

Kai Breuhahn

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3574409/publications.pdf>

Version: 2024-02-01

89
papers

4,496
citations

94269

37
h-index

106150

65
g-index

92
all docs

92
docs citations

92
times ranked

7867
citing authors

#	ARTICLE	IF	CITATIONS
1	Acquired Resistance to Antiangiogenic Therapies in Hepatocellular Carcinoma Is Mediated by Yes-Associated Protein 1 Activation and Transient Expansion of Stem-Like Cancer Cells. <i>Hepatology Communications</i> , 2022, 6, 1140-1156.	2.0	6
2	HELLS Is Negatively Regulated by Wild-Type P53 in Liver Cancer by a Mechanism Involving P21 and FOXM1. <i>Cancers</i> , 2022, 14, 459.	1.7	6
3	Non-canonical NF- κ B signaling induces proliferation in primary liver cancer. <i>Zeitschrift Fur Gastroenterologie</i> , 2022, 60, .	0.2	0
4	Direct interaction of the oncogenes YAP and TAZ with the transcription factor HNF1B in hepatocellular carcinoma. <i>Zeitschrift Fur Gastroenterologie</i> , 2022, 60, .	0.2	0
5	STAT1 and STAT3 Exhibit a Crosstalk and Are Associated with Increased Inflammation in Hepatocellular Carcinoma. <i>Cancers</i> , 2022, 14, 1154.	1.7	11
6	LINC00152 Drives a Competing Endogenous RNA Network in Human Hepatocellular Carcinoma. <i>Cells</i> , 2022, 11, 1528.	1.8	6
7	YAP-induced Ccl2 expression is associated with a switch in hepatic macrophage identity and vascular remodelling in liver cancer. <i>Liver International</i> , 2021, 41, 3011-3023.	1.9	17
8	Histone H3K27 demethylase KDM6A is an epigenetic gatekeeper of mTORC1 signalling in cancer. <i>Gut</i> , 2021, , gutjnl-2021-325405.	6.1	15
9	Co-expression of YAP and TAZ associates with chromosomal instability in human cholangiocarcinoma. <i>BMC Cancer</i> , 2021, 21, 1079.	1.1	14
10	TAZ target gene ITGAV regulates invasion and feeds back positively on YAP and TAZ in liver cancer cells. <i>Cancer Letters</i> , 2020, 473, 164-175.	3.2	39
11	YAP Orchestrates Heterotypic Endothelial Cell Communication via HGF/c-MET Signaling in Liver Tumorigenesis. <i>Cancer Research</i> , 2020, 80, 5502-5514.	0.4	31
12	Yes-associated protein (YAP) induces a secretome phenotype and transcriptionally regulates plasminogen activator inhibitor-1 (PAI-1) expression in hepatocarcinogenesis. <i>Cell Communication and Signaling</i> , 2020, 18, 166.	2.7	21
13	A dual role for hepatocyte-intrinsic canonical NF- κ B signaling in virus control. <i>Journal of Hepatology</i> , 2020, 72, 960-975.	1.8	18
14	Nucleoporin Nup155 is part of the p53 network in liver cancer. <i>Nature Communications</i> , 2019, 10, 2147.	5.8	29
15	YAP-dependent induction of UHMK1 supports nuclear enrichment of the oncogene MYBL2 and proliferation in liver cancer cells. <i>Oncogene</i> , 2019, 38, 5541-5550.	2.6	45
16	Nuclear Translocation of RELB Is Increased in Diseased Human Liver and Promotes Ductular Reaction and Biliary Fibrosis in Mice. <i>Gastroenterology</i> , 2019, 156, 1190-1205.e14.	0.6	19
17	Karyopherin β 2-dependent import of E2F1 and TFDP1 maintains protumorigenic stathmin expression in liver cancer. <i>Cell Communication and Signaling</i> , 2019, 17, 159.	2.7	29
18	Editorial: Systems Biology and Bioinformatics in Gastroenterology and Hepatology. <i>Frontiers in Physiology</i> , 2019, 10, 1438.	1.3	0

