

Daniela Sia

List of Publications by Year in descending order

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Version: 2024-02-01

56
papers

8,599
citations

109321

35
h-index

144013

57
g-index

61
all docs

61
docs citations

61
times ranked

11400
citing authors

#	ARTICLE	IF	CITATIONS
1	Inflamed and non-inflamed classes of HCC: a revised immunogenomic classification. <i>Gut</i> , 2023, 72, 129-140.	12.1	90
2	Novel microenvironment-based classification of intrahepatic cholangiocarcinoma with therapeutic implications. <i>Gut</i> , 2023, 72, 736-748.	12.1	42
3	The portrait of liver cancer is shaped by mitochondrial genetics. <i>Cell Reports</i> , 2022, 38, 110254.	6.4	10
4	Novel insights into molecular and immune subtypes of biliary tract cancers. <i>Advances in Cancer Research</i> , 2022, , 167-199.	5.0	1
5	Transcriptomic profiling of a multiethnic pediatric NAFLD cohort reveals genes and pathways associated with disease. <i>Hepatology Communications</i> , 2022, 6, 1598-1610.	4.3	6
6	NOTCH-YAP1/TEAD-DNMT1 Axis Drives Hepatocyte Reprogramming Into Intrahepatic Cholangiocarcinoma. <i>Gastroenterology</i> , 2022, 163, 449-465.	1.3	23
7	Mutations in circulating tumor DNA predict primary resistance to systemic therapies in advanced hepatocellular carcinoma. <i>Oncogene</i> , 2021, 40, 140-151.	5.9	77
8	Cell of origin in biliary tract cancers and clinical implications. <i>JHEP Reports</i> , 2021, 3, 100226.	4.9	30
9	Molecular markers of response to anti-PD1 therapy in advanced hepatocellular carcinoma.. <i>Journal of Clinical Oncology</i> , 2021, 39, 4100-4100.	1.6	17
10	Promotion of cholangiocarcinoma growth by diverse cancer-associated fibroblast subpopulations. <i>Cancer Cell</i> , 2021, 39, 866-882.e11.	16.8	159
11	Transcriptomic characterization of cancer-testis antigens identifies MAGEA3 as a driver of tumor progression in hepatocellular carcinoma. <i>PLoS Genetics</i> , 2021, 17, e1009589.	3.5	15
12	Immunomodulatory Effects of Lenvatinib Plus Anti-Programmed Cell Death Protein 1 in Mice and Rationale for Patient Enrichment in Hepatocellular Carcinoma. <i>Hepatology</i> , 2021, 74, 2652-2669.	7.3	95
13	Molecular characterisation of hepatocellular carcinoma in patients with non-alcoholic steatohepatitis. <i>Journal of Hepatology</i> , 2021, 75, 865-878.	3.7	111
14	Copy-Number Alteration Burden Differentially Impacts Immune Profiles and Molecular Features of Hepatocellular Carcinoma. <i>Clinical Cancer Research</i> , 2020, 26, 6350-6361.	7.0	35
15	Molecular classification and therapeutic targets in extrahepatic cholangiocarcinoma. <i>Journal of Hepatology</i> , 2020, 73, 315-327.	3.7	164
16	Intratumoral heterogeneity and clonal evolution in liver cancer. <i>Nature Communications</i> , 2020, 11, 291.	12.8	230
17	Epigenetic footprint enables molecular risk stratification of hepatoblastoma with clinical implications. <i>Journal of Hepatology</i> , 2020, 73, 328-341.	3.7	82
18	Molecular predictors of prevention of recurrence in HCC with sorafenib as adjuvant treatment and prognostic factors in the phase 3 STORM trial. <i>Gut</i> , 2019, 68, 1065-1075.	12.1	195

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19	Molecular portrait of high alpha-fetoprotein in hepatocellular carcinoma: implications for biomarker-driven clinical trials. <i>British Journal of Cancer</i> , 2019, 121, 340-343.	6.4	62
20	An Immune Gene Expression Signature Associated With Development of Human Hepatocellular Carcinoma Identifies Mice That Respond to Chemopreventive Agents. <i>Gastroenterology</i> , 2019, 157, 1383-1397.e11.	1.3	62
21	Î²-Catenin Activation Promotes Immune Escape and Resistance to Anti-“PD-1 Therapy in Hepatocellular Carcinoma. <i>Cancer Discovery</i> , 2019, 9, 1124-1141.	9.4	498
22	Immune Exclusion-Wnt/CTNNB1 Class Predicts Resistance to Immunotherapies in HCC. <i>Clinical Cancer Research</i> , 2019, 25, 2021-2023.	7.0	152
23	Translating cancer genomics for precision oncology in biliary tract cancers. <i>Discovery Medicine</i> , 2019, 28, 255-265.	0.5	6
24	A pilot study of ultra-deep targeted sequencing of plasma DNA identifies driver mutations in hepatocellular carcinoma. <i>Oncogene</i> , 2018, 37, 3740-3752.	5.9	89
25	Molecular therapies and precision medicine for hepatocellular carcinoma. <i>Nature Reviews Clinical Oncology</i> , 2018, 15, 599-616.	27.6	1,308
26	Tumour initiating cells and IGF/FGF signalling contribute to sorafenib resistance in hepatocellular carcinoma. <i>Gut</i> , 2017, 66, 530-540.	12.1	161
27	Mixed hepatocellular cholangiocarcinoma tumors: Cholangiolocellular carcinoma is a distinct molecular entity. <i>Journal of Hepatology</i> , 2017, 66, 952-961.	3.7	120
28	Identification of an Immune-specific Class of Hepatocellular Carcinoma, Based on Molecular Features. <i>Gastroenterology</i> , 2017, 153, 812-826.	1.3	650
29	Liver Cancer Cell of Origin, Molecular Class, and Effects on Patient Prognosis. <i>Gastroenterology</i> , 2017, 152, 745-761.	1.3	838
30	Palbociclib (PD-0332991), a selective CDK4/6 inhibitor, restricts tumour growth in preclinical models of hepatocellular carcinoma. <i>Gut</i> , 2017, 66, 1286-1296.	12.1	198
31	Trunk mutational events present minimal intra- and inter-tumoral heterogeneity in hepatocellular carcinoma. <i>Journal of Hepatology</i> , 2017, 67, 1222-1231.	3.7	121
32	Translating ‘-omics’ results into precision medicine for hepatocellular carcinoma. <i>Nature Reviews Gastroenterology and Hepatology</i> , 2017, 14, 571-572.	17.8	28
33	Genetics of Hepatocellular Carcinoma: Risk Stratification, Clinical Outcome, and Implications for Therapy. <i>Digestive Disease Interventions</i> , 2017, 01, 055-065.	0.2	2
34	Advances in cholangiocarcinoma research: report from the third Cholangiocarcinoma Foundation Annual Conference. <i>Journal of Gastrointestinal Oncology</i> , 2016, 7, 819-827.	1.4	17
35	Epigenetic profiling to classify cancer of unknown primary: a multicentre, retrospective analysis. <i>Lancet Oncology</i> , The, 2016, 17, 1386-1395.	10.7	357
36	Molecular Pathogenesis and Targeted Therapies for Intrahepatic Cholangiocarcinoma. <i>Clinical Cancer Research</i> , 2016, 22, 291-300.	7.0	185

