

Peter W Eide

List of Publications by Year in descending order

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17
papers

1,056
citations

759233

12
h-index

888059

17
g-index

17
all docs

17
docs citations

17
times ranked

2239
citing authors

#	ARTICLE	IF	CITATIONS
1	E-cadherin is a robust prognostic biomarker in colorectal cancer and low expression is associated with sensitivity to inhibitors of topoisomerase, aurora, and HSP90 in preclinical models. <i>Molecular Oncology</i> , 2022, 16, 2312-2329.	4.6	4
2	Metastatic heterogeneity of the consensus molecular subtypes of colorectal cancer. <i>Npj Genomic Medicine</i> , 2021, 6, 59.	3.8	29
3	De novo transcriptomic subtyping of colorectal cancer liver metastases in the context of tumor heterogeneity. <i>Genome Medicine</i> , 2021, 13, 143.	8.2	10
4	The expressed mutational landscape of microsatellite stable colorectal cancers. <i>Genome Medicine</i> , 2021, 13, 142.	8.2	4
5	Increased sensitivity to SMAC mimetic LCL161 identified by longitudinal ex vivo pharmacogenomics of recurrent, KRAS mutated rectal cancer liver metastases. <i>Journal of Translational Medicine</i> , 2021, 19, 384.	4.4	6
6	Molecular correlates of sensitivity to PARP inhibition beyond homologous recombination deficiency in pre-clinical models of colorectal cancer point to wild-type TP53 activity. <i>EBioMedicine</i> , 2020, 59, 102923.	6.1	22
7	Patient-Derived Organoids from Multiple Colorectal Cancer Liver Metastases Reveal Moderate Intra-patient Pharmacotranscriptomic Heterogeneity. <i>Clinical Cancer Research</i> , 2020, 26, 4107-4119.	7.0	68
8	Transcriptional and functional consequences of TP53 splice mutations in colorectal cancer. <i>Oncogenesis</i> , 2019, 8, 35.	4.9	19
9	Exploratory analyses of consensus molecular subtype-dependent associations of TP53 mutations with immunomodulation and prognosis in colorectal cancer. <i>ESMO Open</i> , 2019, 4, e000523.	4.5	11
10	Long noncoding RNA <i>MIR31HG</i> is a bona fide prognostic marker with colorectal cancer cell-intrinsic properties. <i>International Journal of Cancer</i> , 2019, 144, 2843-2853.	5.1	33
11	Colorectal Cancer Consensus Molecular Subtypes Translated to Preclinical Models Uncover Potentially Targetable Cancer Cell Dependencies. <i>Clinical Cancer Research</i> , 2018, 24, 794-806.	7.0	177
12	Prognostic, predictive, and pharmacogenomic assessments of <i>CDX2</i> refine stratification of colorectal cancer. <i>Molecular Oncology</i> , 2018, 12, 1639-1655.	4.6	40
13	E3 ubiquitin ligase NEDD4 induces endocytosis and lysosomal sorting of connexin43 to promote loss of gap junctions. <i>Journal of Cell Science</i> , 2017, 130, 2867-2882.	2.0	14
14	CMScaller: an R package for consensus molecular subtyping of colorectal cancer pre-clinical models. <i>Scientific Reports</i> , 2017, 7, 16618.	3.3	229
15	Multi-omics of 34 colorectal cancer cell lines - a resource for biomedical studies. <i>Molecular Cancer</i> , 2017, 16, 116.	19.2	232
16	MicroRNAs as growth regulators, their function and biomarker status in colorectal cancer. <i>Oncotarget</i> , 2016, 7, 6476-6505.	1.8	93
17	NEDD4 is overexpressed in colorectal cancer and promotes colonic cell growth independently of the PI3K/PTEN/AKT pathway. <i>Cellular Signalling</i> , 2013, 25, 12-18.	3.6	65