Gustavo Jacob LourenÃ§o

List of Publications by Year
in descending order

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92	625	687363 13	752698 20
papers	citations	h-index	g-index

96	96	96	888
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Reflections on the COVID-19 Pandemic: Experiences of a Brazilian Cancer Center. Journal of Social Work in End-of-Life and Palliative Care, 2022, 18, 12-16.	0.6	0
2	Addressing psychiatric disorders and genetics: the meaningful use of comics for health information. Journal of Visual Communication in Medicine, 2022, , 1-6.	0.6	0
3	Demographic history differences between Hispanics and Brazilians imprint haplotype features. G3: Genes, Genomes, Genetics, 2022, 12, .	1.8	1
4	<scp>microRNAs</scp> deregulation in head and neck squamous cell carcinoma. Head and Neck, 2021, 43, 645-667.	2.0	12
5	GSTP1andABCB1Polymorphisms Predicting Toxicities and Clinical Management on Carboplatin and Paclitaxelâ€Based Chemotherapy in Ovarian Cancer. Clinical and Translational Science, 2021, 14, 720-728.	3.1	15
6	Single nucleotide variants in immune-response genes and the tumor microenvironment composition predict progression of mantle cell lymphoma. BMC Cancer, 2021, 21, 209.	2.6	5
7	Influence of IL1B (rs16944) and IL1R2 (rs4141134) polymorphisms on aggressiveness and prognosis of cutaneous melanoma. Melanoma Research, 2021, 31, 476-481.	1.2	6
8	Pseudogene Transcripts in Head and Neck Cancer: Literature Review and In Silico Analysis. Genes, 2021, 12, 1254.	2.4	5
9	Intronic variants of MITF (rs7623610) and CREB1 (rs10932201) genes may enhance splicing efficiency in human melanoma cell line. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2021, 823, 111763.	1.0	2
10	Influence of Sociodemographic Characteristics and Inflammation-Related Gene Variants on the Comfort Level of Caregivers of Patients With Head and Neck Cancer. Journal of Holistic Nursing, 2021, , 089801012110467.	1.6	1
11	Interplay between the Mediterranean diet and C-reactive protein genetic polymorphisms towards inflammation in adolescents. Clinical Nutrition, 2020, 39, 1919-1926.	5.0	16
12	Single-nucleotide variants in TGFB1, TGFBR2, IL17A, and IL17F immune response genes contribute to follicular lymphoma susceptibility and aggressiveness. Blood Cancer Journal, 2020, 10, 97.	6.2	3
13	Role of a genetic variation in the microRNA-4421 binding site of ERP29 regarding risk of oropharynx cancer and prognosis. Scientific Reports, 2020, 10, 17039.	3.3	8
14	Inherited variations in human pigmentation-related genes modulate cutaneous melanoma risk and clinicopathological features in Brazilian population. Scientific Reports, 2020, 10, 12129.	3.3	6
15	Influence of CASP9 c.â€1339A>G and CASP3 c.â€1191A>G variants in outcome of patients with head and neck squamous cell carcinoma. Journal of Oral Pathology and Medicine, 2020, 49, 1078-1083.	2.7	1
16	The association of a single-nucleotide variant in the microRNA-146a with advanced colorectal cancer prognosis. Tumor Biology, 2020, 42, 101042832092385.	1.8	5
17	An integrative microenvironment approach for follicular lymphoma: roles of inflammatory cell subsets and immune-response polymorphisms on disease clinical course. Oncotarget, 2020, 11, 3153-3173.	1.8	4
18	Association between polymorphisms in angiogenesisâ€related genes and the prognosis of classical Hodgkin lymphoma. British Journal of Haematology, 2019, 185, 366-370.	2.5	0

