Martha L Slattery

List of Publications by Year in Descending Order

314 papers	15,595 citations	67 h-index	108 g-index
321 ext. papers	17,566 ext. citations	5.2 avg, IF	6.26 L-index

#	Paper	IF	Citations
314	Genome-wide association study identifies tumor anatomical site-specific risk variants for colorectal cancer survival.. *Scientific Reports*, **2022**, 12, 127	4.9	2
313	Energy homeostasis genes modify the association between serum concentrations of IGF-1 and IGFBP-3 and breast cancer risk.. *Scientific Reports*, **2022**, 12, 1837	4.9	1
312	Cumulative menstrual months and breast cancer risk by hormone receptor status and ethnicity: The Breast Cancer Etiology in Minorities Study. *International Journal of Cancer*, **2022**, 150, 208-220	7.5	
311	Beyond GWAS of Colorectal Cancer: Evidence of Interaction with Alcohol Consumption and Putative Causal Variant for the 10q24.2 Region.. *Cancer Epidemiology Biomarkers and Prevention*, **2022**, OF1-OF13	4	0
310	Overall survival is the lowest among young women with postpartum breast cancer.. *European Journal of Cancer*, **2022**, 168, 119-127	7.5	0
309	A Combined Proteomics and Mendelian Randomization Approach to Investigate the Effects of Aspirin-Targeted Proteins on Colorectal Cancer. *Cancer Epidemiology Biomarkers and Prevention*, **2021**, 30, 564-575	4	2
308	Salicylic Acid and Risk of Colorectal Cancer: A Two-Sample Mendelian Randomization Study. *Nutrients*, **2021**, 13,	6.7	1
307	Response to Li and Hopper. *American Journal of Human Genetics*, **2021**, 108, 527-529	11	1
306	Nongenetic Determinants of Risk for Early-Onset Colorectal Cancer. *JNCI Cancer Spectrum*, **2021**, 5, pkab078	4.9	15
305	Genetically Predicted Circulating C-Reactive Protein Concentration and Colorectal Cancer Survival: A Mendelian Randomization Consortium Study. *Cancer Epidemiology Biomarkers and Prevention*, **2021**, 30, 1349-1358	4	1
304	Association Between Smoking and Molecular Subtypes of Colorectal Cancer. *JNCI Cancer Spectrum*, **2021**, 5, pkab056	4.6	2
303	Association of Body Mass Index With Colorectal Cancer Risk by Genome-Wide Variants. *Journal of the National Cancer Institute*, **2021**, 113, 38-47	9.7	6
302	Identifying Novel Susceptibility Genes for Colorectal Cancer Risk From a Transcriptome-Wide Association Study of 125,478 Subjects. *Gastroenterology*, **2021**, 160, 1164-1178.e6	13.3	15
301	Genetically predicted circulating concentrations of micronutrients and risk of colorectal cancer among individuals of European descent: a Mendelian randomization study. *American Journal of Clinical Nutrition*, **2021**, 113, 1490-1502	7	5
300	Genetic architectures of proximal and distal colorectal cancer are partly distinct. *Gut*, **2021**, 70, 1325-1334	19.2	7
299	Smoking Behavior and Prognosis After Colorectal Cancer Diagnosis: A Pooled Analysis of 11 Studies. *JNCI Cancer Spectrum*, **2021**, 5, pkab077	4.6	0
298	Mendelian Randomization of Circulating Polyunsaturated Fatty Acids and Colorectal Cancer Risk. *Cancer Epidemiology Biomarkers and Prevention*, **2020**, 29, 860-870	4	12

297	Functional informed genome-wide interaction analysis of body mass index, diabetes and colorectal cancer risk. *Cancer Medicine*, **2020**, 9, 3563-3573	4.8	4
296	Telomere Maintenance Variants and Survival after Colorectal Cancer: Smoking- and Sex-Specific Associations. *Cancer Epidemiology Biomarkers and Prevention*, **2020**, 29, 1817-1824	4	2
295	Menstrual and reproductive characteristics and breast cancer risk by hormone receptor status and ethnicity: The Breast Cancer Etiology in Minorities study. *International Journal of Cancer*, **2020**, 147, 1808-1822	7.5	4
