

Andre Luiz Vettore

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1689739/publications.pdf>

Version: 2024-02-01

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 mutational landscape of early and typical onset oral tongue squamous cell carcinoma. <i>Cancer</i> , 2021, 127, 544-553.	4.1	27
2	MicroRNA-1252-5p Associated with Extracellular Vesicles Enhances Bortezomib Sensitivity in Multiple Myeloma Cells by Targeting Heparanase. <i>OncoTargets and Therapy</i> , 2021, Volume 14, 455-467.	2.0	16
3	Evaluation of acetylation and methylation in oral rinse of patients with head and neck cancer history exposed to valproic acid. <i>Scientific Reports</i> , 2021, 11, 16415.	3.3	4
4	Assessment of the cytotoxic effects of aporphine prototypes on head and neck cancer cells. <i>Investigational New Drugs</i> , 2020, 38, 70-78.	2.6	6
5	Circulating extracellular vesicle-associated TGF β 3 modulates response to cytotoxic therapy in head and neck squamous cell carcinoma. <i>Carcinogenesis</i> , 2019, 40, 1452-1461.	2.8	9
6	A preliminary investigation of circulating extracellular vesicles and biomarker discovery associated with treatment response in head and neck squamous cell carcinoma. <i>BMC Cancer</i> , 2019, 19, 373.	2.6	20
7	High expression of MLANA in the plasma of patients with head and neck squamous cell carcinoma as a predictor of tumor progression. <i>Head and Neck</i> , 2019, 41, 1199-1205.	2.0	3
8	Downregulation of DCC sensitizes multiple myeloma cells to bortezomib treatment. <i>Molecular Medicine Reports</i> , 2019, 19, 5023-5029.	2.4	2
9	In vitro and in silico validation of CA3 and FHL1 downregulation in oral cancer. <i>BMC Cancer</i> , 2018, 18, 193.	2.6	6
10	Technical challenges of working with extracellular vesicles. <i>Nanoscale</i> , 2018, 10, 881-906.	5.6	366
11	OIP5 Expression Sensitize Glioblastoma Cells to Lomustine Treatment. <i>Journal of Molecular Neuroscience</i> , 2018, 66, 383-389.	2.3	4
12	Expression and Prognostic Relevance of GAGE1 and XAGE1 Cancer/Testis Antigens in Head and Neck Squamous Cell Carcinoma. <i>Current Molecular Medicine</i> , 2018, 17, 707-717.	1.3	10
13	Whole-Genome and Epigenomic Landscapes of Etiologically Distinct Subtypes of Cholangiocarcinoma. <i>Cancer Discovery</i> , 2017, 7, 1116-1135.	9.4	637
14	Exome sequencing reveals recurrent REV3L mutations in cisplatin-resistant squamous cell carcinoma of head and neck. <i>Scientific Reports</i> , 2016, 6, 19552.	3.3	26
15	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
16	Mutational landscapes of tongue carcinoma reveal recurrent mutations in genes of therapeutic and prognostic relevance. <i>Genome Medicine</i> , 2015, 7, 98.	8.2	74
17	Expression of miR-296-5p as predictive marker for radiotherapy resistance in early-stage laryngeal carcinoma. <i>Journal of Translational Medicine</i> , 2015, 13, 262.	4.4	50
18	A Comprehensive Expression Analysis of Cancer Testis Antigens in Head and Neck Squamous Cell Carcinoma Reveals <i>MAGEA3/6</i> as a Marker for Recurrence. <i>Molecular Cancer Therapeutics</i> , 2015, 14, 828-834.	4.1	27

