

# 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	Discovery of common and rare genetic risk variants for colorectal cancer. <i>Nature Genetics</i> , 2019, 51, 76-87.	21.4	377
2	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
3	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
4	Colorectal and Other Cancer Risks for Carriers and Noncarriers From Families With a DNA Mismatch Repair Gene Mutation: A Prospective Cohort Study. <i>Journal of Clinical Oncology</i> , 2012, 30, 958-964.	1.6	286
5	Metachronous colorectal cancer risk for mismatch repair gene mutation carriers: the advantage of more extensive colon surgery. <i>Gut</i> , 2011, 60, 950-957.	12.1	227
6	Cancer Risks for <i>MLH1</i> and <i>MSH2</i> Mutation Carriers. <i>Human Mutation</i> , 2013, 34, 490-497.	2.5	201
7	Risks of Primary Extracolonic Cancers Following Colorectal Cancer in Lynch Syndrome. <i>Journal of the National Cancer Institute</i> , 2012, 104, 1363-1372.	6.3	193
8	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
9	Risk of Colorectal Cancer for Carriers of Mutations in <i>MUTYH</i> , With and Without a Family History of Cancer. <i>Gastroenterology</i> , 2014, 146, 1208-1211.e5.	1.3	180
10	Association analyses identify 31 new risk loci for colorectal cancer susceptibility. <i>Nature Communications</i> , 2019, 10, 2154.	12.8	172
11	Incidence and prevalence of non-melanoma skin cancer in Australia: A systematic review. <i>Australasian Journal of Dermatology</i> , 2015, 56, 258-267.	0.7	164
12	Cancer Risks for <i>PMS2</i> -Associated Lynch Syndrome. <i>Journal of Clinical Oncology</i> , 2018, 36, 2961-2968.	1.6	147
13	Risk Prediction Models for Colorectal Cancer: A Systematic Review. <i>Cancer Prevention Research</i> , 2016, 9, 13-26.	1.5	142
14	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2019, 111, 146-157.	6.3	129
15	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
16	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
17	Colorectal carcinomas with KRAS mutation are associated with distinctive morphological and molecular features. <i>Modern Pathology</i> , 2013, 26, 825-834.	5.5	126
18	BRAFV600E Immunohistochemistry Facilitates Universal Screening of Colorectal Cancers for Lynch Syndrome. <i>American Journal of Surgical Pathology</i> , 2013, 37, 1592-1602.	3.7	125

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19	Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. American Journal of Human Genetics, 2020, 107, 432-444.	6.2	124
20	Rising incidence of early-onset colorectal cancer in Australia over two decades: Report and review. Journal of Gastroenterology and Hepatology (Australia), 2015, 30, 6-13.	2.8	119
21	PIK3CA Activating Mutation in Colorectal Carcinoma: Associations with Molecular Features and Survival. PLoS ONE, 2013, 8, e65479.	2.5	117
22	<i>BRAF</i> Mutation Status and Survival after Colorectal Cancer Diagnosis According to Patient and Tumor Characteristics. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 1792-1798.	2.5	113
23	Cumulative Burden of Colorectal Cancer-Associated Genetic Variants Is More Strongly Associated With Early-Onset vs Late-Onset Cancer. Gastroenterology, 2020, 158, 1274-1286.e12.	1.3	110
24	Risk of breast cancer in Lynch syndrome: a systematic review. Breast Cancer Research, 2013, 15, R27.	5.0	109
25	A new GWAS and meta-analysis with 1000Genomes imputation identifies novel risk variants for colorectal cancer. Scientific Reports, 2015, 5, 10442.	3.3	109
26	A large-scale meta-analysis to refine colorectal cancer risk estimates associated with MUTYH variants. British Journal of Cancer, 2010, 103, 1875-1884.	6.4	107
27	Risk of extracolonic cancers for people with biallelic and monoallelic mutations in <i>MUTYH</i> . International Journal of Cancer, 2016, 139, 1557-1563.	5.1	107
28	Risk of Prostate Cancer in Lynch Syndrome: A Systematic Review and Meta-analysis. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 437-449.	2.5	104
29	Risk of Metachronous Colon Cancer Following Surgery for Rectal Cancer in Mismatch Repair Gene Mutation Carriers. Annals of Surgical Oncology, 2013, 20, 1829-1836.	1.5	103
30	Cancer risks for monoallelic <i>MUTYH</i> mutation carriers with a family history of colorectal cancer. International Journal of Cancer, 2011, 129, 2256-2262.	5.1	93
31	Risks of Colorectal and Other Cancers After Endometrial Cancer for Women With Lynch Syndrome. Journal of the National Cancer Institute, 2013, 105, 274-279.	6.3	93
32	Phenotype and Polyp Landscape in Serrated Polyposis Syndrome. American Journal of Surgical Pathology, 2012, 36, 876-882.	3.7	85
33	Risk Prediction Models for Colorectal Cancer: A Review. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 398-410.	2.5	84
34	Colon and Rectal Cancer Survival by Tumor Location and Microsatellite Instability. Diseases of the Colon and Rectum, 2013, 56, 937-944.	1.3	81
35	Pro-inflammatory fatty acid profile and colorectal cancer risk: A Mendelian randomisation analysis. European Journal of Cancer, 2017, 84, 228-238.	2.8	81
36	Cancer Risks for Relatives of Patients With Serrated Polyposis. American Journal of Gastroenterology, 2012, 107, 770-778.	0.4	80

