

# Graham Casey

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

124 papers	5,926 citations	38 h-index	75 g-index
134 ext. papers	7,629 ext. citations	8.9 avg, IF	4.21 L-index

#	Paper	IF	Citations
124	Genetically proxied therapeutic inhibition of antihypertensive drug targets and risk of common cancers: A mendelian randomization analysis.. <i>PLoS Medicine</i> , <b>2022</b> , 19, e1003897	11.6	2
123	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> , <b>2022</b> , OF1-OF13	4	0
122	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> , <b>2021</b> , 30, 564-575	4	2
121	Salicylic Acid and Risk of Colorectal Cancer: A Two-Sample Mendelian Randomization Study. <i>Nutrients</i> , <b>2021</b> , 13,	6.7	1
120	Response to Li and Hopper. <i>American Journal of Human Genetics</i> , <b>2021</b> , 108, 527-529	11	1
119	Assessment of a Polygenic Risk Score for Colorectal Cancer to Predict Risk of Lynch Syndrome Colorectal Cancer. <i>JNCI Cancer Spectrum</i> , <b>2021</b> , 5, pkab022	4.6	2
118	Circulating Levels of Testosterone, Sex Hormone Binding Globulin and Colorectal Cancer Risk: Observational and Mendelian Randomization Analyses. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2021</b> , 30, 1336-1348	4	3
117	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. <i>American Journal of Human Genetics</i> , <b>2021</b> , 108, 564-582	11	7
116	Controlling for cellular heterogeneity using single-cell deconvolution of gene expression reveals novel markers of colorectal tumors exhibiting microsatellite instability. <i>Oncotarget</i> , <b>2021</b> , 12, 767-782	3.3	3
115	Colon Crypts of Subjects With Familial Adenomatous Polyposis Show an Increased Number of LGR5+ Ectopic Stem Cells. <i>Clinical and Translational Gastroenterology</i> , <b>2021</b> , 12, e00353	4.2	0
114	Nongenetic Determinants of Risk for Early-Onset Colorectal Cancer. <i>JNCI Cancer Spectrum</i> , <b>2021</b> , 5, pkab029	4.8	15
113	Genetically Predicted Circulating C-Reactive Protein Concentration and Colorectal Cancer Survival: A Mendelian Randomization Consortium Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2021</b> , 30, 1349-1358	4	1
112	A Functional Variant on 9p21.3 Related to Glioma Risk Affects Enhancer Activity and Modulates Expression of CDKN2B-AS1. <i>Human Mutation</i> , <b>2021</b> , 42, 1208-1214	4.7	3
111	Association of Body Mass Index With Colorectal Cancer Risk by Genome-Wide Variants. <i>Journal of the National Cancer Institute</i> , <b>2021</b> , 113, 38-47	9.7	6
110	Identifying Novel Susceptibility Genes for Colorectal Cancer Risk From a Transcriptome-Wide Association Study of 125,478 Subjects. <i>Gastroenterology</i> , <b>2021</b> , 160, 1164-1178.e6	13.3	15
109	A functional variant on 20q13.33 related to glioma risk alters enhancer activity and modulates expression of multiple genes. <i>Human Mutation</i> , <b>2021</b> , 42, 77-88	4.7	6
108	Genomic Analysis of Germline Variation Associated with Survival of Patients with Colorectal Cancer Treated with Chemotherapy Plus Biologics in CALGB/SWOG 80405 (Alliance). <i>Clinical Cancer Research</i> , <b>2021</b> , 27, 267-275	12.9	2