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19	Increased risk of Hodgkin lymphoma in males with inherited T lymphocyte receptor programed death-1 deficiency. Leukemia and Lymphoma, 2019, 60, 3552-3556.	1.3	2
20	GSTM1, GSTT1 and GSTP1 Ile105Val polymorphisms in outcomes of head and neck squamous cell carcinoma patients treated with cisplatin chemoradiation. Scientific Reports, 2019, 9, 9312.	3.3	12
21	Variants in the JAK1 and JAK2 genes in the risk and prognosis of patients with cutaneous melanoma. Annals of Oncology, 2019, 30, v14.	1.2	0
22	Modulation of risk of cutaneous melanoma patients by variants in STAT3 gene and functional analysis. Annals of Oncology, 2019, 30, v14-v15.	1.2	0
23	Cytokine genetic variations and worse quality of life among family caregivers of head and neck cancer patients in palliative care. Annals of Oncology, 2019, 30, v668.	1.2	0
24	Associations ofVEGFAandKDRsingle-nucleotide polymorphisms and increased risk and aggressiveness of high-grade gliomas. Tumor Biology, 2019, 41, 101042831987209.	1.8	13
25	CASP8 (rs3834129) and CASP3 (rs4647601) polymorphisms in oropharynx cancer risk, tumor cell differentiation, and prognosis in a cohort of the Brazilian population. Molecular Biology Reports, 2019, 46, 6557-6563.	2.3	3
26	Cost-minimization analysis of GSTP1c.313A>G genotyping for the prevention of cisplatin-induced nausea and vomiting: A Bayesian inference approach. PLoS ONE, 2019, 14, e0213929.	2.5	3
27	Influence of functional variants Asp312Asn and Lys751Gln of Xeroderma Pigmentosum Group D (XPD) and Glutathione Sâ€transferase Mu 1 (GSTM 1) and Theta 1 (GSTT 1) genes on cutaneous melanoma susceptibility and prognosis. Experimental Dermatology, 2019, 28, 631-635.	2.9	6
28	CASP9 c.â€1339A>G and CASP3 c.â€1191A>G polymorphisms alter susceptibility and clinical aspects of head and neck squamous cell carcinoma. Head and Neck, 2019, 41, 2665-2670.	2.0	5
29	Role for <scp>DNA</scp> baseâ€excision repair gene variants in the prognosis of Hodgkin lymphoma. British Journal of Haematology, 2019, 186, 171-175.	2.5	0
30	PDCD1 gene polymorphisms as regulators of Tâ€lymphocyte activity in cutaneous melanoma risk and prognosis. Pigment Cell and Melanoma Research, 2018, 31, 308-317.	3.3	12
31	Polymorphisms in DNA mismatch repair pathway genes predict toxicity and response to cisplatin chemoradiation in head and neck squamous cell carcinoma patients. Oncotarget, 2018, 9, 29538-29547.	1.8	25
32	Polymorphisms in apoptosis-related genes in cutaneous melanoma prognosis: sex disparity. Medical Oncology, 2017, 34, 19.	2.5	4
33	Through translational prospective study, the GSTP1 Ile105Val polymorphism emerges as prognostic marker in de novo large B-cell lymphoma patients. Blood Cancer Journal, 2017, 7, e560-e560.	6.2	0
34	<scp>GSTP</scp>1 c.313A>G, <scp>XPD</scp> c.934G>A, <scp>XPF</scp> c.2505T>C and <scp>CASP</scp>9 c.â€1339A>G Polymorphisms and Severity of Vomiting in Head and Neck Cancer Patients treated with Cisplatin Chemoradiation. Basic and Clinical Pharmacology and Toxicology, 2017, 121, 520-525.	2.5	11
35	VEGF, VEGFR2 and GSTM1 polymorphisms in outcome of multiple myeloma patients treated with thalidomide-based regimens. Blood Cancer Journal, 2017, 7, e580-e580.	6.2	7
36	Modulation of Risk and Prognosis of Cutaneous Melanoma Patients by Genetic Polymorphisms on PDCD1 Gene. Annals of Oncology, 2017, 28, v436.	1.2	1

