

# Gloria M Petersen

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8820344/publications.pdf>

Version: 2024-02-01

130  
papers

18,031  
citations

30070

54  
h-index

15266

126  
g-index

135  
all docs

135  
docs citations

135  
times ranked

22202  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic analyses identify molecular subtypes of pancreatic cancer. <i>Nature</i> , 2016, 531, 47-52.	27.8	2,700
2	Pancreatic cancer genomes reveal aberrations in axon guidance pathway genes. <i>Nature</i> , 2012, 491, 399-405.	27.8	1,741
3	Integrated Genomic Characterization of Pancreatic Ductal Adenocarcinoma. <i>Cancer Cell</i> , 2017, 32, 185-203.e13.	16.8	1,428
4	Very high risk of cancer in familial Peutz-Jeghers syndrome. <i>Gastroenterology</i> , 2000, 119, 1447-1453.	1.3	1,247
5	Prospective Risk of Pancreatic Cancer in Familial Pancreatic Cancer Kindreds. <i>Cancer Research</i> , 2004, 64, 2634-2638.	0.9	595
6	Germline mutations of the gene encoding bone morphogenetic protein receptor 1A in juvenile polyposis. <i>Nature Genetics</i> , 2001, 28, 184-187.	21.4	591
7	A genome-wide association study identifies pancreatic cancer susceptibility loci on chromosomes 13q22.1, 1q32.1 and 5p15.33. <i>Nature Genetics</i> , 2010, 42, 224-228.	21.4	539
8	<i>ATM</i> Mutations in Patients with Hereditary Pancreatic Cancer. <i>Cancer Discovery</i> , 2012, 2, 41-46.	9.4	442
9	Combined circulating tumor DNA and protein biomarker-based liquid biopsy for the earlier detection of pancreatic cancers. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 10202-10207.	7.1	438
10	A renewed model of pancreatic cancer evolution based on genomic rearrangement patterns. <i>Nature</i> , 2016, 538, 378-382.	27.8	418
11	Association Between Telomere Length and Risk of Cancer and Non-Neoplastic Diseases. <i>JAMA Oncology</i> , 2017, 3, 636.	7.1	376
12	Association Between Inherited Germline Mutations in Cancer Predisposition Genes and Risk of Pancreatic Cancer. <i>JAMA - Journal of the American Medical Association</i> , 2018, 319, 2401.	7.4	375
13	Diabetes, Pancreatogenic Diabetes, and Pancreatic Cancer. <i>Diabetes</i> , 2017, 66, 1103-1110.	0.6	311
14	Evaluation of candidate genes MAP2K4, MADH4, ACVR1B, and BRCA2 in familial pancreatic cancer: deleterious BRCA2 mutations in 17%. <i>Cancer Research</i> , 2002, 62, 3789-93.	0.9	308
15	Genome-wide association study identifies multiple susceptibility loci for pancreatic cancer. <i>Nature Genetics</i> , 2014, 46, 994-1000.	21.4	294
16	Whole Genome Sequencing Defines the Genetic Heterogeneity of Familial Pancreatic Cancer. <i>Cancer Discovery</i> , 2016, 6, 166-175.	9.4	282
17	Prevalence of Germline Mutations in Cancer Predisposition Genes in Patients With Pancreatic Cancer. <i>Gastroenterology</i> , 2015, 148, 556-564.	1.3	256
18	The Prevalence of BRCA2 Mutations in Familial Pancreatic Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 342-346.	2.5	255