107	Ethanol exposure drives colon location specific cell composition changes in a normal colon crypt 3D organoid model. <i>Scientific Reports</i> , <b>2021</b> , 11, 432	4.9	7
106	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. <i>Nature Genetics</i> , <b>2021</b> , 53, 65-75	36.3	62
105	Lack of an association between gallstone disease and bilirubin levels with risk of colorectal cancer: a Mendelian randomisation analysis. <i>British Journal of Cancer</i> , <b>2021</b> , 124, 1169-1174	8.7	1
104	Genetically predicted circulating concentrations of micronutrients and risk of colorectal cancer among individuals of European descent: a Mendelian randomization study. <i>American Journal of Clinical Nutrition</i> , <b>2021</b> , 113, 1490-1502	7	5
103	Genetic Effects on Transcriptome Profiles in Colon Epithelium Provide Functional Insights for Genetic Risk Loci. <i>Cellular and Molecular Gastroenterology and Hepatology</i> , <b>2021</b> , 12, 181-197	7.9	4
102	Genetic architectures of proximal and distal colorectal cancer are partly distinct. <i>Gut</i> , <b>2021</b> , 70, 1325-1334	19.2	7
101	SNPs associated with colorectal cancer at 15q13.3 affect risk enhancers that modulate GREM1 gene expression. <i>Human Mutation</i> , <b>2021</b> , 42, 237-245	4.7	5
100	Rare Variants in the DNA Repair Pathway and the Risk of Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2021</b> , 30, 895-903	4	1
99	Transcriptome-wide Effects of Aspirin on Patient-derived Normal Colon Organoids. <i>Cancer Prevention Research</i> , <b>2021</b> , 14, 1089-1100	3.2	3
98	Novel insights into the molecular mechanisms underlying risk of colorectal cancer from smoking and red/processed meat carcinogens by modeling exposure in normal colon organoids. <i>Oncotarget</i> , <b>2021</b> , 12, 1863-1877	3.3	1
97	Prioritization and functional analysis of GWAS risk loci for Barrett's esophagus and esophageal adenocarcinoma. <i>Human Molecular Genetics</i> , <b>2021</b> ,	5.6	2
96	A Germline Variant at 8q24 Contributes to Familial Clustering of Prostate Cancer in Men of African Ancestry. <i>European Urology</i> , <b>2020</b> , 78, 316-320	10.2	13
95	Mendelian Randomization of Circulating Polyunsaturated Fatty Acids and Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2020</b> , 29, 860-870	4	12
94	Functional informed genome-wide interaction analysis of body mass index, diabetes and colorectal cancer risk. <i>Cancer Medicine</i> , <b>2020</b> , 9, 3563-3573	4.8	4
93	Assessment of polygenic architecture and risk prediction based on common variants across fourteen cancers. <i>Nature Communications</i> , <b>2020</b> , 11, 3353	17.4	32
92	Modeling the effect of prolonged ethanol exposure on global gene expression and chromatin accessibility in normal 3D colon organoids. <i>PLoS ONE</i> , <b>2020</b> , 15, e0227116	3.7	12
91	A meta-analysis of genome-wide association studies of multiple myeloma among men and women of African ancestry. <i>Blood Advances</i> , <b>2020</b> , 4, 181-190	7.8	5
90	Racial Disparities in Epigenetic Aging of the Right vs Left Colon. <i>Journal of the National Cancer Institute</i> , <b>2020</b> ,	9.7	9