#	ARTICLE	IF	CITATIONS
19	Epigenetically Regulated Chromosome 14q32 miRNA Cluster Induces Metastasis and Predicts Poor Prognosis in Lung Adenocarcinoma Patients. <i>Molecular Cancer Research</i> , 2018, 16, 390-402.	1.5	63
20	Cytoplasmic localization of the cell polarity factor scribble supports liver tumor formation and tumor cell invasiveness. <i>Hepatology</i> , 2018, 67, 1842-1856.	3.6	48
21	Methylation in MIRLET7A3 Gene Induces the Expression of IGF-II and Its mRNA Binding Proteins IGF2BP-2 and 3 in Hepatocellular Carcinoma. <i>Frontiers in Physiology</i> , 2018, 9, 1918.	1.3	12
22	Induction of Chromosome Instability by Activation of Yes-Associated Protein and Forkhead Box M1 in Liver Cancer. <i>Gastroenterology</i> , 2017, 152, 2037-2051.e22.	0.6	118
23	TTCA: an R package for the identification of differentially expressed genes in time course microarray data. <i>BMC Bioinformatics</i> , 2017, 18, 33.	1.2	16
24	Proteomic Analysis Reveals GMP Synthetase as p53 Repression Target in Liver Cancer. <i>American Journal of Pathology</i> , 2017, 187, 228-235.	1.9	26
25	An individual-based model for collective cancer cell migration explains speed dynamics and phenotype variability in response to growth factors. <i>Npj Systems Biology and Applications</i> , 2017, 3, 5.	1.4	29
26	A20/TNFAIP3 Discriminates Tumor Necrosis Factor (TNF)-Induced NF- κ B from JNK Pathway Activation in Hepatocytes. <i>Frontiers in Physiology</i> , 2017, 8, 610.	1.3	16
27	FOXM1 activates AGR2 and causes progression of lung adenomas into invasive mucinous adenocarcinomas. <i>PLoS Genetics</i> , 2017, 13, e1007097.	1.5	48
28	MicroRNAs are key regulators of hepatocellular carcinoma (HCC) cell dissemination—what we learned from microRNA-494. <i>Hepatobiliary Surgery and Nutrition</i> , 2016, 5, 372-376.	0.7	6
29	Quantitative estimation of tumor cellularity based on histology data. , 2016, , .		1
30	PI3K/AKT/mTOR—dependent stabilization of oncogenic far—upstream element binding proteins in hepatocellular carcinoma cells. <i>Hepatology</i> , 2016, 63, 813-826.	3.6	52
31	Chromosome 8p tumor suppressor genes SH2D4A and SORBS3 cooperate to inhibit interleukin—6 signaling in hepatocellular carcinoma. <i>Hepatology</i> , 2016, 64, 828-842.	3.6	29
32	Inducing Differentiation of Premalignant Hepatic Cells as a Novel Therapeutic Strategy in Hepatocarcinoma. <i>Cancer Research</i> , 2016, 76, 5550-5561.	0.4	15
33	The proto-oncogene Myc drives expression of the NK cell-activating NKp30 ligand B7-H6 in tumor cells. <i>Oncolmmunology</i> , 2016, 5, e1116674.	2.1	39
34	Tumor microvasculature in lung cancer and diffusion-weighted MRI: Preliminary results. , 2016, , .		0
35	Directed random walks and constraint programming reveal active pathways in hepatocyte growth factor signaling. <i>FEBS Journal</i> , 2016, 283, 350-360.	2.2	5
36	Cellular apoptosis susceptibility (CAS) is overexpressed in thyroid carcinoma and maintains tumor cell growth: A potential link to the BRAFV600E mutation. <i>International Journal of Oncology</i> , 2016, 48, 1679-1687.	1.4	11