#	ARTICLE	IF	CITATIONS
19	BRCA1, BRCA2, PALB2, and CDKN2A mutations in familial pancreatic cancer: a PACGENE study. <i>Genetics in Medicine</i> , 2015, 17, 569-577.	2.4	231
20	Common variation at 2p13.3, 3q29, 7p13 and 17q25.1 associated with susceptibility to pancreatic cancer. <i>Nature Genetics</i> , 2015, 47, 911-916.	21.4	224
21	Association of Distinct Mutational Signatures With Correlates of Increased Immune Activity in Pancreatic Ductal Adenocarcinoma. <i>JAMA Oncology</i> , 2017, 3, 774.	7.1	221
22	Detection of early pancreatic ductal adenocarcinoma with thrombospondin-2 and CA19-9 blood markers. <i>Science Translational Medicine</i> , 2017, 9, .	12.4	193
23	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. <i>Nature Communications</i> , 2018, 9, 556.	12.8	188
24	Genetics of Pancreatic Cancer. <i>Surgical Oncology Clinics of North America</i> , 1998, 7, 1-23.	1.5	170
25	Integration of Genomic and Transcriptional Features in Pancreatic Cancer Reveals Increased Cell Cycle Progression in Metastases. <i>Cancer Cell</i> , 2019, 35, 267-282.e7.	16.8	151
26	Pancreatic Cancerâ€™Derived Exosomes Cause Paraneoplastic Î²-cell Dysfunction. <i>Clinical Cancer Research</i> , 2015, 21, 1722-1733.	7.0	147
27	Pancreatic Cancer Genetic Epidemiology Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 704-710.	2.5	133
28	Family history of cancer and risk of pancreatic cancer: A pooled analysis from the Pancreatic Cancer Cohort Consortium (PanScan). <i>International Journal of Cancer</i> , 2010, 127, 1421-1428.	5.1	128
29	Evidence for a major gene influencing risk of pancreatic cancer. <i>Genetic Epidemiology</i> , 2002, 23, 133-149.	1.3	123
30	Prevalence of Pathogenic Mutations in Cancer Predisposition Genes among Pancreatic Cancer Patients. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 207-211.	2.5	116
31	Familial pancreatic cancer. <i>Seminars in Oncology</i> , 2016, 43, 548-553.	2.2	114
32	Prevalence of CDKN2A mutations in pancreatic cancer patients: implications for genetic counseling. <i>European Journal of Human Genetics</i> , 2011, 19, 472-478.	2.8	112
33	Prevalence of germ-line mutations in cancer genes among pancreatic cancer patients with a positive family history. <i>Genetics in Medicine</i> , 2018, 20, 119-127.	2.4	109
34	New DNA Methylation Markers for Pancreatic Cancer: Discovery, Tissue Validation, and Pilot Testing in Pancreatic Juice. <i>Clinical Cancer Research</i> , 2015, 21, 4473-4481.	7.0	108
35	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. <i>American Journal of Human Genetics</i> , 2015, 96, 487-497.	6.2	101
36	Transcriptional regulation by NR5A2 links differentiation and inflammation in the pancreas. <i>Nature</i> , 2018, 554, 533-537.	27.8	101

#	ARTICLE	IF	CITATIONS
37	Winner's Curse Correction and Variable Thresholding Improve Performance of Polygenic Risk Modeling Based on Genome-Wide Association Study Summary-Level Data. <i>PLoS Genetics</i> , 2016, 12, e1006493.	3.5	98
38	Alcohol intake and pancreatic cancer: a pooled analysis from the pancreatic cancer cohort consortium (PanScan). <i>Cancer Causes and Control</i> , 2010, 21, 1213-1225.	1.8	93
39	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> , 2014, 23, 6616-6633.	2.9	90
40	Targeting DNA Damage Response and Replication Stress in Pancreatic Cancer. <i>Gastroenterology</i> , 2021, 160, 362-377.e13.	1.3	90
41	Germ line Fanconi anemia complementation group C mutations and pancreatic cancer. <i>Cancer Research</i> , 2005, 65, 383-6.	0.9	89
42	Three new pancreatic cancer susceptibility signals identified on chromosomes 1q32.1, 5p15.33 and 8q24.21. <i>Oncotarget</i> , 2016, 7, 66328-66343.	1.8	88
43	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016, 7, 11843.	12.8	86
44	Cystic fibrosis transmembrane conductance regulator ( <i>CFTR</i> ) gene mutations and risk for pancreatic adenocarcinoma. <i>Cancer</i> , 2010, 116, 203-209.	4.1	80
45	Risk of malignancy in first-degree relatives of patients with pancreatic carcinoma. <i>Cancer</i> , 2005, 104, 388-394.	4.1	78
46	Exposure to environmental chemicals and heavy metals, and risk of pancreatic cancer. <i>Cancer Causes and Control</i> , 2015, 26, 1583-1591.	1.8	78
47	Fruit and vegetable consumption is inversely associated with having pancreatic cancer. <i>Cancer Causes and Control</i> , 2011, 22, 1613-1625.	1.8	75
48	Assessment of polygenic architecture and risk prediction based on common variants across fourteen cancers. <i>Nature Communications</i> , 2020, 11, 3353.	12.8	75
49	Mitochondrial Genetic Polymorphisms and Pancreatic Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 1455-1459.	2.5	74
50	Metformin Use and Survival of Patients With Pancreatic Cancer: A Cautionary Lesson. <i>Journal of Clinical Oncology</i> , 2016, 34, 1898-1904.	1.6	69
51	Mutations in the pancreatic secretory enzymes <i>CPA1</i> and <i>CPB1</i> are associated with pancreatic cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 4767-4772.	7.1	65
52	Immunosuppressive CD14 <sup>+</sup> HLA-DR <sup>lo/neg</sup> monocytes are elevated in pancreatic cancer and are primed by tumor-derived exosomes. <i>Onc Immunology</i> , 2017, 6, e1252013.	4.6	59
53	A Transcriptome-Wide Association Study Identifies Novel Candidate Susceptibility Genes for Pancreatic Cancer. <i>Journal of the National Cancer Institute</i> , 2020, 112, 1003-1012.	6.3	59
54	EUS-guided fine-needle injection of gemcitabine for locally advanced and metastatic pancreatic cancer. <i>Gastrointestinal Endoscopy</i> , 2017, 86, 161-169.	1.0	58

