

# Michael Poidinger

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2896480/publications.pdf>

Version: 2024-02-01

91  
papers

13,496  
citations

38742

50  
h-index

45317

90  
g-index

93  
all docs

93  
docs citations

93  
times ranked

25008  
citing authors

#	ARTICLE	IF	CITATIONS
1	Histone acetylome-wide associations in immune cells from individuals with active <i>Mycobacterium tuberculosis</i> infection. <i>Nature Microbiology</i> , 2022, 7, 312-326.	13.3	9
2	Inverse association of FCER1A allergy variant in monocytes and plasmacytoid dendritic cells. <i>Journal of Allergy and Clinical Immunology</i> , 2021, 147, 1510-1513.e8.	2.9	4
3	FUT6 deficiency compromises basophil function by selectively abrogating their sialyl-Lewis x expression. <i>Communications Biology</i> , 2021, 4, 832.	4.4	7
4	Endoplasmic reticulum stress response and bile acid signatures associate with multi-strain seroresponsiveness during elderly influenza vaccination. <i>IScience</i> , 2021, 24, 102970.	4.1	5
5	miR-181a Modulation of ERK-MAPK Signaling Sustains DC-SIGN Expression and Limits Activation of Monocyte-Derived Dendritic Cells. <i>Cell Reports</i> , 2020, 30, 3793-3805.e5.	6.4	14
6	Circulating CD1c+ myeloid dendritic cells are potential precursors to LCH lesion CD1a+CD207+ cells. <i>Blood Advances</i> , 2020, 4, 87-99.	5.2	25
7	Resistin expression in human monocytes is controlled by two linked promoter SNPs mediating NFkB p50/p50 binding and C-methylation. <i>Scientific Reports</i> , 2019, 9, 15245.	3.3	8
8	RNA-Seq Signatures Normalized by mRNA Abundance Allow Absolute Deconvolution of Human Immune Cell Types. <i>Cell Reports</i> , 2019, 26, 1627-1640.e7.	6.4	590
9	Two distinct interstitial macrophage populations coexist across tissues in specific subtissular niches. <i>Science</i> , 2019, 363, .	12.6	676
10	A Subset of Type I Conventional Dendritic Cells Controls Cutaneous Bacterial Infections through VEGF±-Mediated Recruitment of Neutrophils. <i>Immunity</i> , 2019, 50, 1069-1083.e8.	14.3	50
11	Multifactorial heterogeneity of virus-specific T cells and association with the progression of human chronic hepatitis B infection. <i>Science Immunology</i> , 2019, 4, .	11.9	57
12	Mapping of $\hat{I}^3/\hat{I}^1$ T cells reveals $\hat{V}^2+$ T cells resistance to senescence. <i>EBioMedicine</i> , 2019, 39, 44-58.	6.1	54
13	The tumour microenvironment creates a niche for the self-renewal of tumour-promoting macrophages in colon adenoma. <i>Nature Communications</i> , 2018, 9, 582.	12.8	76
14	Advantages of meta-total RNA sequencing (MeTRS) over shotgun metagenomics and amplicon-based sequencing in the profiling of complex microbial communities. <i>Npj Biofilms and Microbiomes</i> , 2018, 4, 2.	6.4	65
15	Brief report: Decreased expression of CD244 (SLAMF4) on monocytes and platelets in patients with systemic lupus erythematosus. <i>Clinical Rheumatology</i> , 2018, 37, 811-816.	2.2	12
16	Ezh2 Controls Skin Tolerance through Distinct Mechanisms in Different Subsets of Skin Dendritic Cells. <i>IScience</i> , 2018, 10, 23-39.	4.1	12
17	Streamlining volumetric multi-channel image cytometry using hue-saturation-brightness-based surface creation. <i>Communications Biology</i> , 2018, 1, 136.	4.4	8
18	Induction of Human T-cell and Cytokine Responses Following Vaccination with a Novel Influenza Vaccine. <i>Scientific Reports</i> , 2018, 8, 18007.	3.3	33

