

Michal Kroupa

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1970450/publications.pdf>

Version: 2024-02-01

16
papers

198
citations

1039880

9
h-index

1058333

14
g-index

16
all docs

16
docs citations

16
times ranked

292
citing authors

#	ARTICLE	IF	CITATIONS
1	Relationship of telomere length in colorectal cancer patients with cancer phenotype and patient prognosis. <i>British Journal of Cancer</i> , 2019, 121, 344-350.	2.9	28
2	Oxidative Damage in Sporadic Colorectal Cancer: Molecular Mapping of Base Excision Repair Glycosylases in Colorectal Cancer Patients. <i>International Journal of Molecular Sciences</i> , 2020, 21, 2473.	1.8	28
3	Base excision repair capacity as a determinant of prognosis and therapy response in colon cancer patients. <i>DNA Repair</i> , 2018, 72, 77-85.	1.3	27
4	Interactions of DNA repair gene variants modulate chromosomal aberrations in healthy subjects. <i>Carcinogenesis</i> , 2015, 36, 1299-1306.	1.3	24
5	Genetic variation of acquired structural chromosomal aberrations. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2018, 836, 13-21.	0.9	19
6	Genetic variation in the major mitotic checkpoint genes associated with chromosomal aberrations in healthy humans. <i>Cancer Letters</i> , 2016, 380, 442-446.	3.2	12
7	Bleomycin-induced chromosomal damage and shortening of telomeres in peripheral blood lymphocytes of incident cancer patients. <i>Genes Chromosomes and Cancer</i> , 2018, 57, 61-69.	1.5	12
8	Telomere maintenance in interplay with DNA repair in pathogenesis and treatment of colorectal cancer. <i>Mutagenesis</i> , 2020, 35, 261-271.	1.0	11
9	Telomere length in peripheral blood lymphocytes related to genetic variation in telomerase, prognosis and clinicopathological features in breast cancer patients. <i>Mutagenesis</i> , 2020, 35, 491-497.	1.0	11
10	Genetic variation associated with chromosomal aberration frequency: A genome-wide association study. <i>Environmental and Molecular Mutagenesis</i> , 2019, 60, 17-28.	0.9	9
11	Distinct pathways associated with chromosomal aberration frequency in a cohort exposed to genotoxic compounds compared to general population. <i>Mutagenesis</i> , 2019, 34, 323-330.	1.0	6
12	DNA repair gene polymorphisms and chromosomal aberrations in healthy, nonsmoking population. <i>DNA Repair</i> , 2021, 101, 103079.	1.3	3
13	DNA Repair Gene Polymorphisms and Chromosomal Aberrations in Exposed Populations. <i>Frontiers in Genetics</i> , 2021, 12, 691947.	1.1	3
14	Oxidative Damage in Sporadic Colorectal Cancer: Molecular Mapping of Base Excision Repair Glycosylases MÜTYH and hOGG1 in Colorectal Cancer Patients. <i>International Journal of Molecular Sciences</i> , 2022, 23, 5704.	1.8	3
15	Impact of genetic polymorphisms in kinetochore and spindle assembly genes on chromosomal aberration frequency in healthy humans. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2020, 858-860, 503253.	0.9	2
16	Abstract 2316: Malignant potential of colorectal adenoma based on the telomere length. <i>Cancer Research</i> , 2022, 82, 2316-2316.	0.4	0