Andre Luiz Vettore

List of Publications by Year
in descending order

83 papers	4,244 citations	159585 30 h-index	114465 63 g-index
85 all docs	85 docs citations	85 times ranked	7250 citing authors

#	Article	IF	Citations
1	The genome sequence of the plant pathogen Xylella fastidiosa. Nature, 2000, 406, 151-157.	27.8	827
2	Whole-Genome and Epigenomic Landscapes of Etiologically Distinct Subtypes of Cholangiocarcinoma. Cancer Discovery, 2017, 7, 1116-1135.	9.4	637
3	Technical challenges of working with extracellular vesicles. Nanoscale, 2018, 10, 881-906.	5.6	366
4	Analysis and Functional Annotation of an Expressed Sequence Tag Collection for Tropical Crop Sugarcane. Genome Research, 2003, 13, 2725-2735.	5.5	254
5	The TP53 mutation, R337H, is associated with Li-Fraumeni and Li-Fraumeni-like syndromes in Brazilian families. Cancer Letters, 2007, 245, 96-102.	7.2	170
6	The mitochondrial genome of the blowfly Chrysomya chloropyga (Diptera: Calliphoridae). Gene, 2004, 339, 7-15.	2.2	151
7	The libraries that made SUCEST. Genetics and Molecular Biology, 2001, 24, 1-7.	1.3	146
8	The mitochondrial genome of the primary screwworm fly Cochliomyia hominivorax (Diptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 462 T	2.0	129
9	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
10	Prognostic impact of cancer/testis antigen expression in advanced stage multiple myeloma patients. Cancer Immunity, 2008, 8, 2.	3.2	76
11	Mutational landscapes of tongue carcinoma reveal recurrent mutations in genes of therapeutic and prognostic relevance. Genome Medicine, 2015, 7, 98.	8.2	74
12	The involvement of Opaque 2 on Î²-prolamin gene regulation in maize and Coix suggests a more general role for this transcriptional activator. Plant Molecular Biology, 1995, 27, 1015-1029.	3.9	72
13	Aberrant Promoter Methylation of Multiple Genes during Pathogenesis of Bladder Cancer. Cancer Epidemiology Biomarkers and Prevention, 2008, 17, 2786-2794.	2.5	72
14	Expression of Cancer/Testis Antigens is Correlated with Improved Survival in Glioblastoma. Oncotarget, 2013, 4, 636-646.	1.8	54
15	MT1G Hypermethylation: A Potential Prognostic Marker for Hepatoblastoma. Pediatric Research, 2010, 67, 387-393.	2.3	53
16	Prognostic significance of TIMP3 hypermethylation in post-treatment salivary rinse from head and neck squamous cell carcinoma patients. Carcinogenesis, 2013, 34, 20-27.	2.8	52
17	SAGE analysis highlights the importance of p53csv, ddx5, mapkapk2 and ranbp2 to multiple myeloma tumorigenesis. Cancer Letters, 2009, 278, 41-48.	7.2	51
18	Expression of miR-296-5p as predictive marker for radiotherapy resistance in early-stage laryngeal carcinoma. Journal of Translational Medicine, 2015, 13, 262.	4.4	50

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19	TGFÎ²R2 aberrant methylation is a potential prognostic marker and therapeutic target in multiple myeloma. International Journal of Cancer, 2009, 125, 1985-1991.	5.1	48
20	Glioblastomas: correlation between oligodendroglial components, genetic abnormalities, and prognosis. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2008, 452, 481-490.	2.8	46
21	Identification of proteinâ€coding and intronic noncoding RNAs downâ€regulated in clear cell renal carcinoma. Molecular Carcinogenesis, 2008, 47, 757-767.	2.7	45
22	Clinical significance of molecular alterations in histologically negative surgical margins of head and neck cancer patients. Oral Oncology, 2012, 48, 240-248.	1.5	45
23	The molecular and functional characterization of an Opaque2 homologue gene from Coix and a new classification of plant bZIP proteins. Plant Molecular Biology, 1998, 36, 249-263.	3.9	40
24	Aberrant methylation in pediatric myelodysplastic syndrome. Leukemia Research, 2007, 31, 175-181.	0.8	39
25	Evaluation of Monocot and Eudicot Divergence Using the Sugarcane Transcriptome. Plant Physiology, 2004, 134, 951-959.	4.8	38
26	TIMP3 and CCNA1 hypermethylation in HNSCC is associated with an increased incidence of second primary tumors. Journal of Translational Medicine, 2013, 11, 316.	4.4	36
27	Identification of upregulated genes in oral squamous cell carcinomas. Head and Neck, 2013, 35, 1475-1481.	2.0	35
28	Accuracy of microRNAs as markers for the detection of neck lymph node metastases in patients with head and neck squamous cell carcinoma. BMC Medicine, 2015, 13, 108.	5.5	33
29	Epsteinâ€Šâ€“â€ŠBarr viral load, interleukin-6 and interleukin-10 levels in post-transplant lymphoproliferative disease: A nested caseâ€Šâ€“â€Šcontrol study in a renal transplant cohort. Leukemia and Lymphoma, 2005, 46, 533-539.	1.3	32
30	Cancer/Testis Antigen MAGE-C1/CT7: New Target for Multiple Myeloma Therapy. Clinical and Developmental Immunology, 2012, 2012, 1-7.	3.3	30
31	Physical interaction of two cancer-testis antigens, MAGE-C1 (CT7) and NY-ESO-1 (CT6). Cancer Immunity, 2006, 6, 12.	3.2	29
32	Endosperm-preferred Expression of Maize Genes as Revealed by Transcriptome-wide Analysis of Expressed Sequence Tags. Plant Molecular Biology, 2005, 59, 363-374.	3.9	28
33	A Comprehensive Expression Analysis of Cancer Testis Antigens in Head and Neck Squamous Cell Carcinoma Revels MAGEA3/6 as a Marker for Recurrence. Molecular Cancer Therapeutics, 2015, 14, 828-834.	4.1	27
34	The mutational landscape of earlyâ€and typicalâ€onset oral tongue squamous cell carcinoma. Cancer, 2021, 127, 544-553.	4.1	27
35	Exome sequencing reveals recurrent REV3L mutations in cisplatin-resistant squamous cell carcinoma of head and neck. Scientific Reports, 2016, 6, 19552.	3.3	26
36	Claudinâ€7 downâ€regulation is an important feature in oral squamous cell carcinoma. Histopathology, 2010, 57, 689-698.	2.9	23

