Alison P Klein

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.

16,762 48 129 133 h-index g-index citations papers 6.07 8.9 19,796 141 avg, IF L-index ext. citations ext. papers

#	Paper	IF	Citations
133	Endoplasmic stress-inducing variants in CPB1 and CPA1 and risk of pancreatic cancer: A case-control study and meta-analysis. <i>International Journal of Cancer</i> , 2021 , 150, 1123	7.5	O
132	Novel Models of Genetic Education and Testing for Pancreatic Cancer Interception: Preliminary Results from the GENERATE Study. <i>Cancer Prevention Research</i> , 2021 , 14, 1021-1032	3.2	3
131	A risk prediction tool for individuals with a family history of breast, ovarian, or pancreatic cancer: BRCAPANCPRO. <i>British Journal of Cancer</i> , 2021 , 125, 1712-1717	8.7	1
130	The Role of Inherited Pathogenic CDKN2A Variants in Susceptibility to Pancreatic Cancer. <i>Pancreas</i> , 2021 , 50, 1123-1130	2.6	1
129	Two-Sample Mendelian Randomization Analysis of Associations Between Periodontal Disease and Risk of Cancer. <i>JNCI Cancer Spectrum</i> , 2021 , 5, pkab037	4.6	2
128	Germline sequence analysis of RABL3 in a large series of pancreatic ductal adenocarcinoma patients reveals no evidence of deleterious variants. <i>Genes Chromosomes and Cancer</i> , 2021 , 60, 559-564	5	1
127	Pancreatic cancer epidemiology: understanding the role of lifestyle and inherited risk factors. <i>Nature Reviews Gastroenterology and Hepatology</i> , 2021 , 18, 493-502	24.2	39
126	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, 1408-141	7	2
125	A pooled genome-wide association study identifies pancreatic cancer susceptibility loci on chromosome 19p12 and 19p13.3 in the full-Jewish population. <i>Human Genetics</i> , 2021 , 140, 309-319	6.3	2
124	Challenges of the current precision medicine approach for pancreatic cancer: A single institution experience between 2013 and 2017. <i>Cancer Letters</i> , 2021 , 497, 221-228	9.9	7
123	A multilayered post-GWAS assessment on genetic susceptibility to pancreatic cancer. <i>Genome Medicine</i> , 2021 , 13, 15	14.4	6
122	Smoking Modifies Pancreatic Cancer Risk Loci on 2q21.3. <i>Cancer Research</i> , 2021 , 81, 3134-3143	10.1	2
121	Pancreatic cancer pathology viewed in the light of evolution. <i>Cancer and Metastasis Reviews</i> , 2021 , 40, 661-674	9.6	3
120	Familial pancreatic cancer: who should be considered for genetic testing?. <i>Irish Journal of Medical Science</i> , 2021 , 1	1.9	2
119	Examination of ATM, BRCA1, and BRCA2 promoter methylation in patients with pancreatic cancer. <i>Pancreatology</i> , 2021 , 21, 938-941	3.8	O
118	A 584lbp 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	11	1
117	Risk of Pancreatic Cancer Among Individuals With Pathogenic Variants in the ATM Gene. <i>JAMA Oncology</i> , 2021 , 7, 1664-1668	13.4	7

(2019-2020)

