

# Paige M Bracci

## List of Publications by Year in descending order

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100  
papers

7,029  
citations

94433

37  
h-index

62596

80  
g-index

102  
all docs

102  
docs citations

102  
times ranked

12907  
citing authors

#	ARTICLE	IF	CITATIONS
1	Interactions of Age and Blood Immune Factors and Noninvasive Prediction of Glioma Survival. Journal of the National Cancer Institute, 2022, 114, 446-457.	6.3	11
2	Phase 2B randomized controlled trial of NP001 in amyotrophic lateral sclerosis: Pre-specified and post hoc analyses. Muscle and Nerve, 2022, 66, 39-49.	2.2	16
3	Clinical characteristics of cytomegalovirus-positive pediatric acute lymphoblastic leukemia at diagnosis. American Journal of Hematology, 2022, 97, .	4.1	5
4	The immunogenetics of viral antigen response is associated with subtype-specific glioma risk and survival. American Journal of Human Genetics, 2022, 109, 1105-1116.	6.2	7
5	Pre-surgery immune profiles of adult glioma patients. Journal of Neuro-Oncology, 2022, 159, 103-115.	2.9	7
6	Cognitive impact of lower-grade gliomas and strategies for rehabilitation. Neuro-Oncology Practice, 2021, 8, 117-128.	1.6	21
7	Cancer health disparities in racial/ethnic minorities in the United States. British Journal of Cancer, 2021, 124, 315-332.	6.4	447
8	Genome-wide homozygosity and risk of four non-Hodgkin lymphoma subtypes. , 2021, 5, 200-217.		0
9	A multilayered post-GWAS assessment on genetic susceptibility to pancreatic cancer. Genome Medicine, 2021, 13, 15.	8.2	15
10	Smoking Modifies Pancreatic Cancer Risk Loci on 2q21.3. Cancer Research, 2021, 81, 3134-3143.	0.9	8
11	Hepcidin-regulating iron metabolism genes and pancreatic ductal adenocarcinoma: a pathway analysis of genome-wide association studies. American Journal of Clinical Nutrition, 2021, 114, 1408-1417.	4.7	9
12	The Role of Bacteria in KSHV Infection and KSHV-Induced Cancers. Cancers, 2021, 13, 4269.	3.7	6
13	T2 FLAIR Hyperintensity Volume Is Associated With Cognitive Function and Quality of Life in Clinically Stable Patients With Lower Grade Gliomas. Frontiers in Neurology, 2021, 12, 769345.	2.4	3
14	Abdominal Imaging of Pancreatic Cysts and Cyst-Associated Pancreatic Cancer in BRCA1/2 Mutation Carriers: A Retrospective Cross-Sectional Study. Journal of the American College of Surgeons, 2020, 230, 53-63e1.	0.5	2
15	A Transcriptome-Wide Association Study Identifies Novel Candidate Susceptibility Genes for Pancreatic Cancer. Journal of the National Cancer Institute, 2020, 112, 1003-1012.	6.3	59
16	Pseudomonas aeruginosa Stimulates Inflammation and Enhances Kaposi's Sarcoma Herpesvirus-Induced Cell Proliferation and Cellular Transformation through both Lipopolysaccharide and Flagellin. MBio, 2020, 11, .	4.1	10
17	Mendelian Randomization Analysis of n-6 Polyunsaturated Fatty Acid Levels and Pancreatic Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 2735-2739.	2.5	6
18	Incorporating multiple sets of eQTL weights into gene-by-environment interaction analysis identifies novel susceptibility loci for pancreatic cancer. Genetic Epidemiology, 2020, 44, 880-892.	1.3	0

