M Henar Alonso Aguado

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7317171/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Discovery of common and rare genetic risk variants for colorectal cancer. Nature Genetics, 2019, 51, 76-87.	9.4	377
2	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. Journal of the National Cancer Institute, 2019, 111, 146-157.	3.0	129
3	Low adherence to the western and high adherence to the mediterranean dietary patterns could prevent colorectal cancer. European Journal of Nutrition, 2019, 58, 1495-1505.	1.8	126
4	Comprehensive analysis of copy number aberrations in microsatellite stable colon cancer in view of stromal component. British Journal of Cancer, 2017, 117, 421-431.	2.9	125
5	Lung metastases share common immune features regardless of primary tumor origin. , 2020, 8, e000491.		63
6	Exome Sequencing Reveals <i>AMER1</i> as a Frequently Mutated Gene in Colorectal Cancer. Clinical Cancer Research, 2015, 21, 4709-4718.	3.2	52
7	Genetic architectures of proximal and distal colorectal cancer are partly distinct. Gut, 2021, 70, 1325-1334.	6.1	44
8	Risk Model for Colorectal Cancer in Spanish Population Using Environmental and Genetic Factors: Results from the MCC-Spain study. Scientific Reports, 2017, 7, 43263.	1.6	41
9	Colon-specific eQTL analysis to inform on functional SNPs. British Journal of Cancer, 2018, 119, 971-977.	2.9	25
10	Lymphocytic infiltration in stage II microsatellite stable colorectal tumors: A retrospective prognosis biomarker analysis. PLoS Medicine, 2020, 17, e1003292.	3.9	25
11	Residential proximity to industrial pollution sources and colorectal cancer risk: A multicase-control study (MCC-Spain). Environment International, 2020, 144, 106055.	4.8	24
12	Colorectal cancer, sun exposure and dietary vitamin D and calcium intake in the MCC-Spain study. Environment International, 2018, 121, 428-434.	4.8	23
13	NTHL1 biallelic mutations seldom cause colorectal cancer, serrated polyposis or a multi-tumor phenotype, in absence of colorectal adenomas. Scientific Reports, 2019, 9, 9020.	1.6	23
14	Additive Role of Immune System Infiltration and Angiogenesis in Uveal Melanoma Progression. International Journal of Molecular Sciences, 2021, 22, 2669.	1.8	22
15	DNA methylation events in transcription factors and gene expression changes in colon cancer. Epigenomics, 2020, 12, 1593-1610.	1.0	13
16	Genetically determined telomere length and multiple myeloma risk and outcome. Blood Cancer Journal, 2021, 11, 74.	2.8	10
17	Tumor immune infiltration estimated from gene expression profiles predicts colorectal cancer relapse. Oncolmmunology, 2021, 10, 1862529.	2.1	9
18	Non-Lynch Familial and Early-Onset Colorectal Cancer Explained by Accumulation of Low-Risk Genetic Variants. Cancers, 2021, 13, 3857.	1.7	8

#	Article	IF	CITATIONS
19	Telomere length alterations in microsatellite stable colorectal cancer and association with the immune response. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2018, 1864, 2992-3000.	1.8	7
20	Changes of CD68, CD163, and PD-L1 tumor expression during high-dose-rate and pulsed-dose-rate brachytherapy for cervical cancer. Brachytherapy, 2020, 19, 51-59.	0.2	7
21	Mutanome and expression of immune response genes in microsatellite stable colon cancer. Oncotarget, 2016, 7, 17711-17725.	0.8	6
22	Polygenic risk score across distinct colorectal cancer screening outcomes: from premalignant polyps to colorectal cancer. BMC Medicine, 2021, 19, 261.	2.3	5
23	Abstract 2737: Clinical and epidemiologic predictors of clonal immune responses in colorectal cancer. , 2021, , .		0