Siew Cheng Wong

List of Publications by Year
in descending order

15	2,379	687363 13	940533 16
papers	citations	h-index	g-index

16	16	16	4953
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Increased monocyteâ€platelet aggregates and monocyteâ€endothelial adhesion in healthy individuals with vitamin D deficiency. FASEB Journal, 2020, 34, 11133-11142.	0.5	17
2	Resveratrol attenuates TLR-4 mediated inflammation and elicits therapeutic potential in models of sepsis. Scientific Reports, 2020, 10, 18837.	3.3	14
3	Macrophage polarisation associated with atherosclerosis differentially affects their capacity to handle lipids. Atherosclerosis, 2020, 305, 10-18.	0.8	19
4	A Novel, Five-Marker Alternative to CD16â€“CD14 Gating to Identify the Three Human Monocyte Subsets. Frontiers in Immunology, 2019, 10, 1761.	4.8	77
5	Targeting immune cells for cancer therapy. Redox Biology, 2019, 25, 101174.	9.0	151
6	The pro-inflammatory phenotype of the human non-classical monocyte subset is attributed to senescence. Cell Death and Disease, 2018, 9, 266.	6.3	169
7	Multiplexed Label-Free Fractionation of Peripheral Blood Mononuclear Cells for Identification of Monocyteâ€“Platelet Aggregates. Analytical Chemistry, 2018, 90, 14535-14542.	6.5	15
8	Checkpoint blockade immunotherapy reshapes the high-dimensional phenotypic heterogeneity of murine intratumoural neoantigen-specific CD8+ T cells. Nature Communications, 2017, 8, 562.	12.8	101
9	Warburg metabolism in tumor-conditioned macrophages promotes metastasis in human pancreatic ductal adenocarcinoma. OncoImmunology, 2016, 5, e1191731.	4.6	178
10	MicroRNA-mediated immune modulation as a therapeutic strategy in host-implant integration. Advanced Drug Delivery Reviews, 2015, 88, 92-107.	13.7	17
11	Micro<scp>RNA</scp> expression profiling of human blood monocyte subsets highlights functional differences. Immunology, 2015, 145, 404-416.	4.4	34
12	The three human monocyte subsets: implications for health and disease. Immunologic Research, 2012, 53, 41-57.	2.9	577
13	Gene expression profiling reveals the defining features of the classical, intermediate, and nonclassical human monocyte subsets. Blood, 2011, 118, e16-e31.	1.4	873
14	Study of monocyte membrane proteome perturbation during lipopolysaccharideâ€induced tolerance using iTRAQâ€based quantitative proteomic approach. Proteomics, 2010, 10, 2780-2789.	2.2	45
15	Identification of Novel Functional Differences in Monocyte Subsets Using Proteomic and Transcriptomic Methods. Journal of Proteome Research, 2009, 8, 4028-4038.	3.7	89