KRAS-induced pancreatic cancer. Cellular Oncology (Dordrecht), 2020, 43, 445-460.	4.4	30
10	Short-Term Effects of Sepsis and the Impact of Aging on the Transcriptional Profile of Different Brain Regions. Inflammation, 2019, 42, 1023-1031.	3.8	12
11	Where do we aspire to publish? A position paper on scientific communication in biochemistry and molecular biology. Brazilian Journal of Medical and Biological Research, 2019, 52, e8935.	1.5	1
12	Chromatin Landscape Distinguishes the Genomic Loci of Hundreds of Androgen-Receptor-Associated LincRNAs From the Loci of Non-associated LincRNAs. Frontiers in Genetics, 2018, 9, 132.	2.3	10
13	Evaluating the Stability of mRNAs and Noncoding RNAs. Methods in Molecular Biology, 2017, 1468, 139-153.	0.9	23
14	Insights into the Function of Long Noncoding RNAs in Sepsis Revealed by Gene Co-Expression Network Analysis. Non-coding RNA, 2017, 3, 5.	2.6	30
15	Identification of novel biomarkers associated with poor patient outcomes in invasive breast carcinoma. Tumor Biology, 2016, 37, 13855-13870.	1.8	19
16	Microarray gene expression analysis of neutrophils from elderly septic patients. Genomics Data, 2015, 6, 51-53.	1.3	4
17	Septic Shock in Advanced Age: Transcriptome Analysis Reveals Altered Molecular Signatures in Neutrophil Granulocytes. PLoS ONE, 2015, 10, e0128341.	2.5	27
18	Global analysis of biogenesis, stability and sub-cellular localization of lncRNAs mapping to intragenic regions of the human genome. RNA Biology, 2015, 12, 877-892.	3.1	59

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19	CIP4 promotes metastasis in triple-negative breast cancer and is associated with poor patient prognosis. Oncotarget, 2015, 6, 9397-9408.	1.8	29
20	Loss of Caspase 7 Expression Is Associated With Poor Prognosis in Renal Cell Carcinoma Clear Cell Subtype. Urology, 2013, 82, 974.e1-974.e7.	1.0	9
21	Expression analysis and in silico characterization of intronic long noncoding RNAs in renal cell carcinoma: emerging functional associations. Molecular Cancer, 2013, 12, 140.	19.2	59
22	Transcriptional profiling of Neurospora crassa Δmak-2 reveals that mitogen-activated protein kinase MAK-2 participates in the phosphate signaling pathway. Fungal Genetics and Biology, 2013, 60, 140-149.	2.1	33
23	The Intronic Long Noncoding RNA ANRASSF1 Recruits PRC2 to the RASSF1A Promoter, Reducing the Expression of RASSF1A and Increasing Cell Proliferation. PLoS Genetics, 2013, 9, e1003705.	3.5	180
24	Perspectives of Long Non-Coding RNAs in Cancer Diagnostics. Frontiers in Genetics, 2012, 3, 32.	2.3	131
25	Long noncoding intronic RNAs are differentially expressed in primary and metastatic pancreatic cancer. Molecular Cancer, 2011, 10, 141.	19.2	153
26	Transcriptome analysis of nicotine-exposed cells from the brainstem of neonate spontaneously hypertensive and Wistar Kyoto rats. Pharmacogenomics Journal, 2010, 10, 134-160.	2.0	2
27	A new tick Kunitz type inhibitor, Amblyomin-X, induces tumor cell death by modulating genes related to the cell cycle and targeting the ubiquitin-proteasome system. Toxicon, 2010, 56, 1145-1154.	1.6	58
28	Gene Expression Profiling of Cultured Cells From Brainstem of Newborn Spontaneously Hypertensive and Wistar Kyoto Rats. Cellular and Molecular Neurobiology, 2009, 29, 287-308.	3.3	9
29	Gene expression profiling reveals molecular marker candidates of laryngeal squamous cell carcinoma. Oncology Reports, 2009, , .	2.6	14
30	Identification of proteinâ€coding and intronic noncoding RNAs downâ€regulated in clear cell renal carcinoma. Molecular Carcinogenesis, 2008, 47, 757-767.	2.7	45