#	ARTICLE	IF	CITATIONS
55	<i>TERT</i> gene harbors multiple variants associated with pancreatic cancer susceptibility. <i>International Journal of Cancer</i> , 2015, 137, 2175-2183.	5.1	57
56	Telomere Length and Pancreatic Cancer: A Case-Control Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 2095-2100.	2.5	51
57	CLPTM1L Promotes Growth and Enhances Aneuploidy in Pancreatic Cancer Cells. <i>Cancer Research</i> , 2014, 74, 2785-2795.	0.9	48
58	Quantifying the Genetic Correlation between Multiple Cancer Types. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 1427-1435.	2.5	48
59	Analysis of Heritability and Genetic Architecture of Pancreatic Cancer: A PanC4 Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 1238-1245.	2.5	48
60	Candidate DNA repair susceptibility genes identified by exome sequencing in high-risk pancreatic cancer. <i>Cancer Letters</i> , 2016, 370, 302-312.	7.2	47
61	Transcriptome analysis of pancreatic cancer reveals a tumor suppressor function for HNF1A. <i>Carcinogenesis</i> , 2014, 35, 2670-2678.	2.8	46
62	Familial Pancreatic Adenocarcinoma. <i>Hematology/Oncology Clinics of North America</i> , 2015, 29, 641-653.	2.2	46
63	Functional characterization of a multi-cancer risk locus on chr5p15.33 reveals regulation of TERT by ZNF148. <i>Nature Communications</i> , 2017, 8, 15034.	12.8	40
64	Decreased Skeletal Muscle Volume Is a Predictive Factor for Poorer Survival in Patients Undergoing Surgical Resection for Pancreatic Ductal Adenocarcinoma. <i>Journal of Gastrointestinal Surgery</i> , 2018, 22, 831-839.	1.7	40
65	Methylated DNA in Pancreatic Juice Distinguishes Patients With Pancreatic Cancer From Controls. <i>Clinical Gastroenterology and Hepatology</i> , 2020, 18, 676-683.e3.	4.4	40
66	Quantitative Proteomics Based on Optimized Data-Independent Acquisition in Plasma Analysis. <i>Journal of Proteome Research</i> , 2017, 16, 665-676.	3.7	39
67	Genes-Environment Interactions in Obesity- and Diabetes-Associated Pancreatic Cancer: A GWAS Data Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 98-106.	2.5	32
68	Vitamin D Metabolic Pathway Genes and Pancreatic Cancer Risk. <i>PLoS ONE</i> , 2015, 10, e0117574.	2.5	29
69	Association of Common Susceptibility Variants of Pancreatic Cancer in Higher-Risk Patients: A PACGENE Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1185-1191.	2.5	29
70	Integrated Genomic Analysis of Pancreatic Ductal Adenocarcinomas Reveals Genomic Rearrangement Events as Significant Drivers of Disease. <i>Cancer Research</i> , 2016, 76, 749-761.	0.9	27
71	Impact of Diabetes Mellitus on Clinical Outcomes in Patients Undergoing Surgical Resection for Pancreatic Cancer: A Retrospective, Cohort Study. <i>American Journal of Gastroenterology</i> , 2014, 109, 1484-1492.	0.4	26
72	Characterising <i>cis</i> -regulatory variation in the transcriptome of histologically normal and tumour-derived pancreatic tissues. <i>Gut</i> , 2018, 67, 521-533.	12.1	26