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37	Increased Cancer Risks in Myotonic Dystrophy. Mayo Clinic Proceedings, 2012, 87, 130-135.	3.0	80
38	Aspirin, Ibuprofen, and the Risk of Colorectal Cancer in Lynch Syndrome. Journal of the National Cancer Institute, 2015, 107, djv170.	6.3	80
39	Mendelian randomisation implicates hyperlipidaemia as a risk factor for colorectal cancer. International Journal of Cancer, 2017, 140, 2701-2708.	5.1	76
40	Immunohistochemical testing of conventional adenomas for loss of expression of mismatch repair proteins in Lynch syndrome mutation carriers: a case series from the Australasian site of the colon cancer family registry. Modern Pathology, 2012, 25, 722-730.	5.5	73
41	Family History and Risk of Endometrial Cancer. Obstetrics and Gynecology, 2015, 125, 89-98.	2.4	72
42	Risk Factors for Colorectal Cancer in Patients with Multiple Serrated Polyps: A Cross-Sectional Case Series from Genetics Clinics. PLoS ONE, 2010, 5, e11636.	2.5	68
43	Clinical problems of colorectal cancer and endometrial cancer cases with unknown cause of tumor mismatch repair deficiency (suspected Lynch syndrome). The Application of Clinical Genetics, 2014, 7, 183.	3.0	68
44	Female Hormonal Factors and the Risk of Endometrial Cancer in Lynch Syndrome. JAMA - Journal of the American Medical Association, 2015, 314, 61.	7.4	68
45	Mendelian Randomization Study of Body Mass Index and Colorectal Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1024-1031.	2.5	67
46	Association between monoallelic MUTYH mutation and colorectal cancer risk: a meta-regression analysis. Familial Cancer, 2011, 10, 1-9.	1.9	59
47	Variation in the risk of colorectal cancer in families with Lynch syndrome: a retrospective cohort study. Lancet Oncology, The, 2021, 22, 1014-1022.	10.7	58
48	Mendelian randomisation analysis strongly implicates adiposity with risk of developing colorectal cancer. British Journal of Cancer, 2016, 115, 266-272.	6.4	57
49	Mendelian randomization study of height and risk of colorectal cancer. International Journal of Epidemiology, 2015, 44, 662-672.	1.9	55
50	PMS2 monoallelic mutation carriers: the known unknown. Genetics in Medicine, 2016, 18, 13-19.	2.4	51
51	Body mass index in early adulthood and colorectal cancer risk for carriers and non-carriers of germline mutations in DNA mismatch repair genes. British Journal of Cancer, 2011, 105, 162-169.	6.4	50
52	Risk of colorectal cancer for carriers of a germ-line mutation in POLE or POLD1. Genetics in Medicine, 2018, 20, 890-895.	2.4	49
53	Phenotypic diversity in patients with multiple serrated polyps: a genetics clinic study. International Journal of Colorectal Disease, 2010, 25, 703-712.	2.2	48
54	Tumor testing to identify lynch syndrome in two Australian colorectal cancer cohorts. Journal of Gastroenterology and Hepatology (Australia), 2017, 32, 427-438.	2.8	47

