

# Steven Gallinger

## List of Publications by Citations

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276  
papers

31,036  
citations

66  
h-index

175  
g-index

296  
ext. papers

36,422  
ext. citations

9.8  
avg, IF

6.35  
L-index

#	Paper	IF	Citations
276	A human colon cancer cell capable of initiating tumour growth in immunodeficient mice. <i>Nature</i> , <b>2007</b> , 445, 106-10	50.4	3323
275	Erlotinib plus gemcitabine compared with gemcitabine alone in patients with advanced pancreatic cancer: a phase III trial of the National Cancer Institute of Canada Clinical Trials Group. <i>Journal of Clinical Oncology</i> , <b>2007</b> , 25, 1960-6	2.2	2883
274	Tumor microsatellite-instability status as a predictor of benefit from fluorouracil-based adjuvant chemotherapy for colon cancer. <i>New England Journal of Medicine</i> , <b>2003</b> , 349, 247-57	59.2	1641
273	Pancreatic cancer genomes reveal aberrations in axon guidance pathway genes. <i>Nature</i> , <b>2012</b> , 491, 399-405	50.4	1427
272	Tumor microsatellite instability and clinical outcome in young patients with colorectal cancer. <i>New England Journal of Medicine</i> , <b>2000</b> , 342, 69-77	59.2	1076
271	Defective mismatch repair as a predictive marker for lack of efficacy of fluorouracil-based adjuvant therapy in colon cancer. <i>Journal of Clinical Oncology</i> , <b>2010</b> , 28, 3219-26	2.2	1063
270	Adjuvant chemotherapy with fluorouracil plus folinic acid vs gemcitabine following pancreatic cancer resection: a randomized controlled trial. <i>JAMA - Journal of the American Medical Association</i> , <b>2010</b> , 304, 1073-81	27.4	958
269	Discovery of cross-reactive probes and polymorphic CpGs in the Illumina Infinium HumanMethylation450 microarray. <i>Epigenetics</i> , <b>2013</b> , 8, 203-9	5.7	953
268	MADR2 maps to 18q21 and encodes a TGFbeta-regulated MAD-related protein that is functionally mutated in colorectal carcinoma. <i>Cell</i> , <b>1996</b> , 86, 543-52	56.2	764
267	Convergence of genes and cellular pathways dysregulated in autism spectrum disorders. <i>American Journal of Human Genetics</i> , <b>2014</b> , 94, 677-94	11	635
266	Genome-wide association scan identifies a colorectal cancer susceptibility locus on chromosome 8q24. <i>Nature Genetics</i> , <b>2007</b> , 39, 989-94	36.3	609
265	Variable clonal repopulation dynamics influence chemotherapy response in colorectal cancer. <i>Science</i> , <b>2013</b> , 339, 543-8	33.3	550
264	Genome-wide association scan identifies a colorectal cancer susceptibility locus on 11q23 and replicates risk loci at 8q24 and 18q21. <i>Nature Genetics</i> , <b>2008</b> , 40, 631-7	36.3	486
263	Genome-wide association study identifies variants in the ABO locus associated with susceptibility to pancreatic cancer. <i>Nature Genetics</i> , <b>2009</b> , 41, 986-90	36.3	483
262	A genome-wide association study identifies pancreatic cancer susceptibility loci on chromosomes 13q22.1, 1q32.1 and 5p15.33. <i>Nature Genetics</i> , <b>2010</b> , 42, 224-8	36.3	463
261	Meta-analysis of genome-wide association data identifies four new susceptibility loci for colorectal cancer. <i>Nature Genetics</i> , <b>2008</b> , 40, 1426-35	36.3	457
260	CCAT2, a novel noncoding RNA mapping to 8q24, underlies metastatic progression and chromosomal instability in colon cancer. <i>Genome Research</i> , <b>2013</b> , 23, 1446-61	9.7	442