#	ARTICLE	IF	CITATIONS
73	Barriers to preventive intervention. <i>Gastroenterology Clinics of North America</i> , 2002, 31, 1061-1068.	2.2	25
74	Differential and Joint Effects of Metformin and Statins on Overall Survival of Elderly Patients with Pancreatic Adenocarcinoma: A Large Population-Based Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 1225-1232.	2.5	25
75	Functional characterization of a chr13q22.1 pancreatic cancer risk locus reveals long-range interaction and allele-specific effects on <i>DIS3</i> expression. <i>Human Molecular Genetics</i> , 2016, 25, ddw300.	2.9	24
76	Effect of Germline Mutations in Homologous Recombination Repair Genes on Overall Survival of Patients with Pancreatic Adenocarcinoma. <i>Clinical Cancer Research</i> , 2020, 26, 6505-6512.	7.0	24
77	The Association of Telomere Length with Colorectal Cancer Differs by the Age of Cancer Onset. <i>Clinical and Translational Gastroenterology</i> , 2014, 5, e52.	2.5	23
78	Case-only exome sequencing and complex disease susceptibility gene discovery: study design considerations. <i>Journal of Medical Genetics</i> , 2015, 52, 10-16.	3.2	23
79	Pancreatic cancer risk is modulated by inflammatory potential of diet and ABO genotype: a consortia-based evaluation and replication study. <i>Carcinogenesis</i> , 2018, 39, 1056-1067.	2.8	23
80	<i>CDKN2A</i> Germline Rare Coding Variants and Risk of Pancreatic Cancer in Minority Populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 1364-1370.	2.5	23
81	Inactivation of the Transcription Factor <i>GLI1</i> Accelerates Pancreatic Cancer Progression. <i>Journal of Biological Chemistry</i> , 2014, 289, 16516-16525.	3.4	22
82	Zinc transporter genes and urological cancers: integrated analysis suggests a role for <i>ZIP11</i> in bladder cancer. <i>Tumor Biology</i> , 2015, 36, 7431-7437.	1.8	22
83	Agnostic Pathway/Gene Set Analysis of Genome-Wide Association Data Identifies Associations for Pancreatic Cancer. <i>Journal of the National Cancer Institute</i> , 2019, 111, 557-567.	6.3	21
84	Systemic Proteome Alterations Linked to Early Stage Pancreatic Cancer in Diabetic Patients. <i>Cancers</i> , 2020, 12, 1534.	3.7	18
85	Associations between Genetically Predicted Blood Protein Biomarkers and Pancreatic Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1501-1508.	2.5	18
86	Factors influencing receptivity to future screening options for pancreatic cancer in those with and without pancreatic cancer family history. <i>Hereditary Cancer in Clinical Practice</i> , 2012, 10, 8.	1.5	17
87	A New Strategy to Control and Eradicate "Undruggable" Oncogenic K-RAS-Driven Pancreatic Cancer: Molecular Insights and Core Principles Learned from Developmental and Evolutionary Biology. <i>Cancers</i> , 2018, 10, 142.	3.7	17
88	Should Researchers Offer Results to Family Members of Cancer Biobank Participants? A Mixed-Methods Study of Proband and Family Preferences. <i>AJOB Empirical Bioethics</i> , 2019, 10, 1-22.	1.6	17
89	High Detection Rates of Pancreatic Cancer Across Stages by Plasma Assay of Novel Methylated DNA Markers and CA19-9. <i>Clinical Cancer Research</i> , 2021, 27, 2523-2532.	7.0	17
90	A multilayered post-GWAS assessment on genetic susceptibility to pancreatic cancer. <i>Genome Medicine</i> , 2021, 13, 15.	8.2	15