#	ARTICLE	IF	CITATIONS
19	Accuracy of microRNAs as markers for the detection of neck lymph node metastases in patients with head and neck squamous cell carcinoma. <i>BMC Medicine</i> , 2015, 13, 108.	5.5	33
20	Abstract 3953: HORMAD1 plays an important role in the HNSCC carcinogenesis. , 2015, , .		0
21	Abstract 3874: Mutational landscapes of oral tongue squamous cell carcinoma reveal recurrent mutations in genes of therapeutic and prognostic relevance. , 2015, , .		0
22	Abstract 3968: Identification of microRNAs markers in patients with oral squamous cell carcinomas for the presence of lymph nodes metastasis. , 2015, , .		0
23	Evaluation of the methylation profile of exfoliated cell samples from patients with head and neck squamous cell carcinoma. <i>Head and Neck</i> , 2014, 36, 631-637.	2.0	15
24	Abstract 1481: miR-296 as prognostic and predictive molecular marker for recurrence in early-stage laryngeal carcinoma treated with definitive radiotherapy. , 2014, , .		0
25	Abstract 1495: Identification of markers for the presence of lymph nodes metastasis in patients with oral squamous cell carcinomas. , 2014, , .		0
26	Abstract 2457: Functional Study of DCC Gene in multiple myeloma mell lines. , 2014, , .		0
27	Identification of upregulated genes in oral squamous cell carcinomas. <i>Head and Neck</i> , 2013, 35, 1475-1481.	2.0	35
28	TIMP3 and CCNA1 hypermethylation in HNSCC is associated with an increased incidence of second primary tumors. <i>Journal of Translational Medicine</i> , 2013, 11, 316.	4.4	36
29	Prognostic significance of TIMP3 hypermethylation in post-treatment salivary rinse from head and neck squamous cell carcinoma patients. <i>Carcinogenesis</i> , 2013, 34, 20-27.	2.8	52
30	PP033. <i>Oral Oncology</i> , 2013, 49, S104-S105.	1.5	0
31	Search for mutations in signaling pathways in head and neck squamous cell carcinoma. <i>Oncology Reports</i> , 2013, 30, 334-340.	2.6	18
32	Aberrant DNA methylation of ESR1 and p14ARF genes could be useful as prognostic indicators in osteosarcoma. <i>OncoTargets and Therapy</i> , 2013, 6, 713.	2.0	14
33	Expression of Cancer/Testis Antigens is Correlated with Improved Survival in Glioblastoma. <i>Oncotarget</i> , 2013, 4, 636-646.	1.8	54
34	Cancer/Testis Antigen MAGE-C1/CT7: New Target for Multiple Myeloma Therapy. <i>Clinical and Developmental Immunology</i> , 2012, 2012, 1-7.	3.3	30
35	Clinical significance of molecular alterations in histologically negative surgical margins of head and neck cancer patients. <i>Oral Oncology</i> , 2012, 48, 240-248.	1.5	45
36	Abstract 2104: Molecular alterations in JAK1 and JAK2 genes in head and neck squamous cell carcinoma. , 2012, , .		0

#	ARTICLE	IF	CITATIONS
37	Abstract 5050: MicroRNAs profiling in salivary rinse from patients with head and neck squamous cells carcinoma. , 2012, , .		0
38	Abstract 1142: The expression profile of cancer/testis antigens (CTAs) in head and neck cancer. , 2012, , .		0
39	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
40	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
41	Claudinâ€7 downâ€regulation is an important feature in oral squamous cell carcinoma. Histopathology, 2010, 57, 689-698.	2.9	23
42	Comparative Expression of a Set of Genes to an Internal Housekeeping Control in CDNA Amplified and not Amplified by PolyAPCR in Non-Hodgkinâ€™s Lymphoma Samples Obtained From Fine-Needle Aspiration Cytology. Diagnostic Molecular Pathology, 2010, 19, 40-44.	2.1	3
43	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
44	MT1G Hypermethylation: A Potential Prognostic Marker for Hepatoblastoma. Pediatric Research, 2010, 67, 387-393.	2.3	53
45	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
46	Abstract 4910: Identification of putative epigenetic markers for head and neck squamous cell carcinoma recurrence. , 2010, , .		0
47	Frequency and prognostic relevance of cancer testis antigen 45 expression inâ€multiple myeloma. Experimental Hematology, 2009, 37, 446-449.	0.4	13
48	<i>TGFâ€R2</i> aberrant methylation is a potential prognostic marker and therapeutic target in multiple myeloma. International Journal of Cancer, 2009, 125, 1985-1991.	5.1	48
49	SAGE analysis highlights the importance of p53, ddx5, mapkapk2 and ranbp2 to multiple myeloma tumorigenesis. Cancer Letters, 2009, 278, 41-48.	7.2	51
50	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
51	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
52	Identification of proteinâ€coding and intronic noncoding RNAs downâ€regulated in clear cell renal carcinoma. Molecular Carcinogenesis, 2008, 47, 757-767.	2.7	45
53	Aberrant Promoter Methylation of Multiple Genes during Pathogenesis of Bladder Cancer. Cancer Epidemiology Biomarkers and Prevention, 2008, 17, 2786-2794.	2.5	72
54	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