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37	Influence of an intronic polymorphism in the MITF gene, of melanogenic pathway, in the risk and the prognosis of cutaneous melanoma. Annals of Oncology, 2017, 28, v437.	1.2	0
38	Role of an intronic polymorphism in the CREB1 gene, involved in melanogenesis, with the risk and the aggressiveness of cutaneous melanoma. Annals of Oncology, 2017, 28, v436-v437.	1.2	1
39	Influence of FASL and FAS polymorphisms, enrolled in extrinsic apoptosis pathway, in the inherited increased risk of head and neck squamous cell carcinoma. Annals of Oncology, 2016, 27, vi344.	1.2	0
40	Association between polymorphisms in genes related to DNA base-excision repair with risk and prognosis of oropharyngeal squamous cell carcinoma. Journal of Cancer Research and Clinical Oncology, 2016, 142, 1917-1926.	2.5	19
41	Association between polymorphisms in xenobiotic detoxification-related genes with prognosis of epithelial ovarian cancer. Medical Oncology, 2016, 33, 112.	2.5	4
42	XPC (A2920C), XPF (T30028C), TP53 (Arg72Pro), and GSTP1 (Ile105Val) polymorphisms in prognosis of cutaneous melanoma. Tumor Biology, 2016, 37, 3163-3171.	1.8	2
43	Association of FASL and FAS polymorphisms, enrolled in extrinsic apoptosis pathway, with head and neck squamous cell carcinoma risk and outcomes.. Journal of Clinical Oncology, 2016, 34, 6053-6053.	1.6	0
44	OGG1, APEX1 and XRCC1 polymorphisms in oropharyngeal squamous cell carcinoma risk and prognosis.. Journal of Clinical Oncology, 2016, 34, e17515-e17515.	1.6	0
45	XPD and ERCC1 polymorphisms of the nucleotide excision repair pathway in outcome of larynx squamous cell carcinoma patients.. Journal of Clinical Oncology, 2016, 34, e17523-e17523.	1.6	0
46	VEGF, VEGFR2 and GSTM1 polymorphisms in Outcome of Multiple Myeloma Patients in the Thalidomide Era. Blood, 2016, 128, 4457-4457.	1.4	0
47	3312 XPC (A2920C), XPF (T30028C), TP53 (Arg72Pro) and GSTP1 (Ile105Val) Polymorphisms in Prognosis of Patients with Cutaneous Melanoma. European Journal of Cancer, 2015, 51, S668.	2.8	0
48	Dietary risk factors for colorectal cancer in Brazil: a case control study. Nutrition Journal, 2015, 15, 20.	3.4	13
49	Association between genetic polymorphisms in DNA mismatch repair-related genes with risk and prognosis of head and neck squamous cell carcinoma. International Journal of Cancer, 2015, 137, 810-818.	5.1	47
50	Vascular endothelial growth factor (VEGF) polymorphism and increased risk of epithelial ovarian cancer. Journal of Cancer Research and Clinical Oncology, 2015, 141, 69-73.	2.5	13
51	Association of CYP1A1 A4889G and T6235C polymorphisms with the risk of sporadic breast cancer in Brazilian women. Clinics, 2015, 70, 680-685.	1.5	14
52	XPC, XPF, TP53 and GSTP1 polymorphisms in prognosis of cutaneous melanoma patients.. Journal of Clinical Oncology, 2015, 33, 9038-9038.	1.6	0
53	MLH1, MSH2, MSH3 and EXO1 polymorphisms and head and neck squamous cell carcinoma risk and prognosis.. Journal of Clinical Oncology, 2015, 33, 6063-6063.	1.6	1
54	Association between genetic polymorphisms in apoptosis-related genes and risk of cutaneous melanoma in women and men. Journal of Dermatological Science, 2014, 74, 135-141.	1.9	8

