

Aung Ko Win

List of Publications by Year in descending order

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185
papers

8,545
citations

44069

48
h-index

54911

84
g-index

195
all docs

195
docs citations

195
times ranked

11134
citing authors

#	ARTICLE	IF	CITATIONS
1	Risk Stratification for Early-Onset Colorectal Cancer Using a Combination of Genetic and Environmental Risk Scores: An International Multi-Center Study. <i>Journal of the National Cancer Institute</i> , 2022, , .	6.3	15
2	Genetic Regulation of DNA Methylation Yields Novel Discoveries in GWAS of Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 1068-1076.	2.5	1
3	Analysis in the Prospective Lynch Syndrome Database identifies sarcoma as part of the Lynch syndrome tumor spectrum. <i>International Journal of Cancer</i> , 2021, 148, 512-513.	5.1	9
4	Risk-reducing hysterectomy and bilateral salpingo-oophorectomy in female heterozygotes of pathogenic mismatch repair variants: a Prospective Lynch Syndrome Database report. <i>Genetics in Medicine</i> , 2021, 23, 705-712.	2.4	28
5	Associations of Height With the Risks of Colorectal and Endometrial Cancer in Persons With Lynch Syndrome. <i>American Journal of Epidemiology</i> , 2021, 190, 230-238.	3.4	2
6	Exposure to household air pollution over 10 years is related to asthma and lung function decline. <i>European Respiratory Journal</i> , 2021, 57, 2000602.	6.7	18
7	Evaluating the utility of tumour mutational signatures for identifying hereditary colorectal cancer and polyposis syndrome carriers. <i>Gut</i> , 2021, 70, 2138-2149.	12.1	27
8	Germline and Tumor Sequencing as a Diagnostic Tool To Resolve Suspected Lynch Syndrome. <i>Journal of Molecular Diagnostics</i> , 2021, 23, 358-371.	2.8	12
9	Response to Li and Hopper. <i>American Journal of Human Genetics</i> , 2021, 108, 527-529.	6.2	5
10	Assessment of a Polygenic Risk Score for Colorectal Cancer to Predict Risk of Lynch Syndrome Colorectal Cancer. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab022.	2.9	15
11	Uptake of hysterectomy and bilateral salpingo-oophorectomy in carriers of pathogenic mismatch repair variants: a Prospective Lynch Syndrome Database report. <i>European Journal of Cancer</i> , 2021, 148, 124-133.	2.8	11
12	Nongenetic Determinants of Risk for Early-Onset Colorectal Cancer. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab029.	2.9	39
13	DNA Methylation Signatures and the Contribution of Age-Associated Methylomic Drift to Carcinogenesis in Early-Onset Colorectal Cancer. <i>Cancers</i> , 2021, 13, 2589.	3.7	18
14	A Meta-Analysis of Obesity and Risk of Colorectal Cancer in Patients with Lynch Syndrome: The Impact of Sex and Genetics. <i>Nutrients</i> , 2021, 13, 1736.	4.1	10
15	No Difference in Penetrance between Truncating and Missense/Aberrant Splicing Pathogenic Variants in MLH1 and MSH2: A Prospective Lynch Syndrome Database Study. <i>Journal of Clinical Medicine</i> , 2021, 10, 2856.	2.4	11
16	Variation in the risk of colorectal cancer in families with Lynch syndrome: a retrospective cohort study. <i>Lancet Oncology</i> , The, 2021, 22, 1014-1022.	10.7	58
17	DNA methylation-based signature of CD8+ tumor-infiltrating lymphocytes enables evaluation of immune response and prognosis in colorectal cancer. , 2021, 9, e002671.		37
18	Do the risks of Lynch syndrome-related cancers depend on the parent-of-origin of the mutation?. <i>International Journal of Epidemiology</i> , 2021, 50, .	1.9	0

