

Pilar Mur

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7554628/publications.pdf>

Version: 2024-02-01

27
papers

707
citations

516681

16
h-index

552766

26
g-index

27
all docs

27
docs citations

27
times ranked

1630
citing authors

#	ARTICLE	IF	CITATIONS
1	MLH1 promoter hypermethylation in the analytical algorithm of Lynch syndrome: a cost-effectiveness study. <i>European Journal of Human Genetics</i> , 2012, 20, 762-768.	2.8	76
2	Role of POLE and POLD1 in familial cancer. <i>Genetics in Medicine</i> , 2020, 22, 2089-2100.	2.4	76
3	Codeletion of 1p and 19q determines distinct gene methylation and expression profiles in IDH-mutated oligodendroglial tumors. <i>Acta Neuropathologica</i> , 2013, 126, 277-289.	7.7	49
4	Evidence suggests that germline <i>RNF43</i> mutations are a rare cause of serrated polyposis. <i>Gut</i> , 2018, 67, 2230-2232.	12.1	48
5	Genetic Alterations Associated With Progression and Recurrence in Meningiomas. <i>Journal of Neuropathology and Experimental Neurology</i> , 2012, 71, 882-893.	1.7	47
6	Impact on prognosis of the regional distribution of MGMT methylation with respect to the CpG island methylator phenotype and age in glioma patients. <i>Journal of Neuro-Oncology</i> , 2015, 122, 441-450.	2.9	41
7	Delineating the Phenotypic Spectrum of the NTHL1-Associated Polyposis. <i>Clinical Gastroenterology and Hepatology</i> , 2017, 15, 461-462.	4.4	41
8	The Inherited and Familial Component of Early-Onset Colorectal Cancer. <i>Cells</i> , 2021, 10, 710.	4.1	41
9	Molecular Classification Defines 4 Prognostically Distinct Glioma Groups Irrespective of Diagnosis and Grade. <i>Journal of Neuropathology and Experimental Neurology</i> , 2015, 74, 241-249.	1.7	38
10	MLH1 methylation screening is effective in identifying epimutation carriers. <i>European Journal of Human Genetics</i> , 2012, 20, 1256-1264.	2.8	36
11	Gene expression changes associated with erlotinib response in glioma cell lines. <i>European Journal of Cancer</i> , 2013, 49, 1641-1653.	2.8	35
12	Association Between Germline Mutations in BRF1, a Subunit of the RNA Polymerase III Transcription Complex, and Hereditary Colorectal Cancer. <i>Gastroenterology</i> , 2018, 154, 181-194.e20.	1.3	32
13	NTHL1 biallelic mutations seldom cause colorectal cancer, serrated polyposis or a multi-tumor phenotype, in absence of colorectal adenomas. <i>Scientific Reports</i> , 2019, 9, 9020.	3.3	23
14	Plasma Viral miRNAs Indicate a High Prevalence of Occult Viral Infections. <i>EBioMedicine</i> , 2017, 20, 182-192.	6.1	19
15	Germline mutations in the spindle assembly checkpoint genes BUB1 and BUB3 are infrequent in familial colorectal cancer and polyposis. <i>Molecular Cancer</i> , 2018, 17, 23.	19.2	19
16	miR-543 regulates the epigenetic landscape of myelofibrosis by targeting TET1 and TET2. <i>JCI Insight</i> , 2020, 5, .	5.0	18
17	Germline variation in O6-methylguanine-DNA methyltransferase (MGMT) as cause of hereditary colorectal cancer. <i>Cancer Letters</i> , 2019, 447, 86-92.	7.2	12
18	Germline variation in the oxidative DNA repair genes NUDT1 and OGG1 is not associated with hereditary colorectal cancer or polyposis. <i>Human Mutation</i> , 2018, 39, 1214-1225.	2.5	10

#	ARTICLE	IF	CITATIONS
19	<i>TP53</i> , a gene for colorectal cancer predisposition in the absence of Li-Fraumeni-associated phenotypes. <i>Gut</i> , 2021, 70, 1139-1146.	12.1	10
20	Scarce evidence of the causal role of germline mutations in <i>UNC5C</i> in hereditary colorectal cancer and polyposis. <i>Scientific Reports</i> , 2016, 6, 20697.	3.3	9
21	Non-Lynch Familial and Early-Onset Colorectal Cancer Explained by Accumulation of Low-Risk Genetic Variants. <i>Cancers</i> , 2021, 13, 3857.	3.7	8
22	Approaches to functionally validate candidate genetic variants involved in colorectal cancer predisposition. <i>Molecular Aspects of Medicine</i> , 2019, 69, 27-40.	6.4	5
23	<i>AMER1</i> Is a Frequently Mutated Gene in Colorectal Cancer—Letter. <i>Clinical Cancer Research</i> , 2015, 21, 4985-4985.	7.0	4
24	Reply to: “Development of an MSI-positive colon tumor with aberrant DNA methylation in a PPAP patient” <i>Journal of Human Genetics</i> , 2020, 65, 513-514.	2.3	4
25	Expanding the phenotype of cerebellarâ€facialâ€dental syndrome: Two siblings with a novel variant in <i>BRF1</i> . <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 2742-2745.	1.2	4
26	Solving the enigma of <i>POLD1</i> p.V295M as a potential cause of increased cancer risk. <i>European Journal of Human Genetics</i> , 2022, 30, 485-489.	2.8	2
27	Potential Involvement of <i>NSD1</i> , <i>KRT24</i> and <i>ACACA</i> in the Genetic Predisposition to Colorectal Cancer. <i>Cancers</i> , 2022, 14, 699.	3.7	0