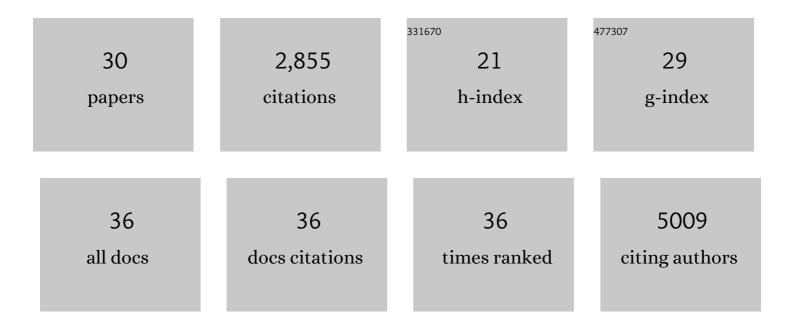
Lukas C Heukamp

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4018058/publications.pdf Version: 2024-02-01



LUKAS C HEUKAMA

#	Article	IF	CITATIONS
1	Frequent and Focal <i>FGFR1</i> Amplification Associates with Therapeutically Tractable FGFR1 Dependency in Squamous Cell Lung Cancer. Science Translational Medicine, 2010, 2, 62ra93.	12.4	761
2	MYCN regulates oncogenic MicroRNAs in neuroblastoma. International Journal of Cancer, 2008, 122, 699-704.	5.1	251
3	Frequent mutations in chromatin-remodelling genes in pulmonary carcinoids. Nature Communications, 2014, 5, 3518.	12.8	239
4	Circulating microRNAs (miRNA) in Serum of Patients With Prostate Cancer. Urology, 2011, 77, 1265.e9-1265.e16.	1.0	210
5	Global levels of histone modifications predict prostate cancer recurrence. Prostate, 2010, 70, 61-69.	2.3	194
6	Diagnostic and Prognostic Information in Prostate Cancer with the Help of a Small Set of Hypermethylated Gene Loci. Clinical Cancer Research, 2005, 11, 4097-4106.	7.0	135
7	Elp3 links tRNA modification to IRES-dependent translation of LEF1 to sustain metastasis in breast cancer. Journal of Experimental Medicine, 2016, 213, 2503-2523.	8.5	128
8	CpG Island hypermethylation in cell-free serum DNA identifies patients with localized prostate cancer. Prostate, 2008, 68, 42-49.	2.3	121
9	CpG Island Hypermethylation at Multiple Gene Sites in Diagnosis and Prognosis of Prostate Cancer. Urology, 2008, 71, 161-167.	1.0	120
10	Prognostic Value of CpG Island Hypermethylation at PTGS2, RAR-beta, EDNRB, and Other Gene Loci in Patients Undergoing Radical Prostatectomy. European Urology, 2007, 51, 665-674.	1.9	72
11	Global histone H4K20 trimethylation predicts cancer-specific survival in patients with muscle-invasive bladder cancer. BJU International, 2011, 108, E290-E296.	2.5	68
12	Elp3 drives Wnt-dependent tumor initiation and regeneration in the intestine. Journal of Experimental Medicine, 2015, 212, 2057-2075.	8.5	67
13	Global Histone H3K27 Methylation Levels are Different in Localized and Metastatic Prostate Cancer. Cancer Investigation, 2012, 30, 92-97.	1.3	51
14	Epidermal growth factor receptor mutations in non-small cell lung cancer influence downstream Akt, MAPK and Stat3 signaling. Journal of Cancer Research and Clinical Oncology, 2009, 135, 723-730.	2.5	47
15	Alterations of global histone H4K20 methylation during prostate carcinogenesis. BMC Urology, 2012, 12, 5.	1.4	46
16	KRAS G12C-mutated advanced non-small cell lung cancer: A real-world cohort from the German prospective, observational, nation-wide CRISP Registry (AIO-TRK-0315). Lung Cancer, 2021, 154, 51-61.	2.0	43
17	Systematic Kinase Inhibitor Profiling Identifies CDK9 as a Synthetic Lethal Target in NUT Midline Carcinoma. Cell Reports, 2017, 20, 2833-2845.	6.4	40
18	MicroRNAs in the pathogenesis of neuroblastoma. Cancer Letters, 2009, 274, 10-15.	7.2	37

LUKAS C HEUKAMP

#	Article	IF	CITATIONS
19	Alterations of Global Histone H3K9 and H3K27 Methylation Levels in Bladder Cancer. Urologia Internationalis, 2014, 93, 113-118.	1.3	31
20	Anti-Proliferative Effect of Cytohesin Inhibition in Gefitinib-Resistant Lung Cancer Cells. PLoS ONE, 2012, 7, e41179.	2.5	29
21	Intermittent high-dose treatment with erlotinib enhances therapeutic efficacy in EGFR-mutant lung cancer. Oncotarget, 2015, 6, 38458-38468.	1.8	19
22	<i>CUL2</i> and <i>STK11</i> as novel response-predictive genes for neoadjuvant radiochemotherapy in esophageal cancer. Pharmacogenomics, 2010, 11, 1105-1113.	1.3	18
23	The LIM-Only Protein FHL2 Reduces Vascular Lesion Formation Involving Inhibition of Proliferation and Migration of Smooth Muscle Cells. PLoS ONE, 2014, 9, e94931.	2.5	17
24	The PDL1-inducible GTPase Arl4d controls T effector function by limiting IL-2 production. Scientific Reports, 2018, 8, 16123.	3.3	13
25	Implementing amplicon-based next generation sequencing in the diagnosis of small cell lung carcinoma metastases. Experimental and Molecular Pathology, 2015, 99, 682-686.	2.1	12
26	The Role of Molecular Diagnostics in Cancer Diagnosis and Treatment. Onkologie, 2012, 35, 8-12.	0.8	11
27	Web-based database for the management of tissue specimens in a transregional histological research facility. Diagnostic Pathology, 2011, 6, 17.	2.0	2
28	CPG ISLAND HYPERMETHYLATION OF CELL-FREE SERUM DNA INDICATES WORSE OUTCOME IN PATIENTS WITH BLADDER CANCER. Journal of Urology, 2008, 179, 315-315.	0.4	1
29	Molekulardiagnostik von Mutationen des epidermalen Wachstumsfaktor-Rezeptors und Aktivierung nachgeschalteter Signalwege in nichtkleinzelligen Lungenkarzinomen. Onkopipeline, 2008, 1, 101-108.	0.0	0
30	HYPERMETHYLATION IN CELL-FREE CIRCULATING SERUM DNA IDENTIFIES PATIENTS WITH LOCALIZED PROSTATE CANCER. Journal of Urology, 2008, 179, 720-721.	0.4	0