## Michael Poidinger

List of Publications by Year in descending order

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Version: 2024-02-01

91 13,496 50 papers citations h-index

h-index g-index

93
25008
times ranked citing authors

90

93 all docs 93 docs citations

#	Article	IF	CITATIONS
1	Bystander CD8+ T cells are abundant and phenotypically distinct in human tumour infiltrates. Nature, 2018, 557, 575-579.	27.8	942
2	C-Myb+ Erythro-Myeloid Progenitor-Derived Fetal Monocytes Give Rise to Adult Tissue-Resident Macrophages. Immunity, 2015, 42, 665-678.	14.3	847
3	IRF4 Transcription Factor-Dependent CD11b+ Dendritic Cells in Human and Mouse Control Mucosal IL-17 Cytokine Responses. Immunity, 2013, 38, 970-983.	14.3	703
4	Unsupervised High-Dimensional Analysis Aligns Dendritic Cells across Tissues and Species. Immunity, 2016, 45, 669-684.	14.3	683
5	Two distinct interstitial macrophage populations coexist across tissues in specific subtissular niches. Science, 2019, 363, .	12.6	676
6	Human Tissues Contain CD141hi Cross-Presenting Dendritic Cells with Functional Homology to Mouse CD103+ Nonlymphoid Dendritic Cells. Immunity, 2012, 37, 60-73.	14.3	643
7	Cistrome: an integrative platform for transcriptional regulation studies. Genome Biology, 2011, 12, R83.	9.6	598
8	RNA-Seq Signatures Normalized by mRNA Abundance Allow Absolute Deconvolution of Human Immune Cell Types. Cell Reports, 2019, 26, 1627-1640.e7.	6.4	590
9	Identification of cDC1- and cDC2-committed DC progenitors reveals early lineage priming at the common DC progenitor stage in the bone marrow. Nature Immunology, 2015, 16, 718-728.	14.5	475
10	Mapping the human DC lineage through the integration of high-dimensional techniques. Science, 2017, 356, .	12.6	429
11	Metformin as adjunct antituberculosis therapy. Science Translational Medicine, 2014, 6, 263ra159.	12.4	404
12	High-dimensional analysis of the murine myeloid cell system. Nature Immunology, 2014, 15, 1181-1189.	14.5	349
13	Human Monocytes Undergo Functional Re-programming during Sepsis Mediated by Hypoxia-Inducible Factor-1α. Immunity, 2015, 42, 484-498.	14.3	340
14	Cytofkit: A Bioconductor Package for an Integrated Mass Cytometry Data Analysis Pipeline. PLoS Computational Biology, 2016, 12, e1005112.	3.2	302
15	An outbreak of Japanese encephalitis in the Torres Strait, Australia, 1995. Medical Journal of Australia, 1996, 165, 256-260.	1.7	298
16	Human Dermal CD14 + Cells Are a Transient Population of Monocyte-Derived Macrophages. Immunity, 2014, 41, 465-477.	14.3	256
17	Induced-Pluripotent-Stem-Cell-Derived Primitive Macrophages Provide a Platform for Modeling Tissue-Resident Macrophage Differentiation and Function. Immunity, 2017, 47, 183-198.e6.	14.3	245
18	Molecular Profiling Reveals a Tumor-Promoting Phenotype of Monocytes and Macrophages in Human Cancer Progression. Immunity, 2014, 41, 815-829.	14.3	240