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19	Cancer risks by gene, age, and gender in 6350 carriers of pathogenic mismatch repair variants: findings from the Prospective Lynch Syndrome Database. <i>Genetics in Medicine</i> , 2020, 22, 15-25.	2.4	365
20	Cumulative Burden of Colorectal Cancer-Associated Genetic Variants Is More Strongly Associated With Early-Onset vs Late-Onset Cancer. <i>Gastroenterology</i> , 2020, 158, 1274-1286.e12.	1.3	110
21	Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 432-444.	6.2	124
22	Pathways to a cancer-free future: a protocol for modelled evaluations to minimise the future burden of colorectal cancer in Australia. <i>BMJ Open</i> , 2020, 10, e036475.	1.9	1
23	Monoallelic NTHL1 Loss-of-Function Variants and Risk of Polyposis and Colorectal Cancer. <i>Gastroenterology</i> , 2020, 159, 2241-2243.e6.	1.3	20
24	Mendelian Randomization of Circulating Polyunsaturated Fatty Acids and Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 860-870.	2.5	26
25	Do the risks of Lynch syndrome-related cancers depend on the parent of origin of the mutation?. <i>Familial Cancer</i> , 2020, 19, 215-222.	1.9	1
26	A New Comprehensive Colorectal Cancer Risk Prediction Model Incorporating Family History, Personal Characteristics, and Environmental Factors. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 549-557.	2.5	25
27	Genetic Predictors of Circulating 25-Hydroxyvitamin D and Prognosis after Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1128-1134.	2.5	1
28	Potential impact of family history-based screening guidelines on the detection of early-onset colorectal cancer. <i>Cancer</i> , 2020, 126, 3013-3020.	4.1	45
29	Early-Age-of-Onset Colorectal Carcinoma: An Emerging Public Health Issue. , 2020, , 1-9.		0
30	Genetic Variants in the Regulatory T cell-Related Pathway and Colorectal Cancer Prognosis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 2719-2728.	2.5	1
31	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2019, 111, 146-157.	6.3	129
32	Type 2 diabetes mellitus, blood cholesterol, triglyceride and colorectal cancer risk in Lynch syndrome. <i>British Journal of Cancer</i> , 2019, 121, 869-876.	6.4	10
33	Ability of known susceptibility SNPs to predict colorectal cancer risk for persons with and without a family history. <i>Familial Cancer</i> , 2019, 18, 389-397.	1.9	23
34	Association analyses identify 31 new risk loci for colorectal cancer susceptibility. <i>Nature Communications</i> , 2019, 10, 2154.	12.8	172
35	Mortality after breast cancer as a function of time since diagnosis by estrogen receptor status and age at diagnosis. <i>International Journal of Cancer</i> , 2019, 145, 3207-3217.	5.1	14
36	Risks of Colorectal Cancer and Cancer-Related Mortality in Familial Colorectal Cancer Type X and Lynch Syndrome Families. <i>Journal of the National Cancer Institute</i> , 2019, 111, 675-683.	6.3	12

