

Lauri A Aaltonen

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/880110/publications.pdf>

Version: 2024-02-01

337
papers

41,972
citations

2970

93
h-index

2624

194
g-index

349
all docs

349
docs citations

349
times ranked

31157
citing authors

#	ARTICLE	IF	CITATIONS
1	Clues to the pathogenesis of familial colorectal cancer. <i>Science</i> , 1993, 260, 812-816.	6.0	2,563
2	Mutations of a mutS homolog in hereditary nonpolyposis colorectal cancer. <i>Cell</i> , 1993, 75, 1215-1225.	13.5	2,195
3	A serine/threonine kinase gene defective in Peutz-Jeghers syndrome. <i>Nature</i> , 1998, 391, 184-187.	13.7	1,451
4	Germline mutations in FH predispose to dominantly inherited uterine fibroids, skin leiomyomata and papillary renal cell cancer. <i>Nature Genetics</i> , 2002, 30, 406-410.	9.4	1,426
5	Controlled 15-year trial on screening for colorectal cancer in families with hereditary nonpolyposis colorectal cancer. <i>Gastroenterology</i> , 2000, 118, 829-834.	0.6	1,259
6	Cancer risk in mutation carriers of DNA-mismatch-repair genes. , 1999, 81, 214-218.		1,061
7	Incidence of Hereditary Nonpolyposis Colorectal Cancer and the Feasibility of Molecular Screening for the Disease. <i>New England Journal of Medicine</i> , 1998, 338, 1481-1487.	13.9	1,048
8	Mutations in the SMAD4/DPC4 Gene in Juvenile Polyposis. <i>Science</i> , 1998, 280, 1086-1088.	6.0	866
9	Genetic mapping of a locus predisposing to human colorectal cancer. <i>Science</i> , 1993, 260, 810-812.	6.0	846
10	Multiple Colorectal Adenomas, Classic Adenomatous Polyposis, and Germ-Line Mutations in MYH. <i>New England Journal of Medicine</i> , 2003, 348, 791-799.	13.9	822
11	Inherited susceptibility to uterine leiomyomas and renal cell cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 3387-3392.	3.3	604
12	Life-time risk of different cancers in hereditary non-polyposis colorectal cancer (hnpcc) syndrome. <i>International Journal of Cancer</i> , 1995, 64, 430-433.	2.3	560
13	Pituitary Adenoma Predisposition Caused by Germline Mutations in the AIP Gene. <i>Science</i> , 2006, 312, 1228-1230.	6.0	557
14	MED12, the Mediator Complex Subunit 12 Gene, Is Mutated at High Frequency in Uterine Leiomyomas. <i>Science</i> , 2011, 334, 252-255.	6.0	547
15	A genome-wide association study identifies colorectal cancer susceptibility loci on chromosomes 10p14 and 8q23.3. <i>Nature Genetics</i> , 2008, 40, 623-630.	9.4	514
16	Meta-analysis of genome-wide association data identifies four new susceptibility loci for colorectal cancer. <i>Nature Genetics</i> , 2008, 40, 1426-1435.	9.4	498
17	Loss-of-Function Mutations in PPAR δ Associated with Human Colon Cancer. <i>Molecular Cell</i> , 1999, 3, 799-804.	4.5	485
18	Population-Based Molecular Detection of Hereditary Nonpolyposis Colorectal Cancer. <i>Journal of Clinical Oncology</i> , 2000, 18, 2193-2200.	0.8	466

#	ARTICLE	IF	CITATIONS
19	The common colorectal cancer predisposition SNP rs6983267 at chromosome 8q24 confers potential to enhanced Wnt signaling. <i>Nature Genetics</i> , 2009, 41, 885-890.	9.4	463
20	Localization of a susceptibility locus for Peutz-Jeghers syndrome to 19p using comparative genomic hybridization and targeted linkage analysis. <i>Nature Genetics</i> , 1997, 15, 87-90.	9.4	444
21	Identification of Lynch Syndrome Among Patients With Colorectal Cancer. <i>JAMA - Journal of the American Medical Association</i> , 2012, 308, 1555.	3.8	443
22	A recurrent mutation in PALB2 in Finnish cancer families. <i>Nature</i> , 2007, 446, 316-319.	13.7	402
23	Metastasis-Associated Gene Expression Changes Predict Poor Outcomes in Patients with Dukes Stage B and C Colorectal Cancer. <i>Clinical Cancer Research</i> , 2009, 15, 7642-7651.	3.2	395
24	CTCF/cohesin-binding sites are frequently mutated in cancer. <i>Nature Genetics</i> , 2015, 47, 818-821.	9.4	383
25	DNA methylation patterns in hereditary human cancers mimic sporadic tumorigenesis. <i>Human Molecular Genetics</i> , 2001, 10, 3001-3007.	1.4	374
26	Early-Onset Renal Cell Carcinoma as a Novel Extraparaganglial Component of SDHB-Associated Heritable Paraganglioma. <i>American Journal of Human Genetics</i> , 2004, 74, 153-159.	2.6	367
27	A role for mitochondrial enzymes in inherited neoplasia and beyond. <i>Nature Reviews Cancer</i> , 2003, 3, 193-202.	12.8	359
28	A TARBP2 mutation in human cancer impairs microRNA processing and DICER1 function. <i>Nature Genetics</i> , 2009, 41, 365-370.	9.4	355
29	Cancer Risk in Hereditary Nonpolyposis Colorectal Cancer Syndrome: Later Age of Onset. <i>Gastroenterology</i> , 2005, 129, 415-421.	0.6	338
30	Mutations in BRIP1 confer high risk of ovarian cancer. <i>Nature Genetics</i> , 2011, 43, 1104-1107.	9.4	338
31	Better survival rates in patients with MLH1-associated hereditary colorectal cancer. <i>Gastroenterology</i> , 1996, 110, 682-687.	0.6	336
32	Meta-analysis of three genome-wide association studies identifies susceptibility loci for colorectal cancer at 1q41, 3q26.2, 12q13.13 and 20q13.33. <i>Nature Genetics</i> , 2010, 42, 973-977.	9.4	335
33	Clinical Characteristics and Therapeutic Responses in Patients with Germ-Line <i>AIP</i> Mutations and Pituitary Adenomas: An International Collaborative Study. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2010, 95, E373-E383.	1.8	323
34	Microsatellite instability is a favorable prognostic indicator in patients with colorectal cancer receiving chemotherapy. <i>Gastroenterology</i> , 2000, 119, 921-928.	0.6	322
35	Molecular Staging for Survival Prediction of Colorectal Cancer Patients. <i>Journal of Clinical Oncology</i> , 2005, 23, 3526-3535.	0.8	313
36	Cancer Risk in Hereditary Nonpolyposis Colorectal Cancer Syndrome: Later Age of Onset. <i>Gastroenterology</i> , 2005, 129, 415-421.	0.6	309