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19	Bayesian copy number detection and association in large-scale studies. <i>BMC Cancer</i> , 2020, 20, 856.	2.6	0
20	Genome-Wide Geneâ€“Diabetes and Geneâ€“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
21	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
22	Infectious mononucleosis, immune genotypes, and non-Hodgkin lymphoma (NHL): an InterLymph Consortium study. <i>Cancer Causes and Control</i> , 2020, 31, 451-462.	1.8	4
23	Lipid Trait Variants and the Risk of Non-Hodgkin Lymphoma Subtypes: A Mendelian Randomization Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1074-1078.	2.5	13
24	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
25	BIOM-13. DNA METHYLATION MARKS GLUCOCORTICOID PATHWAY RESPONSE IN DEXAMETHASONE-TREATED BRAIN TUMOR PATIENTS. <i>Neuro-Oncology</i> , 2020, 22, ii4-ii4.	1.2	0
26	EPID-08. PRE-SURGERY IMMUNE PROFILES OF ADULT GLIOMA PATIENTS. <i>Neuro-Oncology</i> , 2020, 22, ii79-ii80.	1.2	0
27	EPCO-25. AN IMMUNOMETHYLOMIC PLATFORM INTEGRATING SYSTEMIC IMMUNE PROFILES AND EPIGENETIC AGE IN NEURO-ONCOLOGY. <i>Neuro-Oncology</i> , 2020, 22, ii74-ii74.	1.2	0
28	NCOG-21. INTERIM RESULTS OF THREE COGNITIVE REHABILITATION STRATEGIES IN PATIENTS WITH LOWER GRADE GLIOMAS. <i>Neuro-Oncology</i> , 2020, 22, ii133-ii134.	1.2	2
29	Genetic overlap between autoimmune diseases and nonâ€“Hodgkin lymphoma subtypes. <i>Genetic Epidemiology</i> , 2019, 43, 844-863.	1.3	28
30	Nivolumab in patients with advanced hepatocellular carcinoma and Childâ€“Pugh class B cirrhosis: Safety and clinical outcomes in a retrospective case series. <i>Cancer</i> , 2019, 125, 3234-3241.	4.1	73
31	Blood transfusion history and risk of non-Hodgkin lymphoma: an InterLymph pooled analysis. <i>Cancer Causes and Control</i> , 2019, 30, 889-900.	1.8	4
32	Increased neonatal level of arginase 2 in cases of childhood acute lymphoblastic leukemia implicates immunosuppression in the etiology. <i>Haematologica</i> , 2019, 104, e514-e516.	3.5	8
33	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
34	Using germline variants to estimate glioma and subtype risks. <i>Neuro-Oncology</i> , 2019, 21, 451-461.	1.2	23
35	Prognostic implications of 5-hydroxymethylcytosines from circulating cell-free DNA in diffuse large B-cell lymphoma. <i>Blood Advances</i> , 2019, 3, 2790-2799.	5.2	36
36	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

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37	Genetically Determined Height and Risk of Non-hodgkin Lymphoma. <i>Frontiers in Oncology</i> , 2019, 9, 1539.	2.8	6
38	Changes in alpha-fetoprotein (AFP) and systemic therapy outcomes in advanced hepatocellular carcinoma (HCC): A multicenter retrospective analysis.. <i>Journal of Clinical Oncology</i> , 2019, 37, 346-346.	1.6	1
39	Phase II multicenter pilot study of safety, efficacy, and immune cell profiling in advanced hepatocellular carcinoma (HCC) on combination of sorafenib (SOR) plus nivolumab (NIVO).. <i>Journal of Clinical Oncology</i> , 2019, 37, TPS464-TPS464.	1.6	2
40	Association of polygenic risk score with the risk of chronic lymphocytic leukemia and monoclonal B-cell lymphocytosis. <i>Blood</i> , 2018, 131, 2541-2551.	1.4	21
41	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. <i>Nature Communications</i> , 2018, 9, 556.	12.8	188
42	Two high-risk susceptibility loci at 6p25.3 and 14q32.13 for Waldenström macroglobulinemia. <i>Nature Communications</i> , 2018, 9, 4182.	12.8	15
43	Surgical overtreatment of pancreatic intraductal papillary mucinous neoplasms: Do the 2017 International Consensus Guidelines improve clinical decision making?. <i>Surgery</i> , 2018, 164, 1178-1184.	1.9	39
44	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
45	HLA Class I and II Diversity Contributes to the Etiologic Heterogeneity of Non-Hodgkin Lymphoma Subtypes. <i>Cancer Research</i> , 2018, 78, 4086-4096.	0.9	34
46	Association Between Telomere Length and Risk of Cancer and Non-Neoplastic Diseases. <i>JAMA Oncology</i> , 2017, 3, 636.	7.1	376
47	Genome-wide association analysis implicates dysregulation of immunity genes in chronic lymphocytic leukaemia. <i>Nature Communications</i> , 2017, 8, 14175.	12.8	75
48	Adult infiltrating gliomas with WHO 2016 integrated diagnosis: additional prognostic roles of ATRX and TERT. <i>Acta Neuropathologica</i> , 2017, 133, 1001-1016.	7.7	245
49	Immunomethylomic approach to explore the blood neutrophil lymphocyte ratio (NLR) in glioma survival. <i>Clinical Epigenetics</i> , 2017, 9, 10.	4.1	60
50	Lupus-related single nucleotide polymorphisms and risk of diffuse large B-cell lymphoma. <i>Lupus Science and Medicine</i> , 2017, 4, e000187.	2.7	15
51	Fit for genomic and proteomic purposes: Sampling the fitness of nucleic acid and protein derivatives from formalin fixed paraffin embedded tissue. <i>PLoS ONE</i> , 2017, 12, e0181756.	2.5	9
52	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
53	Genetic Variation Associated with Longer Telomere Length Increases Risk of Chronic Lymphocytic Leukemia. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1043-1049.	2.5	61
54	Serum protein profiling in diffuse large B-cell lymphoma. <i>Proteomics - Clinical Applications</i> , 2016, 10, 1113-1121.	1.6	6