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37	Discovery of common and rare genetic risk variants for colorectal cancer. <i>Nature Genetics</i> , 2019, 51, 76-87.	21.4	377
38	Early-life exposure to sibling modifies the relationship between <i>CD14</i> polymorphisms and allergic sensitization. <i>Clinical and Experimental Allergy</i> , 2019, 49, 331-340.	2.9	2
39	Cohort Profile: The Colon Cancer Family Registry Cohort (CCFRC). <i>International Journal of Epidemiology</i> , 2018, 47, 387-388i.	1.9	40
40	Somatic mutations of the coding microsatellites within the beta-2-microglobulin gene in mismatch repair-deficient colorectal cancers and adenomas. <i>Familial Cancer</i> , 2018, 17, 91-100.	1.9	21
41	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.	5.1	26
42	Risk of colorectal cancer for carriers of a germ-line mutation in <i>POLE</i> or <i>POLD1</i> . <i>Genetics in Medicine</i> , 2018, 20, 890-895.	2.4	49
43	Cancer Risks for <i>PMS2</i> -Associated Lynch Syndrome. <i>Journal of Clinical Oncology</i> , 2018, 36, 2961-2968.	1.6	147
44	Genetic susceptibility markers for a breast-colorectal cancer phenotype: Exploratory results from genome-wide association studies. <i>PLoS ONE</i> , 2018, 13, e0196245.	2.5	9
45	The International Mismatch Repair Consortium. , 2018, , 479-495.		0
46	Mendelian randomisation study of age at menarche and age at menopause and the risk of colorectal cancer. <i>British Journal of Cancer</i> , 2018, 118, 1639-1647.	6.4	16
47	284 - Potential Impact of Family History Based Screening Guidelines on Early Cancer Detection Among Individuals at Risk for Young Onset Colorectal Cancer. <i>Gastroenterology</i> , 2018, 154, S-71.	1.3	0
48	Genetic and Environmental Modifiers of Cancer Risk in Lynch Syndrome. , 2018, , 67-89.		4
49	Interaction between polymorphisms in aspirin metabolic pathways, regular aspirin use and colorectal cancer risk: A case-control study in unselected white European populations. <i>PLoS ONE</i> , 2018, 13, e0192223.	2.5	5
50	Physical activity and the risk of colorectal cancer in Lynch syndrome. <i>International Journal of Cancer</i> , 2018, 143, 2250-2260.	5.1	23
51	The Colon Cancer Family Registry Cohort. , 2018, , 427-459.		3
52	Towards personalised risk assessment and clinical management: A worldwide study of age-, sex-, geographic region-, gene- and cancer-specific risks for Lynch syndrome.. <i>Journal of Clinical Oncology</i> , 2018, 36, 1526-1526.	1.6	0
53	Abstract 1238: International comparison of cancer risks for Lynch syndrome. , 2018, , .		0
54	Household Exposures, Glutathione S-Transferase Genes and Asthma Risk in Middle-Age. ISEE Conference Abstracts, 2018, 2018, .	0.0	0

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55	Tumor testing to identify lynch syndrome in two Australian colorectal cancer cohorts. <i>Journal of Gastroenterology and Hepatology (Australia)</i> , 2017, 32, 427-438.	2.8	47
56	Modeling of Successive Cancer Risks in Lynch Syndrome Families in the Presence of Competing Risks Using Copulas. <i>Biometrics</i> , 2017, 73, 271-282.	1.4	5
57	Mendelian randomisation implicates hyperlipidaemia as a risk factor for colorectal cancer. <i>International Journal of Cancer</i> , 2017, 140, 2701-2708.	5.1	76
58	The interaction between farming/rural environment and TLR2, TLR4, TLR6 and CD14 genetic polymorphisms in relation to early- and late-onset asthma. <i>Scientific Reports</i> , 2017, 7, 43681.	3.3	27
59	Prediagnostic alcohol consumption and colorectal cancer survival: The Colon Cancer Family Registry. <i>Cancer</i> , 2017, 123, 1035-1043.	4.1	21
60	Risk factors for metachronous colorectal cancer or polyp: A systematic review and meta-analysis. <i>Journal of Gastroenterology and Hepatology (Australia)</i> , 2017, 32, 301-326.	2.8	13
61	Long-term weight loss after colorectal cancer diagnosis is associated with lower survival: The Colon Cancer Family Registry. <i>Cancer</i> , 2017, 123, 4701-4708.	4.1	20
62	Pro-inflammatory fatty acid profile and colorectal cancer risk: A Mendelian randomisation analysis. <i>European Journal of Cancer</i> , 2017, 84, 228-238.	2.8	81
63	Targeted sequencing of 36 known or putative colorectal cancer susceptibility genes. <i>Molecular Genetics & Genomic Medicine</i> , 2017, 5, 553-569.	1.2	32
64	Testing for Gene-Environment Interactions Using a Prospective Family Cohort Design: Body Mass Index in Early and Later Adulthood and Risk of Breast Cancer. <i>American Journal of Epidemiology</i> , 2017, 185, 487-500.	3.4	5
65	Findings in young adults at colonoscopy from a hospital service database audit. <i>BMC Gastroenterology</i> , 2017, 17, 56.	2.0	14
66	Alcohol Consumption and the Risk of Colorectal Cancer for Mismatch Repair Gene Mutation Carriers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 366-375.	2.5	37
67	Prevalence and Penetrance of Major Genes and Polygenes for Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 404-412.	2.5	341
68	Germline miRNA DNA variants and the risk of colorectal cancer by subtype. <i>Genes Chromosomes and Cancer</i> , 2017, 56, 177-184.	2.8	7
69	Reproductive factors as risk modifiers of breast cancer in <i>BRCA</i> mutation carriers and high-risk non-carriers. <i>Oncotarget</i> , 2017, 8, 102110-102118.	1.8	23
70	Abstract PR05: Does a comprehensive family history of colorectal cancer improve risk prediction?. , 2017, , .		3
71	Abstract PR10: Development of a comprehensive colorectal cancer risk prediction tool (CRiPT) incorporating known and unknown major genes and polygenes. , 2017, , .		0
72	Abstract B04: Development of a comprehensive colorectal cancer risk prediction tool (CRiPT) incorporating known and unknown major genes and polygenes. , 2017, , .		0