General Population. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 999-1008	4	7
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	4	4
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	10.1	1
Assessment of polygenic architecture and risk prediction based on common variants across fourteen cancers. <i>Nature Communications</i> , 2020 , 11, 3353	17.4	32
Recent Trends in the Incidence and Survival of Stage 1A Pancreatic Cancer: A Surveillance, Epidemiology, and End Results Analysis. <i>Journal of the National Cancer Institute</i> , 2020 , 112, 1162-1169	9.7	48
Feasibility of blood testing combined with PET-CT to screen for cancer and guide intervention. <i>Science</i> , 2020 , 369,	33.3	149
Detection of Circulating Tumor DNA in Patients with Pancreatic Cancer Using Digital Next-Generation Sequencing. <i>Journal of Molecular Diagnostics</i> , 2020 , 22, 748-756	5.1	4
Associations between Genetically Predicted Blood Protein Biomarkers and Pancreatic Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020 , 29, 1501-1508	4	9
Gene Variants That Affect Levels of Circulating Tumor Markers Increase Identification of Patients With Pancreatic Cancer. <i>Clinical Gastroenterology and Hepatology</i> , 2020 , 18, 1161-1169.e5	6.9	10
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	9.7	25
Molecular characterization of organoids derived from pancreatic intraductal papillary mucinous neoplasms. <i>Journal of Pathology</i> , 2020 , 252, 252-262	9.4	18
The genetics of ductal adenocarcinoma of the pancreas in the year 2020: dramatic progress, but far to go. <i>Modern Pathology</i> , 2020 , 33, 2544-2563	9.8	9
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	4	2
Bayesian copy number detection and association in large-scale studies. <i>BMC Cancer</i> , 2020 , 20, 856	4.8	
Surgical Outcomes After Pancreatic Resection of Screening-Detected Lesions in Individuals at High Risk for Developing Pancreatic Cancer. <i>Journal of Gastrointestinal Surgery</i> , 2020 , 24, 1101-1110	3.3	26
Pancreatic cancer: a growing burden. <i>The Lancet Gastroenterology and Hepatology</i> , 2019 , 4, 895-896	18.8	11
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	6	7
Analysis of Heritability and Genetic Architecture of Pancreatic Cancer: A PanC4 Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019 , 28, 1238-1245	4	27
	Genome-Wide Gene-Diabetes and Gene-Obesity Interaction Scan in 8,255 Cases and 11,900 Controls from PanScan and PanC4 Consortia. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1784-1791 Genome-Wide Association Study Data Reveal Genetic Susceptibility to Chronic Inflammatory Intestinal Diseases and Pancreatic Ductal Adenocarcinoma Risk. Cancer Research, 2020, 80, 4004-4013 Assessment of polygenic architecture and risk prediction based on common variants across fourteen cancers. Nature Communications, 2020, 11, 3353 Recent Trends in the Incidence and Survival of Stage 1A Pancreatic Cancer: A Surveillance, Epidemiology, and End Results Analysis. Journal of the National Cancer Institute, 2020, 112, 1162-1169 Feasibility of blood testing combined with PET-CT to screen for cancer and guide intervention. Science, 2020, 369, Detection of Circulating Tumor DNA in Patients with Pancreatic Cancer Using Digital Next-Generation Sequencing. Journal of Molecular Diagnostics, 2020, 22, 748-756 Associations between Genetically Predicted Blood Protein Biomarkers and Pancreatic Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1501-1508 Gene Variants That Affect Levels of Circulating Tumor Markers Increase Identification of Patients With Pancreatic Cancer. Clinical Gastroenterology and Hepatology, 2020, 18, 1161-1169-e5 A Transcriptome-Wide Association Study Identifies Novel Candidate Susceptibility Genes for Pancreatic Cancer. Journal of the National Cancer Institute, 2020, 112, 1003-1012 Molecular characterization of organoids derived from pancreatic intraductal papillary mucinous neoplasms. Journal of Pathology, 2020, 252, 252-262 The genetics of ductal adenocarcinoma of the pancreas in the year 2020: dramatic progress, but far to go. Modern Pathology, 2020, 33, 2544-2563 Mendelian Randomization Analysis of n-6 Polyunsaturated Fatty Acid Levels and Pancreatic Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 2735-2739 Bayesian copy number detection and association i	Genome-Wide Gene-Diabetes and Gene-Obesity Interaction Scan in 9,255 Cases and 11,900 Controls from PanScan and PanC4 Consortia. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1784-1791 Genome-Wide Association Study Data Reveal Genetic Susceptibility to Chronic Infiammatory Intestinal Diseases and Pancreatic Ductal Adenocarcinoma Risk. Cancer Research, 2020, 80, 4004-4013 Assessment of polygenic architecture and risk prediction based on common variants across fourteen cancers. Nature Communications, 2020, 11, 3353 7.44 Recent Trends in the Incidence and Survival of Stage 1A Pancreatic Cancer: A Surveillance, Epidemiology, and End Results Analysis. Journal of the National Cancer Institute, 2020, 112, 1162-1169 Peasibility of blood testing combined with PET-CT to screen for cancer and guide intervention. Science, 2020, 369, Detection of Circulating Tumor DNA in Patients with Pancreatic Cancer: Using Digital Next-Generation Sequencing. Journal of Molecular Diagnostics, 2020, 22, 748-756 Associations between Genetically Predicted Blood Protein Biomarkers and Pancreatic Cancer Risk. Acancer Epidemiology Biomarkers and Prevention, 2020, 29, 1501-1508 4 Gene Variants That Affect Levels of Circulating Tumor Markers Increase Identification of Patients With Pancreatic Cancer. Clinical Gastroenterology and Hepatology, 2020, 18, 1161-1169,e5 A Transcriptome-Wide Association Study Identifies Novel Candidate Susceptibility Genes for Pancreatic Cancer. Journal of the National Cancer Institute, 2020, 112, 1003-1012 Molecular characterization of organoids derived from pancreatic intraductal papillary mucinous neoplasms. Journal of Pathology, 2020, 252, 252-262 The genetics of ductal adenocarcinoma of the pancreas in the year 2020: dramatic progress, but far to go. Modern Pathology, 2020, 33, 2544-2563 Mendelian Randomization Analysis of n-6 Polyunsaturated Fatty Acid Levels and Pancreatic Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 2735-2739 Bayesian copy number detection and asso