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55	Mutation Spectrum and Risk of Colorectal Cancer in African American Families with Lynch Syndrome. <i>Gastroenterology</i> , 2015, 149, 1446-1453.	1.3	46
56	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
57	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
58	Should the grading of colorectal adenocarcinoma include microsatellite instability status?. <i>Human Pathology</i> , 2014, 45, 2077-2084.	2.0	44
59	Association between Body Mass Index and Mortality for Colorectal Cancer Survivors: Overall and by Tumor Molecular Phenotype. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1229-1238.	2.5	44
60	Quantifying the utility of single nucleotide polymorphisms to guide colorectal cancer screening. <i>Future Oncology</i> , 2016, 12, 503-513.	2.4	42
61	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
62	Determining the frequency of de novo germline mutations in DNA mismatch repair genes. <i>Journal of Medical Genetics</i> , 2011, 48, 530-534.	3.2	40
63	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
64	Cohort Profile: The Colon Cancer Family Registry Cohort (CCFRC). <i>International Journal of Epidemiology</i> , 2018, 47, 387-388i.	1.9	40
65	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
66	Nongenetic Determinants of Risk for Early-Onset Colorectal Cancer. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab029.	2.9	39
67	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
68	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
69	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
70	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
71	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
72	DNA methylation-based signature of CD8+ tumor-infiltrating lymphocytes enables evaluation of immune response and prognosis in colorectal cancer. , 2021, 9, e002671.		37

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73	Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. <i>Scientific Reports</i> , 2015, 5, 17369.	3.3	35
74	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
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	Targeted sequencing of 36 known or putative colorectal cancer susceptibility genes. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2017, 5, 553-569.	1.2	32
77	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
78	Prediagnostic Physical Activity and Colorectal Cancer Survival: Overall and Stratified by Tumor Characteristics. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1130-1137.	2.5	30
79	Comparison of Prediction Models for Lynch Syndrome Among Individuals With Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2016, 108, .	6.3	29
80	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
81	Criteria and prediction models for mismatch repair gene mutations: a review. <i>Journal of Medical Genetics</i> , 2013, 50, 785-793.	3.2	27
82	Germline Mutations in the Polyposis-Associated Genes BMPR1A, SMAD4, PTEN, MUTYH and GREM1 Are Not Common in Individuals with Serrated Polyposis Syndrome. <i>PLoS ONE</i> , 2013, 8, e66705.	2.5	27
83	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
84	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
85	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
86	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
87	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
88	Genetic variation in <i>UGT</i> 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
89	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
90	Germline HOXB13 p.Gly84Glu mutation and risk of colorectal cancer. <i>Cancer Epidemiology</i> , 2013, 37, 424-427.	1.9	24

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91	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
92	Body Mass Index in Early Adulthood and Endometrial Cancer Risk for Mismatch Repair Gene Mutation Carriers. <i>Obstetrics and Gynecology</i> , 2011, 117, 899-905.	2.4	23
93	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
94	Physical activity and the risk of colorectal cancer in Lynch syndrome. <i>International Journal of Cancer</i> , 2018, 143, 2250-2260.	5.1	23
95	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
96	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
97	Prediagnostic alcohol consumption and colorectal cancer survival: The Colon Cancer Family Registry. <i>Cancer</i> , 2017, 123, 1035-1043.	4.1	21
98	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
99	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
100	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
101	Monoallelic <i>NTHL1</i> Loss-of-Function Variants and Risk of Polyposis and Colorectal Cancer. <i>Gastroenterology</i> , 2020, 159, 2241-2243.e6.	1.3	20
102	Breast cancer risk for Korean women with germline mutations in <i>BRCA1</i> and <i>BRCA2</i> . <i>Breast Cancer Research and Treatment</i> , 2015, 152, 659-665.	2.5	18
103	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
104	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
105	Fertility after young-onset colorectal cancer: a study of subjects with Lynch syndrome. <i>Colorectal Disease</i> , 2015, 17, 787-793.	1.4	17
106	<i>PIK3CA</i> Somatic Mutation Status in Relation to Patient and Tumor Factors in Racial/Ethnic Minorities with Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1046-1051.	2.5	17
107	Colorectal cancer and self-reported tooth agenesis. <i>Hereditary Cancer in Clinical Practice</i> , 2014, 12, 7.	1.5	16
108	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