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73	Abstract 4266: Double somatic mutations as a cause of tumor mismatch repair-deficiency in population-based colorectal and endometrial cancer with Lynch-like syndrome. , 2017, , .		1
74	Reducing the polyp burden in serrated polyposis by serial colonoscopy: the impact of nationally coordinated community surveillance. <i>New Zealand Medical Journal</i> , 2017, 130, 57-67.	0.5	4
75	Risk factors for metachronous colorectal cancer following a primary colorectal cancer: A prospective cohort study. <i>International Journal of Cancer</i> , 2016, 139, 1081-1090.	5.1	32
76	Mendelian randomisation analysis strongly implicates adiposity with risk of developing colorectal cancer. <i>British Journal of Cancer</i> , 2016, 115, 266-272.	6.4	57
77	Cholecystectomy and the risk of colorectal cancer by tumor mismatch repair deficiency status. <i>International Journal of Colorectal Disease</i> , 2016, 31, 1451-1457.	2.2	6
78	The Experience of Extended Bowel Resection in Individuals With a High Metachronous Colorectal Cancer Risk: A Qualitative Study. <i>Oncology Nursing Forum</i> , 2016, 43, 444-452.	1.2	5
79	Su1725 Characteristics of Advanced Neoplasia of the Large Bowel in Young Adults: Results From a Hospital Colonoscopy Service Database. <i>Gastrointestinal Endoscopy</i> , 2016, 83, AB411-AB412.	1.0	0
80	Multivitamin, calcium and folic acid supplements and the risk of colorectal cancer in Lynch syndrome. <i>International Journal of Epidemiology</i> , 2016, 45, 940-953.	1.9	27
81	Common variants in the obesity-associated genes <i>FTO</i> and <i>MC4R</i> are not associated with risk of colorectal cancer. <i>Cancer Epidemiology</i> , 2016, 44, 1-4.	1.9	12
82	Response: Table 1.. <i>Journal of the National Cancer Institute</i> , 2016, 108, djv385.	6.3	1
83	SNP rs16906252C>T Is an Expression and Methylation Quantitative Trait Locus Associated with an Increased Risk of Developing <i>MGMT</i>-Methylated Colorectal Cancer. <i>Clinical Cancer Research</i> , 2016, 22, 6266-6277.	7.0	22
84	Risk of extracolonic cancers for people with biallelic and monoallelic mutations in <i>MUTYH</i>. <i>International Journal of Cancer</i> , 2016, 139, 1557-1563.	5.1	107
85	Determining the familial risk distribution of colorectal cancer: a data mining approach. <i>Familial Cancer</i> , 2016, 15, 241-251.	1.9	6
86	Associations of 5HTTLPR polymorphism with major depressive disorder and alcohol dependence: A systematic review and meta-analysis. <i>Australian and New Zealand Journal of Psychiatry</i> , 2016, 50, 842-857.	2.3	37
87	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.	2.9	37
88	Quantifying the utility of single nucleotide polymorphisms to guide colorectal cancer screening. <i>Future Oncology</i> , 2016, 12, 503-513.	2.4	42
89	Germline mutations in <i>PMS2</i> and <i>MLH1</i> in individuals with solitary loss of PMS2 expression in colorectal carcinomas from the Colon Cancer Family Registry Cohort. <i>BMJ Open</i> , 2016, 6, e010293.	1.9	33
90	Risk Prediction Models for Colorectal Cancer: A Systematic Review. <i>Cancer Prevention Research</i> , 2016, 9, 13-26.	1.5	142