259	Detectable clonal mosaicism and its relationship to aging and cancer. <i>Nature Genetics</i> , <b>2012</b> , 44, 651-8	36.3	409
258	Ductal pancreatic cancer modeling and drug screening using human pluripotent stem cell- and patient-derived tumor organoids. <i>Nature Medicine</i> , <b>2015</b> , 21, 1364-71	50.5	403
257	Hereditary Diffuse Gastric Cancer Syndrome: CDH1 Mutations and Beyond. <i>JAMA Oncology</i> , <b>2015</b> , 1, 23-32	13.4	401
256	DNA mismatch repair status and colon cancer recurrence and survival in clinical trials of 5-fluorouracil-based adjuvant therapy. <i>Journal of the National Cancer Institute</i> , <b>2011</b> , 103, 863-75	9.7	390
255	ATM mutations in patients with hereditary pancreatic cancer. <i>Cancer Discovery</i> , <b>2012</b> , 2, 41-6	24.4	365
254	Self-renewal as a therapeutic target in human colorectal cancer. <i>Nature Medicine</i> , <b>2014</b> , 20, 29-36	50.5	361
253	Sensitive tumour detection and classification using plasma cell-free DNA methylomes. <i>Nature</i> , <b>2018</b> , 563, 579-583	50.4	344
252	Genome-wide association study identifies eight risk loci and implicates metabo-psychiatric origins for anorexia nervosa. <i>Nature Genetics</i> , <b>2019</b> , 51, 1207-1214	36.3	303
251	Identification of Genetic Susceptibility Loci for Colorectal Tumors in a Genome-Wide Meta-analysis. <i>Gastroenterology</i> , <b>2013</b> , 144, 799-807.e24	13.3	250
250	Germline BRCA Mutations in a Large Clinic-Based Cohort of Patients With Pancreatic Adenocarcinoma. <i>Journal of Clinical Oncology</i> , <b>2015</b> , 33, 3124-9	2.2	241
249	Genomics-Driven Precision Medicine for Advanced Pancreatic Cancer: Early Results from the COMPASS Trial. <i>Clinical Cancer Research</i> , <b>2018</b> , 24, 1344-1354	12.9	240
248	Association Between Telomere Length and Risk of Cancer and Non-Neoplastic Diseases: A Mendelian Randomization Study. <i>JAMA Oncology</i> , <b>2017</b> , 3, 636-651	13.4	236
247	Genome-wide association study identifies multiple susceptibility loci for pancreatic cancer. <i>Nature Genetics</i> , <b>2014</b> , 46, 994-1000	36.3	226
246	Whole Genome Sequencing Defines the Genetic Heterogeneity of Familial Pancreatic Cancer. <i>Cancer Discovery</i> , <b>2016</b> , 6, 166-75	24.4	206
245	Prevalence of germline mutations in cancer predisposition genes in patients with pancreatic cancer. <i>Gastroenterology</i> , <b>2015</b> , 148, 556-64	13.3	200
244	Prevalence and Penetrance of Major Genes and Polygenes for Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2017</b> , 26, 404-412	4	185
243	Common variation near CDKN1A, POLD3 and SHROOM2 influences colorectal cancer risk. <i>Nature Genetics</i> , <b>2012</b> , 44, 770-6	36.3	184
242	Germline BRCA2 6174delT mutations in Ashkenazi Jewish pancreatic cancer patients. <i>Nature Genetics</i> , <b>1997</b> , 16, 17-8	36.3	177