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37	A preliminary investigation of circulating extracellular vesicles and biomarker discovery associated with treatment response in head and neck squamous cell carcinoma. BMC Cancer, 2019, 19, 373.	2.6	20
38	Hypermethylation of CpG island in the promoter region of CALCA in acute lymphoblastic leukemia with central nervous system (CNS) infiltration correlates with poorer prognosis. Leukemia Research, 2006, 30, 891-894.	0.8	18
39	Search for mutations in signaling pathways in head and neck squamous cell carcinoma. Oncology Reports, 2013, 30, 334-340.	2.6	18
40	MicroRNA-1252-5p Associated with Extracellular Vesicles Enhances Bortezomib Sensitivity in Multiple Myeloma Cells by Targeting Heparanase. OncoTargets and Therapy, 2021, Volume 14, 455-467.	2.0	16
41	Cooperative DNA Binding and Sequence Discrimination by the Opaque2 bZIP Factor. Plant Cell, 1998, 10, 1941-1955.	6.6	15
42	Evaluation of the methylation profile of exfoliated cell samples from patients with head and neck squamous cell carcinoma. Head and Neck, 2014, 36, 631-637.	2.0	15
43	Aberrant DNA methylation of ESR1 and p14ARF genes could be useful as prognostic indicators in osteosarcoma. OncoTargets and Therapy, 2013, 6, 713.	2.0	14
44	Clinical correlations and prognostic relevance of HGF, VEGF AND FGF expression in Brazilian patients with non-Hodgkin lymphoma. Leukemia and Lymphoma, 2008, 49, 257-264.	1.3	13
45	Frequency and prognostic relevance of cancer testis antigen 45 expression inÂmultiple myeloma. Experimental Hematology, 2009, 37, 446-449.	0.4	13
46	Expression of eight genes of nuclear factor-kappa B pathway in multiple myeloma using bone marrow aspirates obtained at diagnosis. Histology and Histopathology, 2009, 24, 991-7.	0.7	13
47	The Prolamins of Sorghum, Coix and Millets. , 1999, , 141-157.		12
48	Overexpression, purification, biochemical characterization, and molecular modeling of recombinant GDP-mannosyltransferase (GumH) from Xylella fastidiosa. Biochemical and Biophysical Research Communications, 2004, 315, 485-492.	2.1	11
49	Expression and Prognostic Relevance of GAGE1 and XAGE1 Cancer/Testis Antigens in Head and Neck Squamous Cell Carcinoma. Current Molecular Medicine, 2018, 17, 707-717.	1.3	10
50	Quantification of Epstein-Barr viral load and determination of a cut-off value to predict the risk of post-transplant lymphoproliferative disease in a renal transplant cohort. Haematologica, 2004, 89, 366-8.	3.5	10
51	Circulating extracellular vesicle-associated TGFÎ²3 modulates response to cytotoxic therapy in head and neck squamous cell carcinoma. Carcinogenesis, 2019, 40, 1452-1461.	2.8	9
52	Differential expression of apoptosis related proteins and nitric oxide synthases in Epstein Barr associated gastric carcinomas. World Journal of Gastroenterology, 2006, 12, 4959.	3.3	8
53	Overexpression, purification, and biochemical characterization of GumC, an enzyme involved in the biosynthesis of exopolysaccharide by Xylella fastidiosa. Protein Expression and Purification, 2004, 34, 223-228.	1.3	6
54	Number of expressed cancer/testis antigens identifies focal adhesion pathway genes as possible targets for multiple myeloma therapy. Leukemia and Lymphoma, 2010, 51, 1543-1549.	1.3	6