294	Association Between Molecular Subtypes of Colorectal Tumors and Patient Survival, Based on Pooled Analysis of 7 International Studies. *Gastroenterology*, **2020**, 158, 2158-2168.e4	13.3	17
293	Genetic Variants in the Regulatory T cell-Related Pathway and Colorectal Cancer Prognosis. *Cancer Epidemiology Biomarkers and Prevention*, **2020**, 29, 2719-2728	4	
292	Physical activity and risks of breast and colorectal cancer: a Mendelian randomisation analysis. *Nature Communications*, **2020**, 11, 597	17.4	36
291	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	13.3	47
290	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. *Gastroenterology*, **2020**, 158, 1300-1312.e20	13.3	45
289	Postmenopausal Hormone Therapy and Colorectal Cancer Risk by Molecularly Defined Subtypes and Tumor Location. *JNCI Cancer Spectrum*, **2020**, 4, pkaa042	4.6	2
288	Exploratory Genome-Wide Interaction Analysis of Nonsteroidal Anti-inflammatory Drugs and Predicted Gene Expression on Colorectal Cancer Risk. *Cancer Epidemiology Biomarkers and Prevention*, **2020**, 29, 1800-1808	4	1
287	Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. *American Journal of Human Genetics*, **2020**, 107, 432-444	11	31
286	Circulating bilirubin levels and risk of colorectal cancer: serological and Mendelian randomization analyses. *BMC Medicine*, **2020**, 18, 229	11.4	11
285	Intake of Dietary Fruit, Vegetables, and Fiber and Risk of Colorectal Cancer According to Molecular Subtypes: A Pooled Analysis of 9 Studies. *Cancer Research*, **2020**, 80, 4578-4590	10.1	8
284	Adiposity, metabolites, and colorectal cancer risk: Mendelian randomization study. *BMC Medicine*, **2020**, 18, 396	11.4	17
283	DNA repair and cancer in colon and rectum: Novel players in genetic susceptibility. *International Journal of Cancer*, **2020**, 146, 363-372	7.5	13
282	Meta-analysis of 16 studies of the association of alcohol with colorectal cancer. *International Journal of Cancer*, **2020**, 146, 861-873	7.5	39
281	Genetic Predictors of Circulating 25-Hydroxyvitamin D and Prognosis after Colorectal Cancer. *Cancer Epidemiology Biomarkers and Prevention*, **2020**, 29, 1128-1134	4	
280	The functional role of miRNAs in colorectal cancer: insights from a large population-based study. *Cancer Biology and Medicine*, **2019**, 16, 211-219	5.2	4

279	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. *Journal of the National Cancer Institute*, **2019**, 111, 146-157	9.7	67
278	The p53-signaling pathway and colorectal cancer: Interactions between downstream p53 target genes and miRNAs. *Genomics*, **2019**, 111, 762-771	4.3	41
277	Genetic variant predictors of gene expression provide new insight into risk of colorectal cancer. *Human Genetics*, **2019**, 138, 307-326	6.3	17
276	Combined effect of modifiable and non-modifiable risk factors for colorectal cancer risk in a pooled analysis of 11 population-based studies. *BMJ Open Gastroenterology*, **2019**, 6, e000339	3.9	10
275	A Pooled Analysis of Breastfeeding and Breast Cancer Risk by Hormone Receptor Status in Parous Hispanic Women. *Epidemiology*, **2019**, 30, 449-457	3.1	3
274	Mendelian randomization analysis of C-reactive protein on colorectal cancer risk. *International Journal of Epidemiology*, **2019**, 48, 767-780	7.8	18
273	Discovery of common and rare genetic risk variants for colorectal cancer. *Nature Genetics*, **2019**, 51, 76-87	36.3	177
272	Determining Risk of Colorectal Cancer and Starting Age of Screening Based on Lifestyle, Environmental, and Genetic Factors. *Gastroenterology*, **2018**, 154, 2152-2164.e19	13.3	131
271	The MAPK-Signaling Pathway in Colorectal Cancer: Dysregulated Genes and Their Association With MicroRNAs. *Cancer Informatics*, **2018**, 17, 1176935118766522	2.4	29