#	ARTICLE	IF	CITATIONS
37	BRAF screening as a low-cost effective strategy for simplifying HNPCC genetic testing. <i>Journal of Medical Genetics</i> , 2004, 41, 664-668.	1.5	305
38	Loss of the wild type MLH1 gene is a feature of hereditary nonpolyposis colorectal cancer. <i>Nature Genetics</i> , 1994, 8, 405-410.	9.4	304
39	Familial Isolated Pituitary Adenomas (FIPA) and the Pituitary Adenoma Predisposition due to Mutations in the Aryl Hydrocarbon Receptor Interacting Protein (AIP) Gene. <i>Endocrine Reviews</i> , 2013, 34, 239-277.	8.9	289
40	Characterization of Uterine Leiomyomas by Whole-Genome Sequencing. <i>New England Journal of Medicine</i> , 2013, 369, 43-53.	13.9	280
41	Founding mutations and Alu-mediated recombination in hereditary colon cancer. <i>Nature Medicine</i> , 1995, 1, 1203-1206.	15.2	275
42	Germline CDKN1B/p27Kip1 Mutation in Multiple Endocrine Neoplasia. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2007, 92, 3321-3325.	1.8	262
43	A truncating mutation of HDAC2 in human cancers confers resistance to histone deacetylase inhibition. <i>Nature Genetics</i> , 2006, 38, 566-569.	9.4	254
44	Ten Years After Mutation Testing for Lynch Syndrome: Cancer Incidence and Outcome in Mutation-Positive and Mutation-Negative Family Members. <i>Journal of Clinical Oncology</i> , 2009, 27, 4793-4797.	0.8	252
45	Germline Mutations in BMPR1A/ALK3 Cause a Subset of Cases of Juvenile Polyposis Syndrome and of Cowden and Bannayan-Riley-Ruvalcaba Syndromes*. <i>American Journal of Human Genetics</i> , 2001, 69, 704-711.	2.6	236
46	The prevalence of MADH4 and BMPR1A mutations in juvenile polyposis and absence of BMPR2, BMPR1B, and ACVR1 mutations. <i>Journal of Medical Genetics</i> , 2004, 41, 484-491.	1.5	226
47	Aberrant succination of proteins in fumarate hydratase-deficient mice and HLRCC patients is a robust biomarker of mutation status. <i>Journal of Pathology</i> , 2011, 225, 4-11.	2.1	225
48	Molecular Classification of Patients With Unexplained Hamartomatous and Hyperplastic Polyposis. <i>JAMA - Journal of the American Medical Association</i> , 2005, 294, 2465.	3.8	218
49	Differential gene expression in colon cancer of the caecum versus the sigmoid and rectosigmoid. <i>Gut</i> , 2005, 54, 374-384.	6.1	212
50	The DNA repair gene MBD4 (MED1) is mutated in human carcinomas with microsatellite instability. <i>Nature Genetics</i> , 1999, 23, 266-268.	9.4	211
51	Common variation near CDKN1A, POLD3 and SHROOM2 influences colorectal cancer risk. <i>Nature Genetics</i> , 2012, 44, 770-776.	9.4	210
52	Mice Lacking a <i>Myc</i> Enhancer That Includes Human SNP rs6983267 Are Resistant to Intestinal Tumors. <i>Science</i> , 2012, 338, 1360-1363.	6.0	200
53	Familial Cutaneous Leiomyomatosis Is a Two-Hit Condition Associated with Renal Cell Cancer of Characteristic Histopathology. <i>American Journal of Pathology</i> , 2001, 159, 825-829.	1.9	191
54	Multiple Common Susceptibility Variants near BMP Pathway Loci GREM1, BMP4, and BMP2 Explain Part of the Missing Heritability of Colorectal Cancer. <i>PLoS Genetics</i> , 2011, 7, e1002105.	1.5	188

#	ARTICLE	IF	CITATIONS
55	Gene expression in colorectal cancer. <i>Cancer Research</i> , 2002, 62, 4352-63.	0.4	186
56	Increased risk of cancer in patients with fumarate hydratase germline mutation. <i>Journal of Medical Genetics</i> , 2006, 43, 523-526.	1.5	184
57	PTEN Mutational Spectra, Expression Levels, and Subcellular Localization in Microsatellite Stable and Unstable Colorectal Cancers. <i>American Journal of Pathology</i> , 2002, 161, 439-447.	1.9	173
58	Molecular diagnosis of pituitary adenoma predisposition caused by aryl hydrocarbon receptor-interacting protein gene mutations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 4101-4105.	3.3	173
59	SMAD4 as a Prognostic Marker in Colorectal Cancer. <i>Clinical Cancer Research</i> , 2005, 11, 2606-2611.	3.2	172
60	Association analyses identify 31 new risk loci for colorectal cancer susceptibility. <i>Nature Communications</i> , 2019, 10, 2154.	5.8	172
61	Epigenetic inactivation of LKB1 in primary tumors associated with the Peutz-Jeghers syndrome. <i>Oncogene</i> , 2000, 19, 164-168.	2.6	171
62	Biallelic Inactivation of Fumarate Hydratase (FH) Occurs in Nonsyndromic Uterine Leiomyomas but Is Rare in Other Tumors. <i>American Journal of Pathology</i> , 2004, 164, 17-22.	1.9	167
63	Transcription Factor PROX1 Induces Colon Cancer Progression by Promoting the Transition from Benign to Highly Dysplastic Phenotype. <i>Cancer Cell</i> , 2008, 13, 407-419.	7.7	166
64	Integrated data analysis reveals uterine leiomyoma subtypes with distinct driver pathways and biomarkers. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 1315-1320.	3.3	166
65	Genomics of uterine leiomyomas: Insights from high-throughput sequencing. <i>Fertility and Sterility</i> , 2014, 102, 621-629.	0.5	164
66	BRAF-V600E is not involved in the colorectal tumorigenesis of HNPCC in patients with functional MLH1 and MSH2 genes. <i>Oncogene</i> , 2005, 24, 3995-3998.	2.6	155
67	Screening E-cadherin in gastric cancer families reveals germline mutations only in hereditary diffuse gastric cancer kindred. <i>Human Mutation</i> , 2002, 19, 510-517.	1.1	153
68	Explaining the Familial Colorectal Cancer Risk Associated with Mismatch Repair (MMR)-Deficient and MMR-Stable Tumors. <i>Clinical Cancer Research</i> , 2007, 13, 356-361.	3.2	153
69	Features of gastric cancer in hereditary non-polyposis colorectal cancer syndrome. , 1997, 74, 551-555.		152
70	Colibactin DNA-damage signature indicates mutational impact in colorectal cancer. <i>Nature Medicine</i> , 2020, 26, 1063-1069.	15.2	149
71	Loss of SUFU Function in Familial Multiple Meningioma. <i>American Journal of Human Genetics</i> , 2012, 91, 520-526.	2.6	137
72	Serrated carcinomas form a subclass of colorectal cancer with distinct molecular basis. <i>Oncogene</i> , 2007, 26, 312-320.	2.6	136

