

# Apinya Jusakul

## List of Publications by Year in descending order

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Version: 2024-02-01

27  
papers

1,713  
citations

623734

14  
h-index

526287

27  
g-index

28  
all docs

28  
docs citations

28  
times ranked

3185  
citing authors

#	ARTICLE	IF	CITATIONS
1	Whole-Genome and Epigenomic Landscapes of Etiologically Distinct Subtypes of Cholangiocarcinoma. <i>Cancer Discovery</i> , 2017, 7, 1116-1135.	9.4	637
2	Exome sequencing identifies distinct mutational patterns in liver fluke-related and non-infection-related bile duct cancers. <i>Nature Genetics</i> , 2013, 45, 1474-1478.	21.4	426
3	Tissue Microbiome Profiling Identifies an Enrichment of Specific Enteric Bacteria in <i>Opisthorchis viverrini</i> Associated Cholangiocarcinoma. <i>EBioMedicine</i> , 2016, 8, 195-202.	6.1	94
4	Genome-scale mutational signatures of aflatoxin in cells, mice, and human tumors. <i>Genome Research</i> , 2017, 27, 1475-1486.	5.5	90
5	Mechanisms of oxysterol-induced carcinogenesis. <i>Lipids in Health and Disease</i> , 2011, 10, 44.	3.0	69
6	Epigenomic Promoter Alterations Amplify Gene Isoform and Immunogenic Diversity in Gastric Adenocarcinoma. <i>Cancer Discovery</i> , 2017, 7, 630-651.	9.4	48
7	Genetics of <i>Opisthorchis viverrini</i> -related cholangiocarcinoma. <i>Current Opinion in Gastroenterology</i> , 2015, 31, 258-263.	2.3	45
8	Liver fluke-induced hepatic oxysterols stimulate DNA damage and apoptosis in cultured human cholangiocytes. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2012, 731, 48-57.	1.0	36
9	Lack of Targetable FGFR2 Fusions in Endemic Fluke-Associated Cholangiocarcinoma. <i>JCO Global Oncology</i> , 2020, 6, 628-638.	1.8	35
10	Pathogenesis of cholangiocarcinoma: From genetics to signalling pathways. <i>Bailliere's Best Practice and Research in Clinical Gastroenterology</i> , 2015, 29, 233-244.	2.4	34
11	Expression of oxysterol binding protein isoforms in opisthorchiasis-associated cholangiocarcinoma: A potential molecular marker for tumor metastasis. <i>Parasitology International</i> , 2012, 61, 136-139.	1.3	28
12	Establishment of cholangiocarcinoma cell lines from patients in the endemic area of liver fluke infection in Thailand. <i>Tumor Biology</i> , 2017, 39, 101042831772592.	1.8	27
13	Anti-apoptotic phenotypes of cholestan-3 $\beta$ ,5 $\alpha$ ,6 $\beta$ -triol-resistant human cholangiocytes: Characteristics contributing to the genesis of cholangiocarcinoma. <i>Journal of Steroid Biochemistry and Molecular Biology</i> , 2013, 138, 368-375.	2.5	20
14	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. <i>PLoS ONE</i> , 2020, 15, e0231237.	2.5	18
15	Inhibition of FGFR2 enhances chemosensitivity to gemcitabine in cholangiocarcinoma through the AKT/mTOR and EMT signaling pathways. <i>Life Sciences</i> , 2022, 296, 120427.	4.3	14
16	Identification of biliary bile acids in patients with benign biliary diseases, hepatocellular carcinoma and cholangiocarcinoma. <i>Asian Pacific Journal of Cancer Prevention</i> , 2012, 13 Suppl, 77-82.	1.2	13
17	Establishment of a Potential Serum Biomarker Panel for the Diagnosis and Prognosis of Cholangiocarcinoma Using Decision Tree Algorithms. <i>Diagnostics</i> , 2021, 11, 589.	2.6	10
18	Association of CYP39A1, RUNX2 and Oxidized Alpha-1 Antitrypsin Expression in Relation to Cholangiocarcinoma Progression. <i>Asian Pacific Journal of Cancer Prevention</i> , 2015, 15, 10187-10192.	1.2	10

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19	Circulating TGF- $\beta$ 1 as the potential epithelial mesenchymal transition-biomarker for diagnosis of cholangiocarcinoma. <i>Journal of Gastrointestinal Oncology</i> , 2020, 11, 304-318.	1.4	9
20	Vitamin C supplementation reduces expression of circulating miR-451a in subjects with poorly controlled type 2 diabetes mellitus and high oxidative stress. <i>PeerJ</i> , 2021, 9, e10776.	2.0	9
21	<i>ARID1A</i> alterations and their clinical significance in cholangiocarcinoma. <i>PeerJ</i> , 2020, 8, e10464.	2.0	9
22	Diagnostic and Prognostic Value of Circulating Cell-Free DNA for Cholangiocarcinoma. <i>Diagnostics</i> , 2021, 11, 999.	2.6	8
23	Current omics-based biomarkers for cholangiocarcinoma. <i>Expert Review of Molecular Diagnostics</i> , 2019, 19, 997-1005.	3.1	7
24	Serum coiled-coil domain containing 25 protein as a potential screening/diagnostic biomarker for cholangiocarcinoma. <i>Oncology Letters</i> , 2020, 19, 930-942.	1.8	5
25	Therapeutic targeting of ARID1A and PI3K/AKT pathway alterations in cholangiocarcinoma. <i>PeerJ</i> , 2022, 10, e12750.	2.0	5
26	Promoter hypermethylation of early B cell factor 1 (EBF1) is associated with cholangiocarcinoma progression. <i>Journal of Cancer</i> , 2021, 12, 2673-2686.	2.5	4
27	Serum Angiopoietin-Like Protein 4: A Potential Prognostic Biomarker for Prediction of Vascular Invasion and Lymph Node Metastasis in Cholangiocarcinoma Patients. <i>Frontiers in Public Health</i> , 2022, 10, 836985.	2.7	2