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91	GWASeq: targeted re-sequencing follow up to GWAS. BMC Genomics, 2016, 17, 176.	2.8	7
92	Comparison of Prediction Models for Lynch Syndrome Among Individuals With Colorectal Cancer. Journal of the National Cancer Institute, 2016, 108, .	6.3	29
93	PMS2 monoallelic mutation carriers: the known unknown. Genetics in Medicine, 2016, 18, 13-19.	2.4	51
94	Abstract 3425: Prediagnostic alcohol consumption and colorectal cancer survival: the Colon Cancer Family Registry. , 2016, , .		1
95	Abstract 2552: Influence of age, sex, colorectal cancer status, and mutation carrier status on physical activity in families with Lynch syndrome. , 2016, , .		0
96	2079 Risk factors for metachronous colorectal cancer following a primary colorectal cancer: A prospective cohort study. European Journal of Cancer, 2015, 51, S355.	2.8	0
97	1054 Risk of extracolonic cancers for people with biallelic and monoallelic mutations in MUTYH. European Journal of Cancer, 2015, 51, S163.	2.8	0
98	Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. Scientific Reports, 2015, 5, 17369.	3.3	35
99	Incidence and prevalence of non-melanoma skin cancer in Australia: A systematic review. Australasian Journal of Dermatology, 2015, 56, 258-267.	0.7	164
100	Fertility after young-onset colorectal cancer: a study of subjects with Lynch syndrome. Colorectal Disease, 2015, 17, 787-793.	1.4	17
101	Lynch syndrome and cervical cancer. International Journal of Cancer, 2015, 137, 2757-2761.	5.1	13
102	Family History and Risk of Endometrial Cancer. Obstetrics and Gynecology, 2015, 125, 89-98.	2.4	72
103	Mendelian randomization study of height and risk of colorectal cancer. International Journal of Epidemiology, 2015, 44, 662-672.	1.9	55
104	Mendelian Randomization Study of Body Mass Index and Colorectal Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1024-1031.	2.5	67
105	Prediagnostic Physical Activity and Colorectal Cancer Survival: Overall and Stratified by Tumor Characteristics. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1130-1137.	2.5	30
106	PIK3CA Somatic Mutation Status in Relation to Patient and Tumor Factors in Racial/Ethnic Minorities with Colorectal Cancer. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1046-1051.	2.5	17
107	Association between Body Mass Index and Mortality for Colorectal Cancer Survivors: Overall and by Tumor Molecular Phenotype. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1229-1238.	2.5	44
108	Risk of colorectal cancer for people with a mutation in both a MUTYH and a DNA mismatch repair gene. Familial Cancer, 2015, 14, 575-583.	1.9	11