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55	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016, 7, 11843.	12.8	86
56	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. <i>Nature Communications</i> , 2016, 7, 10933.	12.8	94
57	Genetically predicted longer telomere length is associated with increased risk of B-cell lymphoma subtypes. <i>Human Molecular Genetics</i> , 2016, 25, 1663-1676.	2.9	52
58	The potential relevance of the endocannabinoid, 2-arachidonoylglycerol, in diffuse large B-cell lymphoma. <i>Oncoscience</i> , 2016, 3, 31-41.	2.2	17
59	Next-generation sequencing (NGS) in an advanced hepatocellular carcinoma (HCC) cohort: Analyses of TP53 and CTNNB1.. <i>Journal of Clinical Oncology</i> , 2016, 34, 286-286.	1.6	0
60	Metformin use among type 2 diabetics and risk of pancreatic cancer in a clinicâ€based caseâ€control study. <i>International Journal of Cancer</i> , 2015, 136, E646-53.	5.1	29
61	<scp><i>TERT</i></scp> gene harbors multiple variants associated with pancreatic cancer susceptibility. <i>International Journal of Cancer</i> , 2015, 137, 2175-2183.	5.1	57
62	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.	6.3	152
63	Vitamin D Metabolic Pathway Genes and Pancreatic Cancer Risk. <i>PLoS ONE</i> , 2015, 10, e0117574.	2.5	29
64	A genome-wide association study of marginal zone lymphoma shows association to the HLA region. <i>Nature Communications</i> , 2015, 6, 5751.	12.8	58
65	Non-Hodgkin Lymphoma, Body Mass Index, and Cytokine Polymorphisms: A Pooled Analysis from the InterLymph Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1061-1070.	2.5	8
66	Glioma Groups Based on 1p/19q,<i>IDH</i>, and<i>TERT</i>Promoter Mutations in Tumors. <i>New England Journal of Medicine</i> , 2015, 372, 2499-2508.	27.0	1,632
67	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. <i>American Journal of Human Genetics</i> , 2015, 96, 487-497.	6.2	101
68	Development and Validation of a Melanoma Risk Score Based on Pooled Data from 16 Caseâ€Control Studies. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 817-824.	2.5	25
69	Statin use and risk of pancreatic cancer: Results from a large, clinicâ€based caseâ€control study. <i>Cancer</i> , 2015, 121, 1287-1294.	4.1	60
70	Serum macrophage-derived chemokine/CCL22 levels are associated with glioma risk, CD4 T cell lymphopenia and survival time. <i>International Journal of Cancer</i> , 2015, 137, 826-836.	5.1	38
71	Genome-wide association study identifies variants at 16p13 associated with survival in multiple myeloma patients. <i>Nature Communications</i> , 2015, 6, 7539.	12.8	38
72	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