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19	Hyaluronan Receptor LYVE-1-Expressing Macrophages Maintain Arterial Tone through Hyaluronan-Mediated Regulation of Smooth Muscle Cell Collagen. Immunity, 2018, 49, 326-341.e7.	14.3	235
20	Human fetal dendritic cells promote prenatal T-cell immune suppression through arginase-2. Nature, 2017, 546, 662-666.	27.8	199
21	Experimental evolution of a fungal pathogen into a gut symbiont. Science, 2018, 362, 589-595.	12.6	184
22	High Mitochondrial Respiration and Glycolytic Capacity Represent a Metabolic Phenotype of Human Tolerogenic Dendritic Cells. Journal of Immunology, 2015, 194, 5174-5186.	0.8	183
23	flowAl: automatic and interactive anomaly discerning tools for flow cytometry data. Bioinformatics, 2016, 32, 2473-2480.	4.1	166
24	Gene Essentiality Is a Quantitative Property Linked to Cellular Evolvability. Cell, 2015, 163, 1388-1399.	28.9	146
25	IgG1 memory B cells keep the memory of IgE responses. Nature Communications, 2017, 8, 641.	12.8	143
26	Mapping the Diversity of Follicular Helper T Cells in Human Blood and Tonsils Using High-Dimensional Mass Cytometry Analysis. Cell Reports, 2015, 11, 1822-1833.	6.4	140
27	The distinctive germinal center phase of IgE+ B lymphocytes limits their contribution to the classical memory response. Journal of Experimental Medicine, 2013, 210, 2755-2771.	8.5	139
28	Molecular Characterization of the Japanese Encephalitis Serocomplex of the Flavivirus Genus. Virology, 1996, 218, 417-421.	2.4	126
29	The Relationships between West Nile and Kunjin Viruses. Emerging Infectious Diseases, 2001, 7, 697-705.	4.3	126
30	Cell Specific eQTL Analysis without Sorting Cells. PLoS Genetics, 2015, 11, e1005223.	3.5	115
31	CXCR4 identifies transitional bone marrow premonocytes that replenish the mature monocyte pool for peripheral responses. Journal of Experimental Medicine, 2016, 213, 2293-2314.	8.5	108
32	Host sirtuin 1 regulates mycobacterial immunopathogenesis and represents a therapeutic target against tuberculosis. Science Immunology, 2017, $2$ , .	11.9	104
33	Functionally diverse human T cells recognize non-microbial antigens presented by MR1. ELife, 2017, 6, .	6.0	100
34	An Unbiased Approach to Identifying Tau Kinases That Phosphorylate Tau at Sites Associated with Alzheimer Disease. Journal of Biological Chemistry, 2013, 288, 23331-23347.	3.4	99
35	Ubiquitin-conjugating enzyme Ubc13 controls breast cancer metastasis through a TAK1-p38 MAP kinase cascade. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 13870-13875.	7.1	99
36	Rational Design of a Live Attenuated Dengue Vaccine: 2′-O-Methyltransferase Mutants Are Highly Attenuated and Immunogenic in Mice and Macaques. PLoS Pathogens, 2013, 9, e1003521.	4.7	98

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37	The Relationships between West Nile and Kunjin Viruses. Emerging Infectious Diseases, 2001, 7, 697-705.	4.3	90
38	CD103+ Dendritic Cells Control Th17 Cell Function in the Lung. Cell Reports, 2015, 12, 1789-1801.	6.4	89
39	Enhancers Are Major Targets for Murine Leukemia Virus Vector Integration. Journal of Virology, 2014, 88, 4504-4513.	3.4	88
40	Selective Susceptibility of Human Skin Antigen Presenting Cells to Productive Dengue Virus Infection. PLoS Pathogens, 2014, 10, e1004548.	4.7	80
41	The tumour microenvironment creates a niche for the self-renewal of tumour-promoting macrophages in colon adenoma. Nature Communications, 2018, 9, 582.	12.8	76
42	Mpath maps multi-branching single-cell trajectories revealing progenitor cell progression during development. Nature Communications, 2016, 7, 11988.	12.8	67
43	Two contiguous outbreaks of dengue type 2 in north Queensland. Medical Journal of Australia, 1998, 168, 221-225.	1.7	66
44	Advantages of meta-total RNA sequencing (MeTRS) over shotgun metagenomics and amplicon-based sequencing in the profiling of complex microbial communities. Npj Biofilms and Microbiomes, 2018, 4, 2.	6.4	65
45	Plasmablasts During Acute Dengue Infection Represent a Small Subset of a Broader Virus-specific Memory B Cell Pool. EBioMedicine, 2016, 12, 178-188.	6.1	62
46	Neutrophils Self-Regulate Immune Complex-Mediated Cutaneous Inflammation through CXCL2. Journal of Investigative Dermatology, 2016, 136, 416-424.	0.7	62
47	Protumoral role of monocytes in human B-cell precursor acute lymphoblastic leukemia: involvement of the chemokine CXCL10. Blood, 2012, 119, 227-237.	1.4	59
48	Sequence determinants of innate immune activation by short interfering RNAs. BMC Immunology, 2009, 10, 40.	2.2	57
49	Multifactorial heterogeneity of virus-specific T cells and association with the progression of human chronic hepatitis B infection. Science Immunology, 2019, 4, .	11.9	57
50	Involvement of GABA Transporters in Atropine-Treated Myopic Retina As Revealed by iTRAQ Quantitative Proteomics. Journal of Proteome Research, 2014, 13, 4647-4658.	3.7	56
51	Mapping of γ/δT cells reveals Vδ2+T cells resistance to senescence. EBioMedicine, 2019, 39, 44-58.	6.1	54
52	A Subset of Type I Conventional Dendritic Cells Controls Cutaneous Bacterial Infections through VEGFα-Mediated Recruitment of Neutrophils. Immunity, 2019, 50, 1069-1083.e8.	14.3	50
53	RNA sensing by conventional dendritic cells is central to the development of lupus nephritis. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E6195-204.	7.1	49
54	The Transcriptional Stress Response of <i>Candida albicans</i> to Weak Organic Acids. G3: Genes, Genomes, Genetics, 2015, 5, 497-505.	1.8	46