241	Discovery of common and rare genetic risk variants for colorectal cancer. <i>Nature Genetics</i> , <b>2019</b> , 51, 76-83	6.3	177
240	BRCA1, BRCA2, PALB2, and CDKN2A mutations in familial pancreatic cancer: a PACGENE study. <i>Genetics in Medicine</i> , <b>2015</b> , 17, 569-77	8.1	175
239	Meta-analysis of new genome-wide association studies of colorectal cancer risk. <i>Human Genetics</i> , <b>2012</b> , 131, 217-34	6.3	173
238	Common variation at 2p13.3, 3q29, 7p13 and 17q25.1 associated with susceptibility to pancreatic cancer. <i>Nature Genetics</i> , <b>2015</b> , 47, 911-6	36.3	171
237	Molecular characterization of MSI-H colorectal cancer by MLH1 promoter methylation, immunohistochemistry, and mismatch repair germline mutation screening. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2008</b> , 17, 3208-15	4	171
236	Germline MutY human homologue mutations and colorectal cancer: a multisite case-control study. <i>Gastroenterology</i> , <b>2009</b> , 136, 1251-60	13.3	165
235	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> , <b>2020</b> , 22, 15-25	8.1	164
234	Molecular biology of colorectal cancer. <i>Current Problems in Cancer</i> , <b>1997</b> , 21, 233-300	2.3	159
233	Association of Distinct Mutational Signatures With Correlates of Increased Immune Activity in Pancreatic Ductal Adenocarcinoma. <i>JAMA Oncology</i> , <b>2017</b> , 3, 774-783	13.4	157
232	ID1 and ID3 regulate the self-renewal capacity of human colon cancer-initiating cells through p21. <i>Cancer Cell</i> , <b>2012</b> , 21, 777-92	24.3	157
231	Transcription phenotypes of pancreatic cancer are driven by genomic events during tumor evolution. <i>Nature Genetics</i> , <b>2020</b> , 52, 231-240	36.3	148
230	Case-control study of overweight, obesity, and colorectal cancer risk, overall and by tumor microsatellite instability status. <i>Journal of the National Cancer Institute</i> , <b>2010</b> , 102, 391-400	9.7	133
229	Problems of delivery of monoclonal antibodies. Pharmaceutical and pharmacokinetic solutions. <i>Clinical Pharmacokinetics</i> , <b>1995</b> , 28, 126-42	6.2	131
228	Risk of colorectal cancer for carriers of mutations in MUTYH, with and without a family history of cancer. <i>Gastroenterology</i> , <b>2014</b> , 146, 1208-11.e1-5	13.3	128
227	Analysis of the gene coding for the BRCA2-interacting protein PALB2 in familial and sporadic pancreatic cancer. <i>Gastroenterology</i> , <b>2009</b> , 137, 1183-6	13.3	121
226	Characterization of gene-environment interactions for colorectal cancer susceptibility loci. <i>Cancer Research</i> , <b>2012</b> , 72, 2036-44	10.1	119
225	Gastrointestinal cancers and neurofibromatosis type 1 features in children with a germline homozygous MLH1 mutation. <i>Gastroenterology</i> , <b>2004</b> , 126, 576-85	13.3	107
224	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. <i>Nature Communications</i> , <b>2018</b> , 9, 556	17.4	103

