Heikki JÃ¤rvinen

List of Publications by Year
in descending order

24	2,431	394286 19	677027 22
papers	citations	h-index	g-index

24	24	24	3698
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Disease expression in juvenile polyposis syndrome: a retrospective survey on a cohort of 221 European patients and comparison with a literature-derived cohort of 473 SMAD4/BMPR1A pathogenic variant carriers. Genetics in Medicine, 2020, 22, 1524-1532.	1.1	44
2	Genomeâ€wide association study and metaâ€analysis in Northern European populations replicate multiple colorectal cancer risk loci. International Journal of Cancer, 2018, 142, 540-546.	2.3	26
3	No Difference in Colorectal Cancer Incidence or Stage at Detection by Colonoscopy Among 3 Countries With Different Lynch Syndrome Surveillance Policies. Gastroenterology, 2018, 155, 1400-1409.e2.	0.6	112
4	Comprehensive Evaluation of Protein Coding Mononucleotide Microsatellites in Microsatellite-Unstable Colorectal Cancer. Cancer Research, 2017, 77, 4078-4088.	0.4	18
5	Multiple components of PKA and TGF-Î² pathways are mutated in pseudomyxoma peritonei. PLoS ONE, 2017, 12, e0174898.	1.1	15
6	The Role of Chromosomal Instability and Epigenetics in Colorectal Cancers LackingÎ²-Catenin/TCF Regulated Transcription. Gastroenterology Research and Practice, 2016, 2016, 1-11.	0.7	17
7	Mendelian randomisation analysis strongly implicates adiposity with risk of developing colorectal cancer. British Journal of Cancer, 2016, 115, 266-272.	2.9	57
8	Expression of CEA, CA19-9, CA125, and EpCAM in pseudomyxoma peritonei. Human Pathology, 2016, 54, 47-54.	1.1	23
9	Somatic MED12 mutations in prostate cancer and uterine leiomyomas promote tumorigenesis through distinct mechanisms. Prostate, 2016, 76, 22-31.	1.2	33
10	Variation at 2q35 (PNKDandTMBIM1) influences colorectal cancer risk and identifies a pleiotropic effect with inflammatory bowel disease. Human Molecular Genetics, 2016, 25, 2349-2359.	1.4	37
11	CTCF/cohesin-binding sites are frequently mutated in cancer. Nature Genetics, 2015, 47, 818-821.	9.4	383
12	Systematic search for rare variants in Finnish early-onset colorectal cancer patients. Cancer Genetics, 2015, 208, 35-40.	0.2	24
13	DNA hypermethylation appears early and shows increased frequency with dysplasia in Lynch syndrome-associated colorectal adenomas and carcinomas. Clinical Epigenetics, 2015, 7, 71.	1.8	24
14	Response to Dr Braillon. Journal of Medical Screening, 2011, 18, 103-104.	1.1	0
15	Population-Based Molecular Detection of Hereditary Nonpolyposis Colorectal Cancer. Journal of Clinical Oncology, 2000, 18, 2193-2200.	0.8	466
16	Predictive genetic testing for hereditary non-polyposis colorectal cancer: Uptake and long-term satisfaction. International Journal of Cancer, 2000, 89, 44-50.	2.3	116
17	SMAD genes in juvenile polyposis. , 1999, 26, 54-61.		78
18	Intestinal cancer in patients with a germline mutation in the down-regulated in adenoma (DRA) gene. Oncogene, 1998, 16, 681-684.	2.6	37

#	Article	IF	Citations
19	Mutation sharing, predominant involvement of the MLH1 gene and description of four novel mutations in hereditary nonpolyposis colorectal cancer. Human Mutation, 1998, 11, 482-483.	1.1	33
20	Localization of a susceptibility locus for Peutz-Jeghers syndrome to 19p using comparative genomic hybridization and targeted linkage analysis. Nature Genetics, 1997, 15, 87-90.	9.4	444
21	Substance P- and vasoactive intestinal polypeptide-immunoreactive innervation in normal and inflamed pouches after restorative proctocolectomy for ulcerative colitis. Digestive Diseases and Sciences, 1996, 41, 1658-1664.	1.1	33
22	Substance Pâ€"An underlying factor for pouchitis?. Digestive Diseases and Sciences, 1996, 41, 1665-1671.	1.1	27
23	Changes in substance P-immunoreactive innervation of human colon associated with ulcerative colitis. Digestive Diseases and Sciences, 1995, 40, 2250-2258.	1.1	80
24	Loss of the wild type MLH1 gene is a feature of hereditary nonpolyposis colorectal cancer. Nature Genetics, 1994, 8, 405-410.	9.4	304