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55	TLR7 and TLR9 ligands regulate antigen presentation by macrophages. International Immunology, 2016, 28, 223-232.	4.0	43
56	$\hat{l}^2$ -glucan Exposure on the Fungal Cell Wall Tightly Correlates with Competitive Fitness of Candida Species in the Mouse Gastrointestinal Tract. Frontiers in Cellular and Infection Microbiology, 2016, 6, 186.	3.9	41
57	Human Regulatory B Cells Combine Phenotypic and Genetic Hallmarks with a Distinct Differentiation Fate. Journal of Immunology, 2014, 193, 2258-2266.	0.8	40
58	Optimal cellular preservation for high dimensional flow cytometric analysis of multicentre trials. Journal of Immunological Methods, 2012, 385, 79-89.	1.4	38
59	Genetic Stability Among Temporally and Geographically Diverse Isolates of Barmah Forest Virus. American Journal of Tropical Medicine and Hygiene, 1997, 57, 230-234.	1.4	38
60	NFATc2 mediates epigenetic modification of dendritic cell cytokine and chemokine responses to dectin-1 stimulation. Nucleic Acids Research, 2015, 43, 836-847.	14.5	35
61	Micro <scp>RNA</scp> expression profiling of human blood monocyte subsets highlights functional differences. Immunology, 2015, 145, 404-416.	4.4	34
62	Functional variants of 17q12-21 are associated with allergic asthma but not allergic rhinitis. Journal of Allergy and Clinical Immunology, 2016, 137, 758-766.e3.	2.9	34
63	Influenza Vaccine-Induced Antibody Responses Are Not Impaired by Frailty in the Community-Dwelling Elderly With Natural Influenza Exposure. Frontiers in Immunology, 2018, 9, 2465.	4.8	34
64	Automated Identification of Core Regulatory Genes in Human Gene Regulatory Networks. PLoS Computational Biology, 2015, 11, e1004504.	3.2	33
65	Induction of Human T-cell and Cytokine Responses Following Vaccination with a Novel Influenza Vaccine. Scientific Reports, 2018, 8, 18007.	3.3	33
66	Systematic characterization of basophil anergy. Allergy: European Journal of Allergy and Clinical Immunology, 2017, 72, 373-384.	5.7	26
67	Enhanced Neutralizing Antibody Titers and Th1 Polarization from a Novel Escherichia coli Derived Pandemic Influenza Vaccine. PLoS ONE, 2013, 8, e76571.	2.5	25
68	Circulating CD1c+ myeloid dendritic cells are potential precursors to LCH lesion CD1a+CD207+ cells. Blood Advances, 2020, 4, 87-99.	5.2	25
69	Genome-wide analysis of the genetic regulation of gene expression in human neutrophils. Nature Communications, 2015, 6, 7971.	12.8	23
70	NLRP10 Enhances CD4+ T-Cell-Mediated IFN $\hat{I}^3$ Response via Regulation of Dendritic Cell-Derived IL-12 Release. Frontiers in Immunology, 2017, 8, 1462.	4.8	21
71	Calcium and Calcineurin-NFAT Signaling Regulate Granulocyte-Monocyte Progenitor Cell Cycle via Flt3-L. Stem Cells, 2014, 32, 3232-3244.	3.2	20
72	Dengue Serotype Cross-Reactive, Anti-E Protein Antibodies Confound Specific Immune Memory for 1 Year after Infection. Frontiers in Immunology, 2014, 5, 388.	4.8	18