223	Cumulative impact of common genetic variants and other risk factors on colorectal cancer risk in 42,103 individuals. <i>Gut</i> , <b>2013</b> , 62, 871-81	19.2	95
222	Quality assessment and correlation of microsatellite instability and immunohistochemical markers among population- and clinic-based colorectal tumors results from the Colon Cancer Family Registry. <i>Journal of Molecular Diagnostics</i> , <b>2011</b> , 13, 271-81	5.1	95
221	Colorectal carcinomas in mice lacking the catalytic subunit of PI(3)Kgamma. <i>Nature</i> , <b>2000</b> , 406, 897-902	50.4	94
220	Long-range epigenetic regulation is conferred by genetic variation located at thousands of independent loci. <i>Nature Communications</i> , <b>2015</b> , 6, 6326	17.4	90
219	Pathway analysis of genome-wide association study data highlights pancreatic development genes as susceptibility factors for pancreatic cancer. <i>Carcinogenesis</i> , <b>2012</b> , 33, 1384-90	4.6	85
218	An absolute risk model to identify individuals at elevated risk for pancreatic cancer in the general population. <i>PLoS ONE</i> , <b>2013</b> , 8, e72311	3.7	82
217	Variants on 9p24 and 8q24 are associated with risk of colorectal cancer: results from the Colon Cancer Family Registry. <i>Cancer Research</i> , <b>2007</b> , 67, 11128-32	10.1	82
216	Association analyses identify 31 new risk loci for colorectal cancer susceptibility. <i>Nature Communications</i> , <b>2019</b> , 10, 2154	17.4	81
215	Integration of Genomic and Transcriptional Features in Pancreatic Cancer Reveals Increased Cell Cycle Progression in Metastases. <i>Cancer Cell</i> , <b>2019</b> , 35, 267-282.e7	24.3	80
214	Phenotypic and genotypic characterisation of biallelic mismatch repair deficiency (BMMR-D) syndrome. <i>European Journal of Cancer</i> , <b>2015</b> , 51, 977-83	7.5	77
213	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 6616-33	5.6	77
212	Family history characteristics, tumor microsatellite instability and germline MSH2 and MLH1 mutations in hereditary colorectal cancer. <i>Human Genetics</i> , <b>1999</b> , 104, 167-76	6.3	76
211	Essays on science and society. Defining disease in the genomics era. <i>Science</i> , <b>2001</b> , 293, 807-8	33.3	70
210	Aspirin, Ibuprofen, and the Risk of Colorectal Cancer in Lynch Syndrome. <i>Journal of the National Cancer Institute</i> , <b>2015</b> , 107,	9.7	66
209	Genome-wide diet-gene interaction analyses for risk of colorectal cancer. <i>PLoS Genetics</i> , <b>2014</b> , 10, e1004228	4.2	66
208	Prognostic value of microsatellite instability (MSI) and PTEN expression in women with endometrial cancer: results from studies of the NCIC Clinical Trials Group (NCIC CTG). <i>European Journal of Cancer</i> , <b>2010</b> , 46, 1365-73	7.5	66
207	Promoter methylation of Wnt antagonists DKK1 and SFRP1 is associated with opposing tumor subtypes in two large populations of colorectal cancer patients. <i>Carcinogenesis</i> , <b>2011</b> , 32, 741-7	4.6	66
206	Three new pancreatic cancer susceptibility signals identified on chromosomes 1q32.1, 5p15.33 and 8q24.21. <i>Oncotarget</i> , <b>2016</b> , 7, 66328-66343	3.3	66

205	Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. <i>Cancer Research</i> , <b>2016</b> , 76, 5103-14	10.1	66
204	Overall Survival and Clinical Characteristics of BRCA-Associated Cholangiocarcinoma: A Multicenter Retrospective Study. <i>Oncologist</i> , <b>2017</b> , 22, 804-810	5.7	65
203	Identification of Susceptibility Loci and Genes for Colorectal Cancer Risk. <i>Gastroenterology</i> , <b>2016</b> , 150, 1633-1645	13.3	64
202	Hepatic resection of noncolorectal nonneuroendocrine metastases. <i>Liver Transplantation</i> , <b>2000</b> , 6, 97-101	4.5	63
201	Trans-ethnic genome-wide association study of colorectal cancer identifies a new susceptibility locus in VTI1A. <i>Nature Communications</i> , <b>2014</b> , 5, 4613	17.4	62
200	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , <b>2016</b> , 7, 11843	17.4	59
199	Senescent Carcinoma-Associated Fibroblasts Upregulate IL8 to Enhance Prometastatic Phenotypes. <i>Molecular Cancer Research</i> , <b>2017</b> , 15, 3-14	6.6	59
198	Association of the colorectal CpG island methylator phenotype with molecular features, risk factors, and family history. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 512-519	4	59
197	The APC11307K allele and breast cancer risk. <i>Nature Genetics</i> , <b>1998</b> , 20, 13-4	36.3	59
196	Pro-inflammatory fatty acid profile and colorectal cancer risk: A Mendelian randomisation analysis. <i>European Journal of Cancer</i> , <b>2017</b> , 84, 228-238	7.5	56
195	Familial adenomatous polyposis. <i>Journal of Surgical Oncology</i> , <b>2000</b> , 18, 314-23		56
194	Mendelian Randomization Study of Body Mass Index and Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 1024-31	4	54
193	Female Hormonal Factors and the Risk of Endometrial Cancer in Lynch Syndrome. <i>JAMA - Journal of the American Medical Association</i> , <b>2015</b> , 314, 61-71	27.4	53
192	cis-Expression QTL analysis of established colorectal cancer risk variants in colon tumors and adjacent normal tissue. <i>PLoS ONE</i> , <b>2012</b> , 7, e30477	3.7	52
191	Fine-mapping of colorectal cancer susceptibility loci at 8q23.3, 16q22.1 and 19q13.11: refinement of association signals and use of in silico analysis to suggest functional variation and unexpected candidate target genes. <i>Human Molecular Genetics</i> , <b>2011</b> , 20, 2879-88	5.6	51
190	Hereditary colorectal cancer syndromes: familial adenomatous polyposis and lynch syndrome. <i>Surgical Clinics of North America</i> , <b>2008</b> , 88, 819-44, vii	4	51
189	Mendelian randomisation implicates hyperlipidaemia as a risk factor for colorectal cancer. <i>International Journal of Cancer</i> , <b>2017</b> , 140, 2701-2708	7.5	50
188	Overall survival and clinical characteristics of BRCA mutation carriers with stage I/II pancreatic cancer. <i>British Journal of Cancer</i> , <b>2017</b> , 116, 697-702	8.7	49