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55	Associations of VEGF and VEGFR2 polymorphisms with increased risk and aggressiveness of multiple myeloma. Annals of Hematology, 2014, 93, 1363-9.	1.8	15
56	ERP29 genetic polymorphism and breast cancer susceptibility and prognosis.. Journal of Clinical Oncology, 2014, 32, 584-584.	1.6	0
57	Polymorphisms in the apoptosis pathway and prognosis in cutaneous melanoma.. Journal of Clinical Oncology, 2014, 32, 9084-9084.	1.6	0
58	Assessment of the XPC (A2920C), XPF (T30028C), TP53 (Arg72Pro) and GSTP1 (Ile105Val) polymorphisms in the risk of cutaneous melanoma. Journal of Cancer Research and Clinical Oncology, 2013, 139, 1199-1206.	2.5	14
59	Clinical effects of A4889G and T6235C polymorphisms in cytochrome P-450 CYP1A1 for breast cancer patients treated with tamoxifen: implications for tumor aggressiveness and patient survival. Cancer Chemotherapy and Pharmacology, 2013, 72, 529-535.	2.3	5
60	Association of polymorphisms in genes related to cell cycle (ERP29, LEF1, MCC and PTCH1) and DNA transcription factors (IKBKAP and ZNF415) with base of tongue squamous cell carcinoma risk.. Journal of Clinical Oncology, 2013, 31, 6073-6073.	1.6	0
61	Association of CYP1A1 A4889G and T6235C polymorphisms with the risk of breast cancer in Brazilian women.. Journal of Clinical Oncology, 2013, 31, e11551-e11551.	1.6	0
62	Associations Of VEGF and VEGFR2 Polymorphisms With Increased Risk and Aggressiveness Of Multiple Myeloma. Blood, 2013, 122, 1886-1886.	1.4	0
63	Polymorphisms of Glutathione S-Transferase Mu 1 (GSTM1), Theta 1 (GSTT1), and Pi 1 (GSTP1) Genes and Epithelial Ovarian Cancer Risk. Disease Markers, 2012, 33, 155-159.	1.3	21
64	Inherited Abnormalities in Genes That During the Apoptosis Process and Cutaneous Melanoma Risk. Annals of Oncology, 2012, 23, ix365.	1.2	0
65	Association of CYP1A1 A4889G and T6235C Polymorphisms With Increased Risk and Aggressiveness of Breast Cancer. Annals of Oncology, 2012, 23, ix133.	1.2	1
66	Genome-Wide Association Study of Base of Tongue Squamous Cell Carcinoma Risk. Annals of Oncology, 2012, 23, ix339.	1.2	0
67	Association of Genetic Polymorphisms in Tumor Suppressors, ERP29 and PTCH1, and DNA Transcription Factors, IKBKAP and ZNF415, with Cutaneous Melanoma Risk. Annals of Oncology, 2012, 23, ix364-ix365.	1.2	0
68	Polymorphisms of VEGF, GSTM1 and GSTT1 genes in multiple myeloma risk. Hematological Oncology, 2012, 30, 105-107.	1.7	5
69	Base of tongue squamous cell carcinoma susceptibility: Novel candidate genetic polymorphisms identified in genome-wide association study.. Journal of Clinical Oncology, 2012, 30, e16041-e16041.	1.6	0
70	8521 POSTER Novel Candidate Genetic Polymorphisms Identified in Genome-wide Association Study for Base of Tongue Squamous Cell Carcinoma Susceptibility. European Journal of Cancer, 2011, 47, S550.	2.8	0
71	9308 POSTER The XPC A2920C, the XPF T30028C and the P53 Arg72Pro Polymorphisms, Involved in DNA Repair, Alter the Risk for the Malignant Melanoma. European Journal of Cancer, 2011, 47, S654-S655.	2.8	1
72	Polymorphism of cytochrome P450 A2 (CYP1A2) in pure and mixed breed dogs. Journal of Veterinary Pharmacology and Therapeutics, 2011, 34, 184-186.	1.3	9

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73	Polymorphisms in the 5â€²- and 3â€²-untranslated region of the VEGF gene and sporadic breast cancer risk and clinicopathologic characteristics. Tumor Biology, 2011, 32, 295-300.	1.8	30