#	ARTICLE	IF	CITATIONS
73	Mutations and impaired function of LKB1 in familial and non-familial Peutz-Jeghers syndrome and a sporadic testicular cancer. <i>Human Molecular Genetics</i> , 1999, 8, 45-51.	1.4	135
74	LKB1 exonic and whole gene deletions are a common cause of Peutz-Jeghers syndrome. <i>Journal of Medical Genetics</i> , 2005, 43, e18-e18.	1.5	135
75	Low-level microsatellite instability in most colorectal carcinomas. <i>Cancer Research</i> , 2002, 62, 1166-70.	0.4	135
76	Development of Colorectal Tumors in Colonoscopic Surveillance in Lynch Syndrome. <i>Gastroenterology</i> , 2007, 133, 1093-1098.	0.6	131
77	Genetic reconstruction of individual colorectal tumor histories. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 1236-1241.	3.3	130
78	Proportion and Phenotype of MYH-Associated Colorectal Neoplasia in a Population-Based Series of Finnish Colorectal Cancer Patients. <i>American Journal of Pathology</i> , 2003, 163, 827-832.	1.9	129
79	Distinct patterns of KRAS mutations in colorectal carcinomas according to germline mismatch repair defects and hMLH1 methylation status. <i>Human Molecular Genetics</i> , 2004, 13, 2303-2311.	1.4	127
80	Uterine Leiomyoma-Linked MED12 Mutations Disrupt Mediator-Associated CDK Activity. <i>Cell Reports</i> , 2014, 7, 654-660.	2.9	125
81	KSHV-Initiated Notch Activation Leads to Membrane-Type-1 Matrix Metalloproteinase-Dependent Lymphatic Endothelial-to-Mesenchymal Transition. <i>Cell Host and Microbe</i> , 2011, 10, 577-590.	5.1	123
82	Gene-Expression Profiling Predicts Recurrence in Duke's C Colorectal Cancer. <i>Gastroenterology</i> , 2005, 129, 874-884.	0.6	119
83	Few FH mutations in sporadic counterparts of tumor types observed in hereditary leiomyomatosis and renal cell cancer families. <i>Cancer Research</i> , 2002, 62, 4554-7.	0.4	119
84	LKB1 Somatic Mutations in Sporadic Tumors. <i>American Journal of Pathology</i> , 1999, 154, 677-681.	1.9	118
85	Cumulative impact of common genetic variants and other risk factors on colorectal cancer risk in 42,103 individuals. <i>Gut</i> , 2013, 62, 871-881.	6.1	117
86	Induction of cyclooxygenase-2 in a mouse model of Peutz-Jeghers polyposis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 12327-12332.	3.3	113
87	Frequent loss of SMAD4/DPC4 protein in colorectal cancers. <i>Gut</i> , 2002, 51, 56-59.	6.1	111
88	The MDM2 promoter polymorphism SNP309T->G and the risk of uterine leiomyosarcoma, colorectal cancer, and squamous cell carcinoma of the head and neck. <i>Journal of Medical Genetics</i> , 2005, 42, 694-698.	1.5	108
89	A large-scale meta-analysis to refine colorectal cancer risk estimates associated with MUTYH variants. <i>British Journal of Cancer</i> , 2010, 103, 1875-1884.	2.9	107
90	Epigenetic phenotypes distinguish microsatellite-stable and -unstable colorectal cancers. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1999, 96, 12661-12666.	3.3	105

#	ARTICLE	IF	CITATIONS
91	Somatic MED12 mutations in uterine leiomyosarcoma and colorectal cancer. <i>British Journal of Cancer</i> , 2012, 107, 1761-1765.	2.9	105
92	The MDM2 Promoter SNP285C/309G Haplotype Diminishes Sp1 Transcription Factor Binding and Reduces Risk for Breast and Ovarian Cancer in Caucasians. <i>Cancer Cell</i> , 2011, 19, 273-282.	7.7	104
93	MSH2 and MLH1 mutations in sporadic replication error-positive colorectal carcinoma as assessed by two-dimensional DNA electrophoresis. <i>Cancer</i> , 1997, 18, 269-278.		99
94	Candidate driver genes in microsatellite-unstable colorectal cancer. <i>International Journal of Cancer</i> , 2012, 130, 1558-1566.	2.3	99
95	The role of hypermethylation of the hMLH1 promoter region in HNPCC versus MSI+ sporadic colorectal cancers. <i>Journal of Medical Genetics</i> , 2000, 37, 588-592.	1.5	98
96	Whole-Genome Sequencing of Growth Hormone (GH)-Secreting Pituitary Adenomas. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2015, 100, 3918-3927.	1.8	96
97	Does MSI-low exist?. <i>Journal of Pathology</i> , 2002, 197, 6-13.	2.1	95
98	MED12 exon 2 mutations are common in uterine leiomyomas from South African patients. <i>Oncotarget</i> , 2011, 2, 966-969.	0.8	95
99	Dysregulation of the transcription factors SOX4, CBFβ and SMARCC1 correlates with outcome of colorectal cancer. <i>British Journal of Cancer</i> , 2009, 100, 511-523.	2.9	94
100	Microsatellite Instability in Adenomas as a Marker for Hereditary Nonpolyposis Colorectal Cancer. <i>American Journal of Pathology</i> , 1999, 155, 1849-1853.	1.9	89
101	SMAD4 Levels and Response to 5-Fluorouracil in Colorectal Cancer. <i>Clinical Cancer Research</i> , 2005, 11, 6311-6316.	3.2	89
102	MDM2 SNP309 accelerates colorectal tumour formation in women. <i>Journal of Medical Genetics</i> , 2006, 43, 950-952.	1.5	89
103	Segregation of a Missense Variant in Enteric Smooth Muscle Actin β -2 With Autosomal Dominant Familial Visceral Myopathy. <i>Gastroenterology</i> , 2012, 143, 1482-1491.e3.	0.6	89
104	Somatic microsatellite mutations as molecular tumor clocks. <i>Nature Medicine</i> , 1996, 2, 676-681.	15.2	87
105	No Support for Endoscopic Surveillance for Gastric Cancer in Hereditary Non-Polyposis Colorectal Cancer. <i>Scandinavian Journal of Gastroenterology</i> , 2002, 37, 574-577.	0.6	86
106	Close linkage to chromosome 3p and conservation of ancestral founding haplotype in hereditary nonpolyposis colorectal cancer families.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1994, 91, 6054-6058.	3.3	85
107	Mechanisms of Inactivation of the Receptor Tyrosine Kinase EPHB2 in Colorectal Tumors. <i>Cancer Research</i> , 2005, 65, 10170-10173.	0.4	84
108	DNA Copy-Number Alterations Underlie Gene Expression Differences between Microsatellite Stable and Unstable Colorectal Cancers. <i>Clinical Cancer Research</i> , 2008, 14, 8061-8069.	3.2	84

#	ARTICLE	IF	CITATIONS
109	Pro-inflammatory fatty acid profile and colorectal cancer risk: A Mendelian randomisation analysis. <i>European Journal of Cancer</i> , 2017, 84, 228-238.	1.3	81
110	Genetic and Epigenetic Characteristics of Inflammatory Bowel Disease-Associated Colorectal Cancer. <i>Gastroenterology</i> , 2021, 161, 592-607.	0.6	81
111	p53 Codon 72 and MDM2 SNP309 Polymorphisms and Age of Colorectal Cancer Onset in Lynch Syndrome. <i>Clinical Cancer Research</i> , 2005, 11, 6840-6844.	3.2	80
112	Analysis of fumarate hydratase mutations in a population-based series of early onset uterine leiomyosarcoma patients. <i>International Journal of Cancer</i> , 2006, 119, 283-287.	2.3	80
113	EPHB4 and Survival of Colorectal Cancer Patients. <i>Cancer Research</i> , 2006, 66, 8943-8948.	0.4	80
114	<i>Aryl hydrocarbon receptor interacting protein</i> (<i>AIP</i>) gene mutation analysis in children and adolescents with sporadic pituitary adenomas. <i>Clinical Endocrinology</i> , 2008, 69, 621-627.	1.2	80
115	Modifiable pathways for colorectal cancer: a mendelian randomisation analysis. <i>The Lancet Gastroenterology and Hepatology</i> , 2020, 5, 55-62.	3.7	79
116	SMAD genes in juvenile polyposis. , 1999, 26, 54-61.		78
117	Mice with Inactivation of Aryl Hydrocarbon Receptor-Interacting Protein (<i>Aip</i>) Display Complete Penetrance of Pituitary Adenomas with Aberrant <i>ARNT</i> Expression. <i>American Journal of Pathology</i> , 2010, 177, 1969-1976.	1.9	78
118	Exome sequencing reveals germline <i>NPAT</i> mutation as a candidate risk factor for Hodgkin lymphoma. <i>Blood</i> , 2011, 118, 493-498.	0.6	78
119	Unregulated smooth-muscle myosin in human intestinal neoplasia. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 5513-5518.	3.3	77
120	Mutations in the Circadian Gene <i>CLOCK</i> in Colorectal Cancer. <i>Molecular Cancer Research</i> , 2010, 8, 952-960.	1.5	77
121	Mendelian randomisation implicates hyperlipidaemia as a risk factor for colorectal cancer. <i>International Journal of Cancer</i> , 2017, 140, 2701-2708.	2.3	76
122	Direct contact in inviting high-risk members of hereditary colon cancer families to genetic counselling and DNA testing. <i>Journal of Medical Genetics</i> , 2007, 44, 732-738.	1.5	75
123	<i>HOXB13</i> G84E Mutation in Finland: Population-Based Analysis of Prostate, Breast, and Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 452-460.	1.1	75
124	Large Genomic Deletions in <i>AIP</i> in Pituitary Adenoma Predisposition. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2008, 93, 4146-4151.	1.8	74
125	Allelic Variation at the 8q23.3 Colorectal Cancer Risk Locus Functions as a Cis-Acting Regulator of <i>EIF3H</i> . <i>PLoS Genetics</i> , 2010, 6, e1001126.	1.5	74
126	Gene expression signatures for colorectal cancer microsatellite status and HNPCC. <i>British Journal of Cancer</i> , 2005, 92, 2240-2248.	2.9	70

