Luisa Matos do Canto

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Long Non-coding RNAs Involved in Resistance to Chemotherapy in Ovarian Cancer. Frontiers in Oncology, 2019, 9, 1549.	2.8	37
2	CoNVaQ: a web tool for copy number variation-based association studies. BMC Genomics, 2018, 19, 369.	2.8	29
3	Loss of DNA methylation is related to increased expression of miR-21 and miR-146b in papillary thyroid carcinoma. Clinical Epigenetics, 2018, 10, 144.	4.1	27
4	Fusion Potential of Human Osteoclasts In Vitro Reflects Age, Menopause, and In Vivo Bone Resorption Levels of Their Donors—A Possible Involvement of DC-STAMP. International Journal of Molecular Sciences, 2020, 21, 6368.	4.1	27
5	MicroRNA analysis of breast ductal fluid in breast cancer patients. International Journal of Oncology, 2016, 48, 2071-2078.	3.3	24
6	Metabolomic profiling of breast tumors using ductal fluid. International Journal of Oncology, 2016, 49, 2245-2254.	3.3	23
7	Genomic profiling in ovarian cancer retreated with platinum based chemotherapy presented homologous recombination deficiency and copy number imbalances of CCNE1 and RB1 genes. BMC Cancer, 2019, 19, 422.	2.6	22
8	Increased Levels of Genomic Instability and Mutations in Homologous Recombination Genes in Locally Advanced Rectal Carcinomas. Frontiers in Oncology, 2019, 9, 395.	2.8	17
9	Drug Repositioning Based on the Reversal of Gene Expression Signatures Identifies TOP2A as a Therapeutic Target for Rectal Cancer. Cancers, 2021, 13, 5492.	3.7	17
10	Inflammatory Breast Cancer: Clinical Implications of Genomic Alterations and Mutational Profiling. Cancers, 2020, 12, 2816.	3.7	15
11	Locally advanced rectal cancer transcriptomic-based secretome analysis reveals novel biomarkers useful to identify patients according to neoadjuvant chemoradiotherapy response. Scientific Reports, 2019, 9, 8702.	3.3	14
12	Comprehensive Analysis of DNA Methylation and Prediction of Response to NeoadjuvantTherapy in Locally Advanced Rectal Cancer. Cancers, 2020, 12, 3079.	3.7	13
13	Penile Cancer-Derived Cells Molecularly Characterized as Models to Guide Targeted Therapies. Cells, 2021, 10, 814.	4.1	9
14	Rare germline alterations in cancer-related genes associated with the risk of multiple primary tumor development. Journal of Molecular Medicine, 2017, 95, 523-533.	3.9	8
15	Germline variants in DNA repair genes are associated with young-onset head and neck cancer. Oral Oncology, 2021, 122, 105545.	1.5	8
16	Genomic profile of a Li-Fraumeni-like syndrome patient with a 45,X/46,XX karyotype, presenting neither mutations in TP53 nor clinical stigmata of Turner syndrome. Cancer Genetics, 2015, 208, 341-344.	0.4	5
17	Lack of association between interleukin-18 polymorphisms and rheumatoid arthritis. Revista Brasileira De Reumatologia, 2013, 53, 199-205.	0.7	4
18	The Long Non-Coding RNA SNHG12 as a Mediator of Carboplatin Resistance in Ovarian Cancer via Epigenetic Mechanisms. Cancers, 2022, 14, 1664.	3.7	4

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19	Association of PDCD1 polymorphism to systemic lupus erythematosus and rheumatoid arthritis susceptibility. Revista Brasileira De Reumatologia, 2016, 56, 483-489.	0.7	3
20	Associação entre o polimorfismo do gene PDCD1 e a susceptibilidade ao lúpus eritematoso sistêmico e à artrite reumatoide. Revista Brasileira De Reumatologia, 2016, 56, 483-489.	0.8	3
21	Germline large genomic alterations on 7q in patients with multiple primary cancers. Scientific Reports, 2017, 7, 41677.	3.3	2
22	Metabolomic Analysis of Plasma from Breast Cancer Patients Using Ultra-High-Performance Liquid Chromatography Coupled with Mass Spectrometry: An Untargeted Study. Metabolites, 2022, 12, 447.	2.9	2
23	Homologous recombination deficiency and platinum rechallenge in platinum-resistant ovarian cancer patients Journal of Clinical Oncology, 2017, 35, 5576-5576.	1.6	1
24	Genomic profile of Li-Fraumeni syndrome patients with adrenocortical carcinoma in childhood. Annals of Oncology, 2016, 27, vi11.	1.2	0
25	Striking similarities in genetic aberrations between a rectal tumor and its lung recurrence. World Journal of Gastrointestinal Oncology, 2013, 5, 198.	2.0	0
26	Abstract 3465: microRNA profiling of breast ductal lavage fluid , 2013, , .		0
27	Abstract 3459: Metabolomic profiling reveals significant differences between the ductal fluid from cancerous compared to unaffected breasts , 2013, , .		0
28	Abstract 5360: Mutational profile and genomic instability according to response to therapy in rectal carcinomas. , 2018, , .		0