#	ARTICLE	IF	CITATIONS
91	A 584Åbp deletion in CTRB2 inhibits chymotrypsin B2 activity and secretion and confers risk of pancreatic cancer. <i>American Journal of Human Genetics</i> , 2021, 108, 1852-1865.	6.2	15
92	Variants Associated with Susceptibility to Pancreatic Cancer and Melanoma Do Not Reciprocally Affect Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1121-1124.	2.5	14
93	Do Variants Associated with Susceptibility to Pancreatic Cancer and Type 2 Diabetes Reciprocally Affect Risk?. <i>PLoS ONE</i> , 2015, 10, e0117230.	2.5	14
94	Meat-Related Mutagens and Pancreatic Cancer: Null Results from a Clinic-Based Caseâ€“Control Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 1336-1339.	2.5	13
95	A region-based gene association study combined with a leave-one-out sensitivity analysis identifies SMG1 as a pancreatic cancer susceptibility gene. <i>PLoS Genetics</i> , 2019, 15, e1008344.	3.5	13
96	THBS2/CA19-9 Detecting Pancreatic Ductal Adenocarcinoma at Diagnosis Underperforms in Prediagnostic Detection: Implications for Biomarker Advancement. <i>Cancer Prevention Research</i> , 2021, 14, 223-232.	1.5	13
97	Carcinogenesis of pancreatic cancer: Challenges, collaborations, progress. <i>Molecular Carcinogenesis</i> , 2012, 51, 1-2.	2.7	11
98	Receptivity and preferences of pancreatic cancer family members for participating in lifestyle programs to reduce cancer risk. <i>Hereditary Cancer in Clinical Practice</i> , 2013, 11, 3.	1.5	11
99	Genetically Predicted Telomere Length is not Associated with Pancreatic Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 971-974.	2.5	11
100	Attitudes Toward Return of Genetic Research Results to Relatives, Including After Death: Comparison of Cancer Probands, Blood Relatives, and Spouse/Partners. <i>Journal of Empirical Research on Human Research Ethics</i> , 2018, 13, 295-304.	1.3	11
101	Risk of Different Cancers Among First-degree Relatives of Pancreatic Cancer Patients: Influence of Probandsâ€™ Susceptibility Gene Mutation Status. <i>Journal of the National Cancer Institute</i> , 2019, 111, 264-271.	6.3	10
102	Gene-by-Environment Interactions in Pancreatic Cancer: Implications for Prevention. <i>Yale Journal of Biology and Medicine</i> , 2015, 88, 115-26.	0.2	10
103	Association between Alcohol Consumption, Folate Intake, and Risk of Pancreatic Cancer: A Case-Control Study. <i>Nutrients</i> , 2017, 9, 0448.	4.1	9
104	Leukocyte Telomere Length and Pancreatic Cancer Risk. <i>Pancreas</i> , 2018, 47, 265-271.	1.1	9
105	Hepcidin-regulating iron metabolism genes and pancreatic ductal adenocarcinoma: a pathway analysis of genome-wide association studies. <i>American Journal of Clinical Nutrition</i> , 2021, 114, 1408-1417.	4.7	9
106	Smoking Modifies Pancreatic Cancer Risk Loci on 2q21.3. <i>Cancer Research</i> , 2021, 81, 3134-3143.	0.9	8
107	Psychological Impact of Learning & CDKN2A Variant Status as a Genetic Research Result. <i>Public Health Genomics</i> , 2018, 21, 154-163.	1.0	7
108	Pancreatic cancer and melanoma related perceptions and behaviors following disclosure of CDKN2A variant status as a research result. <i>Genetics in Medicine</i> , 2019, 21, 2468-2477.	2.4	6