270	Mutation analysis of adenomas and carcinomas of the colon: Early and late drivers. *Genes Chromosomes and Cancer*, **2018**, 57, 366-376	5	33
269	Associations between ALDH1A1 polymorphisms, alcohol consumption, and mortality among Hispanic and non-Hispanic white women diagnosed with breast cancer: the Breast Cancer Health Disparities Study. *Breast Cancer Research and Treatment*, **2018**, 168, 443-455	4.4	1
268	Reproductive history, breast-feeding and risk of triple negative breast cancer: The Breast Cancer Etiology in Minorities (BEM) study. *International Journal of Cancer*, **2018**, 142, 2273-2285	7.5	35
267	Association of family history and survival in patients with colorectal cancer: a pooled analysis of eight epidemiologic studies. *Cancer Medicine*, **2018**, 7, 2192-2199	4.8	6
266	Dysregulated genes and miRNAs in the apoptosis pathway in colorectal cancer patients. *Apoptosis: an International Journal on Programmed Cell Death*, **2018**, 23, 237-250	5.4	56
265	The PI3K/AKT signaling pathway: Associations of miRNAs with dysregulated gene expression in colorectal cancer. *Molecular Carcinogenesis*, **2018**, 57, 243-261	5	54
264	The TGF signaling pathway and colorectal cancer: associations between dysregulated genes and miRNAs. *Journal of Translational Medicine*, **2018**, 16, 191	8.5	22
263	Influence of Smoking, Body Mass Index, and Other Factors on the Preventive Effect of Nonsteroidal Anti-Inflammatory Drugs on Colorectal Cancer Risk. *Cancer Research*, **2018**, 78, 4790-4799	10.1	19
262	miRNA involvement in cell cycle regulation in colorectal cancer cases. *Genes and Cancer*, **2018**, 9, 53-65	2.9	24

261	MicroRNA-messenger RNA interactions involving JAK-STAT signaling genes in colorectal cancer. *Genes and Cancer*, **2018**, 9, 232-246	2.9	4
260	Expression of Wnt-signaling pathway genes and their associations with miRNAs in colorectal cancer. *Oncotarget*, **2018**, 9, 6075-6085	3.3	13
259	The NF-B signalling pathway in colorectal cancer: associations between dysregulated gene and miRNA expression. *Journal of Cancer Research and Clinical Oncology*, **2018**, 144, 269-283	4.9	44
258	MicroRNA-transcription factor interactions and their combined effect on target gene expression in colon cancer cases. *Genes Chromosomes and Cancer*, **2018**, 57, 192-202	5	26
257	Power in pairs: assessing the statistical value of paired samples in tests for differential expression. *BMC Genomics*, **2018**, 19, 953	4.5	12
256	Leptin gene variants and colorectal cancer risk: Sex-specific associations. *PLoS ONE*, **2018**, 13, e0206519	3.7	9
255	Response to Conner et al. Re: "Cigarette Smoking and Breast Cancer Risk in Hispanic and Non-Hispanic White Women: The Breast Cancer Health Disparities Study". *Journal of Womenls Health*, **2017**, 26, 92-93	3	1
254	Infrequently expressed miRNAs in colorectal cancer tissue and tumor molecular phenotype. *Modern Pathology*, **2017**, 30, 1152-1169	9.8	15
253	Alterations in microRNA expression associated with alcohol consumption in rectal cancer subjects. *Cancer Causes and Control*, **2017**, 28, 545-555	2.8	5
252	The miRNA landscape of colorectal polyps. *Genes Chromosomes and Cancer*, **2017**, 56, 347-353	5	7
251	Projecting Individualized Absolute Invasive Breast Cancer Risk in US Hispanic Women. *Journal of the National Cancer Institute*, **2017**, 109,	9.7	31
250	The Interaction between Genetic Ancestry and Breast Cancer Risk Factors among Hispanic Women: The Breast Cancer Health Disparities Study. *Cancer Epidemiology Biomarkers and Prevention*, **2017**, 26, 692-701	4	14
249	Identifying factors associated with the direction and significance of microRNA tumor-normal expression differences in colorectal cancer. *BMC Cancer*, **2017**, 17, 707	4.8	4
