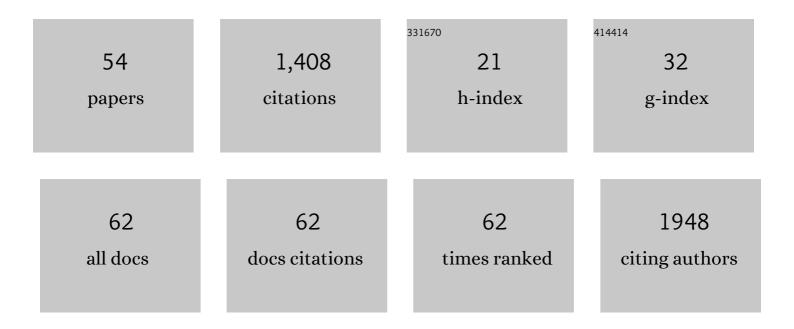
Darawan Rinchai

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1845618/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Fasting-Mimicking Diet Is Safe and Reshapes Metabolism and Antitumor Immunity in Patients with Cancer. Cancer Discovery, 2022, 12, 90-107.	9.4	124
2	Interleukin 10 inhibits pro-inflammatory cytokine responses and killing of Burkholderia pseudomallei. Scientific Reports, 2017, 7, 42791.	3.3	63
3	Flt3 ligand augments immune responses to anti-DEC-205-NY-ESO-1 vaccine through expansion of dendritic cell subsets. Nature Cancer, 2020, 1, 1204-1217.	13.2	58
4	Oncogenic states dictate the prognostic and predictive connotations of intratumoral immune response. , 2020, 8, e000617.		57
5	Exploiting the Burkholderia pseudomallei Acute Phase Antigen BPSL2765 for Structure-Based Epitope Discovery/Design in Structural Vaccinology. Chemistry and Biology, 2013, 20, 1147-1156.	6.0	50
6	A Structure-Based Strategy for Epitope Discovery in Burkholderia pseudomallei OppA Antigen. Structure, 2013, 21, 167-175.	3.3	49
7	An interactive web application for the dissemination of human systems immunology data. Journal of Translational Medicine, 2015, 13, 196.	4.4	49
8	Glibenclamide reduces pro-inflammatory cytokine production by neutrophils of diabetes patients in response to bacterial infection. Scientific Reports, 2013, 3, 3363.	3.3	47
9	Superoxide dismutase C is required for intracellular survival and virulence of Burkholderia pseudomallei. Microbiology (United Kingdom), 2011, 157, 2392-2400.	1.8	46
10	T Cell Immunity to the Alkyl Hydroperoxide Reductase of <i>Burkholderia pseudomallei</i> : A Correlate of Disease Outcome in Acute Melioidosis. Journal of Immunology, 2015, 194, 4814-4824.	0.8	44
11	Three Copies of Four Interferon Receptor Genes Underlie a Mild Type I Interferonopathy in Down Syndrome. Journal of Clinical Immunology, 2020, 40, 807-819.	3.8	44
12	Sequence- and Structure-Based Immunoreactive Epitope Discovery for Burkholderia pseudomallei Flagellin. PLoS Neglected Tropical Diseases, 2015, 9, e0003917.	3.0	40
13	Production of interleukinâ€⊋7 by human neutrophils regulates their function during bacterial infection. European Journal of Immunology, 2012, 42, 3280-3290.	2.9	37
14	A transcriptomic reporter assay employing neutrophils to measure immunogenic activity of septic patients' plasma. Journal of Translational Medicine, 2014, 12, 65.	4.4	34
15	Long-Chain Acyl-CoA Synthetase 1 Role in Sepsis and Immunity: Perspectives From a Parallel Review of Public Transcriptome Datasets and of the Literature. Frontiers in Immunology, 2019, 10, 2410.	4.8	33
16	Blood Interferon Signatures Putatively Link Lack of Protection Conferred by the RTS,S Recombinant Malaria Vaccine to an Antigen-specific IgE Response. F1000Research, 2015, 4, 919.	1.6	33
17	Annexin A3 in sepsis: novel perspectives from an exploration of public transcriptome data. Immunology, 2020, 161, 291-302.	4.4	32
18	Programmed Death Ligand 1 onBurkholderia pseudomallei–Infected Human Polymorphonuclear Neutrophils Impairs T Cell Functions. Journal of Immunology, 2015, 194, 4413-4421.	0.8	29

