

Tabitha A Harrison

List of Publications by Year in descending order

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Version: 2024-02-01

84
papers

6,655
citations

156536

32
h-index

87275

74
g-index

92
all docs

92
docs citations

92
times ranked

16602
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular and Pathology Features of Colorectal Tumors and Patient Outcomes Are Associated with <i>Fusobacterium nucleatum</i> and Its Subspecies <i>animalis</i> . <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 210-220.	1.1	19
2	Genome-wide association study identifies tumor anatomical site-specific risk variants for colorectal cancer survival. <i>Scientific Reports</i> , 2022, 12, 127.	1.6	6
3	Genetic variants associated with circulating C-reactive protein levels and colorectal cancer survival: Sex-specific and lifestyle factors specific associations. <i>International Journal of Cancer</i> , 2022, 150, 1447-1454.	2.3	2
4	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, , .	3.0	15
5	Diabetes mellitus in relation to colorectal tumor molecular subtypes – a pooled analysis of more than 9,000 cases. <i>International Journal of Cancer</i> , 2022, , .	2.3	2
6	Beyond GWAS of Colorectal Cancer: Evidence of Interaction with Alcohol Consumption and Putative Causal Variant for the 10q24.2 Region. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 1077-1089.	1.1	6
7	Genetic Regulation of DNA Methylation Yields Novel Discoveries in GWAS of Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 1068-1076.	1.1	1
8	OUP accepted manuscript. <i>Journal of the National Cancer Institute</i> , 2022, , .	3.0	0
9	Association between germline variants and somatic mutations in colorectal cancer. <i>Scientific Reports</i> , 2022, 12, .	1.6	1
10	Identifying colorectal cancer caused by biallelic MUTYH pathogenic variants using tumor mutational signatures. <i>Nature Communications</i> , 2022, 13, .	5.8	15
11	A Statistical Method for Association Analysis of Cell Type Compositions. <i>Statistics in Biosciences</i> , 2021, 13, 373-385.	0.6	0
12	Association of Body Mass Index With Colorectal Cancer Risk by Genome-Wide Variants. <i>Journal of the National Cancer Institute</i> , 2021, 113, 38-47.	3.0	14
13	Identifying Novel Susceptibility Genes for Colorectal Cancer Risk From a Transcriptome-Wide Association Study of 125,478 Subjects. <i>Gastroenterology</i> , 2021, 160, 1164-1178.e6.	0.6	36
14	Genetic Predictors of Severe Skin Toxicity in Patients with Stage III Colon Cancer Treated with Cetuximab: NCCTG N0147 (Alliance). <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 404-411.	1.1	1
15	Genetic architectures of proximal and distal colorectal cancer are partly distinct. <i>Gut</i> , 2021, 70, 1325-1334.	6.1	44
16	Association of <i>Fusobacterium nucleatum</i> with Specific T-cell Subsets in the Colorectal Carcinoma Microenvironment. <i>Clinical Cancer Research</i> , 2021, 27, 2816-2826.	3.2	36
17	Response to Li and Hopper. <i>American Journal of Human Genetics</i> , 2021, 108, 527-529.	2.6	5
18	Nongenetic Determinants of Risk for Early-Onset Colorectal Cancer. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab029.	1.4	39

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19	Genetically Predicted Circulating C-Reactive Protein Concentration and Colorectal Cancer Survival: A Mendelian Randomization Consortium Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 1349-1358.	1.1	6
20	Association between Smoking and Molecular Subtypes of Colorectal Cancer. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab056.	1.4	8
21	A Combined Proteomics and Mendelian Randomization Approach to Investigate the Effects of Aspirin-Targeted Proteins on Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 564-575.	1.1	10
22	Salicylic Acid and Risk of Colorectal Cancer: A Two-Sample Mendelian Randomization Study. <i>Nutrients</i> , 2021, 13, 4164.	1.7	3
23	DNA repair and cancer in colon and rectum: Novel players in genetic susceptibility. <i>International Journal of Cancer</i> , 2020, 146, 363-372.	2.3	40
24	Meta-analysis of 16 studies of the association of alcohol with colorectal cancer. <i>International Journal of Cancer</i> , 2020, 146, 861-873.	2.3	89
25	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.	0.6	110
26	Circulating Levels of Insulin-like Growth Factor 1 and Insulin-like Growth Factor Binding Protein 3 Associate With Risk of Colorectal Cancer Based on Serologic and Mendelian Randomization Analyses. <i>Gastroenterology</i> , 2020, 158, 1300-1312.e20.	0.6	90
27	Identification of Novel Loci and New Risk Variant in Known Loci for Colorectal Cancer Risk in East Asians. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 477-486.	1.1	25
28	Genetic Variant Associated With Survival of Patients With Stage II-III Colon Cancer. <i>Clinical Gastroenterology and Hepatology</i> , 2020, 18, 2717-2723.e3.	2.4	7
29	Postmenopausal Hormone Therapy and Colorectal Cancer Risk by Molecularly Defined Subtypes and Tumor Location. <i>JNCI Cancer Spectrum</i> , 2020, 4, pkaa042.	1.4	8
30	Landscape of somatic single nucleotide variants and indels in colorectal cancer and impact on survival. <i>Nature Communications</i> , 2020, 11, 3644.	5.8	55
31	Exploratory Genome-Wide Interaction Analysis of Nonsteroidal Anti-inflammatory Drugs and Predicted Gene Expression on Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1800-1808.	1.1	1
32	Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 432-444.	2.6	124
33	A general framework for functionally informed set-based analysis: Application to a large-scale colorectal cancer study. <i>PLoS Genetics</i> , 2020, 16, e1008947.	1.5	6
34	Intake of Dietary Fruit, Vegetables, and Fiber and Risk of Colorectal Cancer According to Molecular Subtypes: A Pooled Analysis of 9 Studies. <i>Cancer Research</i> , 2020, 80, 4578-4590.	0.4	26
35	Adiposity, metabolites, and colorectal cancer risk: Mendelian randomization study. <i>BMC Medicine</i> , 2020, 18, 396.	2.3	76
36	Hemochromatosis risk genotype is not associated with colorectal cancer or age at its diagnosis. <i>Human Genetics and Genomics Advances</i> , 2020, 1, 100010.	1.0	3