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73	Cigarette smoking and risk of pancreatic cancer: a clinic-based case-control study in the San Francisco Bay Area. <i>Annals of Epidemiology</i> , 2015, 25, 816-823.	1.9	5
74	Longer genotypically-estimated leukocyte telomere length is associated with increased adult glioma risk. <i>Oncotarget</i> , 2015, 6, 42468-42477.	1.8	87
75	Rationale and Design of the International Lymphoma Epidemiology Consortium (InterLymph) Non-Hodgkin Lymphoma Subtypes Project. <i>Journal of the National Cancer Institute Monographs</i> , 2014, 2014, 1-14.	2.1	52
76	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
77	A comprehensive evaluation of the role of genetic variation in follicular lymphoma survival. <i>BMC Medical Genetics</i> , 2014, 15, 113.	2.1	17
78	Medical History, Lifestyle, Family History, and Occupational Risk Factors for Marginal Zone Lymphoma: The InterLymph Non-Hodgkin Lymphoma Subtypes Project. <i>Journal of the National Cancer Institute Monographs</i> , 2014, 2014, 52-65.	2.1	70
79	Serum Immunoglobulin E and Risk of Pancreatic Cancer in the Prostate, Lung, Colorectal, and Ovarian Cancer Screening Trial. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1414-1420.	2.5	11
80	Etiologic Heterogeneity Among Non-Hodgkin Lymphoma Subtypes: The InterLymph Non-Hodgkin Lymphoma Subtypes Project. <i>Journal of the National Cancer Institute Monographs</i> , 2014, 2014, 130-144.	2.1	265
81	Genome-wide association study identifies multiple susceptibility loci for diffuse large B cell lymphoma. <i>Nature Genetics</i> , 2014, 46, 1233-1238.	21.4	147
82	Genome-wide Association Study Identifies Five Susceptibility Loci for Follicular Lymphoma outside the HLA Region. <i>American Journal of Human Genetics</i> , 2014, 95, 462-471.	6.2	96
83	Genome-wide association study identifies multiple susceptibility loci for pancreatic cancer. <i>Nature Genetics</i> , 2014, 46, 994-1000.	21.4	294
84	Medical History, Lifestyle, Family History, and Occupational Risk Factors for Mantle Cell Lymphoma: The InterLymph Non-Hodgkin Lymphoma Subtypes Project. <i>Journal of the National Cancer Institute Monographs</i> , 2014, 2014, 76-86.	2.1	31
85	Variants near TERT and TERC influencing telomere length are associated with high-grade glioma risk. <i>Nature Genetics</i> , 2014, 46, 731-735.	21.4	161
86	Copy Number Variation Analysis on a Non-Hodgkin Lymphoma Case-Control Study Identifies an 11q25 Duplication Associated with Diffuse Large B-Cell Lymphoma. <i>PLoS ONE</i> , 2014, 9, e105382.	2.5	17
87	A Meta-Analysis Of Genome-Wide Association Studies Of Multiple Myeloma In Cases and Controls Of European Origin Identifies a Risk Locus In 12q23.1. <i>Blood</i> , 2013, 122, 3111-3111.	1.4	2
88	Obesity and Hepatitis B Infection and Risk Of Primary CNS Lymphoma. <i>Blood</i> , 2013, 122, 4298-4298.	1.4	0
89	Serum autoantibodies to pancreatic cancer antigens as biomarkers of pancreatic cancer in a San Francisco Bay Area case-control study. <i>Cancer</i> , 2012, 118, 5384-5394.	4.1	43
90	Obesity and survival in population-based patients with pancreatic cancer in the San Francisco Bay Area. <i>Cancer Causes and Control</i> , 2012, 23, 1929-1937.	1.8	22

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91	Obesity and pancreatic cancer: Overview of epidemiologic evidence and biologic mechanisms. <i>Molecular Carcinogenesis</i> , 2012, 51, 53-63.	2.7	179
92	Cigarette Smoking Associated With Lung Adenocarcinoma In Situ in a Large Case-Control Study (SFBALCS). <i>Journal of Thoracic Oncology</i> , 2012, 7, 1352-1360.	1.1	14
93	Intake of fatty acids and antioxidants and pancreatic cancer in a large population-based case-control study in the San Francisco Bay Area. <i>International Journal of Cancer</i> , 2010, 127, 1893-1904.	5.1	47
94	Chemokine polymorphisms and lymphoma: a pooled analysis. <i>Leukemia and Lymphoma</i> , 2010, 51, 497-506.	1.3	22
95	Intake of folate, vitamins B6, B12 and methionine and risk of pancreatic cancer in a large population-based case-control study. <i>Cancer Causes and Control</i> , 2009, 20, 1317-1325.	1.8	40
96	Pancreatitis and pancreatic cancer in two large pooled case-control studies. <i>Cancer Causes and Control</i> , 2009, 20, 1723-1731.	1.8	58
97	Hepatitis C and Non-Hodgkin Lymphoma Among 4784 Cases and 6269 Controls From the International Lymphoma Epidemiology Consortium. <i>Clinical Gastroenterology and Hepatology</i> , 2008, 6, 451-458.	4.4	313
98	Screening the human serum proteome for genotype-phenotype associations: An analysis of the L6-174G>C polymorphism. <i>Proteomics</i> , 2007, 7, 548-557.	2.2	18
99	Residential History, Family Characteristics and Non-Hodgkin Lymphoma, A Population-Based Case-Control Study in the San Francisco Bay Area. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 1287-1294.	2.5	29
100	Tobacco use and non-Hodgkin lymphoma: results from a population-based case-control study in the San Francisco Bay Area, California. <i>Cancer Causes and Control</i> , 2005, 16, 333-346.	1.8	15