98	A New Fast Phasing Method Based On Haplotype Subtraction. <i>Journal of Molecular Diagnostics</i> , 2019 , 21, 427-436	5.1	
97	Prevalence of Germline Mutations Associated With Cancer Risk in Patients With Intraductal Papillary Mucinous Neoplasms. <i>Gastroenterology</i> , 2019 , 156, 1905-1913	13.3	27
96	Deleterious Germline Mutations Are a Risk Factor for Neoplastic Progression Among High-Risk Individuals Undergoing Pancreatic Surveillance. <i>Journal of Clinical Oncology</i> , 2019 , 37, 1070-1080	2.2	36
95	Screening for Pancreatic Cancer-Is There Hope?. <i>JAMA Internal Medicine</i> , 2019 , 179, 1313-1315	11.5	6
94	Histomorphology of pancreatic cancer in patients with inherited ATM serine/threonine kinase pathogenic variants. <i>Modern Pathology</i> , 2019 , 32, 1806-1813	9.8	11
93	A multimodality test to guide the management of patients with a pancreatic cyst. <i>Science Translational Medicine</i> , 2019 , 11,	17.5	71
92	A Pathway Analysis of Hereditary Hemochromatosis-related Genes and Pancreatic Ductal Adenocarcinoma Risk (FS11-05-19). <i>Current Developments in Nutrition</i> , 2019 , 3,	0.4	78
91	Abstract 1591: Large-scale transcriptome-wide association study (TWAS) identifies novel candidate susceptibility genes for pancreatic cancer 2019 ,		3
90	Abstract 1591: Large-scale transcriptome-wide association study (TWAS) identifies novel candidate susceptibility genes for pancreatic cancer 2019 ,		3
89	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	9.7	16
88	Determinants and prognostic value of quality of life in patients with pancreatic ductal adenocarcinoma. <i>European Journal of Cancer</i> , 2018 , 92, 20-32	7.5	14
87	Detection and localization of surgically resectable cancers with a multi-analyte blood test. <i>Science</i> , 2018 , 359, 926-930	33.3	1204
86	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. <i>Nature Communications</i> , 2018 , 9, 556	17.4	103
85	Epidemiology of Pancreatic Cancer 2018 , 665-672		
84	Familial Pancreatic Cancer 2018 , 688-692		
83	Familial Pancreatic Cancer 2018 , 553-572		1
82	Long-term analysis of 2 prospective studies that incorporate mitomycin C into an adjuvant chemoradiation regimen for pancreatic and periampullary cancers. <i>Advances in Radiation Oncology</i> , 2018 , 3, 42-51	3.3	1
81	Exome-Wide Association Study of Pancreatic Cancer Risk. <i>Gastroenterology</i> , 2018 , 154, 719-722.e3	13.3	27

