

Saiful Effendi Syafruddin

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

20 papers	219 citations	9 h-index	14 g-index
24 ext. papers	390 ext. citations	8.5 avg, IF	3.39 L-index

#	Paper	IF	Citations
20	A KLF6-driven transcriptional network links lipid homeostasis and tumour growth in renal carcinoma. <i>Nature Communications</i> , 2019 , 10, 1152	17.4	36
19	MicroRNA-200c and microRNA-31 regulate proliferation, colony formation, migration and invasion in serous ovarian cancer. <i>Journal of Ovarian Research</i> , 2015 , 8, 56	5.5	35
18	Revisiting the Roles of Pro-Metastatic EpCAM in Cancer. <i>Biomolecules</i> , 2020 , 10,	5.9	24
17	NF- κ B-Dependent Lymphoid Enhancer Co-option Promotes Renal Carcinoma Metastasis. <i>Cancer Discovery</i> , 2018 , 8, 850-865	24.4	23
16	Two Sides of the Same Coin: The Roles of KLF6 in Physiology and Pathophysiology. <i>Biomolecules</i> , 2020 , 10,	5.9	15
15	VHL-Mediated Regulation of CHCHD4 and Mitochondrial Function. <i>Frontiers in Oncology</i> , 2018 , 8, 388	5.3	14
14	CD44: A Multifunctional Mediator of Cancer Progression.. <i>Biomolecules</i> , 2021 , 11,	5.9	13
13	Molecular characterization of serous ovarian carcinoma using a multigene next generation sequencing cancer panel approach. <i>BMC Research Notes</i> , 2014 , 7, 805	2.3	12
12	Recent progress in mass spectrometry-based strategies for elucidating protein-protein interactions. <i>Cellular and Molecular Life Sciences</i> , 2021 , 78, 5325-5339	10.3	12
11	Predicting type 2 diabetes using genetic and environmental risk factors in a multi-ethnic Malaysian cohort. <i>Public Health</i> , 2017 , 149, 31-38	4	7
10	Characterizing the genetic risk for Type 2 diabetes in a Malaysian multi-ethnic cohort. <i>Diabetic Medicine</i> , 2015 , 32, 1377-84	3.5	6
9	Secretion of pro-oncogenic AGR2 protein in cancer. <i>Heliyon</i> , 2020 , 6, e05000	3.6	6
8	Molecular Characterization of Somatic Alterations in DukesVB and C Colorectal Cancers by Targeted Sequencing. <i>Frontiers in Pharmacology</i> , 2017 , 8, 465	5.6	4
7	Epigenome-Wide DNA Methylation Profiling in Colorectal Cancer and Normal Adjacent Colon Using Infinium Human Methylation 450K.. <i>Diagnostics</i> , 2022 , 12,	3.8	3
6	Integration of RNA-Seq and proteomics data identifies glioblastoma multiforme surfaceome signature. <i>BMC Cancer</i> , 2021 , 21, 850	4.8	3
5	Targeted Next-Generation Sequencing Identifies Actionable Targets in Estrogen Receptor Positive and Estrogen Receptor Negative Endometriod Endometrial Cancer. <i>Frontiers in Pharmacology</i> , 2018 , 9, 750	5.6	2
4	Reverting TP53 Mutation in Breast Cancer Cells: Prime Editing Workflow and Technical Considerations. <i>Cells</i> , 2022 , 11, 1612	7.9	1

3	Endogenous HIF2A reporter systems for high-throughput functional screening. <i>Scientific Reports</i> , 2018 , 8, 12063	4.9	○
2	CRISPR Gene-Editing Models Geared Toward Therapy for Hereditary and Developmental Neurological Disorders. <i>Frontiers in Pediatrics</i> , 2021 , 9, 592571	3.4	○
1	Short open reading frames (sORFs) and microproteins: an update on their identification and validation measures.. <i>Journal of Biomedical Science</i> , 2022 , 29, 19	13.3	○