31	Detailed molecular characterization of cord blood–derived endothelial progenitors. Experimental Hematology, 2008, 36, 193.e1-193.e15.	0.4	33
32	Conserved tissue expression signatures of intronic noncoding RNAs transcribed from human and mouse loci. Genomics, 2008, 92, 18-25.	2.9	66
33	Splice variants of TLE family genes and up-regulation of a TLE3 isoform in prostate tumors. Biochemical and Biophysical Research Communications, 2007, 364, 918-923.	2.1	16
34	Genome mapping and expression analyses of human intronic noncoding RNAs reveal tissue-specific patterns and enrichment in genes related to regulation of transcription. Genome Biology, 2007, 8, R43.	9.6	209
35	Concepts on Microarray Design for Genome and Transcriptome Analyses. , 2007, , 265-307.		3
36	Apert p.Ser252Trp Mutation in FGFR2 Alters Osteogenic Potential and Gene Expression of Cranial Periosteal Cells. Molecular Medicine, 2007, 13, 422-442.	4.4	28

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37	Androgen responsive intronic non-coding RNAs. BMC Biology, 2007, 5, 4.	3.8	73
38	Auto-antibodies in prostate cancer: Humoral immune response to antigenic determinants coded by the differentially expressed transcripts FLJ23438 and VAMP3. Prostate, 2006, 66, 1463-1473.	2.3	11
39	Evaluation of reference-based two-color methods for measurement of gene expression ratios using spotted cDNA microarrays. BMC Genomics, 2006, 7, 35.	2.8	17
40	Gene expression arrays in cancer research: methods and applications. Critical Reviews in Oncology/Hematology, 2005, 54, 95-105.	4.4	72
41	Large-scale Transcriptome Analyses Reveal New Genetic Marker Candidates of Head, Neck, and Thyroid Cancer. Cancer Research, 2005, 65, 1693-1699.	0.9	55
42	As Antisense RNA Gets Intronic. OMICS A Journal of Integrative Biology, 2005, 9, 2-12.	2.0	37
43	Antisense intronic non-coding RNA levels correlate to the degree of tumor differentiation in prostate cancer. Oncogene, 2004, 23, 6684-6692.	5.9	150
44	RASL11A, member of a novel small monomeric GTPase gene family, is down-regulated in prostate tumors. Biochemical and Biophysical Research Communications, 2004, 316, 618-627.	2.1	29
45	Transcriptome analysis of the acoelomate human parasite Schistosoma mansoni. Nature Genetics, 2003, 35, 148-157.	21.4	433
46	Zerg: a very fast BLAST parser library. Bioinformatics, 2003, 19, 1035-1036.	4.1	17
47	The generation and utilization of a cancer-oriented representation of the human transcriptome by using expressed sequence tags. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 13418-13423.	7.1	105
48	ESTWeb: bioinformatics services for EST sequencing projects. Bioinformatics, 2003, 19, 1587-1588.	4.1	36
49	The contribution of 700,000 ORF sequence tags to the definition of the human transcriptome. Proceedings of the National Academy of Sciences of the United States of America, 2001, 98, 12103-12108.	7.1	123
50	N-terminal chimeric constructs improve the expression of sarcoplasmic reticulum Ca2+-ATPase in yeast. Biochimica Et Biophysica Acta - Biomembranes, 1999, 1461, 83-95.	2.6	11
51	Mitochondrial superoxide dismutase is essential for ethanol tolerance of Saccharomyces cerevisiae in the post-diauxic phase. Microbiology (United Kingdom), 1997, 143, 1649-1656.	1.8	116
52	Heterologous expression of sarcoplasmic reticulum Ca2+-ATPase. Bioscience Reports, 1996, 16, 107-113.	2.4	5
53	Acquisition of Ethanol Tolerance in Saccharomyces cerevisiae: The Key Role of the Mitochondrial Superoxide Dismutase. Archives of Biochemistry and Biophysics, 1993, 300, 608-614.	3.0	75