#	ARTICLE	IF	CITATIONS
37	Cellular apoptosis susceptibility (CAS) is linked to integrin β 1 and required for tumor cell migration and invasion in hepatocellular carcinoma (HCC). <i>Oncotarget</i> , 2016, 7, 22883-22892.	0.8	18
38	Implementation of systems theory in liver cancer research. <i>Hepatic Oncology</i> , 2015, 2, 9-11.	4.2	0
39	Concomitant expression of far upstream element (FUSE) binding protein (FBP) interacting repressor (FIR) and its splice variants induce migration and invasion of non-small cell lung cancer (NSCLC) cells. <i>Journal of Pathology</i> , 2015, 237, 390-401.	2.1	32
40	Curcumin effectively inhibits oncogenic NF- κ B signaling and restrains stemness features in liver cancer. <i>Journal of Hepatology</i> , 2015, 63, 661-669.	1.8	237
41	SKP2 cooperates with N-Ras or AKT to induce liver tumor development in mice. <i>Oncotarget</i> , 2015, 6, 2222-2234.	0.8	27
42	Insulin-like growth factor 2 mRNA-binding protein 1 (IGF2BP1) is an important protumorigenic factor in hepatocellular carcinoma. <i>Hepatology</i> , 2014, 59, 1900-1911.	3.6	155
43	Overexpression of far upstream element (FUSE) binding protein (FBP)-interacting repressor (FIR) supports growth of hepatocellular carcinoma. <i>Hepatology</i> , 2014, 60, 1241-1250.	3.6	39
44	Global alterations of DNA methylation in cholangiocarcinoma target the Wnt signaling pathway. <i>Hepatology</i> , 2014, 59, 544-554.	3.6	97
45	Human and Mouse VEGFA-Amplified Hepatocellular Carcinomas Are Highly Sensitive to Sorafenib Treatment. <i>Cancer Discovery</i> , 2014, 4, 730-743.	7.7	165
46	Prosurvival function of the cellular apoptosis susceptibility/importin-1 transport cycle is repressed by p53 in liver cancer. <i>Hepatology</i> , 2014, 60, 884-895.	3.6	29
47	Nuclear Expression of the Deubiquitinase CYLD Is Associated with Improved Survival in Human Hepatocellular Carcinoma. <i>PLoS ONE</i> , 2014, 9, e110591.	1.1	12
48	Downregulation of the activating NKp30 ligand B7-H6 by HDAC inhibitors impairs tumor cell recognition by NK cells. <i>Blood</i> , 2013, 122, 684-693.	0.6	109
49	Cytosolic and nuclear caspase-8 have opposite impact on survival after liver resection for hepatocellular carcinoma. <i>BMC Cancer</i> , 2013, 13, 532.	1.1	23
50	Posttranscriptional destabilization of the liver-specific long noncoding RNA HULC by the IGF2 mRNA-binding protein 1 (IGF2BP1). <i>Hepatology</i> , 2013, 58, 1703-1712.	3.6	208
51	Yes-Associated Protein Up-regulates Jagged-1 and Activates the NOTCH Pathway in Human Hepatocellular Carcinoma. <i>Gastroenterology</i> , 2013, 144, 1530-1542.e12.	0.6	278
52	Endothelial transdifferentiation in hepatocellular carcinoma: loss of Stabilin-2 expression in peritumorous liver correlates with increased survival. <i>Liver International</i> , 2013, 33, 1428-1440.	1.9	49
53	Endothelial plasticity governs the site-specific leukocyte recruitment in hepatocellular cancer. <i>International Journal of Cancer</i> , 2013, 133, 2372-2382.	2.3	4
54	Comparative Analysis of TGF- β 2/Smad Signaling Dependent Cytostasis in Human Hepatocellular Carcinoma Cell Lines. <i>PLoS ONE</i> , 2013, 8, e72252.	1.1	59

#	ARTICLE	IF	CITATIONS
55	Stathmin Regulates Keratinocyte Proliferation and Migration during Cutaneous Regeneration. PLoS ONE, 2013, 8, e75075.	1.1	16
56	Molecular Pathology of Liver Tumors. , 2013, , 43-63.		3
57	A Systems Biology Study on NF κ B Signaling in Primary Mouse Hepatocytes. Frontiers in Physiology, 2012, 3, 466.	1.3	9
58	Nuclear Pore Component Nup98 Is a Potential Tumor Suppressor and Regulates Posttranscriptional Expression of Select p53 Target Genes. Molecular Cell, 2012, 48, 799-810.	4.5	57
59	Transcriptional regulators in hepatocarcinogenesis â€œ Key integrators of malignant transformation. Journal of Hepatology, 2012, 57, 186-195.	1.8	20
60	Nuclear accumulation of seven in absentia homologueâ€2 supports motility and proliferation of liver cancer cells. International Journal of Cancer, 2012, 131, 2016-2026.	2.3	21
61	Insulin/IGF signaling drives cell proliferation in part via Yorkie/YAP. Developmental Biology, 2012, 367, 187-196.	0.9	126
62	Nuclear expression of the ubiquitin ligase seven in absentia homolog (SIAH)-1 induces proliferation and migration of liver cancer cells. Journal of Hepatology, 2011, 55, 1049-1057.	1.8	52
63	Strategies for hepatocellular carcinoma therapy and diagnostics: Lessons learned from high throughput and profiling approaches. Hepatology, 2011, 53, 2112-2121.	3.6	49
64	Expression of cyclooxygenase-2 (COX-2) in an advanced metastasized hypopharyngeal carcinoma and cultured tumor cells. Oral and Maxillofacial Surgery, 2010, 14, 53-57.	0.6	6
65	Down-regulation of tumor suppressor a kinase anchor protein 12 in human hepatocarcinogenesis by epigenetic mechanisms. Hepatology, 2010, 52, 2023-2033.	3.6	61
66	Lipid droplet-associated PAT-proteins show frequent and differential expression in neoplastic steatogenesis. Modern Pathology, 2010, 23, 480-492.	2.9	131
67	A Cellular View of Nf2 in Liver Homeostasis and Tumorigenesis. Developmental Cell, 2010, 19, 363-364.	3.1	5
68	The cyclin E regulator cullin 3 prevents mouse hepatic progenitor cells from becoming tumor-initiating cells. Journal of Clinical Investigation, 2010, 120, 3820-3833.	3.9	45
69	Coordinated Expression of Stathmin Family Members by Far Upstream Sequence Element-Binding Protein-1 Increases Motility in Nonâ€Small Cell Lung Cancer. Cancer Research, 2009, 69, 2234-2243.	0.4	85
70	Overexpression of far upstream element binding proteins: A mechanism regulating proliferation and migration in liver cancer cells. Hepatology, 2009, 50, 1130-1139.	3.6	92
71	S100A8 and S100A9 are novel nuclear factor kappa B target genes during malignant progression of murine and human liver carcinogenesis. Hepatology, 2009, 50, 1251-1262.	3.6	129
72	AP-1-Controlled Hepatocyte Growth Factor Activation Promotes Keratinocyte Migration via CEACAM1 and Urokinase Plasminogen Activator/Urokinase Plasminogen Receptor. Journal of Investigative Dermatology, 2009, 129, 1140-1148.	0.3	17