248	Multiple Gene-Environment Interactions on the Angiogenesis Gene-Pathway Impact Rectal Cancer Risk and Survival. *International Journal of Environmental Research and Public Health*, **2017**, 14,	4.6	2
247	Transcription factor-microRNA associations and their impact on colorectal cancer survival. *Molecular Carcinogenesis*, **2017**, 56, 2512-2526	5	12
246	The co-regulatory networks of tumor suppressor genes, oncogenes, and miRNAs in colorectal cancer. *Genes Chromosomes and Cancer*, **2017**, 56, 769-787	5	49
245	Pre-diagnostic breastfeeding, adiposity, and mortality among parous Hispanic and non-Hispanic white women with invasive breast cancer: the Breast Cancer Health Disparities Study. *Breast Cancer Research and Treatment*, **2017**, 161, 321-331	4.4	4
244	Single nucleotide polymorphisms within MicroRNAs, MicroRNA targets, and MicroRNA biogenesis genes and their impact on colorectal cancer survival. *Genes Chromosomes and Cancer*, **2017**, 56, 285-295	5	13

243	An Assessment of Database-Validated microRNA Target Genes in Normal Colonic Mucosa: Implications for Pathway Analysis. *Cancer Informatics*, **2017**, 16, 1176935117716405	2.4	9
242	Genetic variants in the TGF signaling pathway influence expression of miRNAs in colon and rectal normal mucosa and tumor tissue. *Oncotarget*, **2017**, 8, 16765-16783	3.3	23
241	Infrequently expressed miRNAs influence survival after diagnosis with colorectal cancer. *Oncotarget*, **2017**, 8, 83845-83859	3.3	20
240	Diet and lifestyle factors associated with miRNA expression in colorectal tissue. *Pharmacogenomics and Personalized Medicine*, **2017**, 10, 1-16	2.1	23
239	Energy homeostasis genes and survival after breast cancer diagnosis: the Breast Cancer Health Disparities Study. *Cancer Causes and Control*, **2016**, 27, 47-57	2.8	5
238	Association of cigarette smoking and microRNA expression in rectal cancer: Insight into tumor phenotype. *Cancer Epidemiology*, **2016**, 45, 98-107	2.8	29
237	Impact of polymorphisms in microRNA biogenesis genes on colon cancer risk and microRNA expression levels: a population-based, case-control study. *BMC Medical Genomics*, **2016**, 9, 21	3.7	26
236	CYP24A1 variant modifies the association between use of oestrogen plus progestogen therapy and colorectal cancer risk. *British Journal of Cancer*, **2016**, 114, 221-9	8.7	16
235	MicroRNA profiles in colorectal carcinomas, adenomas and normal colonic mucosa: variations in miRNA expression and disease progression. *Carcinogenesis*, **2016**, 37, 245-261	4.6	84
234	Accounting for Missing Data in Clinical Research. *JAMA - Journal of the American Medical Association* , **2016**, 315, 517-8	27.4	3
233	Identification of Susceptibility Loci and Genes for Colorectal Cancer Risk. *Gastroenterology*, **2016**, 150, 1633-1645	13.3	64
232	Genome-Wide Interaction Analyses between Genetic Variants and Alcohol Consumption and Smoking for Risk of Colorectal Cancer. *PLoS Genetics*, **2016**, 12, e1006296	6	30
231	MicroRNA Seed Region Length Impact on Target Messenger RNA Expression and Survival in Colorectal Cancer. *PLoS ONE*, **2016**, 11, e0154177	3.7	42
230	Telomere Length, TERT, and miRNA Expression. *PLoS ONE*, **2016**, 11, e0162077	3.7	9
229	Site-specific associations between miRNA expression and survival in colorectal cancer cases. *Oncotarget*, **2016**, 7, 60193-60205	3.3	35
228	Genes, environment and gene expression in colon tissue: a pathway approach to determining functionality. *International Journal of Molecular Epidemiology and Genetics*, **2016**, 7, 45-57	0.9	6
227	Bone Mineral Density in Navajo Men and Women and Comparison to Non-Hispanic Whites from NHANES (2005-2008). *Journal of Health Care for the Poor and Underserved*, **2016**, 27, 644-62	1.4	