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#	Article	IF	CITATIONS
19	Development of a fixed module repertoire for the analysis and interpretation of blood transcriptome data. Nature Communications, 2021, 12, 4385.	12.8	29
20	CD4+ T Cell Epitopes of FliC Conserved between Strains of <i>Burkholderia</i> : Implications for Vaccines against Melioidosis and Cepacia Complex in Cystic Fibrosis. Journal of Immunology, 2014, 193, 6041-6049.	0.8	27
21	Macroautophagy is essential for killing of intracellular <i>Burkholderia pseudomallei</i> in human neutrophils. Autophagy, 2015, 11, 748-755.	9.1	27
22	CDX2 as a Prognostic Biomarker in Colon Cancer. New England Journal of Medicine, 2016, 374, 2182-2184.	27.0	23
23	Definition of erythroid cellâ€positive blood transcriptome phenotypes associated with severe respiratory syncytial virus infection. Clinical and Translational Medicine, 2020, 10, e244.	4.0	22
24	<i>Burkholderia pseudomallei</i> Proteins Presented by Monocyte-Derived Dendritic Cells Stimulate Human Memory T Cells <i>In Vitro</i> . Infection and Immunity, 2011, 79, 305-313.	2.2	21
25	A curated compendium of monocyte transcriptome datasets of relevance to human monocyte immunobiology research. F1000Research, 2016, 5, 291.	1.6	20
26	A Neutrophil-Driven Inflammatory Signature Characterizes the Blood Transcriptome Fingerprint of Psoriasis. Frontiers in Immunology, 2020, 11, 587946.	4.8	19
27	Emerging dynamics pathways of response and resistance to PD-1 and CTLA-4 blockade: tackling uncertainty by confronting complexity. Journal of Experimental and Clinical Cancer Research, 2021, 40, 74.	8.6	19
28	Increased abundance of ADAM9 transcripts in the blood is associated with tissue damage. F1000Research, 2015, 4, 89.	1.6	19
29	Blood Interferon Signatures Putatively Link Lack of Protection Conferred by the RTS,S Recombinant Malaria Vaccine to an Antigen-specific IgE Response. F1000Research, 2015, 4, 919.	1.6	19
30	BloodGen3Module: blood transcriptional module repertoire analysis and visualization using R. Bioinformatics, 2021, 37, 2382-2389.	4.1	18
31	Finger stick blood collection for gene expression profiling and storage of tempus blood RNA tubes. F1000Research, 2016, 5, 1385.	1.6	17
32	Transketolase and vitamin B1 influence on ROS-dependent neutrophil extracellular traps (NETs) formation. PLoS ONE, 2019, 14, e0221016.	2.5	16
33	Finger stick blood collection for gene expression profiling and storage of tempus blood RNA tubes. F1000Research, 2016, 5, 1385.	1.6	16
34	Glibenclamide impairs responses of neutrophils against Burkholderia pseudomallei by reduction of intracellular glutathione. Scientific Reports, 2016, 6, 34794.	3.3	15
35	Immune Control of Burkholderia pseudomallei––Common, High-Frequency T-Cell Responses to a Broad Repertoire of Immunoprevalent Epitopes. Frontiers in Immunology, 2018, 9, 484.	4.8	15
36	Using â€~collective omics data' for biomedical research training. Immunology, 2018, 155, 18-23.	4.4	15

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#	Article	IF	CITATIONS
37	Abundance of ADAM9 transcripts increases in the blood in response to tissue damage. F1000Research, 2015, 4, 89.	1.6	15
38	A collection of annotated and harmonized human breast cancer transcriptome datasets, including immunologic classification. F1000Research, 2017, 6, 296.	1.6	14
39	SLFN11 captures cancer-immunity interactions associated with platinum sensitivity in high-grade serous ovarian cancer. JCl Insight, 2021, 6, .	5.0	14
40	Prospective validation study of prognostic biomarkers to predict adverse outcomes in patients with COVID-19: a study protocol. BMJ Open, 2021, 11, e044497.	1.9	14
41	A collection of annotated and harmonized human breast cancer transcriptome datasets, including immunologic classification. F1000Research, 2017, 6, 296.	1.6	14
42	A modular framework for the development of targeted Covid-19 blood transcript profiling panels. Journal of Translational Medicine, 2020, 18, 291.	4.4	13
43	The immune landscape of solid pediatric tumors. Journal of Experimental and Clinical Cancer Research, 2022, 41, .	8.6	13
44	Network-based identification of key master regulators associated with an immune-silent cancer phenotype. Briefings in Bioinformatics, 2021, 22, .	6.5	11
45	A proteasome inhibitor produced by Burkholderia pseudomallei modulates intracellular growth. Microbial Pathogenesis, 2017, 107, 175-180.	2.9	7
46	Myeloid Cells Are Enriched in Tonsillar Crypts, Providing Insight into the Viral Tropism of Human Papillomavirus. American Journal of Pathology, 2021, 191, 1774-1786.	3.8	7
47	Transcriptomic profile investigations highlight a putative role for NUDT16 in sepsis. Journal of Cellular and Molecular Medicine, 2022, 26, 1714-1721.	3.6	5
48	A compendium of monocyte transcriptome datasets to foster biomedical knowledge discovery. F1000Research, 2016, 5, 291.	1.6	4
49	Integrated transcriptionalâ€phenotypic analysis captures systemic immunomodulation following antiangiogenic therapy in renal cell carcinoma patients. Clinical and Translational Medicine, 2021, 11, e434.	4.0	3
50	SysInflam HuDB, a Web Resource for Mining Human Blood Cells Transcriptomic Data Associated with Systemic Inflammatory Responses to Sepsis. Journal of Immunology, 2021, 207, 2195-2202.	0.8	3
51	Organizing gene literature retrieval,Âprofiling, and visualization training workshops for early career researchers. F1000Research, 0, 10, 275.	1.6	2
52	Mapping the Road of Gvhd and GVT: A Longitudinal Study of Immune-Transcriptome Signatures As Novel Approach to Solve Post-Allogeneic Hematopoietic Cell Transplantation Dilemmas. Blood, 2019, 134, 4550-4550.	1.4	1
53	A Web-Based Systems Immunology Toolkit Allows the Visualization and Analysis of Public Collective Data to Decipher Immunity in Early Life. , 2016, , .		0
54	Big Data as the Foundation of a Novel Training Platform for Biomedical Researchers in Qatar. , 2016, , .		0

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