

Bruno Costa Gomes

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

373
citations

758635

12
h-index

839053

18
g-index

21
all docs

21
docs citations

21
times ranked

548
citing authors

#	ARTICLE	IF	CITATIONS
1	Association Between miR-148a and DNA Methylation Profile in Individuals Exposed to Lead (Pb). <i>Frontiers in Genetics</i> , 2021, 12, 620744.	1.1	12
2	Male and female breast cancer: the two faces of the same genetic susceptibility coin. <i>Breast Cancer Research and Treatment</i> , 2021, 188, 295-305.	1.1	7
3	Biomarkers of effect as determined in human biomonitoring studies on hexavalent chromium and cadmium in the period 2008â€“2020. <i>Environmental Research</i> , 2021, 197, 110998.	3.7	22
4	Micronuclei Formation upon Radioiodine Therapy for Well-Differentiated Thyroid Cancer: The Influence of DNA Repair Genes Variants. <i>Genes</i> , 2020, 11, 1083.	1.0	7
5	ABC Efflux Transporters and the Circuitry of miRNAs: Kinetics of Expression in Cancer Drug Resistance. <i>International Journal of Molecular Sciences</i> , 2020, 21, 2985.	1.8	12
6	The Role of the FMN-Domain of Human Cytochrome P450 Oxidoreductase in Its Promiscuous Interactions With Structurally Diverse Redox Partners. <i>Frontiers in Pharmacology</i> , 2020, 11, 299.	1.6	22
7	Thyroid Cancer: The Quest for Genetic Susceptibility Involving DNA Repair Genes. <i>Genes</i> , 2019, 10, 586.	1.0	11
8	Regulation of ABCB1 activity by microRNA-200c and microRNA-203a in breast cancer cells: the quest for microRNAsâ€™ involvement in cancer drug resistance. , 2019, 2, 897-911.		3
9	MicroRNAs and cancer drug resistance: over two thousand characters in search of a role. , 2019, 2, 618-633.		3
10	Probing the Role of the Hinge Segment of Cytochrome P450 Oxidoreductase in the Interaction with Cytochrome P450. <i>International Journal of Molecular Sciences</i> , 2018, 19, 3914.	1.8	16
11	Genetic Susceptibility in Acute Pancreatitis. <i>Pancreas</i> , 2017, 46, 71-76.	0.5	12
12	Prognostic value of microRNA-203a expression in breast cancer. <i>Oncology Reports</i> , 2016, 36, 1748-1756.	1.2	18
13	MicroRNAs and Cancer Drug Resistance. <i>Methods in Molecular Biology</i> , 2016, 1395, 137-162.	0.4	34
14	Methods for Studying MicroRNA Expression and Their Targets in Formalin-Fixed, Paraffin-Embedded (FFPE) Breast Cancer Tissues. <i>Methods in Molecular Biology</i> , 2016, 1395, 189-205.	0.4	7
15	Induction of sister chromatid exchange by acrylamide and glycidamide in human lymphocytes: Role of polymorphisms in detoxification and DNA-repair genes in the genotoxicity of glycidamide. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2013, 752, 1-7.	0.9	18
16	The role of CCNH Val270Ala (rs2230641) and other nucleotide excision repair polymorphisms in individual susceptibility to well-differentiated thyroid cancer. <i>Oncology Reports</i> , 2013, 30, 2458-2466.	1.2	14
17	A Data Mining Approach for the Detection of High-Risk Breast Cancer Groups. <i>Advances in Intelligent and Soft Computing</i> , 2010, , 43-51.	0.2	27
18	Breast cancer risk and common single nucleotide polymorphisms in homologous recombination DNA repair pathway genes XRCC2, XRCC3, NBS1 and RAD51. <i>Cancer Epidemiology</i> , 2010, 34, 85-92.	0.8	86

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19	The role of common variants of non-homologous end-joining repair genes XRCC4, LIG4 and Ku80 in thyroid cancer risk. <i>Oncology Reports</i> , 2010, , .	1.2	13
20	The role of common variants of non-homologous end-joining repair genes XRCC4, LIG4 and Ku80 in thyroid cancer risk. <i>Oncology Reports</i> , 2010, 24, 1079-85.	1.2	28