226	Dietary intake alters gene expression in colon tissue: possible underlying mechanism for the influence of diet on disease. *Pharmacogenetics and Genomics*, **2016**, 26, 294-306	1.9	15

225	Cigarette Smoking and Breast Cancer Risk in Hispanic and Non-Hispanic White Women: The Breast Cancer Health Disparities Study. *Journal of Womenls Health*, **2016**, 25, 299-310	3	9
224	Colorectal tumor molecular phenotype and miRNA: expression profiles and prognosis. *Modern Pathology*, **2016**, 29, 915-27	9.8	33
223	Red meat, poultry, and fish intake and breast cancer risk among Hispanic and Non-Hispanic white women: The Breast Cancer Health Disparities Study. *Cancer Causes and Control*, **2016**, 27, 527-43	2.8	13
222	Ethnic differences in the relationships between diabetes, early age adiposity and mortality among breast cancer survivors: the Breast Cancer Health Disparities Study. *Breast Cancer Research and Treatment*, **2016**, 157, 167-78	4.4	13
221	Expression Profiles of miRNA Subsets Distinguish Human Colorectal Carcinoma and Normal Colonic Mucosa. *Clinical and Translational Gastroenterology*, **2016**, 7, e152	4.2	69
220	Telomere structure and maintenance gene variants and risk of five cancer types. *International Journal of Cancer*, **2016**, 139, 2655-2670	7.5	30
219	A Candidate-Pathway Approach to Identify Gene-Environment Interactions: Analyses of Colon Cancer Risk and Survival. *Journal of the National Cancer Institute*, **2015**, 107,	9.7	17
218	Genome-wide association study of colorectal cancer identifies six new susceptibility loci. *Nature Communications*, **2015**, 6, 7138	17.4	106
217	Body size throughout adult life influences postmenopausal breast cancer risk among hispanic women: the breast cancer health disparities study. *Cancer Epidemiology Biomarkers and Prevention*, **2015**, 24, 128-37	4	18
216	Improved survival among colon cancer patients with increased differentially expressed pathways. *BMC Medicine*, **2015**, 13, 75	11.4	16
215	Association of aspirin and NSAID use with risk of colorectal cancer according to genetic variants. *JAMA - Journal of the American Medical Association*, **2015**, 313, 1133-42	27.4	135
214	Overall and abdominal adiposity and premenopausal breast cancer risk among hispanic women: the breast cancer health disparities study. *Cancer Epidemiology Biomarkers and Prevention*, **2015**, 24, 138-47	4	23
213	Active and passive cigarette smoking and mortality among Hispanic and non-Hispanic white women diagnosed with invasive breast cancer. *Annals of Epidemiology*, **2015**, 25, 824-31	6.4	19
212	A genome-wide association study for colorectal cancer identifies a risk locus in 14q23.1. *Human Genetics*, **2015**, 134, 1249-1262	6.3	25
211	Cross Cancer Genomic Investigation of Inflammation Pathway for Five Common Cancers: Lung, Ovary, Prostate, Breast, and Colorectal Cancer. *Journal of the National Cancer Institute*, **2015**, 107,	9.7	47
210	Interaction between common breast cancer susceptibility variants, genetic ancestry, and nongenetic risk factors in Hispanic women. *Cancer Epidemiology Biomarkers and Prevention*, **2015**, 24, 1731-8	4	12
209	An evaluation and replication of miRNAs with disease stage and colorectal cancer-specific mortality. *International Journal of Cancer*, **2015**, 137, 428-38	7.5	110
208	Incorporation of subject-level covariates in quantile normalization of miRNA data. *BMC Genomics*, **2015**, 16, 1045	4.5	5

207	Associations between ALOX, COX, and CRP polymorphisms and breast cancer among Hispanic and non-Hispanic white women: The breast cancer health disparities study. *Molecular Carcinogenesis*, **2015**, 54, 1541-53	5	17
206	Gene expression in colon cancer: A focus on tumor site and molecular phenotype. *Genes Chromosomes and Cancer*, **2015**, 54, 527-41	5	34