74	CYP1A1, GSTM1 and GSTT1 polymorphisms, tobacco and alcohol status and risk of head and neck squamous cell carcinoma. Tumor Biology, 2011, 32, 1209-1215.	1.8	30
75	The GSTT1 polymorphism of the glutathione S-transferase system in the intratumoral microvessel density of breast cancer patients. Tumor Biology, 2010, 31, 489-493.	1.8	3
76	Glutathione S-transferase mu 1 (GSTM1) and theta 1 (GSTT1) genetic polymorphisms and atopic asthma in children from Southeastern Brazil. Genetics and Molecular Biology, 2010, 33, 438-441.	1.3	19
77	Increased Risk for Acute Lymphoblastic Leukemia in Children with Cytochrome P₄₅₀ A₁<i>(CYP1A1)-</i> and NAD(P)H:Quinone Oxidoreductase 1 <i>(NQO1)</i>-Inherited Gene Variants. Acta Haematologica, 2010, 124, 182-184.	1.4	23
78	Polymorphisms of glutathione S-transferase mu 1, theta 1, and pi 1 genes and prognosis in Hodgkin lymphoma. Leukemia and Lymphoma, 2010, 51, 2215-2221.	1.3	15
79	No contribution of GSTM1 and GSTT1 null genotypes to the risk of neutropenia due to benzene exposure in Southeastern Brazil. Genetics and Molecular Biology, 2009, 32, 709-711.	1.3	2
80	High risk of â€˜de novoâ€™ acute myeloid leukaemia in individuals with cytochrome P450 A1 (CYP1A1) and NAD(P)H:quinone oxidoreductase 1 (NQO1) gene defects. European Journal of Haematology, 2009, 83, 270-272.	2.2	14
81	Polymorphisms of glutathione S-transferase Mu 1, glutathione S-transferase theta 1 and glutathione S-transferase Pi 1 genes in Hodgkin's lymphoma susceptibility and progression. Leukemia and Lymphoma, 2009, 50, 1005-1009.	1.3	9
82	9229 Polymorphisms of glutathione S-transferase mu 1(GSTM1), theta 1 (GSTT1) and pi 1 (GSTP1) in outcome of Hodgkin's lymphoma patients. European Journal of Cancer, Supplement, 2009, 7, 568.	2.2	0
83	Clinical and pathological implications of GSTM1 and GSTT1 gene deletions in sporadic breast cancer. Oncology Reviews, 2008, 2, 36-43.	1.8	2
84	A polymorphism in the angiogenesis inhibitor, endostatin, in lung cancer susceptibility. Lung Cancer, 2008, 59, 276-278.	2.0	8
85	The rare t(6;8) (q27;p11) translocation in a case of chronic myeloid neoplasm mimicking polycythemia vera. Leukemia and Lymphoma, 2008, 49, 1832-1835.	1.3	7
86	D104N polymorphism in endostatin, an angiogenesis inhibitor, in acute and chronic myeloid leukaemia. Leukemia Research, 2007, 31, 1158-1159.	0.8	3
87	Inherited pericentric inversion of chromosome 9 in acquired hematological disorders. Annals of Hematology, 2007, 86, 465-467.	1.8	7
88	High Risk of Acute Myeloid Leukemia in Individuals with NAD(P)H:Quinone Oxidoreductase 1 (NQO1) and Cytochrome P450 A1 (CYP1A1) Gene Defects.. Blood, 2007, 110, 2843-2843.	1.4	0
89	Inherited pericentric inversion of chromosome 16 in chronic phase of chronic myeloid leukaemia. Leukemia Research, 2006, 30, 115-117.	0.8	3
90	A high risk of occurrence of sporadic breast cancer in individuals with the 104NN polymorphism of the COL18A1 gene. Breast Cancer Research and Treatment, 2006, 100, 335-338.	2.5	30

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91	Polymorphisms of glutathione S-transferase mu1 (GSTM1) and theta 1 (GSTT1) genes in chronic myeloid leukaemia. European Journal of Haematology, 2005, 75, 530-531.	2.2	19
92	Molecular analysis of the most prevalent mutations of the FANCA and FANCC genes in Brazilian patients with Fanconi anaemia. Genetics and Molecular Biology, 2005, 28, 205-209.	1.3	4