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109	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
110	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
111	Findings in young adults at colonoscopy from a hospital service database audit. <i>BMC Gastroenterology</i> , 2017, 17, 56.	2.0	14
112	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
113	The Australasian Colorectal Cancer Family Registry. <i>Medical Journal of Australia</i> , 2012, 197, 480-481.	1.7	13
114	Lynch syndrome and cervical cancer. <i>International Journal of Cancer</i> , 2015, 137, 2757-2761.	5.1	13
115	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
116	Cancer Risks for the Relatives of Colorectal Cancer Cases with a Methylated <i>MLH1</i> Promoter Region: Data from the Colorectal Cancer Family Registry. <i>Cancer Prevention Research</i> , 2012, 5, 328-335.	1.5	12
117	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
118	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
119	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
120	Risk of colorectal cancer for people with a mutation in both a <i>MUTYH</i> and a DNA mismatch repair gene. <i>Familial Cancer</i> , 2015, 14, 575-583.	1.9	11
121	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
122	No Difference in Penetrance between Truncating and Missense/Aberrant Splicing Pathogenic Variants in <i>MLH1</i> and <i>MSH2</i> : A Prospective Lynch Syndrome Database Study. <i>Journal of Clinical Medicine</i> , 2021, 10, 2856.	2.4	11
123	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
124	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
125	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
126	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

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127	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
128	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
129	Childhood cancers in families with and without Lynch syndrome. <i>Familial Cancer</i> , 2015, 14, 545-551.	1.9	8
130	Using tumour pathology to identify people at high genetic risk of breast and colorectal cancers. <i>Pathology</i> , 2012, 44, 89-98.	0.6	7
131	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
132	GWASeq: targeted re-sequencing follow up to GWAS. <i>BMC Genomics</i> , 2016, 17, 176.	2.8	7
133	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
134	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
135	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
136	Determining the familial risk distribution of colorectal cancer: a data mining approach. <i>Familial Cancer</i> , 2016, 15, 241-251.	1.9	6
137	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
138	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
139	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
140	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
141	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
142	Response to Li and Hopper. <i>American Journal of Human Genetics</i> , 2021, 108, 527-529.	6.2	5
143	Genetic and Environmental Modifiers of Cancer Risk in Lynch Syndrome. , 2018, , 67-89.		4
144	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

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145	Estimates of familial risks from family data are biased when ascertainment of families is not independent of family history. <i>Gut</i> , 2011, 60, 1162-1163.	12.1	3
146	Fertility and apparent genetic anticipation in Lynch syndrome. <i>Familial Cancer</i> , 2014, 13, 369-374.	1.9	3
147	The Colon Cancer Family Registry Cohort. , 2018, , 427-459.		3
148	Abstract PR05: Does a comprehensive family history of colorectal cancer improve risk prediction?. , 2017, , .		3
149	Metachronous colorectal cancer risk for mismatch repair gene mutation carriers – the advantage of more extensive surgery. <i>Hereditary Cancer in Clinical Practice</i> , 2011, 9, O1.	1.5	2
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	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
152	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
153	Colorectal cancer cases with de novo germ-line mutations in <i>MLH1</i> , <i>MSH2</i> , and <i>MSH6</i> from the Colon Cancer Family Registry.. <i>Journal of Clinical Oncology</i> , 2011, 29, 3538-3538.	1.6	2
154	Is prostate cancer a Lynch syndrome cancer?. <i>Asian Journal of Andrology</i> , 2013, 15, 588-589.	1.6	2
155	Response. <i>Journal of the National Cancer Institute</i> , 2013, 105, 1837-1838.	6.3	1
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