187	Microsatellite instability as a prognostic factor in resected colorectal cancer liver metastases. <i>Annals of Surgical Oncology</i> , <b>2004</b> , 11, 977-82	3.1	49
186	Genotype-environment interactions in microsatellite stable/microsatellite instability-low colorectal cancer: results from a genome-wide association study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2011</b> , 20, 758-66	4	47
185	A pooled analysis of smoking and colorectal cancer: timing of exposure and interactions with environmental factors. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2012</b> , 21, 1974-85	4	47
184	Molecular Events in the Natural History of Pancreatic Cancer. <i>Trends in Cancer</i> , <b>2017</b> , 3, 336-346	12.5	45
183	Improved long-term outcomes after resection of pancreatic adenocarcinoma: a comparison between two time periods. <i>Annals of Surgical Oncology</i> , <b>2015</b> , 22, 1160-7	3.1	43
182	Candidate DNA repair susceptibility genes identified by exome sequencing in high-risk pancreatic cancer. <i>Cancer Letters</i> , <b>2016</b> , 370, 302-12	9.9	42
181	Association between colonic screening, subject characteristics, and stage of colorectal cancer. <i>American Journal of Gastroenterology</i> , <b>2005</b> , 100, 2531-9	0.7	42
180	Recurrent noncoding regulatory mutations in pancreatic ductal adenocarcinoma. <i>Nature Genetics</i> , <b>2017</b> , 49, 825-833	36.3	41
179	STK11/LKB1 germline mutations are not identified in most Peutz-Jeghers syndrome patients. <i>Clinical Genetics</i> , <b>1999</b> , 56, 136-41	4	40
178	Hypoxia provokes base excision repair changes and a repair-deficient, mutator phenotype in colorectal cancer cells. <i>Molecular Cancer Research</i> , <b>2014</b> , 12, 1407-15	6.6	39
177	Microsatellite instability, mismatch repair deficiency, and colorectal cancer. <i>Surgery</i> , <b>2001</b> , 130, 17-20	3.6	39
176	Intraductal Transplantation Models of Human Pancreatic Ductal Adenocarcinoma Reveal Progressive Transition of Molecular Subtypes. <i>Cancer Discovery</i> , <b>2020</b> , 10, 1566-1589	24.4	39
175	Association between body mass index and mortality for colorectal cancer survivors: overall and by tumor molecular phenotype. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 1229-38	4	38
174	A Four-Chemokine Signature Is Associated with a T-cell-Inflamed Phenotype in Primary and Metastatic Pancreatic Cancer. <i>Clinical Cancer Research</i> , <b>2020</b> , 26, 1997-2010	12.9	37
173	Axonal guidance signaling pathway interacting with smoking in modifying the risk of pancreatic cancer: a gene- and pathway-based interaction analysis of GWAS data. <i>Carcinogenesis</i> , <b>2014</b> , 35, 1039-45	4.6	36
172	Lack of evidence for germline mutations in patients with serrated polyposis syndrome from a large multinational study. <i>Gut</i> , <b>2017</b> , 66, 1170-1172	19.2	35
171	Allergies are associated with reduced pancreas cancer risk: A population-based case-control study in Ontario, Canada. <i>International Journal of Cancer</i> , <b>2007</b> , 121, 2241-5	7.5	35
170	Mutations in the pancreatic secretory enzymes and are associated with pancreatic cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2018</b> , 115, 4767-4772	11.5	34