80	Exome Array Analysis of Nuclear Lens Opacity. <i>Ophthalmic Epidemiology</i> , 2018 , 25, 215-219	1.9	1
79	Genomic analysis identifies frequent deletions of Dystrophin in olfactory neuroblastoma. <i>Nature Communications</i> , 2018 , 9, 5410	17.4	14
78	Refraction and Change in Refraction Over a 20-Year Period in the Beaver Dam Eye Study 2018 , 59, 4518	3-4524	8
77	Association analysis of exome variants and refraction, axial length, and corneal curvature in a European-American population. <i>Human Mutation</i> , 2018 , 39, 1973-1979	4.7	2
76	Risk of Neoplastic Progression in Individuals at High Risk for Pancreatic Cancer Undergoing Long-term Surveillance. <i>Gastroenterology</i> , 2018 , 155, 740-751.e2	13.3	154
75	Association Between Telomere Length and Risk of Cancer and Non-Neoplastic Diseases: A Mendelian Randomization Study. <i>JAMA Oncology</i> , 2017 , 3, 636-651	13.4	236
74	Alterations of type II classical cadherin, cadherin-10 (CDH10), is associated with pancreatic ductal adenocarcinomas. <i>Genes Chromosomes and Cancer</i> , 2017 , 56, 427-435	5	6
73	Haplotype Counting for Sensitive Chimerism Testing: Potential for Early Leukemia Relapse Detection. <i>Journal of Molecular Diagnostics</i> , 2017 , 19, 427-436	5.1	9
72	Deleterious Germline Mutations in Patients With Apparently Sporadic Pancreatic Adenocarcinoma. Journal of Clinical Oncology, 2017 , 35, 3382-3390	2.2	207
71	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	11.5	303
70	Impact of Sixteen Established Pancreatic Cancer Susceptibility Loci in American Jews. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017 , 26, 1540-1548	4	5
69	Quantifying the Genetic Correlation between Multiple Cancer Types. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017 , 26, 1427-1435	4	25
68	Inherited pancreatic cancer. Chinese Clinical Oncology, 2017, 6, 58	2.3	15
67	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016 , 7, 11843	17.4	59
66	Whole Genome Sequencing Defines the Genetic Heterogeneity of Familial Pancreatic Cancer. <i>Cancer Discovery</i> , 2016 , 6, 166-75	24.4	206
65	Using Quantitative Seroproteomics to Identify Antibody Biomarkers in Pancreatic Cancer. <i>Cancer Immunology Research</i> , 2016 , 4, 225-33	12.5	19
64	Winner& 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, e10	06493	67
63	Variation in PTCHD2, CRISP3, NAP1L4, FSCB, and AP3B2 associated with spherical equivalent. <i>Molecular Vision</i> , 2016 , 22, 783-96	2.3	5