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37	Incorporating Participant and Clinical Feedback into a Community-Based Participatory Research Study of Colorectal Cancer Among Alaska Native People. <i>Journal of Community Health</i> , 2020, 45, 803-811.	1.9	2
38	Telomere Maintenance Variants and Survival after Colorectal Cancer: Smoking- and Sex-Specific Associations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1817-1824.	1.1	5
39	Association Between Molecular Subtypes of Colorectal Tumors and Patient Survival, Based on Pooled Analysis of 7 International Studies. <i>Gastroenterology</i> , 2020, 158, 2158-2168.e4.	0.6	34
40	Physical activity and risks of breast and colorectal cancer: a Mendelian randomisation analysis. <i>Nature Communications</i> , 2020, 11, 597.	5.8	193
41	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2019, 111, 146-157.	3.0	129
42	Genetic variant predictors of gene expression provide new insight into risk of colorectal cancer. <i>Human Genetics</i> , 2019, 138, 307-326.	1.8	44
43	Combined effect of modifiable and non-modifiable risk factors for colorectal cancer risk in a pooled analysis of 11 population-based studies. <i>BMJ Open Gastroenterology</i> , 2019, 6, e000339.	1.1	28
44	Discovery of common and rare genetic risk variants for colorectal cancer. <i>Nature Genetics</i> , 2019, 51, 76-87.	9.4	377
45	Determining Risk of Colorectal Cancer and Starting Age of Screening Based on Lifestyle, Environmental, and Genetic Factors. <i>Gastroenterology</i> , 2018, 154, 2152-2164.e19.	0.6	226
46	A Mixed-Effects Model for Powerful Association Tests in Integrative Functional Genomics. <i>American Journal of Human Genetics</i> , 2018, 102, 904-919.	2.6	30
47	Association of family history and survival in patients with colorectal cancer: a pooled analysis of eight epidemiologic studies. <i>Cancer Medicine</i> , 2018, 7, 2192-2199.	1.3	9
48	Genetic susceptibility markers for a breast-colorectal cancer phenotype: Exploratory results from genome-wide association studies. <i>PLoS ONE</i> , 2018, 13, e0196245.	1.1	9
49	Inherited variation in circadian rhythm genes and risks of prostate cancer and three other cancer sites in combined cancer consortia. <i>International Journal of Cancer</i> , 2017, 141, 1794-1802.	2.3	28
50	Enrichment of colorectal cancer associations in functional regions: Insight for using epigenomics data in the analysis of whole genome sequence-imputed GWAS data. <i>PLoS ONE</i> , 2017, 12, e0186518.	1.1	8
51	Fine-Mapping of Common Genetic Variants Associated with Colorectal Tumor Risk Identified Potential Functional Variants. <i>PLoS ONE</i> , 2016, 11, e0157521.	1.1	8
52	Relationship of prediagnostic body mass index with survival after colorectal cancer: Stage-specific associations. <i>International Journal of Cancer</i> , 2016, 139, 1065-1072.	2.3	26
53	A reference panel of 64,976 haplotypes for genotype imputation. <i>Nature Genetics</i> , 2016, 48, 1279-1283.	9.4	2,421
54	CYP24A1 variant modifies the association between use of oestrogen plus progestogen therapy and colorectal cancer risk. <i>British Journal of Cancer</i> , 2016, 114, 221-229.	2.9	18