205	Powerful Set-Based Gene-Environment Interaction Testing Framework for Complex Diseases. *Genetic Epidemiology*, **2015**, 39, 609-18	2.6	11
204	Differential Gene Expression in Colon Tissue Associated With Diet, Lifestyle, and Related Oxidative Stress. *PLoS ONE*, **2015**, 10, e0134406	3.7	22
203	SNP Regulation of microRNA Expression and Subsequent Colon Cancer Risk. *PLoS ONE*, **2015**, 10, e0143894	3.7	24
202	Effectiveness and Usability of Bioinformatics Tools to Analyze Pathways Associated with miRNA Expression. *Cancer Informatics*, **2015**, 14, 121-30	2.4	5
201	Mendelian randomization study of height and risk of colorectal cancer. *International Journal of Epidemiology*, **2015**, 44, 662-72	7.8	44
200	Energy homeostasis genes and breast cancer risk: The influence of ancestry, body size, and menopausal status, the breast cancer health disparities study. *Cancer Epidemiology*, **2015**, 39, 1113-22	2.8	14
199	A model to determine colorectal cancer risk using common genetic susceptibility loci. *Gastroenterology*, **2015**, 148, 1330-9.e14	13.3	89
198	A pathway approach to evaluating the association between the CHIEF pathway and risk of colorectal cancer. *Carcinogenesis*, **2015**, 36, 49-59	4.6	11
197	MAPK genes interact with diet and lifestyle factors to alter risk of breast cancer: the Breast Cancer Health Disparities Study. *Nutrition and Cancer*, **2015**, 67, 292-304	2.8	15
196	Accounting for dependence induced by weighted KNN imputation in paired samples, motivated by a colorectal cancer study. *PLoS ONE*, **2015**, 10, e0119876	3.7	21
195	Estimating the heritability of colorectal cancer. *Human Molecular Genetics*, **2014**, 23, 3898-905	5.6	85
194	The influence of genetic ancestry and ethnicity on breast cancer survival associated with genetic variation in the TGF-β signaling pathway: The Breast Cancer Health Disparities Study. *Cancer Causes and Control*, **2014**, 25, 293-307	2.8	10
193	Large-scale genetic study in East Asians identifies six new loci associated with colorectal cancer risk. *Nature Genetics*, **2014**, 46, 533-42	36.3	175
192	Angiogenesis genes, dietary oxidative balance and breast cancer risk and progression: the Breast Cancer Health Disparities Study. *International Journal of Cancer*, **2014**, 134, 629-44	7.5	32
191	Diet and lifestyle factors modify immune/inflammation response genes to alter breast cancer risk and prognosis: the Breast Cancer Health Disparities Study. *Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis*, **2014**, 770, 19-28	3.3	21
190	Genetic variation in the JAK/STAT/SOCS signaling pathway influences breast cancer-specific mortality through interaction with cigarette smoking and use of aspirin/NSAIDs: the Breast Cancer Health Disparities Study. *Breast Cancer Research and Treatment*, **2014**, 147, 145-58	4.4	28

189	Diet and lifestyle factors interact with MAPK genes to influence survival: the Breast Cancer Health Disparities Study. *Cancer Causes and Control*, **2014**, 25, 1211-25	2.8	13
188	An analysis of genetic factors related to risk of inflammatory bowel disease and colon cancer. *Cancer Epidemiology*, **2014**, 38, 583-90	2.8	22
187	Genetic variants in interleukin genes are associated with breast cancer risk and survival in a genetically admixed population: the Breast Cancer Health Disparities Study. *Carcinogenesis*, **2014**, 35, 1750-9	4.6	29
186	VEGFA, FLT1, KDR and colorectal cancer: assessment of disease risk, tumor molecular phenotype, and survival. *Molecular Carcinogenesis*, **2014**, 53 Suppl 1, E140-50	5	31
185	The influence of the CHIEF pathway on colorectal cancer-specific mortality. *PLoS ONE*, **2014**, 9, e116169	3.7	13
