

# Na You

## List of Publications by Year in descending order

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Version: 2024-02-01

20  
papers

238  
citations

1307594

7  
h-index

996975

15  
g-index

21  
all docs

21  
docs citations

21  
times ranked

473  
citing authors

#	ARTICLE	IF	CITATIONS
1	A Dietary Intervention in Urban African Americans. <i>American Journal of Preventive Medicine</i> , 2016, 50, 87-95.	3.0	50
2	Prognostic Value of a BCSC-associated MicroRNA Signature in Hormone Receptor-Positive HER2-Negative Breast Cancer. <i>EBioMedicine</i> , 2016, 11, 199-209.	6.1	43
3	Survival impact of waiting time for radical radiotherapy in nasopharyngeal carcinoma: A large institution-based cohort study from an endemic area. <i>European Journal of Cancer</i> , 2017, 73, 48-60.	2.8	32
4	SNP calling using genotype model selection on high-throughput sequencing data. <i>Bioinformatics</i> , 2012, 28, 643-650.	4.1	22
5	Effect of younger age on survival outcomes in T1N0M0 breast cancer: A propensity score matching analysis. <i>Journal of Surgical Oncology</i> , 2019, 119, 1039-1046.	1.7	22
6	Undetectable circulating tumor DNA levels correlate with low risk of recurrence/metastasis in postoperative pathologic stage I lung adenocarcinoma patients. <i>Lung Cancer</i> , 2020, 146, 327-334.	2.0	14
7	Prognostic immune-related gene models for breast cancer: a pooled analysis. <i>OncoTargets and Therapy</i> , 2017, Volume 10, 4423-4433.	2.0	10
8	Comparison of beta diversity measures in clustering the high-dimensional microbial data. <i>PLoS ONE</i> , 2021, 16, e0246893.	2.5	9
9	Association of polymorphisms in C1orf106, IL1RN, and IL10 with post-induction infliximab trough level in Crohn's disease patients. <i>Gastroenterology Report</i> , 2020, 8, 367-373.	1.3	8
10	Neoadjuvant everolimus plus letrozole versus fluorouracil, epirubicin and cyclophosphamide for ER-positive, HER2-negative breast cancer: study protocol for a randomized pilot trial. <i>Trials</i> , 2017, 18, 497.	1.6	6
11	An empirical Bayes method for genotyping and SNP detection using multi-sample next-generation sequencing data. <i>Bioinformatics</i> , 2016, 32, 3240-3245.	4.1	5
12	Subtype Classification and Heterogeneous Prognosis Model Construction in Precision Medicine. <i>Biometrics</i> , 2018, 74, 814-822.	1.4	5
13	An empirical Bayes method for robust variance estimation in detecting DEGs using microarray data. <i>Journal of Bioinformatics and Computational Biology</i> , 2017, 15, 1750020.	0.8	4
14	MultiGeMS: detection of SNVs from multiple samples using model selection on high-throughput sequencing data. <i>Bioinformatics</i> , 2016, 32, 1486-1492.	4.1	3
15	B-CAN: a resource sharing platform to improve the operation, visualization and integrated analysis of TCGA breast cancer data. <i>Oncotarget</i> , 2017, 8, 108778-108785.	1.8	3
16	Single-sample SNP detection by empirical Bayes method using next-generation sequencing data. <i>Statistics and Its Interface</i> , 2015, 8, 457-462.	0.3	1
17	Application of Clustering Method to Explore the Correlation Between Dominant Flora and the Autism Spectrum Disorder Clinical Phenotype in Chinese Children. <i>Frontiers in Neuroscience</i> , 2021, 15, 760779.	2.8	1
18	Gene Expression Network Reconstruction by LEP Method Using Microarray Data. <i>Scientific World Journal</i> , The, 2012, 2012, 1-6.	2.1	0

#	ARTICLE	IF	CITATIONS
19	Prognostic value of a BCSC-associated microRNA signature in hormone receptor-positive HER2-negative breast cancer.. Journal of Clinical Oncology, 2016, 34, 532-532.	1.6	0
20	Circulating tumor DNA dynamics to predict cancer recurrence/metastasis in Chinese pathologic stage I lung adenocarcinoma.. Journal of Clinical Oncology, 2020, 38, 3537-3537.	1.6	0