#	ARTICLE	IF	CITATIONS
55	Prognostic impact of cancer/testis antigen expression in advanced stage multiple myeloma patients. <i>Cancer Immunity</i> , 2008, 8, 2.	3.2	76
56	The TP53 mutation, R337H, is associated with Li-Fraumeni and Li-Fraumeni-like syndromes in Brazilian families. <i>Cancer Letters</i> , 2007, 245, 96-102.	7.2	170
57	Response to germline TP53 R337H mutation is not sufficient to establish Li-Fraumeni or Li-Fraumeni-like syndrome, by Ribeiro et al.. <i>Cancer Letters</i> , 2007, 247, 356-358.	7.2	1
58	Aberrant methylation in pediatric myelodysplastic syndrome. <i>Leukemia Research</i> , 2007, 31, 175-181.	0.8	39
59	Expression of Nuclear Factor-kappa B Pathway Genes and Their Correlation with Clinical Features in Multiple Myeloma.. <i>Blood</i> , 2007, 110, 4735-4735.	1.4	0
60	Prognostic Impact of Cancer Testis Antigens Expression in Advanced Stage Multiple Myeloma Patients.. <i>Blood</i> , 2007, 110, 4733-4733.	1.4	6
61	Clinical, Prognostic and Possible Therapeutic Relevance of Angiogenesis in Non-Hodgkin's Lymphoma.. <i>Blood</i> , 2007, 110, 3583-3583.	1.4	0
62	Hypermethylation of CpG island in the promoter region of CALCA in acute lymphoblastic leukemia with central nervous system (CNS) infiltration correlates with poorer prognosis. <i>Leukemia Research</i> , 2006, 30, 891-894.	0.8	18
63	Differential expression of apoptosis related proteins and nitric oxide synthases in Epstein Barr associated gastric carcinomas. <i>World Journal of Gastroenterology</i> , 2006, 12, 4959.	3.3	8
64	Frequent Expression of Cancer/Testis Antigens CT7 and LAGE-1 in Advanced Stage Multiple Myeloma: Are They Possible Targets for Immunotherapy?.. <i>Blood</i> , 2006, 108, 5034-5034.	1.4	0
65	Physical interaction of two cancer-testis antigens, MAGE-C1 (CT7) and NY-ESO-1 (CT6). <i>Cancer Immunity</i> , 2006, 6, 12.	3.2	29
66	Endosperm-preferred Expression of Maize Genes as Revealed by Transcriptome-wide Analysis of Expressed Sequence Tags. <i>Plant Molecular Biology</i> , 2005, 59, 363-374.	3.9	28
67	Epstein-Barr viral load, interleukin-6 and interleukin-10 levels in post-transplant lymphoproliferative disease: A nested case-control study in a renal transplant cohort. <i>Leukemia and Lymphoma</i> , 2005, 46, 533-539.	1.3	32
68	Evaluation of Monocot and Eudicot Divergence Using the Sugarcane Transcriptome. <i>Plant Physiology</i> , 2004, 134, 951-959.	4.8	38
69	The mitochondrial genome of the blowfly <i>Chrysomya chloropyga</i> (Diptera: Calliphoridae). <i>Gene</i> , 2004, 339, 7-15.	2.2	151
70	Overexpression, purification, biochemical characterization, and molecular modeling of recombinant GDP-mannosyltransferase (GumH) from <i>Xylella fastidiosa</i> . <i>Biochemical and Biophysical Research Communications</i> , 2004, 315, 485-492.	2.1	11
71	Overexpression, purification, and biochemical characterization of GumC, an enzyme involved in the biosynthesis of exopolysaccharide by <i>Xylella fastidiosa</i> . <i>Protein Expression and Purification</i> , 2004, 34, 223-228.	1.3	6
72	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. <i>Haematologica</i> , 2004, 89, 366-8.	3.5	10

#	ARTICLE	IF	CITATIONS
73	Analysis and Functional Annotation of an Expressed Sequence Tag Collection for Tropical Crop Sugarcane. <i>Genome Research</i> , 2003, 13, 2725-2735.	5.5	254
74	The generation and utilization of a cancer-oriented representation of the human transcriptome by using expressed sequence tags. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 13418-13423.	7.1	105
75	The libraries that made SUCEST. <i>Genetics and Molecular Biology</i> , 2001, 24, 1-7.	1.3	146
76	The mitochondrial genome of the primary screwworm fly <i>Cochliomyia hominivorax</i> (Diptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 622 T	2.0	129
77	The genome sequence of the plant pathogen <i>Xylella fastidiosa</i> . <i>Nature</i> , 2000, 406, 151-157.	27.8	827
78	The Prolamins of Sorghum, Coix and Millets. , 1999, , 141-157.		12
79	Identification of a DNA-binding factor that recognizes an alpha-coixin promoter and interacts with a Coix Opaque-2 like protein. <i>Plant Molecular Biology</i> , 1999, 39, 95-104.	3.9	4
80	The molecular and functional characterization of an Opaque2 homologue gene from Coix and a new classification of plant bZIP proteins. <i>Plant Molecular Biology</i> , 1998, 36, 249-263.	3.9	40
81	Cooperative DNA Binding and Sequence Discrimination by the Opaque2 bZIP Factor. <i>Plant Cell</i> , 1998, 10, 1941-1955.	6.6	15
82	Cooperative DNA Binding and Sequence Discrimination by the Opaque2 bZIP Factor. <i>Plant Cell</i> , 1998, 10, 1941.	6.6	2
83	The involvement of Opaque 2 on \hat{I}^2 -prolamin gene regulation in maize and Coix suggests a more general role for this transcriptional activator. <i>Plant Molecular Biology</i> , 1995, 27, 1015-1029.	3.9	72