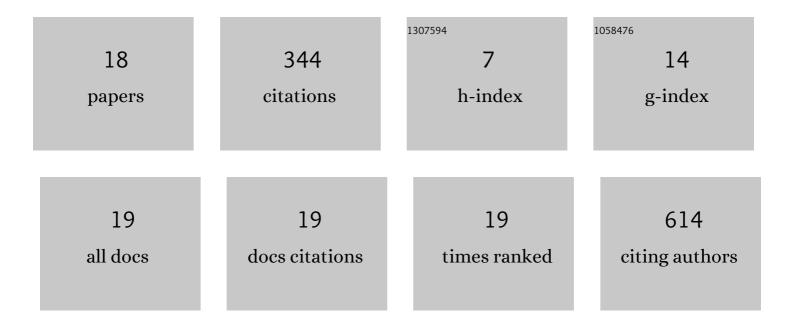
Jian Bai

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Common Postzygotic Mutational Signatures in Healthy Adult Tissues Related to Embryonic Hypoxia. Genomics, Proteomics and Bioinformatics, 2022, 20, 177-191.	6.9	1
2	Genome-scale profiling of circulating cell-free DNA signatures for early detection of hepatocellular carcinoma in cirrhotic patients. Cell Research, 2021, 31, 589-592.	12.0	59
3	Influence of low tumor content on tumor mutational burden estimation by wholeâ€exome sequencing and targeted panel sequencing. Clinical and Translational Medicine, 2021, 11, e415.	4.0	7
4	The Landscape of Cell-Free HBV Integrations and Mutations in Cirrhosis and Hepatocellular Carcinoma Patients. Clinical Cancer Research, 2021, 27, 3772-3783.	7.0	21
5	Durable Response to the Combination of Atezolizumab With Platinum-Based Chemotherapy in an Untreated Non-Smoking Lung Adenocarcinoma Patient With BRAF V600E Mutation: A Case Report. Frontiers in Oncology, 2021, 11, 634920.	2.8	7
6	Identification of Somatic Mutations in Papanicolaou Smear DNA and Plasma Circulating Cell-Free DNA for Detection of Endometrial and Epithelial Ovarian Cancers: A Pilot Study. Frontiers in Oncology, 2020, 10, 582546.	2.8	4
7	Neoantigen Tracking: Tracking Neoantigens by Personalized Circulating Tumor DNA Sequencing during Checkpoint Blockade Immunotherapy in Nonâ€5mall Cell Lung Cancer (Adv. Sci. 9/2020). Advanced Science, 2020, 7, 2070047.	11.2	Ο
8	Tracking Neoantigens by Personalized Circulating Tumor DNA Sequencing during Checkpoint Blockade Immunotherapy in Non‣mall Cell Lung Cancer. Advanced Science, 2020, 7, 1903410.	11.2	30
9	Abstract 4597: 5-Hydroxymethylcytosine signatures in cell-free DNA as a potential biomarker for colorectal cancer andprecancerous adenoma. , 2020, , .		Ο
10	A Reference System for BRCA Mutation Detection Based on Next-Generation Sequencing in the Chinese Population. Journal of Molecular Diagnostics, 2019, 21, 677-686.	2.8	7
11	Germline Mutations in Cancer Susceptibility Genes in a Large Series of Unselected Breast Cancer Patients. Clinical Cancer Research, 2017, 23, 6113-6119.	7.0	159
12	Retinol dehydrogenase-10 promotes development and progression of human glioma via the TWEAK-NF-κB axis. Oncotarget, 2017, 8, 105262-105275.	1.8	6
13	Discovery of potential colorectal cancer serum biomarkers through quantitative proteomics on the colonic tissue interstitial fluids from the AOM–DSS mouse model. Journal of Proteomics, 2016, 132, 31-40.	2.4	28
14	Abstract 2397: Mutation signatures and intratumor heterogeneity of esophageal squamous cell carcinoma in a Chinese cohort. , 2016, , .		0
15	An Old Story Retold: Loss of G1 Control Defines A Distinct Genomic Subtype of Esophageal Squamous Cell Carcinoma. Genomics, Proteomics and Bioinformatics, 2015, 13, 258-270.	6.9	7
16	Abstract 4797: Mutation signature and intratumor heterogeneity of esophageal squamous cell carcinoma in a Chinese cohort. Cancer Research, 2015, 75, 4797-4797.	0.9	1
17	Positive selection signals of hepatitis B virus and their association with disease stages and viral genotypes. Infection, Genetics and Evolution, 2013, 19, 176-187.	2.3	7
18	Abstract C03: A comprehensive study of genomic alterations reveals deregulation of the cell cycle in most esophageal squamous cell carcinomas. , 2013, , .		0