169	Genetic variants in vitamin d pathway genes and risk of pancreas cancer; results from a population-based case-control study in ontario, Canada. <i>PLoS ONE</i> , <b>2013</b> , 8, e66768	3.7	34
168	Red meat intake, NAT2, and risk of colorectal cancer: a pooled analysis of 11 studies. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 198-205	4	32
167	Patients with both pancreatic adenocarcinoma and melanoma may harbor germline CDKN2A mutations <b>2000</b> , 27, 358-361		32
166	Specific variants in the MLH1 gene region may drive DNA methylation, loss of protein expression, and MSI-H colorectal cancer. <i>PLoS ONE</i> , <b>2010</b> , 5, e13314	3.7	32
165	Role of tumour molecular and pathology features to estimate colorectal cancer risk for first-degree relatives. <i>Gut</i> , <b>2015</b> , 64, 101-10	19.2	31
164	The New Era of Transplant Oncology: Liver Transplantation for Nonresectable Colorectal Cancer Liver Metastases. <i>Canadian Journal of Gastroenterology and Hepatology</i> , <b>2018</b> , 2018, 9531925	2.8	31
163	Genome-Wide Interaction Analyses between Genetic Variants and Alcohol Consumption and Smoking for Risk of Colorectal Cancer. <i>PLoS Genetics</i> , <b>2016</b> , 12, e1006296	6	30
162	Genomic Features and Classification of Homologous Recombination Deficient Pancreatic Ductal Adenocarcinoma. <i>Gastroenterology</i> , <b>2021</b> , 160, 2119-2132.e9	13.3	30
161	Genome-wide search for gene-gene interactions in colorectal cancer. <i>PLoS ONE</i> , <b>2012</b> , 7, e52535	3.7	29
160	Spatially confined sub-tumor microenvironments in pancreatic cancer. <i>Cell</i> , <b>2021</b> , 184, 5577-5592.e18	56.2	29
159	Whole genomes define concordance of matched primary, xenograft, and organoid models of pancreas cancer. <i>PLoS Computational Biology</i> , <b>2019</b> , 15, e1006596	5	29
158	Exome sequencing identifies nonsegregating nonsense ATM and PALB2 variants in familial pancreatic cancer. <i>Human Genomics</i> , <b>2013</b> , 7, 11	6.8	28
157	Characterization, Detection, and Treatment Approaches for Homologous Recombination Deficiency in Cancer. <i>Trends in Molecular Medicine</i> , <b>2017</b> , 23, 1121-1137	11.5	28
156	Alcohol Consumption and the Risk of Colorectal Cancer for Mismatch Repair Gene Mutation Carriers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2017</b> , 26, 366-375	4	28
155	Identification of genes expressed by immune cells of the colon that are regulated by colorectal cancer-associated variants. <i>International Journal of Cancer</i> , <b>2014</b> , 134, 2330-41	7.5	28
154	Diagnosis and management of pancreatic cancer. <i>Cmaj</i> , <b>2013</b> , 185, 1219-26	3.5	28
153	Analysis of Heritability and Genetic Architecture of Pancreatic Cancer: A PanC4 Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2019</b> , 28, 1238-1245	4	27
152	Pleiotropic effects of genetic risk variants for other cancers on colorectal cancer risk: PAGE, GECCO and CCFR consortia. <i>Gut</i> , <b>2014</b> , 63, 800-7	19.2	27