89	Modifiable pathways for colorectal cancer: a mendelian randomisation analysis. <i>The Lancet Gastroenterology and Hepatology</i> , <b>2020</b> , 5, 55-62	18.8	31
88	Cumulative Burden of Colorectal Cancer-Associated Genetic Variants Is More Strongly Associated With Early-Onset vs Late-Onset Cancer. <i>Gastroenterology</i> , <b>2020</b> , 158, 1274-1286.e12	13.3	47
87	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> , <b>2020</b> , 158, 1300-1312.e20	13.3	45
86	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> , <b>2020</b> , 29, 1800-1808	4	1
85	Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. <i>American Journal of Human Genetics</i> , <b>2020</b> , 107, 432-444	11	31
84	Circulating bilirubin levels and risk of colorectal cancer: serological and Mendelian randomization analyses. <i>BMC Medicine</i> , <b>2020</b> , 18, 229	11.4	11
83	Oncogenic Features in Histologically Normal Mucosa: Novel Insights Into Field Effect From a Mega-Analysis of Colorectal Transcriptomes. <i>Clinical and Translational Gastroenterology</i> , <b>2020</b> , 11, e00210	4.2	5
82	Adiposity, metabolites, and colorectal cancer risk: Mendelian randomization study. <i>BMC Medicine</i> , <b>2020</b> , 18, 396	11.4	17
81	DNA repair and cancer in colon and rectum: Novel players in genetic susceptibility. <i>International Journal of Cancer</i> , <b>2020</b> , 146, 363-372	7.5	13
80	A latent unknown clustering integrating multi-omics data (LUCID) with phenotypic traits. <i>Bioinformatics</i> , <b>2020</b> , 36, 842-850	7.2	9
79	Type 2 diabetes mellitus, blood cholesterol, triglyceride and colorectal cancer risk in Lynch syndrome. <i>British Journal of Cancer</i> , <b>2019</b> , 121, 869-876	8.7	4
78	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , <b>2019</b> , 10, 431	17.4	45
77	Ability of known susceptibility SNPs to predict colorectal cancer risk for persons with and without a family history. <i>Familial Cancer</i> , <b>2019</b> , 18, 389-397	3	17
76	Association analyses identify 31 new risk loci for colorectal cancer susceptibility. <i>Nature Communications</i> , <b>2019</b> , 10, 2154	17.4	81
75	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , <b>2019</b> , 111, 146-157	9.7	67
74	Genome-wide association study of circulating folate one-carbon metabolites. <i>Genetic Epidemiology</i> , <b>2019</b> , 43, 1030-1045	2.6	1
73	Large-Scale Genome-Wide Association Study of East Asians Identifies Loci Associated With Risk for Colorectal Cancer. <i>Gastroenterology</i> , <b>2019</b> , 156, 1455-1466	13.3	55
72	Mendelian randomization analysis of C-reactive protein on colorectal cancer risk. <i>International Journal of Epidemiology</i> , <b>2019</b> , 48, 767-780	7.8	18

71	Discovery of common and rare genetic risk variants for colorectal cancer. <i>Nature Genetics</i> , <b>2019</b> , 51, 76-83	6.3	177
70	Cohort Profile: The Colon Cancer Family Registry Cohort (CCFRC). <i>International Journal of Epidemiology</i> , <b>2018</b> , 47, 387-388	7.8	23
69	A Mixed-Effects Model for Powerful Association Tests in Integrative Functional Genomics. <i>American Journal of Human Genetics</i> , <b>2018</b> , 102, 904-919	11	20
68	Genome-wide association study and meta-analysis in Northern European populations replicate multiple colorectal cancer risk loci. <i>International Journal of Cancer</i> , <b>2018</b> , 142, 540-546	7.5	21
67	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> , <b>2018</b> , 13, e0192223	3.7	4
66	Genetic susceptibility markers for a breast-colorectal cancer phenotype: Exploratory results from genome-wide association studies. <i>PLoS ONE</i> , <b>2018</b> , 13, e0196245	3.7	2
65	Mendelian randomisation study of age at menarche and age at menopause and the risk of colorectal cancer. <i>British Journal of Cancer</i> , <b>2018</b> , 118, 1639-1647	8.7	7
64	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
63	Cross-Cancer Analysis Reveals Novel Pleiotropic Associations-Response. <i>Cancer Research</i> , <b>2017</b> , 77, 6045-6046	10.4	1
62	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
61	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
60	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
59	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2017</b> , 26, 126-135	4	183
58	Germline miRNA DNA variants and the risk of colorectal cancer by subtype. <i>Genes Chromosomes and Cancer</i> , <b>2017</b> , 56, 177-184	5	6
57	Two Novel Susceptibility Loci for Prostate Cancer in Men of African Ancestry. <i>Journal of the National Cancer Institute</i> , <b>2017</b> , 109,	9.7	38
56	Common variants in the obesity-associated genes FTO and MC4R are not associated with risk of colorectal cancer. <i>Cancer Epidemiology</i> , <b>2016</b> , 44, 1-4	2.8	9
55	Atlas of prostate cancer heritability in European and African-American men pinpoints tissue-specific regulation. <i>Nature Communications</i> , <b>2016</b> , 7, 10979	17.4	37
54	Risk of extracolonic cancers for people with biallelic and monoallelic mutations in MUTYH. <i>International Journal of Cancer</i> , <b>2016</b> , 139, 1557-63	7.5	67