#	ARTICLE	IF	CITATIONS
127	Allelic Imbalance at <i>rs6983267</i> Suggests Selection of the Risk Allele in Somatic Colorectal Tumor Evolution. <i>Cancer Research</i> , 2008, 68, 14-17.	0.4	69
128	Eleven Candidate Susceptibility Genes for Common Familial Colorectal Cancer. <i>PLoS Genetics</i> , 2013, 9, e1003876.	1.5	69
129	Mismatch repair genes and mononucleotide tracts as mutation targets in colorectal tumors with different degrees of microsatellite instability. <i>Oncogene</i> , 1998, 17, 157-163.	2.6	68
130	Distinct expression profile in fumarate-hydratase-deficient uterine fibroids. <i>Human Molecular Genetics</i> , 2006, 15, 97-103.	1.4	67
131	Impact of constitutional TET2 haploinsufficiency on molecular and clinical phenotype in humans. <i>Nature Communications</i> , 2019, 10, 1252.	5.8	67
132	MED12 exon 2 mutations in histopathological uterine leiomyoma variants. <i>European Journal of Human Genetics</i> , 2013, 21, 1300-1303.	1.4	66
133	Genomic profile of pseudomyxoma peritonei analyzed using next-generation sequencing and immunohistochemistry. <i>International Journal of Cancer</i> , 2015, 136, E282-9.	2.3	66
134	The I1307K polymorphism of the APC gene in colorectal cancer. <i>Gastroenterology</i> , 1999, 116, 58-63.	0.6	65
135	CHEK2 I157T associates with familial and sporadic colorectal cancer. <i>Journal of Medical Genetics</i> , 2005, 43, e34-e34.	1.5	65
136	Differential expression of DHHC9 in microsatellite stable and unstable human colorectal cancer subgroups. <i>British Journal of Cancer</i> , 2007, 96, 1896-1903.	2.9	65
137	Identification of Candidate Oncogenes in Human Colorectal Cancers With Microsatellite Instability. <i>Gastroenterology</i> , 2013, 145, 540-543.e22.	0.6	65
138	Comprehensive Analysis of SMAD4 Mutations and Protein Expression in Juvenile Polyposis. <i>American Journal of Pathology</i> , 2001, 159, 1293-1300.	1.9	64
139	Comparison of Serous and Mucinous Ovarian Carcinomas: Distinct Pattern of Allelic Loss at Distal 8p and Expression of Transcription Factor GATA-4. <i>Laboratory Investigation</i> , 2001, 81, 517-526.	1.7	64
140	Subsequent primary malignancies after endometrial carcinoma and ovarian carcinoma. <i>Cancer</i> , 2003, 97, 2432-2439.	2.0	63
141	Expression Profiling in Progressive Stages of Fumarate-Hydratase Deficiency: The Contribution of Metabolic Changes to Tumorigenesis. <i>Cancer Research</i> , 2010, 70, 9153-9165.	0.4	63
142	MED12 mutation frequency in unselected sporadic uterine leiomyomas. <i>Fertility and Sterility</i> , 2014, 102, 1137-1142.	0.5	62
143	Exome-wide somatic mutation characterization of small bowel adenocarcinoma. <i>PLoS Genetics</i> , 2018, 14, e1007200.	1.5	62
144	Clusterin Expression in Normal Mucosa and Colorectal Cancer. <i>Molecular and Cellular Proteomics</i> , 2007, 6, 1039-1048.	2.5	61

#	ARTICLE	IF	CITATIONS
145	Refinement of the basis and impact of common 11q23.1 variation to the risk of developing colorectal cancer. <i>Human Molecular Genetics</i> , 2008, 17, 3720-3727.	1.4	61
146	The Expression of AIP-Related Molecules in Elucidation of Cellular Pathways in Pituitary Adenomas. <i>American Journal of Pathology</i> , 2009, 175, 2501-2507.	1.9	61
147	Prevalence of Germline PTEN, BMPR1A, SMAD4, STK11, and ENG Mutations in Patients With Moderate-Load Colorectal Polyps. <i>Gastroenterology</i> , 2013, 144, 1402-1409.e5.	0.6	61
148	Brush border Myosin Ia has tumor suppressor activity in the intestine. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 1530-1535.	3.3	60
149	Frequent L1 retrotranspositions originating from <i>TTC28</i> in colorectal cancer. <i>Oncotarget</i> , 2014, 5, 853-859.	0.8	60
150	Lessons from Functional Analysis of Genome-Wide Association Studies. <i>Cancer Research</i> , 2013, 73, 4180-4184.	0.4	58
151	Genetic predisposition to uterine leiomyoma is determined by loci for genitourinary development and genome stability. <i>ELife</i> , 2018, 7, .	2.8	58
152	Diagnostic Cancer Genome Sequencing and the Contribution of Germline Variants. <i>Science</i> , 2013, 339, 1559-1562.	6.0	57
153	Mendelian randomisation analysis strongly implicates adiposity with risk of developing colorectal cancer. <i>British Journal of Cancer</i> , 2016, 115, 266-272.	2.9	57
154	Colorectal Adenoma and Cancer Divergence. <i>American Journal of Pathology</i> , 1999, 154, 1815-1824.	1.9	56
155	Mutation analysis of aryl hydrocarbon receptor interacting protein (AIP) gene in colorectal, breast, and prostate cancers. <i>British Journal of Cancer</i> , 2007, 96, 352-356.	2.9	56
156	Retrotransposon insertions can initiate colorectal cancer and are associated with poor survival. <i>Nature Communications</i> , 2019, 10, 4022.	5.8	53
157	Deficient H2A.Z deposition is associated with genesis of uterine leiomyoma. <i>Nature</i> , 2021, 596, 398-403.	13.7	53
158	High Familial Risk in Nodular Lymphocyte-Predominant Hodgkin Lymphoma. <i>Journal of Clinical Oncology</i> , 2013, 31, 938-943.	0.8	51
159	Exome sequencing reveals three novel candidate predisposition genes for diffuse gastric cancer. <i>Familial Cancer</i> , 2015, 14, 241-246.	0.9	50
160	Analysis of a Finnish family confirms RHBDF2 mutations as the underlying factor in tylosis with esophageal cancer. <i>Familial Cancer</i> , 2012, 11, 525-528.	0.9	49
161	No evidence of somatic aryl hydrocarbon receptor interacting protein mutations in sporadic endocrine neoplasia. <i>Endocrine-Related Cancer</i> , 2007, 14, 901-906.	1.6	48
162	Recessively inherited right atrial isomerism caused by mutations in growth/differentiation factor 1 (GDF1). <i>Human Molecular Genetics</i> , 2010, 19, 2747-2753.	1.4	48

