

Peter J Bailey

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.

17
papers

6,305
citations

14
h-index

19
g-index

19
ext. papers

8,077
ext. citations

16
avg, IF

3.75
L-index

#	Paper	IF	Citations
17	Clinical Impact of Molecular Subtyping of Pancreatic Cancer. <i>Frontiers in Cell and Developmental Biology</i> , 2021 , 9, 743908	5.7	4
16	HNF4A and GATA6 Loss Reveals Therapeutically Actionable Subtypes in Pancreatic Cancer. <i>Cell Reports</i> , 2020 , 31, 107625	10.6	34
15	Altered RNA Splicing by Mutant p53 Activates Oncogenic RAS Signaling in Pancreatic Cancer. <i>Cancer Cell</i> , 2020 , 38, 198-211.e8	24.3	38
14	PDAC Subtypes/Stratification. <i>Molecular and Translational Medicine</i> , 2020 , 117-128	0.4	0
13	Using Chromatin Accessibility to Delineate Therapeutic Subtypes in Pancreatic Cancer Patient-Derived Cell Lines. <i>STAR Protocols</i> , 2020 , 1, 100079	1.4	
12	Prolactin Promotes Fibrosis and Pancreatic Cancer Progression. <i>Cancer Research</i> , 2019 , 79, 5316-5327	10.1	19
11	CSF1R Macrophages Sustain Pancreatic Tumor Growth through T Cell Suppression and Maintenance of Key Gene Programs that Define the Squamous Subtype. <i>Cell Reports</i> , 2018 , 23, 1448-1460	10.6	105
10	BRCA2 secondary mutation-mediated resistance to platinum and PARP inhibitor-based therapy in pancreatic cancer. <i>British Journal of Cancer</i> , 2017 , 116, 1021-1026	8.7	48
9	Integrated Genomic Characterization of Pancreatic Ductal Adenocarcinoma. <i>Cancer Cell</i> , 2017 , 32, 185-203.e13	20.5	13896
8	Hypermutation In Pancreatic Cancer. <i>Gastroenterology</i> , 2017 , 152, 68-74.e2	13.3	130
7	Exploiting the neoantigen landscape for immunotherapy of pancreatic ductal adenocarcinoma. <i>Scientific Reports</i> , 2016 , 6, 35848	4.9	87
6	CXCR2 Inhibition Profoundly Suppresses Metastases and Augments Immunotherapy in Pancreatic Ductal Adenocarcinoma. <i>Cancer Cell</i> , 2016 , 29, 832-845	24.3	442
5	Genomic analyses identify molecular subtypes of pancreatic cancer. <i>Nature</i> , 2016 , 531, 47-52	50.4	1785
4	Whole genomes redefine the mutational landscape of pancreatic cancer. <i>Nature</i> , 2015 , 518, 495-501	50.4	1579
3	Integrated genomic and transcriptomic analysis of human brain metastases identifies alterations of potential clinical significance. <i>Journal of Pathology</i> , 2015 , 237, 363-78	9.4	72
2	Whole-genome characterization of chemoresistant ovarian cancer. <i>Nature</i> , 2015 , 521, 489-94	50.4	890
1	Genomic catastrophes frequently arise in esophageal adenocarcinoma and drive tumorigenesis. <i>Nature Communications</i> , 2014 , 5, 5224	17.4	176