53	CYP24A1 variant modifies the association between use of oestrogen plus progestogen therapy and colorectal cancer risk. <i>British Journal of Cancer</i> , <b>2016</b> , 114, 221-9	8.7	16
52	Determining the familial risk distribution of colorectal cancer: a data mining approach. <i>Familial Cancer</i> , <b>2016</b> , 15, 241-51	3	5
51	Variation at 2q35 (PNKD and TMBIM1) influences colorectal cancer risk and identifies a pleiotropic effect with inflammatory bowel disease. <i>Human Molecular Genetics</i> , <b>2016</b> , 25, 2349-2359	5.6	27
50	Prostate Cancer Susceptibility in Men of African Ancestry at 8q24. <i>Journal of the National Cancer Institute</i> , <b>2016</b> , 108,	9.7	72
49	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
48	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
47	Genetically Predicted Body Mass Index and Breast Cancer Risk: Mendelian Randomization Analyses of Data from 145,000 Women of European Descent. <i>PLoS Medicine</i> , <b>2016</b> , 13, e1002105	11.6	80
46	Fine-Mapping of Common Genetic Variants Associated with Colorectal Tumor Risk Identified Potential Functional Variants. <i>PLoS ONE</i> , <b>2016</b> , 11, e0157521	3.7	5
45	Risk factors for metachronous colorectal cancer following a primary colorectal cancer: A prospective cohort study. <i>International Journal of Cancer</i> , <b>2016</b> , 139, 1081-90	7.5	19
44	Mendelian randomisation analysis strongly implicates adiposity with risk of developing colorectal cancer. <i>British Journal of Cancer</i> , <b>2016</b> , 115, 266-72	8.7	39
43	Cholecystectomy and the risk of colorectal cancer by tumor mismatch repair deficiency status. <i>International Journal of Colorectal Disease</i> , <b>2016</b> , 31, 1451-7	3	5
42	Association of genetic susceptibility variants for type 2 diabetes with breast cancer risk in women of European ancestry. <i>Cancer Causes and Control</i> , <b>2016</b> , 27, 679-93	2.8	15
41	A Meta-analysis of Multiple Myeloma Risk Regions in African and European Ancestry Populations Identifies Putatively Functional Loci. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2016</b> , 25, 1609-1618	4	13
40	Telomere structure and maintenance gene variants and risk of five cancer types. <i>International Journal of Cancer</i> , <b>2016</b> , 139, 2655-2670	7.5	30
39	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , <b>2015</b> , 47, 373-80	36.3	406
38	Risk of colorectal cancer for people with a mutation in both a MUTYH and a DNA mismatch repair gene. <i>Familial Cancer</i> , <b>2015</b> , 14, 575-83	3	8
37	Genome-wide association study of colorectal cancer identifies six new susceptibility loci. <i>Nature Communications</i> , <b>2015</b> , 6, 7138	17.4	106
36	A new GWAS and meta-analysis with 1000Genomes imputation identifies novel risk variants for colorectal cancer. <i>Scientific Reports</i> , <b>2015</b> , 5, 10442	4.9	94