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37	YAP Inhibition Restores Hepatocyte Differentiation in Advanced HCC, Leading to Tumor Regression. <i>Cell Reports</i> , 2015, 10, 1692-1707.	6.4	213
38	Massive parallel sequencing uncovers actionable FGFR2-PPHLN1 fusion and ARAF mutations in intrahepatic cholangiocarcinoma. <i>Nature Communications</i> , 2015, 6, 6087.	12.8	240
39	Progenitor cell markers predict outcome of patients with hepatocellular carcinoma beyond Milan criteria undergoing liver transplantation. <i>Journal of Hepatology</i> , 2015, 63, 1368-1377.	3.7	64
40	The future of patient-derived tumor xenografts in cancer treatment. <i>Pharmacogenomics</i> , 2015, 16, 1671-1683.	1.3	43
41	Mutant IDH inhibits HNF-4 α to block hepatocyte differentiation and promote biliary cancer. <i>Nature</i> , 2014, 513, 110-114.	27.8	367
42	VEGF Signaling in Cancer Treatment. <i>Current Pharmaceutical Design</i> , 2014, 20, 2834-2842.	1.9	74
43	Sex bias occurrence of hepatocellular carcinoma in Poly7 molecular subclass is associated with EGFR. <i>Hepatology</i> , 2013, 57, 120-130.	7.3	52
44	Integrative Molecular Analysis of Intrahepatic Cholangiocarcinoma Reveals 2 Classes That Have Different Outcomes. <i>Gastroenterology</i> , 2013, 144, 829-840.	1.3	438
45	Reply to: "Network-based discovery of gene signature for vascular invasion prediction in HCC". <i>Journal of Hepatology</i> , 2012, 56, 1424-1425.	3.7	0
46	microRNAs: New ways to block tumor angiogenesis?. <i>Journal of Hepatology</i> , 2012, 57, 490-491.	3.7	9
47	Telomere loss in Philadelphia-negative hematopoiesis after successful treatment of chronic myeloid leukemia: Evidence for premature aging of the myeloid compartment. <i>Mechanisms of Ageing and Development</i> , 2012, 133, 479-488.	4.6	9
48	Combining Clinical, Pathology, and Gene Expression Data to Predict Recurrence of Hepatocellular Carcinoma. <i>Gastroenterology</i> , 2011, 140, 1501-1512.e2.	1.3	389
49	microRNAs and the MYC Network: A Major Piece in the Puzzle of Liver Cancer. <i>Gastroenterology</i> , 2011, 140, 2138-2140.	1.3	5
50	Signaling Pathways in Hepatocellular Carcinoma. <i>Oncology</i> , 2011, 81, 18-23.	1.9	39
51	Gene-expression signature of vascular invasion in hepatocellular carcinoma. <i>Journal of Hepatology</i> , 2011, 55, 1325-1331.	3.7	133
52	Human CD34+ cells engineered to express membrane-bound tumor necrosis factor-related apoptosis-inducing ligand target both tumor cells and tumor vasculature. <i>Blood</i> , 2010, 115, 2231-2240.	1.4	32
53	Cancer gene discovery in hepatocellular carcinoma. <i>Journal of Hepatology</i> , 2010, 52, 921-929.	3.7	173
54	A computational approach to compare microvessel distributions in tumors following antiangiogenic treatments. <i>Laboratory Investigation</i> , 2009, 89, 1063-1070.	3.7	12

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55	IFN- γ Enhances the Antimyeloma Activity of the Fully Human Anti- α -Human Leukocyte Antigen-DR Monoclonal Antibody 1D09C3. <i>Cancer Research</i> , 2007, 67, 3269-3275.	0.9	18
56	The Portrait of Liver Cancer is Shaped by Mitochondrial Genetics. <i>SSRN Electronic Journal</i> , 0, , .	0.4	0