#	ARTICLE	IF	CITATIONS
163	Conventional renal cancer in a patient with fumarate hydratase mutation. <i>Human Pathology</i> , 2007, 38, 793-796.	1.1	47
164	Carbonic Anhydrase IX Is Highly Expressed in Hereditary Nonpolyposis Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 1760-1766.	1.1	46
165	Germline and Somatic Mutation Analysis of MLH3 in MSI-Positive Colorectal Cancer. <i>American Journal of Pathology</i> , 2000, 157, 347-352.	1.9	45
166	Common Deletion of SMAD4 in Juvenile Polyposis Is a Mutational Hotspot. <i>American Journal of Human Genetics</i> , 2002, 70, 1357-1362.	2.6	45
167	Aggressive pituitary adenomas occurring in young patients in a large Polynesian kindred with a germline R271W mutation in the AIP gene. <i>European Journal of Endocrinology</i> , 2009, 161, 799-804.	1.9	45
168	Multiple clinical characteristics separate MED12-mutation-positive and -negative uterine leiomyomas. <i>Scientific Reports</i> , 2017, 7, 1015.	1.6	44
169	Oncogenic exon 2 mutations in Mediator subunit MED12 disrupt allosteric activation of cyclin C-CDK8/19. <i>Journal of Biological Chemistry</i> , 2018, 293, 4870-4882.	1.6	44
170	COGENT (COlorectal cancer GENEtics): an international consortium to study the role of polymorphic variation on the risk of colorectal cancer. <i>British Journal of Cancer</i> , 2010, 102, 447-454.	2.9	43
171	MED12 mutations and FH inactivation are mutually exclusive in uterine leiomyomas. <i>British Journal of Cancer</i> , 2016, 114, 1405-1411.	2.9	43
172	Little evidence for involvement of MLH3 in colorectal cancer predisposition. <i>International Journal of Cancer</i> , 2003, 106, 292-296.	2.3	42
173	A Finnish founder mutation in <i>RAD51D</i> : analysis in breast, ovarian, prostate, and colorectal cancer: Table 1. <i>Journal of Medical Genetics</i> , 2012, 49, 429-432.	1.5	41
174	Background Mutation Frequency in Microsatellite-Unstable Colorectal Cancer. <i>Cancer Research</i> , 2007, 67, 5691-5698.	0.4	38
175	High Frequency of <i>RPL22</i> Mutations in Microsatellite-Unstable Colorectal and Endometrial Tumors. <i>Human Mutation</i> , 2014, 35, 1442-1445.	1.1	38
176	Intestinal cancer in patients with a germline mutation in the down-regulated in adenoma (DRA) gene. <i>Oncogene</i> , 1998, 16, 681-684.	2.6	37
177	Variation at 2q35 (<i>PNKD</i> and <i>TMBIM1</i>) influences colorectal cancer risk and identifies a pleiotropic effect with inflammatory bowel disease. <i>Human Molecular Genetics</i> , 2016, 25, 2349-2359.	1.4	37
178	CHEK2 1100delC and colorectal cancer. <i>Journal of Medical Genetics</i> , 2003, 40, 110e-110.	1.5	36
179	Exomic landscape of <i>MED12</i> mutation-negative and -positive uterine leiomyomas. <i>International Journal of Cancer</i> , 2014, 134, 1008-1012.	2.3	36
180	Germline mutations in young non-smoking women with lung adenocarcinoma. <i>Lung Cancer</i> , 2018, 122, 76-82.	0.9	36

#	ARTICLE	IF	CITATIONS
181	No evidence of RET germline mutations in familial pituitary adenoma. <i>Journal of Molecular Endocrinology</i> , 2011, 46, 1-8.	1.1	35
182	Complexity of 12q13â€“22 amplicon in liposarcoma: Microsatellite repeat analysis. , 1997, 18, 66-79.		34
183	No SMAD4 hypermethylation in colorectal cancer. <i>British Journal of Cancer</i> , 2000, 83, 1015-1019.	2.9	34
184	NOD2 3020insC Alone Is Not Sufficient for Colorectal Cancer Predisposition. <i>Cancer Research</i> , 2004, 64, 7245-7247.	0.4	34
185	An A13 Repeat within the 3â€“Untranslated Region of Epidermal Growth Factor Receptor (EGFR) Is Frequently Mutated in Microsatellite Instability Colon Cancers and Is Associated with Increased EGFR Expression. <i>Cancer Research</i> , 2009, 69, 7811-7818.	0.4	34
186	Screening for microsatellite instability target genes in colorectal cancers. <i>Journal of Medical Genetics</i> , 2002, 39, 785-789.	1.5	33
187	7q deletion mapping and expression profiling in uterine fibroids. <i>Oncogene</i> , 2005, 24, 6545-6554.	2.6	33
188	Preferential amplification of AURKA 91A (Ile31) in familial colorectal cancers. <i>International Journal of Cancer</i> , 2006, 118, 505-508.	2.3	33
189	Functional Diversity of Human Protection of Telomeres 1 Isoforms in Telomere Protection and Cellular Senescence. <i>Cancer Research</i> , 2007, 67, 11677-11686.	0.4	33
190	Hereditary intestinal cancer. <i>Seminars in Cancer Biology</i> , 2000, 10, 289-298.	4.3	32
191	Allelic Analysis of Serous Ovarian Carcinoma Reveals Two Putative Tumor Suppressor Loci at 18q22-q23 Distal to SMAD4, SMAD2, and DCC. <i>American Journal of Pathology</i> , 2001, 159, 35-42.	1.9	32
192	The C/C-13910 genotype of adult-type hypolactasia is associated with an increased risk of colorectal cancer in the Finnish population. <i>Gut</i> , 2005, 54, 643-647.	6.1	32
193	Downregulation of SRFâ€“FOSâ€“JUNB pathway in fumarate hydratase deficiency and in uterine leiomyomas. <i>Oncogene</i> , 2009, 28, 1261-1273.	2.6	31
194	Phosphoprotein Keratin 23 accumulates in MSS but not MSI colon cancers in vivo and impacts viability and proliferation in vitro. <i>Molecular Oncology</i> , 2007, 1, 181-195.	2.1	30
195	Mutation screening of fumarate hydratase by multiplex ligation-dependent probe amplification: detection of exonic deletion in a patient with leiomyomatosis and renal cell cancer. <i>Cancer Genetics and Cytogenetics</i> , 2008, 183, 83-88.	1.0	30
196	Identification of 33 candidate oncogenes by screening for base-specific mutations. <i>British Journal of Cancer</i> , 2014, 111, 1657-1662.	2.9	30
197	E-cadherin is not frequently mutated in hereditary gastric cancer. <i>Journal of Medical Genetics</i> , 2001, 38, 49-52.	1.5	30
198	Mutation analysis of three genes encoding novel LKB1-interacting proteins, BRG1, STRAD1, and MO251, in Peutzâ€“Jeghers syndrome. <i>British Journal of Cancer</i> , 2005, 92, 1126-1129.	2.9	29