184	CYP19A1 single nucleotide polymorphism associations with CYP19A1, NFB1, and IL6 gene expression in human normal colon and normal liver samples. *Pharmacogenomics and Personalized Medicine*, **2014**, 7, 163-71	2.1	2
183	Gene-environment interaction involving recently identified colorectal cancer susceptibility Loci. *Cancer Epidemiology Biomarkers and Prevention*, **2014**, 23, 1824-33	4	40
182	Genome-wide diet-gene interaction analyses for risk of colorectal cancer. *PLoS Genetics*, **2014**, 10, e1004228	8	66
181	Associations between CYP19A1 polymorphisms, Native American ancestry, and breast cancer risk and mortality: the Breast Cancer Health Disparities Study. *Cancer Causes and Control*, **2014**, 25, 1461-71	2.8	7
180	Genetic variants and non-genetic factors predict circulating vitamin D levels in Hispanic and non-Hispanic White women: the Breast Cancer Health Disparities Study. *International Journal of Molecular Epidemiology and Genetics*, **2014**, 5, 31-46	0.9	15
179	Influence of CHIEF pathway genes on gene expression: a pathway approach to functionality. *International Journal of Molecular Epidemiology and Genetics*, **2014**, 5, 100-11	0.9	8
178	Interleukin genes and associations with colon and rectal cancer risk and overall survival. *International Journal of Cancer*, **2013**, 132, 905-15	7.5	60
177	JAK/STAT/SOCS-signaling pathway and colon and rectal cancer. *Molecular Carcinogenesis*, **2013**, 52, 155-66	5	134
176	TERTs role in colorectal carcinogenesis. *Molecular Carcinogenesis*, **2013**, 52, 507-13	5	17
175	Dietary influence on MAPK-signaling pathways and risk of colon and rectal cancer. *Nutrition and Cancer*, **2013**, 65, 729-38	2.8	35
174	Associations with growth factor genes (FGF1, FGF2, PDGFB, FGFR2, NRG2, EGF, ERBB2) with breast cancer risk and survival: the Breast Cancer Health Disparities Study. *Breast Cancer Research and Treatment*, **2013**, 140, 587-601	4.4	49
173	Telomere length, telomere-related genes, and breast cancer risk: the breast cancer health disparities study. *Genes Chromosomes and Cancer*, **2013**, 52, 595-609	5	66
172	Genetic ancestry modifies the association between genetic risk variants and breast cancer risk among Hispanic and non-Hispanic white women. *Carcinogenesis*, **2013**, 34, 1787-93	4.6	22

171	Genetic variation in bone morphogenetic proteins and breast cancer risk in hispanic and non-hispanic white women: The breast cancer health disparities study. *International Journal of Cancer*, **2013**, 132, 2928-39	7.5	15
170	Identification of Genetic Susceptibility Loci for Colorectal Tumors in a Genome-Wide Meta-analysis. *Gastroenterology*, **2013**, 144, 799-807.e24	13.3	250
169	Associations between genetic variants in the TGF- signaling pathway and breast cancer risk among Hispanic and non-Hispanic white women. *Breast Cancer Research and Treatment*, **2013**, 141, 287-97	4.4	10
168	IBK and NFB1, NSAID use and risk of colorectal cancer in the Colon Cancer Family Registry. *Carcinogenesis*, **2013**, 34, 79-85	4.6	33
167	Genetic variation in the lipoxygenase pathway and risk of colorectal neoplasia. *Genes Chromosomes and Cancer*, **2013**, 52, 437-49	5	25
166	Dietary intake of folate, B-vitamins and methionine and breast cancer risk among Hispanic and non-Hispanic white women. *PLoS ONE*, **2013**, 8, e54495	3.7	22
165	Matrix metalloproteinase genes are associated with breast cancer risk and survival: the Breast Cancer Health Disparities Study. *PLoS ONE*, **2013**, 8, e63165	3.7	60
164	SEPP1 influences breast cancer risk among women with greater native american ancestry: the breast cancer health disparities study. *PLoS ONE*, **2013**, 8, e80554	3.7	29
