Azeet Narayan

List of Publications by Year
in descending order

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Version: 2024-02-01

18	713	1040056 9	1058476 14
papers	citations	h-index	g-index

18	18	18	1614
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Tumor DNA Mutations From Intraparenchymal Brain Metastases Are Detectable in CSF. JCO Precision Oncology, 2021, 5, 163-172.	3.0	9
2	Dynamics of circulating tumor DNA in patients with advanced solid tumors treated with cediranib and olaparib.. Journal of Clinical Oncology, 2021, 39, 3035-3035.	1.6	0
3	35. EVALUATING CSF CIRCULATING TUMOR DNA IN INTRAPARENCHYMAL BRAIN METASTASIS. Neuro-Oncology Advances, 2020, 2, ii6-ii6.	0.7	1
4	META RNA profiling: Multiplexed quantitation of targeted RNAs across large numbers of samples. Methods, 2019, 152, 41-47.	3.8	1
5	Early Assessment of Lung Cancer Immunotherapy Response via Circulating Tumor DNA. Clinical Cancer Research, 2018, 24, 1872-1880.	7.0	319
6	Oligosaccharyltransferase Inhibition Overcomes Therapeutic Resistance to EGFR Tyrosine Kinase Inhibitors. Cancer Research, 2018, 78, 5094-5106.	0.9	47
7	Circulating tumor DNA (ctDNA) to monitor treatment response and progression in patients treated with tyrosine kinase inhibitors (TKIs) and immunotherapy for EGFR-mutant non-small cell lung cancer (NSCLC).. Journal of Clinical Oncology, 2017, 35, e20652-e20652.	1.6	1
8	High-throughput RNA profiling via up-front sample parallelization. Nature Methods, 2015, 12, 343-346.	19.0	18
9	Measurement of circulating tumor DNA as a cancer biomarker in gastrointestinal malignancies using a novel next-generation sequencing method.. Journal of Clinical Oncology, 2014, 32, 217-217.	1.6	0
10	Detection of sensitizing and resistance EGFR mutations from circulating tumor DNA (ctDNA) in blood using multiplexed next-generation sequencing in patients with advanced EGFR-mutant lung adenocarcinoma.. Journal of Clinical Oncology, 2014, 32, 8093-8093.	1.6	0
11	Monitoring changes in circulating tumor DNA in gastrointestinal malignancies using a novel next-generation sequencing method.. Journal of Clinical Oncology, 2014, 32, 3645-3645.	1.6	0
12	Ultrasensitive Measurement of Hotspot Mutations in Tumor DNA in Blood Using Error-Suppressed Multiplexed Deep Sequencing. Cancer Research, 2012, 72, 3492-3498.	0.9	159
13	Phenylalanine-Rich Peptides Potently Bind ESAT6, a Virulence Determinant of Mycobacterium tuberculosis, and Concurrently Affect the Pathogen's Growth. PLoS ONE, 2009, 4, e7615.	2.5	16
14	Comparative genomic study of spo0E family genes and elucidation of the role of Spo0E in BacillusÂanthracis. Archives of Microbiology, 2009, 191, 241-253.	2.2	6
15	Spo0B of Bacillusâ€ƒanthracis â€“ a protein with pleiotropic functions. FEBS Journal, 2008, 275, 739-752.	4.7	14
16	Loss of kinase activity in Mycobacteriumâ€ƒtuberculosis multidomain protein Rv1364c. FEBS Journal, 2008, 275, 6295-6308.	4.7	15
17	Serine threonine protein kinases of mycobacterial genus: phylogeny to function. Physiological Genomics, 2007, 29, 66-75.	2.3	76
18	PknH, a transmembrane Hank's type serine/threonine kinase fromMycobacterium tuberculosisis differentially expressed under stress conditions. FEMS Microbiology Letters, 2004, 233, 107-113.	1.8	31