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55	Identification of Susceptibility Loci and Genes for Colorectal Cancer Risk. <i>Gastroenterology</i> , 2016, 150, 1633-1645.	0.6	97
56	Common genetic variation and survival after colorectal cancer diagnosis: a genome-wide analysis. <i>Carcinogenesis</i> , 2016, 37, 87-95.	1.3	62
57	Genome-Wide Interaction Analyses between Genetic Variants and Alcohol Consumption and Smoking for Risk of Colorectal Cancer. <i>PLoS Genetics</i> , 2016, 12, e1006296.	1.5	38
58	Mendelian randomization study of height and risk of colorectal cancer. <i>International Journal of Epidemiology</i> , 2015, 44, 662-672.	0.9	55
59	Mendelian Randomization Study of Body Mass Index and Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1024-1031.	1.1	67
60	A Model to Determine Colorectal Cancer Risk Using Common Genetic Susceptibility Loci. <i>Gastroenterology</i> , 2015, 148, 1330-1339.e14.	0.6	129
61	Red Meat Intake, NAT2, and Risk of Colorectal Cancer: A Pooled Analysis of 11 Studies. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 198-205.	1.1	38
62	Genome-wide association study of colorectal cancer identifies six new susceptibility loci. <i>Nature Communications</i> , 2015, 6, 7138.	5.8	138
63	Association of Aspirin and NSAID Use With Risk of Colorectal Cancer According to Genetic Variants. <i>JAMA - Journal of the American Medical Association</i> , 2015, 313, 1133.	3.8	171
64	A genome-wide association study for colorectal cancer identifies a risk locus in 14q23.1. <i>Human Genetics</i> , 2015, 134, 1249-1262.	1.8	28
65	Identification of a common variant with potential pleiotropic effect on risk of inflammatory bowel disease and colorectal cancer. <i>Carcinogenesis</i> , 2015, 36, 999-1007.	1.3	28
66	Genetic variants of adiponectin and risk of colorectal cancer. <i>International Journal of Cancer</i> , 2015, 137, 154-164.	2.3	16
67	Gene-Environment Interaction Involving Recently Identified Colorectal Cancer Susceptibility Loci. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1824-1833.	1.1	48
68	No Evidence of Gene-Calcium Interactions from Genome-Wide Analysis of Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 2971-2976.	1.1	9
69	Pleiotropic effects of genetic risk variants for other cancers on colorectal cancer risk: PAGE, GECCO and CCFR consortia. <i>Gut</i> , 2014, 63, 800-807.	6.1	35
70	Genome-Wide Diet-Gene Interaction Analyses for Risk of Colorectal Cancer. <i>PLoS Genetics</i> , 2014, 10, e1004228.	1.5	81
71	No association between germline variation in catechol-O-methyltransferase and colorectal cancer survival in postmenopausal women. <i>Menopause</i> , 2014, 21, 415-420.	0.8	5
72	Estimating the heritability of colorectal cancer. <i>Human Molecular Genetics</i> , 2014, 23, 3898-3905.	1.4	114

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73	Identification of Genetic Susceptibility Loci for Colorectal Tumors in a Genome-Wide Meta-analysis. <i>Gastroenterology</i> , 2013, 144, 799-807.e24.	0.6	292
74	Genetic Predictors of Circulating 25-Hydroxyvitamin D and Risk of Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 2037-2046.	1.1	30
75	Confirmation of the Reported Association of Clonal Chromosomal Mosaicism with an Increased Risk of Incident Hematologic Cancer. <i>PLoS ONE</i> , 2013, 8, e59823.	1.1	26
76	Genome-Wide Association Study of Serum Selenium Concentrations. <i>Nutrients</i> , 2013, 5, 1706-1718.	1.7	14
77	A Pooled Analysis of Smoking and Colorectal Cancer: Timing of Exposure and Interactions with Environmental Factors. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 1974-1985.	1.1	54
78	Characterization of Gene-Environment Interactions for Colorectal Cancer Susceptibility Loci. <i>Cancer Research</i> , 2012, 72, 2036-2044.	0.4	140
79	Genome-Wide Search for Gene-Gene Interactions in Colorectal Cancer. <i>PLoS ONE</i> , 2012, 7, e52535.	1.1	35
80	Billing for Medical Genetics and Genetic Counseling Services: A National Survey. <i>Journal of Genetic Counseling</i> , 2010, 19, 38-43.	0.9	29
81	The asthma consultative process: a collaborative approach to integrating genomics into public health practice. <i>Preventing Chronic Disease</i> , 2005, 2, A27.	1.7	1
82	Trade-related Infections: Global Traffic and Microbial Travel. <i>EcoHealth</i> , 2004, 1, 39-49.	0.9	5
83	Family history of diabetes as a potential public health tool. <i>American Journal of Preventive Medicine</i> , 2003, 24, 152-159.	1.6	263
84	Identification of members of the Wnt signaling pathway in the embryonic pituitary gland. <i>Mammalian Genome</i> , 2001, 12, 843-851.	1.0	63