62	Three new pancreatic cancer susceptibility signals identified on chromosomes 1q32.1, 5p15.33 and 8q24.21. <i>Oncotarget</i> , 2016 , 7, 66328-66343	3.3	66
61	Familial Pancreatic Cancer 2016 , 1-20		
60	Association of Common Susceptibility Variants of Pancreatic Cancer in Higher-Risk Patients: A PACGENE Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016 , 25, 1185-91	4	22
59	Functional characterization of a chr13q22.1 pancreatic cancer risk locus reveals long-range interaction and allele-specific effects on DIS3 expression. <i>Human Molecular Genetics</i> , 2016 , 25, 4726-47	73 § .6	17
58	Family history as a marker of platinum sensitivity in pancreatic adenocarcinoma. <i>Cancer Chemotherapy and Pharmacology</i> , 2015 , 76, 489-498	3.5	45
57	Common variation at 2p13.3, 3q29, 7p13 and 17q25.1 associated with susceptibility to pancreatic cancer. <i>Nature Genetics</i> , 2015 , 47, 911-6	36.3	171
56	Intraductal papillary mucinous neoplasm in a neonate with congenital hyperinsulinism and a de novo germline SKIL gene mutation. <i>Pancreatology</i> , 2015 , 15, 194-6	3.8	5
55	BRCA1, BRCA2, PALB2, and CDKN2A mutations in familial pancreatic cancer: a PACGENE study. <i>Genetics in Medicine</i> , 2015 , 17, 569-77	8.1	175
54	Transflip mutations produce deletions in pancreatic cancer. <i>Genes Chromosomes and Cancer</i> , 2015 , 54, 472-481	5	6
53	TERT gene harbors multiple variants associated with pancreatic cancer susceptibility. <i>International Journal of Cancer</i> , 2015 , 137, 2175-83	7.5	46
52	Analysis of Heritability and Shared Heritability Based on Genome-Wide Association Studies for Thirteen Cancer Types. <i>Journal of the National Cancer Institute</i> , 2015 , 107, djv279	9.7	107
51	A histomorphologic comparison of familial and sporadic pancreatic cancers. <i>Pancreatology</i> , 2015 , 15, 387-391	3.8	22
50	Familial and sporadic pancreatic cancer share the same molecular pathogenesis. <i>Familial Cancer</i> , 2015 , 14, 95-103	3	43
49	Genome-wide association study identifies multiple susceptibility loci for pancreatic cancer. <i>Nature Genetics</i> , 2014 , 46, 994-1000	36.3	226
48	The early detection of pancreatic cancer: what will it take to diagnose and treat curable pancreatic neoplasia?. <i>Cancer Research</i> , 2014 , 74, 3381-9	10.1	162
47	Association of PD-1, PD-1 ligands, and other features of the tumor immune microenvironment with response to anti-PD-1 therapy. <i>Clinical Cancer Research</i> , 2014 , 20, 5064-74	12.9	1661
46	Exome array analysis identifies CAV1/CAV2 as a susceptibility locus for intraocular pressure. <i>Investigative Ophthalmology and Visual Science</i> , 2014 , 56, 544-51		37
45	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, 6616-33	5.6	77

(2009-2014)