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73	Visualization of bone marrow monocyte mobilization using <i>Cx3cr1gfp/+Flt3Lâ^'/â^'</i> reporter mouse by multiphoton intravital microscopy. Journal of Leukocyte Biology, 2015, 97, 611-619.	3.3	15
74	miR-181a Modulation of ERK-MAPK Signaling Sustains DC-SIGN Expression and Limits Activation of Monocyte-Derived Dendritic Cells. Cell Reports, 2020, 30, 3793-3805.e5.	6.4	14
75	Brief report: Decreased expression of CD244 (SLAMF4) on monocytes and platelets in patients with systemic lupus erythematosus. Clinical Rheumatology, 2018, 37, 811-816.	2.2	12
76	Ezh2 Controls Skin Tolerance through Distinct Mechanisms in Different Subsets of Skin Dendritic Cells. IScience, 2018, 10, 23-39.	4.1	12
77	Complete genomic sequence of the Australian south-west genotype of Sindbis virus: comparisons with other Sindbis strains and identification of a unique deletion in the 3'-untranslated region. Virus Genes, 2003, 26, 317-327.	1.6	11
78	Transcriptional and functional characterization of CD137L-dendritic cells identifies a novel dendritic cell phenotype. Scientific Reports, 2016, 6, 29712.	3.3	10
79	Healthy elderly Singaporeans show no age-related humoral hyporesponsiveness nor diminished plasmablast generation in response to influenza vaccine. Immunity and Ageing, 2018, 15, 28.	4.2	10
80	Histone acetylome-wide associations in immune cells from individuals with active Mycobacterium tuberculosis infection. Nature Microbiology, 2022, 7, 312-326.	13.3	9
81	Deep Sequencing in Infectious Diseases: Immune and Pathogen Repertoires for the Improvement of Patient Outcomes. Frontiers in Immunology, 2017, 8, 593.	4.8	8
82	Streamlining volumetric multi-channel image cytometry using hue-saturation-brightness-based surface creation. Communications Biology, 2018, 1, 136.	4.4	8
83	Resistin expression in human monocytes is controlled by two linked promoter SNPs mediating NFKB p50/p50 binding and C-methylation. Scientific Reports, 2019, 9, 15245.	3.3	8
84	FUT6 deficiency compromises basophil function by selectively abrogating their sialyl-Lewis x expression. Communications Biology, 2021, 4, 832.	4.4	7
85	Endoplasmic reticulum stress response and bile acid signatures associate with multi-strain seroresponsiveness during elderly influenza vaccination. IScience, 2021, 24, 102970.	4.1	5
86	Genetic variants of inducible costimulator are associated with allergic asthma susceptibility. Journal of Allergy and Clinical Immunology, 2015, 135, 556-558.e13.	2.9	4
87	Complete human CD1a deficiency on Langerhans cells due to a rare point mutation in the coding sequence. Journal of Allergy and Clinical Immunology, 2016, 138, 1709-1712.e11.	2.9	4
88	Inverse association of FCER1A allergy variant in monocytes and plasmacytoid dendritic cells. Journal of Allergy and Clinical Immunology, 2021, 147, 1510-1513.e8.	2.9	4
89	Interfering ribonucleic acids that suppress expression of multiple unrelated genes. BMC Biotechnology, 2009, 9, 57.	3.3	3
90	RNA-Seq Signatures Normalized by mRNA Abundance Allow Absolute Deconvolution of Human Immune Cells. SSRN Electronic Journal, 0, , .	0.4	3

# ARTICLE IF CITATIONS

91 Molecular Epidemiology and Evolution of Mosquito-Borne Flaviviruses and Alphaviruses Enzootic in Australia., 1996,, 153-165.