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109	Mutation Spectrum and Risk of Colorectal Cancer in African American Families with Lynch Syndrome. <i>Gastroenterology</i> , 2015, 149, 1446-1453.	1.3	46
110	Aspirin, Ibuprofen, and the Risk of Colorectal Cancer in Lynch Syndrome. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv170.	6.3	80
111	A new GWAS and meta-analysis with 1000Genomes imputation identifies novel risk variants for colorectal cancer. <i>Scientific Reports</i> , 2015, 5, 10442.	3.3	109
112	Female Hormonal Factors and the Risk of Endometrial Cancer in Lynch Syndrome. <i>JAMA - Journal of the American Medical Association</i> , 2015, 314, 61.	7.4	68
113	Childhood cancers in families with and without Lynch syndrome. <i>Familial Cancer</i> , 2015, 14, 545-551.	1.9	8
114	Rising incidence of early-onset colorectal cancer in Australia over two decades: Report and review. <i>Journal of Gastroenterology and Hepatology (Australia)</i> , 2015, 30, 6-13.	2.8	119
115	Breast cancer risk for Korean women with germline mutations in BRCA1 and BRCA2. <i>Breast Cancer Research and Treatment</i> , 2015, 152, 659-665.	2.5	18
116	Short-Term Risk of Colorectal Cancer in Individuals With Lynch Syndrome: A Meta-Analysis. <i>Journal of Clinical Oncology</i> , 2015, 33, 326-331.	1.6	37
117	Role of tumour molecular and pathology features to estimate colorectal cancer risk for first-degree relatives. <i>Gut</i> , 2015, 64, 101-110.	12.1	40
118	Genetic variants within the hTERT gene and the risk of colorectal cancer in Lynch syndrome. <i>Genes and Cancer</i> , 2015, 6, 445-451.	1.9	6
119	Exposure to siblings in early life modifies the association between CD14 polymorphisms and allergic sensitization in adult life. , 2015, . .		0
120	Clinical problems of colorectal cancer and endometrial cancer cases with unknown cause of tumor mismatch repair deficiency (suspected Lynch syndrome). <i>The Application of Clinical Genetics</i> , 2014, 7, 183.	3.0	68
121	Risk of Prostate Cancer in Lynch Syndrome: A Systematic Review and Meta-analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 437-449.	2.5	104
122	High prevalence of mismatch repair deficiency in prostate cancers diagnosed in mismatch repair gene mutation carriers from the colon cancer family registry. <i>Familial Cancer</i> , 2014, 13, 573-582.	1.9	44
123	Re: Microsatellite Instability and BRAF Mutation Testing in Colorectal Cancer Prognostication. <i>Journal of the National Cancer Institute</i> , 2014, 106, dju180-dju180.	6.3	6
124	Family History of Colorectal Cancer Is Not Associated with Colorectal Cancer Survival Regardless of Microsatellite Instability Status. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1700-1704.	2.5	9
125	Genetic variation in UGT genes modify the associations of NSAIDs with risk of colorectal cancer: Colon cancer family registry. <i>Genes Chromosomes and Cancer</i> , 2014, 53, 568-578.	2.8	25
126	Perceived Versus Predicted Risks of Colorectal Cancer and Self-Reported Colonoscopies by Members of Mismatch Repair Gene Mutation-Carrying Families Who Have Declined Genetic Testing. <i>Journal of Genetic Counseling</i> , 2014, 23, 79-88.	1.6	8

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127	Risk of Colorectal Cancer for Carriers of Mutations in MUTYH, With and Without a Family History of Cancer. <i>Gastroenterology</i> , 2014, 146, 1208-1211.e5.	1.3	180
128	Should the grading of colorectal adenocarcinoma include microsatellite instability status?. <i>Human Pathology</i> , 2014, 45, 2077-2084.	2.0	44
129	Characterisation of Familial Colorectal Cancer Type X, Lynch syndrome, and non-familial colorectal cancer. <i>British Journal of Cancer</i> , 2014, 111, 598-602.	6.4	38
130	Identification of susceptibility loci for colorectal cancer in a genome-wide meta-analysis. <i>Human Molecular Genetics</i> , 2014, 23, 4729-4737.	2.9	128
131	Fertility and apparent genetic anticipation in Lynch syndrome. <i>Familial Cancer</i> , 2014, 13, 369-374.	1.9	3
132	Colorectal cancer and self-reported tooth agenesis. <i>Hereditary Cancer in Clinical Practice</i> , 2014, 12, 7.	1.5	16
133	Does risk of endometrial cancer for women without a germline mutation in a DNA mismatch repair gene depend on family history of endometrial cancer or colorectal cancer?. <i>Gynecologic Oncology</i> , 2014, 133, 287-292.	1.4	20
134	Abstract LB-276: Prospective study of body mass index and adult weight change with colorectal cancer survival, overall and by tumor microsatellite instability status. , 2014, , .		0
135	Risk of breast cancer in Lynch syndrome: a systematic review. <i>Breast Cancer Research</i> , 2013, 15, R27.	5.0	109
136	Are the common genetic variants associated with colorectal cancer risk for DNA mismatch repair gene mutation carriers?. <i>European Journal of Cancer</i> , 2013, 49, 1578-1587.	2.8	31
137	Expression of MUC2, MUC5AC, MUC5B, and MUC6 mucins in colorectal cancers and their association with the CpG island methylator phenotype. <i>Modern Pathology</i> , 2013, 26, 1642-1656.	5.5	127
138	Differential effects of phosphate binders on pre-dialysis serum bicarbonate in end-stage kidney disease patients on maintenance haemodialysis. <i>BMC Nephrology</i> , 2013, 14, 205.	1.8	7
139	Risks of Colorectal and Other Cancers After Endometrial Cancer for Women With Lynch Syndrome. <i>Journal of the National Cancer Institute</i> , 2013, 105, 274-279.	6.3	93
140	Cancer Risks for <i>MLH1</i> and <i>MSH2</i> Mutation Carriers. <i>Human Mutation</i> , 2013, 34, 490-497.	2.5	201
141	KRAS-mutation status in relation to colorectal cancer survival: the joint impact of correlated tumour markers. <i>British Journal of Cancer</i> , 2013, 108, 1757-1764.	6.4	191
142	Risk of Metachronous Colon Cancer Following Surgery for Rectal Cancer in Mismatch Repair Gene Mutation Carriers. <i>Annals of Surgical Oncology</i> , 2013, 20, 1829-1836.	1.5	103
143	Colorectal carcinomas with KRAS mutation are associated with distinctive morphological and molecular features. <i>Modern Pathology</i> , 2013, 26, 825-834.	5.5	126
144	Association between hypermethylation of DNA repetitive elements in white blood cell DNA and early-onset colorectal cancer. <i>Epigenetics</i> , 2013, 8, 748-755.	2.7	41