#	ARTICLE	IF	CITATIONS
199	Susceptibility to pituitary neoplasia related to MEN-1, CDKN1B and AIP mutations: an update. <i>Human Molecular Genetics</i> , 2007, 16, R73-R79.	1.4	29
200	Global metabolomic profiling of uterine leiomyomas. <i>British Journal of Cancer</i> , 2017, 117, 1855-1864.	2.9	29
201	Germline fumarate hydratase mutations in patients with ovarian mucinous cystadenoma. <i>European Journal of Human Genetics</i> , 2006, 14, 880-883.	1.4	28
202	Variants in the Netrin-1 Receptor UNC5C Prevent Apoptosis and Increase Risk of Familial Colorectal Cancer. <i>Gastroenterology</i> , 2011, 141, 2039-2046.	0.6	28
203	Strong family history of uterine leiomyomatosis warrants fumarate hydratase mutation screening. <i>Human Reproduction</i> , 2012, 27, 1865-1869.	0.4	28
204	Villin Expression Is Frequently Lost in Poorly Differentiated Colon Cancer. <i>American Journal of Pathology</i> , 2012, 180, 1509-1521.	1.9	28
205	COGENT (COlorectal cancer GENEtics) revisited. <i>Mutagenesis</i> , 2012, 27, 143-151.	1.0	27
206	Discovery of potential causative mutations in human coding and noncoding genome with the interactive software BasePlayer. <i>Nature Protocols</i> , 2018, 13, 2580-2600.	5.5	27
207	Mutational processes of distinct POLE exonuclease domain mutants drive an enrichment of a specific TP53 mutation in colorectal cancer. <i>PLoS Genetics</i> , 2020, 16, e1008572.	1.5	27
208	Somatic mutation analysis of MYH11 in breast and prostate cancer. <i>BMC Cancer</i> , 2008, 8, 263.	1.1	26
209	No evidence for a genetic modifier for renal cell cancer risk in HLRCC syndrome. <i>Familial Cancer</i> , 2010, 9, 245-251.	0.9	26
210	Somatic mutations and germline sequence variants in patients with familial colorectal cancer. <i>International Journal of Cancer</i> , 2010, 127, 2974-2980.	2.3	26
211	Mixed lineage kinase 3 gene mutations in mismatch repair deficient gastrointestinal tumours. <i>Human Molecular Genetics</i> , 2010, 19, 697-706.	1.4	26
212	Genome-wide association study and meta-analysis in Northern European populations replicate multiple colorectal cancer risk loci. <i>International Journal of Cancer</i> , 2018, 142, 540-546.	2.3	26
213	CDX2 mutations do not account for juvenile polyposis or Peutz-Jeghers syndrome and occur infrequently in sporadic colorectal cancers. <i>British Journal of Cancer</i> , 2001, 84, 1314-1316.	2.9	25
214	Germline deletions of EXO1 do not cause colorectal tumors and lesions which are null for EXO1 do not have microsatellite instability. <i>Cancer Genetics and Cytogenetics</i> , 2003, 147, 121-127.	1.0	25
215	Colorectal Pretumor Progression Before and After Loss of DNA Mismatch Repair. <i>American Journal of Pathology</i> , 2004, 164, 1447-1453.	1.9	25
216	Tumour selection advantage of non-dominant negative P53 mutations in homozygotic MDM2-SNP309 colorectal cancer cells. <i>Journal of Medical Genetics</i> , 2006, 44, 75-80.	1.5	25

#	ARTICLE	IF	CITATIONS
217	Array comparative genomic hybridization identifies a distinct DNA copy number profile in renal cell cancer associated with hereditary leiomyomatosis and renal cell cancer. <i>Genes Chromosomes and Cancer</i> , 2009, 48, 544-551.	1.5	25
218	Identification of homozygous deletion in <i>ACAN</i> and other candidate variants in familial classical Hodgkin lymphoma by exome sequencing. <i>British Journal of Haematology</i> , 2015, 170, 428-431.	1.2	25
219	Whole-Genome Sequencing Identifies <i>STAT4</i> as a Putative Susceptibility Gene in Classic Kaposi Sarcoma. <i>Journal of Infectious Diseases</i> , 2015, 211, 1842-1851.	1.9	25
220	Contribution of allelic imbalance to colorectal cancer. <i>Nature Communications</i> , 2018, 9, 3664.	5.8	25
221	Characterization of the colorectal cancer-associated enhancer MYC-335 at 8q24: the role of rs67491583. <i>Cancer Genetics</i> , 2012, 205, 25-33.	0.2	24
222	Systematic search for rare variants in Finnish early-onset colorectal cancer patients. <i>Cancer Genetics</i> , 2015, 208, 35-40.	0.2	24
223	Detection of subclonal L1 transductions in colorectal cancer by long-distance inverse-PCR and Nanopore sequencing. <i>Scientific Reports</i> , 2017, 7, 14521.	1.6	24
224	Tracing Cell Fates in Human Colorectal Tumors from Somatic Microsatellite Mutations. <i>American Journal of Pathology</i> , 1998, 153, 1189-1200.	1.9	23
225	Characterization of the 17p amplicon in human sarcomas: Microsatellite marker analysis. , 1999, 82, 329-333.		23
226	Smad4 haploinsufficiency: a matter of dosage. <i>PathoGenetics</i> , 2008, 1, 2.	5.7	23
227	Candidate susceptibility variants for esophageal squamous cell carcinoma. <i>Genes Chromosomes and Cancer</i> , 2017, 56, 453-459.	1.5	23
228	Low-Penetrance Susceptibility Variants in Familial Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 1478-1483.	1.1	22
229	Cancer risk in mutation carriers of DNA-mismatch-repair genes. , 1999, 81, 214.		22
230	Population distribution and ancestry of the cancer protective MDM2 SNP285 (rs117039649). <i>Oncotarget</i> , 2014, 5, 8223-8234.	0.8	22
231	No evidence for association of NOD2 R702W and G908R with colorectal cancer. <i>International Journal of Cancer</i> , 2007, 121, 76-79.	2.3	21
232	Increased HIF1 α in SDH and FH deficient tumors does not cause microsatellite instability. <i>International Journal of Cancer</i> , 2007, 121, 1386-1389.	2.3	21
233	Primary mediastinal large B-cell lymphoma segregating in a family: exome sequencing identifies MLL as a candidate predisposition gene. <i>Blood</i> , 2013, 121, 3428-3430.	0.6	21
234	A novel functionally deficient MYH variant in individuals with colorectal adenomatous polyposis. <i>Human Mutation</i> , 2005, 26, 393-393.	1.1	19

#	ARTICLE	IF	CITATIONS
235	EPHB2 germline variants in patients with colorectal cancer or hyperplastic polyposis. <i>BMC Cancer</i> , 2006, 6, 145.	1.1	19
236	Refinement of the associations between risk of colorectal cancer and polymorphisms on chromosomes 1q41 and 12q13.13. <i>Human Molecular Genetics</i> , 2012, 21, 934-946.	1.4	19
237	Clinical characterization, genetic mapping and whole-genome sequence analysis of a novel autosomal recessive intellectual disability syndrome. <i>European Journal of Medical Genetics</i> , 2014, 57, 543-551.	0.7	19
238	Mutation analysis of components of the Mediator kinase module in MED12 mutation-negative uterine leiomyomas. <i>British Journal of Cancer</i> , 2014, 110, 2246-2249.	2.9	19
239	Clonally related uterine leiomyomas are common and display branched tumor evolution. <i>Human Molecular Genetics</i> , 2015, 24, 4407-4416.	1.4	19
240	Biallelic loss-of-function P4HTM gene variants cause hypotonia, hypoventilation, intellectual disability, dysautonomia, epilepsy, and eye abnormalities (HIDEA syndrome). <i>Genetics in Medicine</i> , 2019, 21, 2355-2363.	1.1	19
241	No evidence of microsatellite instability in bone tumours. <i>British Journal of Cancer</i> , 1996, 74, 453-455.	2.9	18
242	No evidence for dual role of loss of heterozygosity in hereditary non-polyposis colorectal cancer. <i>Oncogene</i> , 2007, 26, 2513-2517.	2.6	18
243	Nationwide Registry-Based Analysis of Cancer Clustering Detects Strong Familial Occurrence of Kaposi Sarcoma. <i>PLoS ONE</i> , 2013, 8, e55209.	1.1	18
244	Comprehensive Evaluation of Protein Coding Mononucleotide Microsatellites in Microsatellite-Unstable Colorectal Cancer. <i>Cancer Research</i> , 2017, 77, 4078-4088.	0.4	18
245	PPP2R1B Gene in Chronic Lymphocytic Leukemias and Mantle Cell Lymphomas. <i>Leukemia and Lymphoma</i> , 2001, 41, 177-183.	0.6	16
246	Genetic predisposition and somatic diversification in tumor development and progression. <i>Advances in Cancer Research</i> , 2001, 80, 83-114.	1.9	16
247	Genome-wide allelotyping of 104 Finnish colorectal cancers reveals an excess of allelic imbalance in chromosome 20q in familial cases. <i>Oncogene</i> , 2003, 22, 2206-2214.	2.6	16
248	Numbers of mutations to different types of colorectal cancer. <i>BMC Cancer</i> , 2005, 5, 126.	1.1	16
249	Mutations in Two Short Noncoding Mononucleotide Repeats in Most Microsatellite-Unstable Colorectal Cancers. <i>Cancer Research</i> , 2005, 65, 4607-4613.	0.4	16
250	Systematic search for enhancer elements and somatic allelic imbalance at seven low-penetrance colorectal cancer predisposition loci. <i>BMC Medical Genetics</i> , 2011, 12, 23.	2.1	16
251	Genetic and Epigenetic Characterization of Growth Hormone- α -Secreting Pituitary Tumors. <i>Molecular Cancer Research</i> , 2019, 17, 2432-2443.	1.5	16
252	From <i>APC</i> to the genetics of hereditary and familial colon cancer syndromes. <i>Human Molecular Genetics</i> , 2021, 30, R206-R224.	1.4	15