44	Having pancreatic cancer with tumoral loss of ATM and normal TP53 protein expression is associated with a poorer prognosis. <i>Clinical Cancer Research</i> , 2014 , 20, 1865-72	12.9	61
43	Identifying people at a high risk of developing pancreatic cancer. <i>Nature Reviews Cancer</i> , 2013 , 13, 66-74	431.3	108
42	An absolute risk model to identify individuals at elevated risk for pancreatic cancer in the general population. <i>PLoS ONE</i> , 2013 , 8, e72311	3.7	82
41	Genetic susceptibility to pancreatic cancer. <i>Molecular Carcinogenesis</i> , 2012 , 51, 14-24	5	150
40	Frequent detection of pancreatic lesions in asymptomatic high-risk individuals. <i>Gastroenterology</i> , 2012 , 142, 796-804; quiz e14-5	13.3	455
39	ATM mutations in patients with hereditary pancreatic cancer. Cancer Discovery, 2012, 2, 41-6	24.4	365
38	Pathway analysis of genome-wide association study data highlights pancreatic development genes as susceptibility factors for pancreatic cancer. <i>Carcinogenesis</i> , 2012 , 33, 1384-90	4.6	85
37	Clinical significance of the genetic landscape of pancreatic cancer and implications for identification of potential long-term survivors. <i>Clinical Cancer Research</i> , 2012 , 18, 6339-47	12.9	163
36	Identification of functional genetic variation in exome sequence analysis. <i>BMC Proceedings</i> , 2011 , 5 Suppl 9, S13	2.3	7
35	Linkage analysis of quantitative refraction and refractive errors in the Beaver Dam Eye Study 2011 , 52, 5220-5		15
35		17.5	15 599
	S2, 5220-5 Recurrent GNAS mutations define an unexpected pathway for pancreatic cyst development.	17.5	599
34	Science Translational Medicine, 2011, 3, 92ra66 Whole-exome sequencing of neoplastic cysts of the pancreas reveals recurrent mutations in components of ubiquitin-dependent pathways. Proceedings of the National Academy of Sciences of		599
34	Recurrent GNAS mutations define an unexpected pathway for pancreatic cyst development. Science Translational Medicine, 2011, 3, 92ra66 Whole-exome sequencing of neoplastic cysts of the pancreas reveals recurrent mutations in components of ubiquitin-dependent pathways. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 21188-93 Personalizing cancer treatment in the age of global genomic analyses: PALB2 gene mutations and the response to DNA damaging agents in pancreatic cancer. Molecular Cancer Therapeutics, 2011,	11.5	599 484
34 33 32	Recurrent GNAS mutations define an unexpected pathway for pancreatic cyst development. Science Translational Medicine, 2011, 3, 92ra66 Whole-exome sequencing of neoplastic cysts of the pancreas reveals recurrent mutations in components of ubiquitin-dependent pathways. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 21188-93 Personalizing cancer treatment in the age of global genomic analyses: PALB2 gene mutations and the response to DNA damaging agents in pancreatic cancer. Molecular Cancer Therapeutics, 2011, 10, 3-8 A genome-wide association study identifies pancreatic cancer susceptibility loci on chromosomes	11.5	599 484 208
34 33 32 31	Recurrent GNAS mutations define an unexpected pathway for pancreatic cyst development. Science Translational Medicine, 2011, 3, 92ra66 Whole-exome sequencing of neoplastic cysts of the pancreas reveals recurrent mutations in components of ubiquitin-dependent pathways. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 21188-93 Personalizing cancer treatment in the age of global genomic analyses: PALB2 gene mutations and the response to DNA damaging agents in pancreatic cancer. Molecular Cancer Therapeutics, 2011, 10, 3-8 A genome-wide association study identifies pancreatic cancer susceptibility loci on chromosomes 13q22.1, 1q32.1 and 5p15.33. Nature Genetics, 2010, 42, 224-8 Importance of age of onset in pancreatic cancer kindreds. Journal of the National Cancer Institute,	11.5 6.1 36.3	599 484 208 463
34 33 32 31 30	Recurrent GNAS mutations define an unexpected pathway for pancreatic cyst development. <i>Science Translational Medicine</i> , 2011 , 3, 92ra66 Whole-exome sequencing of neoplastic cysts of the pancreas reveals recurrent mutations in components of ubiquitin-dependent pathways. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 21188-93 Personalizing cancer treatment in the age of global genomic analyses: PALB2 gene mutations and the response to DNA damaging agents in pancreatic cancer. <i>Molecular Cancer Therapeutics</i> , 2011 , 10, 3-8 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, 224-8 Importance of age of onset in pancreatic cancer kindreds. <i>Journal of the National Cancer Institute</i> , 2010 , 102, 119-26	11.5 6.1 36.3 9.7	599 484 208 463

