Nicholas Light

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/10396346/publications.pdf

Version: 2024-02-01

11	1,017	1040056 9	1372567 10
papers	citations	h-index	g-index

12	12	12	3184
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Pathogenic Germline Variants in Cancer Susceptibility Genes in Children and Young Adults With Rhabdomyosarcoma. JCO Precision Oncology, 2021, 5, 75-87.	3.0	27
2	Lineage-defined leiomyosarcoma subtypes emerge years before diagnosis and determine patient survival. Nature Communications, 2021, 12, 4496.	12.8	28
3	DNA Polymerase and Mismatch Repair Exert Distinct Microsatellite Instability Signatures in Normal and Malignant Human Cells. Cancer Discovery, 2021, 11, 1176-1191.	9.4	46
4	Myc and Loss of p53 Cooperate to Drive Formation of Choroid Plexus Carcinoma. Cancer Research, 2019, 79, 2208-2219.	0.9	15
5	Ewingâ€like sarcoma: An emerging family of round cell sarcomas. Journal of Cellular Physiology, 2019, 234, 7999-8007.	4.1	68
6	HGG-17. TUMOR MUTATIONAL BURDEN ANALYSIS OF PEDIATRIC TUMORS PROVIDES A DIAGNOSTIC TOOL FOR GERMLINE REPLICATION REPAIR DEFICIENCY AND PREDICT RESPONSE TO IMMUNE CHECKPOINT INHIBITION. Neuro-Oncology, 2018, 20, i92-i92.	1.2	0
7	Explosive mutation accumulation triggered by heterozygous human Pol Îµ proofreading-deficiency is driven by suppression of mismatch repair. ELife, 2018, 7, .	6.0	33
8	Comprehensive Analysis of Hypermutation in Human Cancer. Cell, 2017, 171, 1042-1056.e10.	28.9	596
9	Interrogation of allelic chromatin states in human cells by high-density ChIP-genotyping. Epigenetics, 2014, 9, 1238-1251.	2.7	9
10	Allelic expression mapping across cellular lineages to establish impact of nonâ€coding <scp>SNP</scp>s. Molecular Systems Biology, 2014, 10, 754.	7.2	21
11	Analysis of expressed SNPs identifies variable extents of expression from the human inactive X chromosome. Genome Biology, 2013, 14, R122.	9.6	174