Miroslav Levy

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/8770741/publications.pdf

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27	907	430874	501196
papers	citations	18 h-index	28 g-index

28	28	28	1829
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Local Immune Changes in Early Stages of Inflammation and Carcinogenesis Correlate with the Collagen Scaffold Changes of the Colon Mucosa. Cancers, 2021, 13, 2463.	3.7	3
2	Analysis of MicroRNA Expression Changes During the Course of Therapy In Rectal Cancer Patients. Frontiers in Oncology, 2021, 11, 702258.	2.8	11
3	Association of circulating short chain fatty acid levels with colorectal adenomas and colorectal cancer. Clinical Nutrition ESPEN, 2021, 46, 297-304.	1.2	10
4	Expression quantitative trait loci in ABC transporters are associated with survival in 5-FU treated colorectal cancer patients. Mutagenesis, 2020, 35, 273-281.	2.6	2
5	Epistatic effect of TLR3 and cGASâ€STINGâ€IKKÎµâ€TBK1â€IFN signaling variants on colorectal cancer risk. Cancer Medicine, 2020, 9, 1473-1484.	2.8	10
6	Fusobacterium nucleatum tumor DNA levels are associated with survival in colorectal cancer patients. European Journal of Clinical Microbiology and Infectious Diseases, 2019, 38, 1891-1899.	2.9	33
7	Functional Polymorphisms in DNA Repair Genes Are Associated with Sporadic Colorectal Cancer Susceptibility and Clinical Outcome. International Journal of Molecular Sciences, 2019, 20, 97.	4.1	20
8	Significance of postoperative follow-up of patients with metastatic colorectal cancer using circulating tumor DNA. World Journal of Gastroenterology, 2019, 25, 6939-6948.	3.3	24
9	Base excision repair capacity as a determinant of prognosis and therapy response in colon cancer patients. DNA Repair, 2018, 72, 77-85.	2.8	27
10	Expression profile of miR-17/92 cluster is predictive of treatment response in rectal cancer. Carcinogenesis, 2018, 39, 1359-1367.	2.8	29
11	Short article: Influence of regulatory NLRC5 variants on colorectal cancer survival and 5-fluorouracil-based chemotherapy. European Journal of Gastroenterology and Hepatology, 2018, 30, 838-842.	1.6	6
12	Investigation of single and synergic effects of NLRC5 and PD-L1 variants on the risk of colorectal cancer. PLoS ONE, 2018, 13, e0192385.	2.5	20
13	MicroRNA-binding site polymorphisms in genes involved in colorectal cancer etiopathogenesis and their impact on disease prognosis. Mutagenesis, 2017, 32, 533-542.	2.6	20
14	The focus on sample quality: Influence of colon tissue collection on reliability of qPCR data. Scientific Reports, 2016, 6, 29023.	3.3	7
15	Epigenome-wide analysis of DNA methylation reveals a rectal cancer-specific epigenomic signature. Epigenomics, 2016, 8, 1193-1207.	2.1	22
16	Monitoring of Circulating Tumor Cells by a Combination of Immunomagnetic Enrichment and RT-PCR in Colorectal Cancer Patients Undergoing Surgery. Advances in Clinical and Experimental Medicine, 2016, 25, 1273-1279.	1.4	11
17	Double-strand break repair and colorectal cancer: gene variants within 3â€² UTRs and microRNAs binding as modulators of cancer risk and clinical outcome. Oncotarget, 2016, 7, 23156-23169.	1.8	40
18	Post-treatment recovery of suboptimal DNA repair capacity and gene expression levels in colorectal cancer patients. Molecular Carcinogenesis, 2015, 54, 769-778.	2.7	16

#	Article	IF	Citations
19	Circulating miRNAs miR-34a and miR-150 associated with colorectal cancer progression. BMC Cancer, 2015, 15, 329.	2.6	77
20	Elevated levels of 14-3-3 proteins, serotonin, gamma enolase and pyruvate kinase identified in clinical samples from patients diagnosed with colorectal cancer. Clinica Chimica Acta, 2015, 441, 133-141.	1.1	28
21	HOTAIR long non-coding RNA is a negative prognostic factor not only in primary tumors, but also in the blood of colorectal cancer patients. Carcinogenesis, 2014, 35, 1510-1515.	2.8	227
22	Variations in mismatch repair genes and colorectal cancer risk and clinical outcome. Mutagenesis, 2014, 29, 259-265.	2.6	20
23	Variation within 3â€²-UTRs of Base Excision Repair Genes and Response to Therapy in Colorectal Cancer Patients: A Potential Modulation of microRNAs Binding. Clinical Cancer Research, 2013, 19, 6044-6056.	7.0	56
24	Polymorphisms in miRNA-binding sites of nucleotide excision repair genes and colorectal cancer risk. Carcinogenesis, 2012, 33, 1346-1351.	2.8	59
25	Differences in nucleotide excision repair capacity between newly diagnosed colorectal cancer patients and healthy controls. Mutagenesis, 2012, 27, 225-232.	2.6	35
26	Functional, Genetic, and Epigenetic Aspects of Base and Nucleotide Excision Repair in Colorectal Carcinomas. Clinical Cancer Research, 2012, 18, 5878-5887.	7.0	66
27	Utility of cell-free tumour DNA for post-surgical follow-up of colorectal cancer patients. Anticancer Research, 2012, 32, 1621-6.	1.1	18