26	Elevated cancer mortality in the relatives of patients with pancreatic cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009 , 18, 2829-34	4	51
25	Absence of germline BRCA1 mutations in familial pancreatic cancer patients. <i>Cancer Biology and Therapy</i> , 2009 , 8, 131-5	4.6	40
24	Increased Prevalence of Precursor Lesions in Familial Pancreatic Cancer Patients. <i>Clinical Cancer Research</i> , 2009 , 15, 7737-7743	12.9	172
23	Genome-wide association study identifies variants in the ABO locus associated with susceptibility to pancreatic cancer. <i>Nature Genetics</i> , 2009 , 41, 986-90	36.3	483
22	Heritability analysis of spherical equivalent, axial length, corneal curvature, and anterior chamber depth in the Beaver Dam Eye Study. <i>JAMA Ophthalmology</i> , 2009 , 127, 649-55		71
21	Familial pancreatic cancer. Archives of Pathology and Laboratory Medicine, 2009, 133, 365-74	5	131
20	Incorporating tumor immunohistochemical markers in BRCA1 and BRCA2 carrier prediction. <i>Breast Cancer Research</i> , 2008 , 10, 401	8.3	20
19	Core signaling pathways in human pancreatic cancers revealed by global genomic analyses. <i>Science</i> , 2008 , 321, 1801-6	33.3	3223
18	Prevalence of unsuspected pancreatic cysts on MDCT. <i>American Journal of Roentgenology</i> , 2008 , 191, 802-7	5.4	636
17	The prevalence of BRCA2 mutations in familial pancreatic cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007 , 16, 342-6	4	217
16	Copy-number variants in patients with a strong family history of pancreatic cancer. <i>Cancer Biology and Therapy</i> , 2007 , 6, 1592-9	4.6	33
15	Linkage analysis of chromosome 4 in families with familial pancreatic cancer. <i>Cancer Biology and Therapy</i> , 2007 , 6, 320-3	4.6	19
14	PancPRO: risk assessment for individuals with a family history of pancreatic cancer. <i>Journal of Clinical Oncology</i> , 2007 , 25, 1417-22	2.2	150
13	Confirmation of linkage to ocular refraction on chromosome 22q and identification of a novel linkage region on 1q. <i>JAMA Ophthalmology</i> , 2007 , 125, 80-5		43
12	Familial pancreatic cancer: from genes to improved patient care. <i>Expert Review of Gastroenterology and Hepatology</i> , 2007 , 1, 81-8	4.2	14
11	DNA methylation alterations in the pancreatic juice of patients with suspected pancreatic disease. <i>Cancer Research</i> , 2006 , 66, 1208-17	10.1	181
10	Pancreatic cancer genetic epidemiology consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006 , 15, 704-10	4	116
9	Overview of linkage analysis: application to pancreatic cancer. <i>Methods in Molecular Medicine</i> , 2005 , 103, 329-41		3

LIST OF PUBLICATIONS

8	Investigation of altering single-nucleotide polymorphism density on the power to detect trait loci and frequency of false positive in nonparametric linkage analyses of qualitative traits. <i>BMC Genetics</i> , 2005 , 6 Suppl 1, S20	2.6	5
7	Support for polygenic influences on ocular refractive error. <i>Investigative Ophthalmology and Visual Science</i> , 2005 , 46, 442-6		46
6	Polygenic effects and cigarette smoking account for a portion of the familial aggregation of nuclear sclerosis. <i>American Journal of Epidemiology</i> , 2005 , 161, 707-13	3.8	12
5	Prospective risk of pancreatic cancer in familial pancreatic cancer kindreds. <i>Cancer Research</i> , 2004 , 64, 2634-8	10.1	473
4	Evidence for a major gene influencing risk of pancreatic cancer. <i>Genetic Epidemiology</i> , 2002 , 23, 133-49	2.6	108
3	Multipoint linkage analysis under heterogeneity: incorporation of parametric and nonparametric approaches. <i>Genetic Epidemiology</i> , 2001 , 21 Suppl 1, S55-60	2.6	
2	Environmental covariates: effects on the power of sib-pair linkage methods. <i>Genetic Epidemiology</i> , 1999 , 17 Suppl 1, S643-8	2.6	5
1	Assessment of Polygenic Architecture and Risk Prediction based on Common Variants Across Fourteen Cancers		1