#	ARTICLE	IF	CITATIONS
73	Autocrine insulin-like growth factor-II stimulation of tumor cell migration is a progression step in human hepatocarcinogenesis. <i>Hepatology</i> , 2008, 48, 146-156.	3.6	100
74	Quantitative Analysis of Gene Expression Relative to 18S rRNA in Carcinoma Samples Using the LightCycler® Instrument and a SYBR GreenI-based Assay: Determining FAT10 mRNA Levels in Hepatocellular Carcinoma. <i>Methods in Molecular Biology</i> , 2008, 429, 59-72.	0.4	11
75	Reactivation of the insulin-like growth factor-II signaling pathway in human hepatocellular carcinoma. <i>World Journal of Gastroenterology</i> , 2008, 14, 1690.	1.4	67
76	Protumorigenic overexpression of stathmin/Op18 by gain-of-function mutation in p53 in human hepatocarcinogenesis. <i>Hepatology</i> , 2007, 46, 759-768.	3.6	103
77	Non-specific effects of siRNAs on tumor cells with implications on therapeutic applicability using RNA interference. <i>Pathology and Oncology Research</i> , 2007, 13, 84-90.	0.9	18
78	Ex vivo analysis of antineoplastic agents in precision-cut tissue slices of human origin: effects of cyclooxygenase-2 inhibition in hepatocellular carcinoma. <i>Liver International</i> , 2006, 26, 604-612.	1.9	23
79	Expression of epithelial cellular adhesion molecule (Ep-CAM) in chronic (necro-)inflammatory liver diseases and hepatocellular carcinoma. <i>Hepatology Research</i> , 2006, 34, 50-56.	1.8	22
80	The Insulin-Like Growth Factor (IGF) Signaling Pathway: Strategies for Successful Therapeutic Tasks in Cancer Treatment. <i>Current Cancer Therapy Reviews</i> , 2006, 2, 157-167.	0.2	5
81	Cyclooxygenase-2 Inhibition Induces Apoptosis Signaling via Death Receptors and Mitochondria in Hepatocellular Carcinoma. <i>Cancer Research</i> , 2006, 66, 7059-7066.	0.4	151
82	Cyclooxygenase-2 inhibitors suppress the growth of human hepatocellular carcinoma implants in nude mice. <i>Carcinogenesis</i> , 2004, 25, 1193-1199.	1.3	75
83	Factors of transforming growth factor β signalling are co-regulated in human hepatocellular carcinoma. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2004, 445, 589-596.	1.4	29
84	Molecular Profiling of Human Hepatocellular Carcinoma Defines Mutually Exclusive Interferon Regulation and Insulin-Like Growth Factor II Overexpression. <i>Cancer Research</i> , 2004, 64, 6058-6064.	0.4	119
85	Beta-catenin accumulation in the progression of human hepatocarcinogenesis correlates with loss of E-cadherin and accumulation of p53, but not with expression of conventional WNT-1 target genes. <i>Journal of Pathology</i> , 2003, 201, 250-259.	2.1	107
86	Molecular pathogenesis of human hepatocellular carcinoma. <i>Advances in Cancer Research</i> , 2002, 86, 67-112.	1.9	48
87	Proapoptotic and antiproliferative potential of selective cyclooxygenase-2 inhibitors in human liver tumor cells. <i>Hepatology</i> , 2002, 36, 885-894.	3.6	143
88	Keratinocyte-Derived Granulocyte-Macrophage Colony Stimulating Factor Accelerates Wound Healing: Stimulation of Keratinocyte Proliferation, Granulation Tissue Formation, and Vascularization. <i>Journal of Investigative Dermatology</i> , 2001, 117, 1382-1390.	0.3	142
89	The designer cytokine hyper-IL-6 mediates growth inhibition and GM-CSF-dependent rejection of B16 melanoma cells. <i>Oncogene</i> , 2001, 20, 972-979.	2.6	27