Apinya Jusakul

List of Publications by Year in descending order

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Δρινινα Ιμελαμι

#	Article	IF	CITATIONS
1	Therapeutic targeting of ARID1A and PI3K/AKT pathway alterations in cholangiocarcinoma. PeerJ, 2022, 10, e12750.	2.0	5
2	Serum Angiopoietin-Like Protein 4: A Potential Prognostic Biomarker for Prediction of Vascular Invasion and Lymph Node Metastasis in Cholangiocarcinoma Patients. Frontiers in Public Health, 2022, 10, 836985.	2.7	2
3	Inhibition of FGFR2 enhances chemosensitivity to gemcitabine in cholangiocarcinoma through the AKT/mTOR and EMT signaling pathways. Life Sciences, 2022, 296, 120427.	4.3	14
4	Promoter hypermethylation of early B cell factor 1 (EBF1) is associated with cholangiocarcinoma progression. Journal of Cancer, 2021, 12, 2673-2686.	2.5	4
5	Vitamin C supplementation reduces expression of circulating miR-451a in subjects with poorly controlled type 2 diabetes mellitus and high oxidative stress. PeerJ, 2021, 9, e10776.	2.0	9
6	Establishment of a Potential Serum Biomarker Panel for the Diagnosis and Prognosis of Cholangiocarcinoma Using Decision Tree Algorithms. Diagnostics, 2021, 11, 589.	2.6	10
7	Diagnostic and Prognostic Value of Circulating Cell-Free DNA for Cholangiocarcinoma. Diagnostics, 2021, 11, 999.	2.6	8
8	Circulating TGF-β1 as the potential epithelial mesenchymal transition-biomarker for diagnosis of cholangiocarcinoma. Journal of Gastrointestinal Oncology, 2020, 11, 304-318.	1.4	9
9	Lack of Targetable FGFR2 Fusions in Endemic Fluke-Associated Cholangiocarcinoma. JCO Global Oncology, 2020, 6, 628-638.	1.8	35
10	A combination of monosodium glutamate and high-fat and high-fructose diets increases the risk of kidney injury, gut dysbiosis and host-microbial co-metabolism. PLoS ONE, 2020, 15, e0231237.	2.5	18
11	Serum coiled‑coil domain containing 25 protein as a potential screening/diagnostic biomarker for cholangiocarcinoma. Oncology Letters, 2020, 19, 930-942.	1.8	5
12	<i>ARID1A</i> alterations and their clinical significance in cholangiocarcinoma. PeerJ, 2020, 8, e10464.	2.0	9
13	Current omics-based biomarkers for cholangiocarcinoma. Expert Review of Molecular Diagnostics, 2019, 19, 997-1005.	3.1	7
14	Epigenomic Promoter Alterations Amplify Gene Isoform and Immunogenic Diversity in Gastric Adenocarcinoma. Cancer Discovery, 2017, 7, 630-651.	9.4	48
15	Genome-scale mutational signatures of aflatoxin in cells, mice, and human tumors. Genome Research, 2017, 27, 1475-1486.	5.5	90
16	Establishment of cholangiocarcinoma cell lines from patients in the endemic area of liver fluke infection in Thailand. Tumor Biology, 2017, 39, 101042831772592.	1.8	27
17	Whole-Genome and Epigenomic Landscapes of Etiologically Distinct Subtypes of Cholangiocarcinoma. Cancer Discovery, 2017, 7, 1116-1135	9.4	637
18	Tissue Microbiome Profiling Identifies an Enrichment of Specific Enteric Bacteria in Opisthorchis viverrini Associated Cholangiocarcinoma. EBioMedicine, 2016, 8, 195-202.	6.1	94

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19	Genetics of Opisthorchis viverrini-related cholangiocarcinoma. Current Opinion in Gastroenterology, 2015, 31, 258-263.	2.3	45
20	Pathogenesis of cholangiocarcinoma: From genetics to signalling pathways. Bailliere's Best Practice and Research in Clinical Gastroenterology, 2015, 29, 233-244.	2.4	34
21	Association of CYP39A1, RUNX2 and Oxidized Alpha-1 Antitrypsin Expression in Relation to Cholangiocarcinoma Progression. Asian Pacific Journal of Cancer Prevention, 2015, 15, 10187-10192.	1.2	10
22	Anti-apoptotic phenotypes of cholestan-3β,5α,6β-triol-resistant human cholangiocytes: Characteristics contributing to the genesis of cholangiocarcinoma. Journal of Steroid Biochemistry and Molecular Biology, 2013, 138, 368-375.	2.5	20
23	Exome sequencing identifies distinct mutational patterns in liver fluke–related and non-infection-related bile duct cancers. Nature Genetics, 2013, 45, 1474-1478.	21.4	426
24	Expression of oxysterol binding protein isoforms in opisthorchiasis-associated cholangiocarcinoma: A potential molecular marker for tumor metastasis. Parasitology International, 2012, 61, 136-139.	1.3	28
25	Liver fluke-induced hepatic oxysterols stimulate DNA damage and apoptosis in cultured human cholangiocytes. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2012, 731, 48-57.	1.0	36
26	Identification of biliary bile acids in patients with benign biliary diseases, hepatocellular carcinoma and cholangiocarcinoma. Asian Pacific Journal of Cancer Prevention, 2012, 13 Suppl, 77-82.	1.2	13
27	Mechanisms of oxysterol-induced carcinogenesis. Lipids in Health and Disease, 2011, 10, 44.	3.0	69