

# Su Ting Tay

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8908637/publications.pdf>

Version: 2024-02-01

14  
papers

911  
citations

840776

11  
h-index

996975

15  
g-index

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15  
docs citations

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times ranked

1785  
citing authors

#	ARTICLE	IF	CITATIONS
1	Single-Cell Atlas of Lineage States, Tumor Microenvironment, and Subtype-Specific Expression Programs in Gastric Cancer. <i>Cancer Discovery</i> , 2022, 12, 670-691.	9.4	165
2	Chromatin Rewiring by Mismatch Repair Protein MSH2 Alters Cell Adhesion Pathways and Sensitivity to BET Inhibition in Gastric Cancer. <i>Cancer Research</i> , 2022, 82, 2538-2551.	0.9	7
3	Long-read transcriptome sequencing reveals abundant promoter diversity in distinct molecular subtypes of gastric cancer. <i>Genome Biology</i> , 2021, 22, 44.	8.8	46
4	â€œ3Gâ€• Trial: An RNA Editing Signature to Guide Gastric Cancer Chemotherapy. <i>Cancer Research</i> , 2021, 81, 2788-2798.	0.9	9
5	<i>HNF4Î±</i> pathway mapping identifies wild-type <i>IDH1</i> as a targetable metabolic node in gastric cancer. <i>Gut</i> , 2020, 69, 231-242.	12.1	27
6	Genomic and epigenomic EBF1 alterations modulate TERT expression in gastric cancer. <i>Journal of Clinical Investigation</i> , 2020, 130, 3005-3020.	8.2	12
7	Activation of Transforming Growth Factor Beta 1 Signaling in Gastric Cancer-associated Fibroblasts Increases Their Motility, via Expression of Rho GTPase 2, and Ability to Induce Invasiveness of Gastric Cancer Cells. <i>Gastroenterology</i> , 2017, 153, 191-204.e16.	1.3	158
8	Epigenomic Promoter Alterations Amplify Gene Isoform and Immunogenic Diversity in Gastric Adenocarcinoma. <i>Cancer Discovery</i> , 2017, 7, 630-651.	9.4	48
9	<i>VHL</i> Deficiency Drives Enhancer Activation of Oncogenes in Clear Cell Renal Cell Carcinoma. <i>Cancer Discovery</i> , 2017, 7, 1284-1305.	9.4	111
10	An intrinsic mechanism controls reactivation of neural stem cells by spindle matrix proteins. <i>Nature Communications</i> , 2017, 8, 122.	12.8	25
11	Epigenomic profiling of primary gastric adenocarcinoma reveals super-enhancer heterogeneity. <i>Nature Communications</i> , 2016, 7, 12983.	12.8	123
12	<i>SETD2</i> histone modifier loss in aggressive GI stromal tumours. <i>Gut</i> , 2016, 65, 1960-1972.	12.1	49
13	Exome-wide Sequencing Shows Low Mutation Rates and Identifies Novel Mutated Genes in Seminomas. <i>European Urology</i> , 2015, 68, 77-83.	1.9	56
14	Nanoscale chromatin profiling of gastric adenocarcinoma reveals cancer-associated cryptic promoters and somatically acquired regulatory elements. <i>Nature Communications</i> , 2014, 5, 4361.	12.8	72