163	Epidermal growth factor receptor (EGFR) polymorphisms and breast cancer among Hispanic and non-Hispanic white women: the Breast Cancer Health Disparities Study. *International Journal of Molecular Epidemiology and Genetics*, **2013**, 4, 235-49	0.9	7
162	Oxidative balance and colon and rectal cancer: interaction of lifestyle factors and genes. *Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis*, **2012**, 734, 30-40	3.3	32
161	Glutathione peroxidase tagSNPs: associations with rectal cancer but not with colon cancer. *Genes Chromosomes and Cancer*, **2012**, 51, 598-605	5	15
160	Genetic variation in bone morphogenetic protein and colon and rectal cancer. *International Journal of Cancer*, **2012**, 130, 653-64	7.5	23
159	Meta-analysis of new genome-wide association studies of colorectal cancer risk. *Human Genetics*, **2012**, 131, 217-34	6.3	173
158	MAP kinase genes and colon and rectal cancer. *Carcinogenesis*, **2012**, 33, 2398-408	4.6	70
157	Associations between TCF7L2 polymorphisms and risk of breast cancer among Hispanic and non-Hispanic white women: the Breast Cancer Health Disparities Study. *Breast Cancer Research and Treatment*, **2012**, 136, 593-602	4.4	23
156	ADRB2 G-G haplotype associated with breast cancer risk among Hispanic and non-Hispanic white women: interaction with type 2 diabetes and obesity. *Cancer Causes and Control*, **2012**, 23, 1653-63	2.8	10
155	Characterization of gene-environment interactions for colorectal cancer susceptibility loci. *Cancer Research*, **2012**, 72, 2036-44	10.1	119
154	Genetic variability in IL23R and risk of colorectal adenoma and colorectal cancer. *Cancer Epidemiology*, **2012**, 36, e104-10	2.8	16

153	Genetic variation in selenoprotein genes, lifestyle, and risk of colon and rectal cancer. *PLoS ONE*, **2012**, 7, e37312	3.7	34
152	Construct validity of the SF-12 among American Indian and Alaska Native people using two known scoring methods. *Journal of Health Care for the Poor and Underserved*, **2012**, 23, 1123-36	1.4	13
151	Toll-like receptor genes and their association with colon and rectal cancer development and prognosis. *International Journal of Cancer*, **2012**, 130, 2974-80	7.5	81
150	Genetic variation in genes involved in hormones, inflammation and energetic factors and breast cancer risk in an admixed population. *Carcinogenesis*, **2012**, 33, 1512-21	4.6	63
149	A pooled analysis of smoking and colorectal cancer: timing of exposure and interactions with environmental factors. *Cancer Epidemiology Biomarkers and Prevention*, **2012**, 21, 1974-85	4	47
148	Genetic variation in the transforming growth factor-β signaling pathway, lifestyle factors, and risk of colon or rectal cancer. *Diseases of the Colon and Rectum*, **2012**, 55, 532-40	3.1	9
147	Genome-wide search for gene-gene interactions in colorectal cancer. *PLoS ONE*, **2012**, 7, e52535	3.7	29
146	Genetic and lifestyle influence on telomere length and subsequent risk of colon cancer in a case control study. *International Journal of Molecular Epidemiology and Genetics*, **2012**, 3, 184-94	0.9	32
145	Nutrients in folate-mediated, one-carbon metabolism and the risk of rectal tumors in men and women. *Nutrition and Cancer*, **2011**, 63, 357-66	2.8	23
144	Interferon-signaling pathway: associations with colon and rectal cancer risk and subsequent survival. *Carcinogenesis*, **2011**, 32, 1660-7	4.6	78
143	CpG island methylation in colorectal cancer: past, present and future. *Pathology Research International*, **2011**, 2011, 902674		91
142	Genetic variation in RPS6KA1, RPS6KA2, RPS6KB1, RPS6KB2, and PDK1 and risk of colon or rectal cancer. *Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis*, **2011**, 706, 13-20	3.3	40
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