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55	In vitro and in silico validation of CA3 and FHL1 downregulation in oral cancer. BMC Cancer, 2018, 18, 193.	2.6	6
56	Assessment of the cytotoxic effects of aporphine prototypes on head and neck cancer cells. Investigational New Drugs, 2020, 38, 70-78.	2.6	6
57	Prognostic Impact of Cancer Testis Antigens Expression in Advanced Stage Multiple Myeloma Patients.. Blood, 2007, 110, 4733-4733.	1.4	6
58	Identification of a DNA-binding factor that recognizes an alpha-coixin promoter and interacts with a Coix Opaque-2 like protein. Plant Molecular Biology, 1999, 39, 95-104.	3.9	4
59	OIP5 Expression Sensitize Glioblastoma Cells to Lomustine Treatment. Journal of Molecular Neuroscience, 2018, 66, 383-389.	2.3	4
60	Evaluation of acetylation and methylation in oral rinse of patients with head and neck cancer history exposed to valproic acid. Scientific Reports, 2021, 11, 16415.	3.3	4
61	Overexpression of CTLA-4 in the Bone Marrow of Patients with Multiple Myeloma As a Sign of Local Accumulation of Immunosuppressive Tregs â€“ Perspectives for Novel Treatment Strategies. Blood, 2011, 118, 1829-1829.	1.4	4
62	Comparative Expression of a Set of Genes to an Internal Housekeeping Control in CDNA Amplified and not Amplified by PolyAPCR in Non-HodgkinÊ¼s Lymphoma Samples Obtained From Fine-Needle Aspiration Cytology. Diagnostic Molecular Pathology, 2010, 19, 40-44.	2.1	3
63	High expression of MLANA in the plasma of patients with head and neck squamous cell carcinoma as a predictor of tumor progression. Head and Neck, 2019, 41, 1199-1205.	2.0	3
64	Cooperative DNA Binding and Sequence Discrimination by the Opaque2 bZIP Factor. Plant Cell, 1998, 10, 1941.	6.6	2
65	Downregulation of DCC sensitizes multiple myeloma cells to bortezomib treatment. Molecular Medicine Reports, 2019, 19, 5023-5029.	2.4	2
66	Response to â€œGermline TP53 R337H mutation is not sufficient to establish Li-Fraumeni or Li-Fraumeni-like syndromeâ€, by Ribeiro et al.. Cancer Letters, 2007, 247, 356-358.	7.2	1
67	PP033. Oral Oncology, 2013, 49, S104-S105.	1.5	0
68	Frequent Expression of Cancer/Testis Antigens CT7 and LAGE-1 in Advanced Stage Multiple Myeloma: Are They Possible Targets for Immunotherapy?.. Blood, 2006, 108, 5034-5034.	1.4	0
69	Expression of Nuclear Factor-kappa B Pathway Genes and Their Correlation with Clinical Features in Multiple Myeloma.. Blood, 2007, 110, 4735-4735.	1.4	0
70	Clinical, Prognostic and Possible Therapeutic Relevance of Angiogenesis in Non-Hodgkinâ€™s Lymphoma.. Blood, 2007, 110, 3583-3583.	1.4	0
71	Abstract 2954: Overexpression of specific genes in surgical margins of head and neck squamous cell carcinoma patients may predict a significantly increased risk of recurrence. , 2010, , .		0
72	Abstract 4910: Identification of putative epigenetic markers for head and neck squamous cell carinoma recurrence. , 2010, , .		0

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73	Abstract 4808: Detection of aberrant DNA methylation in saliva samples as a predictor of recurrence in head and neck squamous cell carcinoma patients. , 2011, , .		0
74	Abstract 2104: Molecular alterations in JAK1 and JAK2 genes in head and neck squamous cell carcinoma. , 2012, , .		0
75	Abstract 5050: MicroRNAs profiling in salivary rinse from patients with head and neck squamous cells carcinoma. , 2012, , .		0
76	Abstract 1142: The expression profile of cancer/testis antigens (CTAs) in head and neck cancer. , 2012, , .		0
77	Abstract 1481: miR-296 as prognostic and predictive molecular marker for recurrence in early-stage laryngeal carcinoma treated with definitive radiotherapy. , 2014, , .		0
78	Abstract 1495: Identification of markers for the presence of lymph nodes metastasis in patients with oral squamous cell carcinomas. , 2014, , .		0
79	Abstract 2457: Functional Study of DCC Gene in multiple myeloma mell lines. , 2014, , .		0
80	Abstract 3953: HORMAD1 plays an important role in the HNSCC carcinogenesis. , 2015, , .		0
81	Abstract 3874: Mutational landscapes of oral tongue squamous cell carcinoma reveal recurrent mutations in genes of therapeutic and prognostic relevance. , 2015, , .		0
82	Abstract 3968: Identification of microRNAs markers in patients with oral squamous cell carcinomas for the presence of lymph nodes metastasis. , 2015, , .		0
83	Abstract 3150: Analysis of the extracellular vesicles content present in the plasma of patients with head and neck squamous cell carcinoma for identification of molecular markers for treatment response. , 2016, , .		0