35	Integration of multiethnic fine-mapping and genomic annotation to prioritize candidate functional SNPs at prostate cancer susceptibility regions. <i>Human Molecular Genetics</i> , <b>2015</b> , 24, 5603-18	5.6	35
34	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
33	Childhood cancers in families with and without Lynch syndrome. <i>Familial Cancer</i> , <b>2015</b> , 14, 545-51	3	4
32	Association of aspirin and NSAID use with risk of colorectal cancer according to genetic variants. <i>JAMA - Journal of the American Medical Association</i> , <b>2015</b> , 313, 1133-42	27.4	135
31	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
30	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
29	Common colorectal cancer risk alleles contribute to the multiple colorectal adenoma phenotype, but do not influence colonic polyposis in FAP. <i>European Journal of Human Genetics</i> , <b>2015</b> , 23, 260-3	5.3	14
28	Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. <i>Scientific Reports</i> , <b>2015</b> , 5, 17369	4.9	27
27	Methodological Considerations in Estimation of Phenotype Heritability Using Genome-Wide SNP Data, Illustrated by an Analysis of the Heritability of Height in a Large Sample of African Ancestry Adults. <i>PLoS ONE</i> , <b>2015</b> , 10, e0131106	3.7	2
26	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
25	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
24	Identification and characterization of functional risk variants for colorectal cancer mapping to chromosome 11q23.1. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 2198-209	5.6	31
23	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
22	Large-scale genetic study in East Asians identifies six new loci associated with colorectal cancer risk. <i>Nature Genetics</i> , <b>2014</b> , 46, 533-42	36.3	175
21	Identification of susceptibility loci for colorectal cancer in a genome-wide meta-analysis. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 4729-37	5.6	107
20	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. <i>Nature Genetics</i> , <b>2014</b> , 46, 1103-9	36.3	331
19	Genome-wide scan of 29,141 African Americans finds no evidence of directional selection since admixture. <i>American Journal of Human Genetics</i> , <b>2014</b> , 95, 437-44	11	46
18	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

17	Gene-environment interaction involving recently identified colorectal cancer susceptibility Loci. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2014</b> , 23, 1824-33	4	40
16	No evidence of gene-calcium interactions from genome-wide analysis of colorectal cancer risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2014</b> , 23, 2971-6	4	9
15	Genome-wide diet-gene interaction analyses for risk of colorectal cancer. <i>PLoS Genetics</i> , <b>2014</b> , 10, e1004228	4.28	66
14	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
13	Multiple functional risk variants in a SMAD7 enhancer implicate a colorectal cancer risk haplotype. <i>PLoS ONE</i> , <b>2014</b> , 9, e111914	3.7	24
12	DCC and RET pathway analysis to identify factors associated with advanced colorectal cancer.. <i>Journal of Clinical Oncology</i> , <b>2014</b> , 32, 457-457	2.2	
11	Next generation sequencing and a new era of medicine. <i>Gut</i> , <b>2013</b> , 62, 920-32	19.2	19
10	Deciphering the genetic architecture of low-penetrance susceptibility to colorectal cancer. <i>Human Molecular Genetics</i> , <b>2013</b> , 22, 5075-82	5.6	17
9	Genome-wide association analyses in East Asians identify new susceptibility loci for colorectal cancer. <i>Nature Genetics</i> , <b>2013</b> , 45, 191-6	36.3	155
8	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
7	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
6	Common variation near CDKN1A, POLD3 and SHROOM2 influences colorectal cancer risk. <i>Nature Genetics</i> , <b>2012</b> , 44, 770-6	36.3	184
5	Principles for the post-GWAS functional characterization of cancer risk loci. <i>Nature Genetics</i> , <b>2011</b> , 43, 513-8	36.3	326
4	Multiple common susceptibility variants near BMP pathway loci GREM1, BMP4, and BMP2 explain part of the missing heritability of colorectal cancer. <i>PLoS Genetics</i> , <b>2011</b> , 7, e1002105	6	169
3	Principles for the post-GWAS functional characterisation of risk loci. <i>Nature Precedings</i> , <b>2010</b> ,		1
2	A RecQ Protein-like 5 Haplotype is Associated With Colon Cancer. <i>Gastroenterology Research</i> , <b>2010</b> , 3, 101-105	1.8	1
1	No Association of Obesity and Type 2 Diabetes Mellitus Related Genetic Variants With Colon Cancer. <i>Gastroenterology Research</i> , <b>2009</b> , 2, 311-316	1.8	