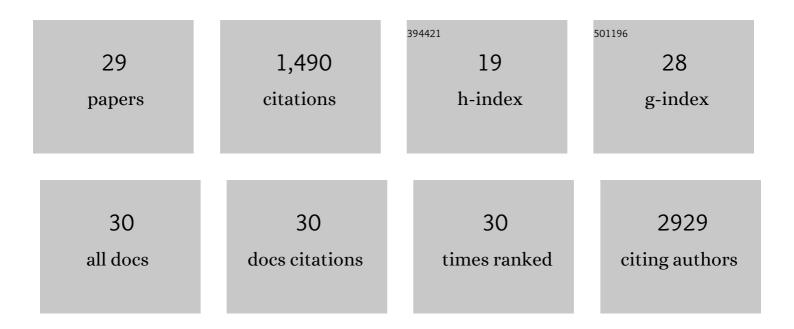
Hwee Tong Tan

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Serum autoantibodies as biomarkers for early cancer detection. FEBS Journal, 2009, 276, 6880-6904.	4.7	272
2	Membrane proteins and membrane proteomics. Proteomics, 2008, 8, 3924-3932.	2.2	257
3	Enhancing gold recovery from electronic waste via lixiviant metabolic engineering in Chromobacterium violaceum. Scientific Reports, 2013, 3, 2236.	3.3	100
4	Subcellular fractionation methods and strategies for proteomics. Proteomics, 2010, 10, 3935-3956.	2.2	91
5	Identification of Key Players for Colorectal Cancer Metastasis by iTRAQ Quantitative Proteomics Profiling of Isogenic SW480 and SW620 Cell Lines. Journal of Proteome Research, 2011, 10, 4373-4387.	3.7	72
6	Prognostic biomarkers for prediction of recurrence of hepatocellular carcinoma: Current status and future prospects. World Journal of Gastroenterology, 2014, 20, 3112.	3.3	72
7	Mining the Gastric Cancer Secretome: Identification of GRN as a Potential Diagnostic Marker for Early Gastric Cancer. Journal of Proteome Research, 2012, 11, 1759-1772.	3.7	71
8	Quantitative and Temporal Proteome Analysis of Butyrate-treated Colorectal Cancer Cells. Molecular and Cellular Proteomics, 2008, 7, 1174-1185.	3.8	66
9	Cancer proteomics. Mass Spectrometry Reviews, 2012, 31, 583-605.	5.4	60
10	Proteomic Analysis of Colorectal Cancer Metastasis: Stathmin-1 Revealed as a Player in Cancer Cell Migration and Prognostic Marker. Journal of Proteome Research, 2012, 11, 1433-1445.	3.7	51
11	Identification and Functional Validation of Caldesmon as a Potential Gastric Cancer Metastasis-associated Protein. Journal of Proteome Research, 2013, 12, 980-990.	3.7	50
12	Analysis of colorectal cancer glycoâ€secretome identifies laminin βâ€1 (LAMB1) as a potential serological biomarker for colorectal cancer. Proteomics, 2015, 15, 3905-3920.	2.2	45
13	Novel Proteomic Biomarker Panel for Prediction of Aggressive Metastatic Hepatocellular Carcinoma Relapse in Surgically Resectable Patients. Journal of Proteome Research, 2014, 13, 4833-4846.	3.7	40
14	Sieving through the cancer secretome. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2013, 1834, 2360-2371.	2.3	28
15	iTRAQ analysis of colorectal cancer cell lines suggests Drebrin (DBN1) is overexpressed during liver metastasis. Proteomics, 2014, 14, 1434-1443.	2.2	28
16	Unbiased Proteomic and Transcript Analyses Reveal that Stathmin-1 Silencing Inhibits Colorectal Cancer Metastasis and Sensitizes to 5-Fluorouracil Treatment. Molecular Cancer Research, 2014, 12, 1717-1728.	3.4	24
17	S-Nitrosylation of Divalent Metal Transporter 1 Enhances Iron Uptake to Mediate Loss of Dopaminergic Neurons and Motoric Deficit. Journal of Neuroscience, 2018, 38, 8364-8377.	3.6	24
18	Proteomics discovery of biomarkers for mitral regurgitation caused by mitral valve prolapse. Journal of Proteomics, 2013, 94, 337-345.	2.4	22

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19	2-D DIGE Analysis of Butyrate-Treated HCT-116 Cells after Enrichment with Heparin Affinity Chromatography. Journal of Proteome Research, 2006, 5, 1098-1106.	3.7	19
20	Unravelling the proteome of degenerative human mitral valves. Proteomics, 2015, 15, 2934-2944.	2.2	17
21	A comprehensive CHO SWATH-MS spectral library for robust quantitative profiling of 10,000 proteins. Scientific Data, 2020, 7, 263.	5.3	17
22	The prognostic value of the stem-like group in colorectal cancer using a panel of immunohistochemistry markers. Oncotarget, 2015, 6, 12763-12773.	1.8	14
23	Identification of Potential Pathways Involved in Induction of Apoptosis by Butyrate and 4-Benzoylbutyrate in HT29 Colorectal Cancer Cells. Journal of Proteome Research, 2012, 11, 6019-6029.	3.7	13
24	Proteomic analysis of human gastric juice: A shotgun approach. Proteomics, 2010, 10, 3928-3931.	2.2	12
25	Labelâ€Free Quantitative Phosphoproteomics Reveals Regulation of Vasodilatorâ€Stimulated Phosphoprotein upon Stathminâ€I Silencing in a Pair of Isogenic Colorectal Cancer Cell Lines. Proteomics, 2018, 18, e1700242.	2.2	10
26	iTRAQâ,,¢ Labeling Coupled with LC-MALDI Mass Spectrometry for Monitoring Temporal Response of Colorectal Cancer Cells to Butyrate Treatment. Methods in Molecular Biology, 2011, 716, 207-224.	0.9	6
27	Next Generation Proteomics for Clinical Biomarker Detection Using SWATH-MS. Methods in Molecular Biology, 2019, 1977, 3-15.	0.9	5
28	Global analysis of RNA-binding proteins identifies a positive feedback loop between LARP1 and MYC that promotes tumorigenesis. Cellular and Molecular Life Sciences, 2022, 79, 147.	5.4	4
29	Biomarkers for Recurrence of Hepatocellular Carcinoma. Biomarkers in Disease, 2017, , 167-191.	0.1	0