#	ARTICLE	IF	CITATIONS
253	Multiple components of PKA and TGF- β pathways are mutated in pseudomyxoma peritonei. PLoS ONE, 2017, 12, e0174898.	1.1	15
254	PolyA Deletions in Hereditary Nonpolyposis Colorectal Cancer. American Journal of Pathology, 2002, 160, 1503-1506.	1.9	14
255	High frequency of TTK mutations in microsatellite-unstable colorectal cancer and evaluation of their effect on spindle assembly checkpoint. Carcinogenesis, 2011, 32, 305-311.	1.3	14
256	Exome sequencing in diagnostic evaluation of colorectal cancer predisposition in young patients. Scandinavian Journal of Gastroenterology, 2013, 48, 672-678.	0.6	14
257	No MSH6 germline mutations in breast cancer families with colorectal and/or endometrial cancer. Journal of Medical Genetics, 2005, 42, e22-e22.	1.5	13
258	Chromothripsis in Uterine Leiomyomas. New England Journal of Medicine, 2013, 369, 2160-2161.	13.9	13
259	ARLTS1 germline variants and the risk for breast, prostate, and colorectal cancer. European Journal of Human Genetics, 2008, 16, 983-991.	1.4	12
260	Human cell transformation by combined lineage conversion and oncogene expression. Oncogene, 2021, 40, 5533-5547.	2.6	12
261	Whole-exome sequencing identifies novel candidate predisposition genes for familial polycythemia vera. Human Genomics, 2017, 11, 6.	1.4	11
262	A missense mutation in the BRCA2 gene in three siblings with ovarian cancer. British Journal of Cancer, 1998, 77, 1199-1202.	2.9	10
263	Definition of a minimal region of deletion of chromosome 7 in uterine leiomyomas by tiling-path microarray CGH and mutation analysis of known genes in this region. Genes Chromosomes and Cancer, 2007, 46, 451-458.	1.5	10
264	New Target Genes in Endometrial Tumors Show a Role for the Estrogen-Receptor Pathway in Microsatellite-Unstable Cancers. Human Mutation, 2014, 35, 1514-1523.	1.1	10
265	Comprehensive evaluation of coding region point mutations in microsatellite-unstable colorectal cancer. EMBO Molecular Medicine, 2018, 10, .	3.3	10
266	Microsatellite markers as tools for characterization of DNA amplifications evaluated by comparative genomic hybridization. Cancer Genetics and Cytogenetics, 1997, 93, 33-38.	1.0	9
267	Mutation and LOH analysis of ACO2 in colorectal cancer: no evidence of biallelic genetic inactivation. Journal of Medical Genetics, 2003, 40, 73e-73.	1.5	9
268	No fumarate hydratase (FH) mutations in hereditary prostate cancer. Journal of Medical Genetics, 2003, 40, 19e-19.	1.5	8
269	No germline FH mutations in familial breast cancer patients. European Journal of Human Genetics, 2005, 13, 506-509.	1.4	8
270	Screening of Finnish RAD51C founder mutations in prostate and colorectal cancer patients. BMC Cancer, 2012, 12, 552.	1.1	8

#	ARTICLE	IF	CITATIONS
271	Candidate susceptibility variants in angioimmunoblastic T-cell lymphoma. <i>Familial Cancer</i> , 2019, 18, 113-119.	0.9	8
272	Comparison of 2SC, AKR1B10, and FH Antibodies as Potential Biomarkers for FH-deficient Uterine Leiomyomas. <i>American Journal of Surgical Pathology</i> , 2022, 46, 537-546.	2.1	8
273	Histopathologic and Molecular Characterization of Uterine Leiomyoma-like Inflammatory Myofibroblastic Tumor. <i>American Journal of Surgical Pathology</i> , 2022, 46, 1126-1136.	2.1	8
274	Mutation analysis of SMAD2, SMAD3, and SMAD4 genes in hereditary non-polyposis colorectal cancer. <i>Journal of Medical Genetics</i> , 2000, 37, 298-301.	1.5	7
275	Modeling tumor predisposing FH mutations in yeast: Effects on fumarase activity, growth phenotype and gene expression profile. <i>International Journal of Cancer</i> , 2006, 118, 1340-1345.	2.3	7
276	Aryl Hydrocarbon Receptor-Interacting Protein and Acromegaly. <i>Hormone Research in Paediatrics</i> , 2007, 68, 127-131.	0.8	7
277	Analysis of <i>KLHDC8B</i> in familial nodular lymphocyte predominant Hodgkin lymphoma. <i>British Journal of Haematology</i> , 2011, 154, 413-415.	1.2	7
278	Blood-derived gene-expression profiling in unravelling susceptibility to recessive disease. <i>Journal of Medical Genetics</i> , 2007, 44, 718-720.	1.5	6
279	Lack of an association between gallstone disease and bilirubin levels with risk of colorectal cancer: a Mendelian randomisation analysis. <i>British Journal of Cancer</i> , 2021, 124, 1169-1174.	2.9	6
280	<i>WNT2</i> activation through proximal germline deletion predisposes to small intestinal neuroendocrine tumors and intestinal adenocarcinomas. <i>Human Molecular Genetics</i> , 2021, 30, 2429-2440.	1.4	6
281	A 17p11.2 germline deletion in a patient with Smith-Magenis syndrome and neuroblastoma. <i>Journal of Medical Genetics</i> , 2005, 42, e3-e3.	1.5	5
282	Downregulation of the hedgehog receptor PTCH1 in colorectal serrated adenocarcinomas is not caused by PTCH1 mutations. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2011, 458, 213-219.	1.4	5
283	Candidate Predisposition Variants in Kaposi Sarcoma as Detected by Whole-Genome Sequencing. <i>Open Forum Infectious Diseases</i> , 2019, 6, ofz337.	0.4	5
284	Prognostic Value of Immune Environment Analysis in Small Bowel Adenocarcinomas with Verified Mutational Landscape and Predisposing Conditions. <i>Cancers</i> , 2020, 12, 2018.	1.7	5
285	No evidence of EMAST in whole genome sequencing data from 248 colorectal cancers. <i>Genes Chromosomes and Cancer</i> , 2021, 60, 463-473.	1.5	5
286	Identification of ZBTB18 as a novel colorectal tumor suppressor gene through genome-wide promoter hypermethylation analysis. <i>Clinical Epigenetics</i> , 2021, 13, 88.	1.8	5
287	Uterine leiomyomas in hereditary leiomyomatosis and renal cell cancer (HLRCC) syndrome can be identified through distinct clinical characteristics and typical morphology. <i>Acta Obstetrica Et Gynecologica Scandinavica</i> , 2021, 100, 2066-2075.	1.3	5
288	Defining eligible patients for allele-selective chemotherapies targeting NAT2 in colorectal cancer. <i>Scientific Reports</i> , 2020, 10, 22436.	1.6	5