151	Genes-environment interactions in obesity- and diabetes-associated pancreatic cancer: a GWAS data analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2014</b> , 23, 98-106	4	26
150	Genetic predictors of circulating 25-hydroxyvitamin d and risk of colorectal cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2013</b> , 22, 2037-46	4	26
149	Meta-analysis of 8q24 for seven cancers reveals a locus between NOV and ENPP2 associated with cancer development. <i>BMC Medical Genetics</i> , <b>2011</b> , 12, 156	2.1	26
148	Clinical and genomic characterisation of mismatch repair deficient pancreatic adenocarcinoma. <i>Gut</i> , <b>2021</b> , 70, 1894-1903	19.2	26
147	A genome-wide association study for colorectal cancer identifies a risk locus in 14q23.1. <i>Human Genetics</i> , <b>2015</b> , 134, 1249-1262	6.3	25
146	A novel colorectal cancer risk locus at 4q32.2 identified from an international genome-wide association study. <i>Carcinogenesis</i> , <b>2014</b> , 35, 2512-9	4.6	25
145	A Transcriptome-Wide Association Study Identifies Novel Candidate Susceptibility Genes for Pancreatic Cancer. <i>Journal of the National Cancer Institute</i> , <b>2020</b> , 112, 1003-1012	9.7	25
144	Germline mutations in PMS2 and MLH1 in individuals with solitary loss of PMS2 expression in colorectal carcinomas from the Colon Cancer Family Registry Cohort. <i>BMJ Open</i> , <b>2016</b> , 6, e010293	3	24
143	Association between Variants in Atopy-Related Immunologic Candidate Genes and Pancreatic Cancer Risk. <i>PLoS ONE</i> , <b>2015</b> , 10, e0125273	3.7	24
142	Glypican-1 and glycoprotein 2 bearing extracellular vesicles do not discern pancreatic cancer from benign pancreatic diseases. <i>Oncotarget</i> , <b>2019</b> , 10, 1045-1055	3.3	24
141	Hereditary Pancreatic Cancer Syndromes. <i>Surgical Oncology Clinics of North America</i> , <b>2015</b> , 24, 733-64	2.7	23
140	Cohort Profile: The Colon Cancer Family Registry Cohort (CCFRC). <i>International Journal of Epidemiology</i> , <b>2018</b> , 47, 387-388i	7.8	23
139	Colorectal cancer linkage on chromosomes 4q21, 8q13, 12q24, and 15q22. <i>PLoS ONE</i> , <b>2012</b> , 7, e38175	3.7	23
138	Association between alcohol consumption and pancreatic cancer risk: a case-control study. <i>PLoS ONE</i> , <b>2015</b> , 10, e0124489	3.7	22
137	Association of Common Susceptibility Variants of Pancreatic Cancer in Higher-Risk Patients: A PACGENE Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2016</b> , 25, 1185-91	4	22
136	Identification of a common variant with potential pleiotropic effect on risk of inflammatory bowel disease and colorectal cancer. <i>Carcinogenesis</i> , <b>2015</b> , 36, 999-1007	4.6	21
135	Pattern of Invasion in Human Pancreatic Cancer Organoids Is Associated with Loss of SMAD4 and Clinical Outcome. <i>Cancer Research</i> , <b>2020</b> , 80, 2804-2817	10.1	21
134	Germline HOXB13 p.Gly84Glu mutation and risk of colorectal cancer. <i>Cancer Epidemiology</i> , <b>2013</b> , 37, 424-278		21

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