

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	eQTL mapping using allele-specific count data is computationally feasible, powerful, and provides individual-specific estimates of genetic effects. PLoS Genetics, 2022, 18, e1010076.	3.5	13
2	A Statistical Method for Association Analysis of Cell Type Compositions. Statistics in Biosciences, 2021, 13, 373-385.	1.2	0
3	Correlation of alterations in the <i><scp>KEAP1</scp>/<scp>CUL3</scp>/<scp>NFE2L2</scp></i> pathway with radiation failure in larynx squamous cell carcinoma. Laryngoscope Investigative Otolaryngology, 2021, 6, 699-707.	1.5	11
4	UNMASC: tumor-only variant calling with unmatched normal controls. NAR Cancer, 2021, 3, zcab040.	3.1	4
5	Associating somatic mutations to clinical outcomes: a pan-cancer study of survival time. Genome Medicine, 2019, 11, 37.	8.2	8
6	The association between copy number aberration, DNA methylation and gene expression in tumor samples. Nucleic Acids Research, 2018, 46, 3009-3018.	14.5	51
7	LCCC 1025: a phase II study of everolimus, trastuzumab, and vinorelbine to treat progressive HER2-positive breast cancer brain metastases. Breast Cancer Research and Treatment, 2018, 171, 637-648.	2.5	40
8	The Prognostic Significance of Low-Frequency Somatic Mutations in Metastatic Cutaneous Melanoma. Frontiers in Oncology, 2018, 8, 584.	2.8	14
9	LCCC 1025: Phase II study of everolimus, trastuzumab, and vinorelbine for HER2+ breast cancer brain metastases (BCBM) Journal of Clinical Oncology, 2017, 35, 1011-1011.	1.6	3
10	Characterization of the number and site of APC mutations in sporadic colorectal cancer Journal of Clinical Oncology, 2017, 35, 630-630.	1.6	5
11	Abstract 3584: Statistical considerations in tumor only variant calling. , 2017, , .		0
12	Myeloid-derived suppressor cells regulate T cell and B cell responses during autoimmune disease. Journal of Leukocyte Biology, 2015, 97, 573-582.	3.3	90