#	ARTICLE	IF	CITATIONS
289	Role of TP53 P72R polymorphism in human papillomavirus associated premalignant laryngeal neoplasm. <i>Journal of Medical Genetics</i> , 2001, 38, 327-327.	1.5	5
290	Trilateral Retinoblastoma in a Patient With Peutz-Jeghers Syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 1096-1100.	0.7	4
291	Germline MSH6 Mutation in a Patient With Two Independent Primary Glioblastomas. <i>Journal of Neuropathology and Experimental Neurology</i> , 2017, 76, 848-853.	0.9	4
292	Exome and immune cell score analyses reveal great variation within synchronous primary colorectal cancers. <i>British Journal of Cancer</i> , 2019, 120, 922-930.	2.9	4
293	Cancer risk in mutation carriers of DNA-mismatch-repair genes. , 1999, 81, 214.		4
294	MSH2 and MLH1 mutations in sporadic replication error-positive colorectal carcinoma as assessed by two-dimensional DNA electrophoresis. <i>Genes Chromosomes and Cancer</i> , 1997, 18, 269-278.	1.5	4
295	Multiple colorectal adenomas, familial adenomatous polyposis and germline mutations in MYH. <i>Gastroenterology</i> , 2003, 124, A45.	0.6	2
296	Computational identification of candidate loci for recessively inherited mutation using high-throughput SNP arrays. <i>Bioinformatics</i> , 2007, 23, 1952-1961.	1.8	2
297	3'-UTR poly(T/U) repeat of EWSR1 is altered in microsatellite unstable colorectal cancer with nearly perfect sensitivity. <i>Familial Cancer</i> , 2015, 14, 449-453.	0.9	2
298	Novel germline variant in the histone demethylase and transcription regulator KDM4C induces a multi-cancer phenotype. <i>Journal of Medical Genetics</i> , 2021, , jmedgenet-2021-107747.	1.5	2
299	Next-generation sequencing in a large pedigree segregating visceral artery aneurysms suggests potential role of COL4A1/COL4A2 in disease etiology. <i>Vascular</i> , 2021, , 170853812110331.	0.4	2
300	Parity associates with chromosomal damage in uterine leiomyomas. <i>Nature Communications</i> , 2021, 12, 5448.	5.8	2
301	How Many Colon Cancer Genes?. <i>Annals of Medicine</i> , 1995, 27, 287-288.	1.5	1
302	Stress-Induced Expression of a Novel Variant of Human Fumarate Hydratase (FH). <i>Gene Expression</i> , 2007, 14, 59-69.	0.5	1
303	Mutation analysis of MYH11 in acute myeloid leukemia. <i>Leukemia and Lymphoma</i> , 2008, 49, 1829-1831.	0.6	1
304	Abstract 4435: Genome-scale DNA methylation changes delineate uterine leiomyoma subgroups. , 2016, , .		1
305	Abstract 5193: Novel candidate oncogenes with mutation hot spots in microsatellite unstable colorectal cancer. , 2014, , .		1
306	Direct Sequencing for Peutz-Jeghers Gene LKB1 (STK11) Mutations. , 2001, 50, 175-183.		0

#	ARTICLE	IF	CITATIONS
307	Reply to the Letter to the Editor by Watanabe et al.. Clinical Cancer Research, 2006, 12, 1654.1-1655.	3.2	0
308	Rule-based induction method for haplotype comparison and identification of candidate disease loci. Genome Medicine, 2012, 4, 21.	3.6	0
309	Abstract 5756: The receptor tyrosine kinase EPHB4 has tumor suppressor activities in intestinal tumorigenesis. , 2010, , .		0
310	Abstract 3864: Low-penetrance variants underlying familial colorectal cancer. , 2010, , .		0
311	Abstract 1844: No evidence for genetic factor increasing renal cell cancer risk in Hereditary Leiomyomatosis and Renal Cell Cancer syndrome. , 2010, , .		0
312	Abstract LB-121: Serrated colorectal adenocarcinoma: Specific copy number alterations. , 2010, , .		0
313	Abstract 1161: The role of MED12 exon 2 mutations in histopathological variants of uterine leiomyoma. , 2012, , .		0
314	Abstract 105: Characterizing the exomic profile of MSI colorectal cancer. , 2012, , .		0
315	Abstract 2976: KSHV-initiated Notch activation leads to membrane-type-1 matrix metalloproteinase-dependent lymphatic endothelial-to-mesenchymal transition. , 2012, , .		0
316	Abstract 4316: Villin expression is frequently lost in colon cancers with microsatellite instability.. , 2012, , .		0
317	Abstract 3514: MED12 and FH mutations in HLRCC associated uterine leiomyomas. , 2014, , .		0
318	Abstract 2401: Identification of new target genes in microsatellite unstable colorectal cancer by exome sequencing. , 2014, , .		0
319	Abstract 2744: Familial multiple metastatic small intestine neuroendocrine tumors: searching for genetic susceptibility. , 2015, , .		0
320	Abstract 4800: Identification of candidate predisposition genes for familial uterine leiomyomas. , 2015, , .		0
321	Abstract 2176: Joint structural variant analysis of colorectal cancer whole genome sequencing data. , 2015, , .		0
322	Abstract 1079: Transcriptional profiling reveals uterine leiomyoma subtypes with distinct pathways and biomarkers of tumorigenesis. , 2015, , .		0
323	Abstract 171: Comparative genomic analyses of synchronous colorectal cancers by exome sequencing. , 2016, , .		0
324	Abstract 5281: Fast and scalable software for comparative variant analysis and visualization of massive next-generation sequencing data. , 2016, , .		0

#	ARTICLE	IF	CITATIONS
325	Abstract 120: Estrogen and progesterone receptor expression in different molecular uterine leiomyoma subclasses. , 2016, , .		0
326	Abstract 4381: The mobile genome of colorectal cancer: Characterization of retrotransposon insertions in 202 colorectal cancer whole genomes. , 2017, , .		0
327	Abstract 4379: Somatic exomic landscape of small intestinal adenocarcinomas. , 2017, , .		0
328	Abstract 1440: Germline loss-of-function alleles in Finnish colorectal cancer patients. , 2017, , .		0
329	Abstract 2461: Molecular classification and clinical characterization of a large uterine leiomyoma patient cohort. , 2017, , .		0
330	Abstract 1457: Somatic biallelic inactivation offumarate hydratase(FH) in uterine leiomyomas. , 2017, , .		0
331	Abstract 1159: Development of uterine leiomyoma 3D in vitro models for high-throughput drug and chemical compound screenings: Towards personalized medicine. , 2018, , .		0
332	Abstract LB-375: The landscape of somatic mutations in uterine adenomyomas. , 2018, , .		0
333	Abstract LB-382: Identification of predisposing genes for small bowel adenocarcinoma by exome sequencing. , 2018, , .		0
334	Title is missing!. , 2020, 16, e1008572.		0
335	Title is missing!. , 2020, 16, e1008572.		0
336	Title is missing!. , 2020, 16, e1008572.		0
337	Title is missing!. , 2020, 16, e1008572.		0