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145	Germline HOXB13 p.Gly84Glu mutation and risk of colorectal cancer. <i>Cancer Epidemiology</i> , 2013, 37, 424-427.	1.9	24
146	Family History of Colorectal Cancer in <i>BRAF</i> p.V600E-Mutated Colorectal Cancer Cases. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 917-926.	2.5	24
147	Colon and Rectal Cancer Survival by Tumor Location and Microsatellite Instability. <i>Diseases of the Colon and Rectum</i> , 2013, 56, 937-944.	1.3	81
148	<i>BRAF</i> V600E Immunohistochemistry Facilitates Universal Screening of Colorectal Cancers for Lynch Syndrome. <i>American Journal of Surgical Pathology</i> , 2013, 37, 1592-1602.	3.7	125
149	Response. <i>Journal of the National Cancer Institute</i> , 2013, 105, 1837-1838.	6.3	1
150	Is the reported modifying effect of 8q23.3 and 11q23.1 on colorectal cancer risk for <i>MLH1</i> mutation carriers valid?. <i>International Journal of Cancer</i> , 2013, 133, 1762-1763.	5.1	2
151	Criteria and prediction models for mismatch repair gene mutations: a review. <i>Journal of Medical Genetics</i> , 2013, 50, 785-793.	3.2	27
152	Multiplicity and Molecular Heterogeneity of Colorectal Carcinomas in Individuals With Serrated Polyposis. <i>American Journal of Surgical Pathology</i> , 2013, 37, 434-442.	3.7	39
153	<i>PIK3CA</i> Activating Mutation in Colorectal Carcinoma: Associations with Molecular Features and Survival. <i>PLoS ONE</i> , 2013, 8, e65479.	2.5	117
154	Germline Mutations in the Polyposis-Associated Genes <i>BMPRI1A</i> , <i>SMAD4</i> , <i>PTEN</i> , <i>MUTYH</i> and <i>GREM1</i> Are Not Common in Individuals with Serrated Polyposis Syndrome. <i>PLoS ONE</i> , 2013, 8, e66705.	2.5	27
155	Abstract 4831: Additive and multiplicative gene-environment interactions for colorectal cancer risk.. , 2013, , .		0
156	Abstract 4842: Hazard ratio for colorectal cancer risk in Lynch syndrome is inversely associated with age.. , 2013, , .		0
157	Is prostate cancer a Lynch syndrome cancer?. <i>Asian Journal of Andrology</i> , 2013, 15, 588-589.	1.6	2
158	Kaplan-Meier failure estimate for metachronous colorectal cancer risk is clinically relevant. <i>Gut</i> , 2012, 61, 783.2-784.	12.1	0
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