#	ARTICLE	IF	CITATIONS
109	Mendelian Randomization Analysis of n-6 Polyunsaturated Fatty Acid Levels and Pancreatic Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 2735-2739.	2.5	6
110	Detection of DNA damage in peripheral blood mononuclear cells from pancreatic cancer patients. <i>Molecular Carcinogenesis</i> , 2015, 54, 1220-1226.	2.7	5
111	Role of Surgery and Perioperative Therapy in Older Patients with Resectable Pancreatic Ductal Adenocarcinoma. <i>Oncologist</i> , 2020, 25, e1681-e1690.	3.7	5
112	Survival Benefit of Combination Chemotherapy in Elderly Patients With Metastatic Pancreatic Ductal Adenocarcinoma. <i>American Journal of Clinical Oncology: Cancer Clinical Trials</i> , 2020, 43, 586-590.	1.3	5
113	Genome-Wide Gene-€Diabetes and Gene-€Obesity Interaction Scan in 8,255 Cases and 11,900 Controls from PanScan and PanC4 Consortia. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1784-1791.	2.5	5
114	Genome-Wide Association Study Data Reveal Genetic Susceptibility to Chronic Inflammatory Intestinal Diseases and Pancreatic Ductal Adenocarcinoma Risk. <i>Cancer Research</i> , 2020, 80, 4004-4013.	0.9	5
115	Leukocyte Telomere Length and Its Interaction with Germline Variation in Telomere-Related Genes in Relation to Pancreatic Adenocarcinoma Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1492-1500.	2.5	5
116	Aspirin, Statins, Non-aspirin NSAIDs, Metformin, and the Risk of Biliary Cancer: A Swedish Population-Based Cohort Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 804-810.	2.5	5
117	Intact SMAD-4 is a predictor of increased locoregional recurrence in upfront resected pancreas cancer receiving adjuvant therapy. <i>Journal of Gastrointestinal Oncology</i> , 2021, 12, 2275-2286.	1.4	4
118	A risk prediction tool for individuals with a family history of breast, ovarian, or pancreatic cancer: BRCAPANPRO. <i>British Journal of Cancer</i> , 2021, 125, 1712-1717.	6.4	4
119	Pancreatic cancer risk to siblings of probands in bilineal cancer settings. <i>Genetics in Medicine</i> , 2022, 24, 1008-1016.	2.4	4
120	Influence of Cancer Susceptibility Gene Mutations and ABO Blood Group of Pancreatic Cancer Proband on Concomitant Risk to First-Degree Relatives. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 372-381.	2.5	3
121	A rare germline CDKN2A variant (47T>G; p16-L16R) predisposes carriers to pancreatic cancer by reducing cell cycle inhibition. <i>Journal of Biological Chemistry</i> , 2021, 296, 100634.	3.4	2
122	Shorter Treatment-Na-ve Leukocyte Telomere Length is Associated with Poorer Overall Survival of Patients with Pancreatic Ductal Adenocarcinoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 210-216.	2.5	2
123	Intercepting Pancreatic Cancer. <i>Pancreas</i> , 2018, 47, 1175-1176.	1.1	1
124	Gallbladder disease, cholecystectomy, and pancreatic cancer risk in the International Pancreatic Cancer Case-Control Consortium (PanC4). <i>European Journal of Cancer Prevention</i> , 2020, 29, 408-415.	1.3	1
125	Accuracy of Smoking Status Reporting. <i>Mayo Clinic Proceedings Innovations, Quality &amp; Outcomes</i> , 2020, 4, 801-809.	2.4	1
126	Telomere Length and Pancreatic Cancer Risk-Reply. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 1158-1159.	2.5	0



#	ARTICLE	IF	CITATIONS
127	A Pathway Analysis of Hereditary Hemochromatosis-related Genes and Pancreatic Ductal Adenocarcinoma Risk (FS11-05-19). <i>Current Developments in Nutrition</i> , 2019, 3, nzz037.FS11-05-19.	0.3	0
128	Bayesian copy number detection and association in large-scale studies. <i>BMC Cancer</i> , 2020, 20, 856.	2.6	0
129	Abstract A26: Attacking the most downstream "gatekeeper," the SIAH-dependent proteolytic machinery, in the oncogenic ERBB/K-RAS signaling pathway to block tumorigenesis and control metastasis in human cancer. , 2014, , .		0
130	A Pilot Study of Blood-Based Methylation Markers Associated With Pancreatic Cancer. <i>Frontiers in Genetics</i> , 2022, 13, 849839.	2.3	0