#	ARTICLE	IF	CITATIONS
19	Influenza Vaccine-Induced Antibody Responses Are Not Impaired by Frailty in the Community-Dwelling Elderly With Natural Influenza Exposure. <i>Frontiers in Immunology</i> , 2018, 9, 2465.	4.8	34
20	Healthy elderly Singaporeans show no age-related humoral hyporesponsiveness nor diminished plasmablast generation in response to influenza vaccine. <i>Immunity and Ageing</i> , 2018, 15, 28.	4.2	10
21	Experimental evolution of a fungal pathogen into a gut symbiont. <i>Science</i> , 2018, 362, 589-595.	12.6	184
22	Bystander CD8+ T cells are abundant and phenotypically distinct in human tumour infiltrates. <i>Nature</i> , 2018, 557, 575-579.	27.8	942
23	Hyaluronan Receptor LYVE-1-Expressing Macrophages Maintain Arterial Tone through Hyaluronan-Mediated Regulation of Smooth Muscle Cell Collagen. <i>Immunity</i> , 2018, 49, 326-341.e7.	14.3	235
24	Systematic characterization of basophil anergy. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2017, 72, 373-384.	5.7	26
25	Mapping the human DC lineage through the integration of high-dimensional techniques. <i>Science</i> , 2017, 356, .	12.6	429
26	Human fetal dendritic cells promote prenatal T-cell immune suppression through arginase-2. <i>Nature</i> , 2017, 546, 662-666.	27.8	199
27	Host sirtuin 1 regulates mycobacterial immunopathogenesis and represents a therapeutic target against tuberculosis. <i>Science Immunology</i> , 2017, 2, .	11.9	104
28	IgG1 memory B cells keep the memory of IgE responses. <i>Nature Communications</i> , 2017, 8, 641.	12.8	143
29	Induced-Pluripotent-Stem-Cell-Derived Primitive Macrophages Provide a Platform for Modeling Tissue-Resident Macrophage Differentiation and Function. <i>Immunity</i> , 2017, 47, 183-198.e6.	14.3	245
30	Functionally diverse human T cells recognize non-microbial antigens presented by MR1. <i>ELife</i> , 2017, 6, .	6.0	100
31	Deep Sequencing in Infectious Diseases: Immune and Pathogen Repertoires for the Improvement of Patient Outcomes. <i>Frontiers in Immunology</i> , 2017, 8, 593.	4.8	8
32	NLRP10 Enhances CD4+ T-Cell-Mediated IFN $\gamma$ Response via Regulation of Dendritic Cell-Derived IL-12 Release. <i>Frontiers in Immunology</i> , 2017, 8, 1462.	4.8	21
33	$\beta$ -glucan Exposure on the Fungal Cell Wall Tightly Correlates with Competitive Fitness of <i>Candida</i> Species in the Mouse Gastrointestinal Tract. <i>Frontiers in Cellular and Infection Microbiology</i> , 2016, 6, 186.	3.9	41
34	Mpath maps multi-branching single-cell trajectories revealing progenitor cell progression during development. <i>Nature Communications</i> , 2016, 7, 11988.	12.8	67
35	flowAI: automatic and interactive anomaly discerning tools for flow cytometry data. <i>Bioinformatics</i> , 2016, 32, 2473-2480.	4.1	166
36	CXCR4 identifies transitional bone marrow premonocytes that replenish the mature monocyte pool for peripheral responses. <i>Journal of Experimental Medicine</i> , 2016, 213, 2293-2314.	8.5	108

#	ARTICLE	IF	CITATIONS
37	Complete human CD1a deficiency on Langerhans cells due to a rare point mutation in the coding sequence. <i>Journal of Allergy and Clinical Immunology</i> , 2016, 138, 1709-1712.e11.	2.9	4
38	Plasmablasts During Acute Dengue Infection Represent a Small Subset of a Broader Virus-specific Memory B Cell Pool. <i>EBioMedicine</i> , 2016, 12, 178-188.	6.1	62
39	Transcriptional and functional characterization of CD137L-dendritic cells identifies a novel dendritic cell phenotype. <i>Scientific Reports</i> , 2016, 6, 29712.	3.3	10
40	Unsupervised High-Dimensional Analysis Aligns Dendritic Cells across Tissues and Species. <i>Immunity</i> , 2016, 45, 669-684.	14.3	683
41	Functional variants of 17q12-21 are associated with allergic asthma but not allergic rhinitis. <i>Journal of Allergy and Clinical Immunology</i> , 2016, 137, 758-766.e3.	2.9	34
42	TLR7 and TLR9 ligands regulate antigen presentation by macrophages. <i>International Immunology</i> , 2016, 28, 223-232.	4.0	43
43	Neutrophils Self-Regulate Immune Complex-Mediated Cutaneous Inflammation through CXCL2. <i>Journal of Investigative Dermatology</i> , 2016, 136, 416-424.	0.7	62
44	Cytofkit: A Bioconductor Package for an Integrated Mass Cytometry Data Analysis Pipeline. <i>PLoS Computational Biology</i> , 2016, 12, e1005112.	3.2	302
45	Automated Identification of Core Regulatory Genes in Human Gene Regulatory Networks. <i>PLoS Computational Biology</i> , 2015, 11, e1004504.	3.2	33
46	Identification of cDC1- and cDC2-committed DC progenitors reveals early lineage priming at the common DC progenitor stage in the bone marrow. <i>Nature Immunology</i> , 2015, 16, 718-728.	14.5	475
47	Visualization of bone marrow monocyte mobilization using <i>Cx3cr1gfp/+Flt3L<sup>Δ</sup>/Δ</i> reporter mouse by multiphoton intravital microscopy. <i>Journal of Leukocyte Biology</i> , 2015, 97, 611-619.	3.3	15
48	The Transcriptional Stress Response of <i>Candida albicans</i> to Weak Organic Acids. <i>G3: Genes, Genomes, Genetics</i> , 2015, 5, 497-505.	1.8	46
49	Genetic variants of inducible costimulator are associated with allergic asthma susceptibility. <i>Journal of Allergy and Clinical Immunology</i> , 2015, 135, 556-558.e13.	2.9	4
50	NFATc2 mediates epigenetic modification of dendritic cell cytokine and chemokine responses to dectin-1 stimulation. <i>Nucleic Acids Research</i> , 2015, 43, 836-847.	14.5	35
51	MicroRNA expression profiling of human blood monocyte subsets highlights functional differences. <i>Immunology</i> , 2015, 145, 404-416.	4.4	34
52	Cell Specific eQTL Analysis without Sorting Cells. <i>PLoS Genetics</i> , 2015, 11, e1005223.	3.5	115
53	Mapping the Diversity of Follicular Helper T Cells in Human Blood and Tonsils Using High-Dimensional Mass Cytometry Analysis. <i>Cell Reports</i> , 2015, 11, 1822-1833.	6.4	140
54	C-Myb+ Erythro-Myeloid Progenitor-Derived Fetal Monocytes Give Rise to Adult Tissue-Resident Macrophages. <i>Immunity</i> , 2015, 42, 665-678.	14.3	847

#	ARTICLE	IF	CITATIONS
55	High Mitochondrial Respiration and Glycolytic Capacity Represent a Metabolic Phenotype of Human Tolerogenic Dendritic Cells. <i>Journal of Immunology</i> , 2015, 194, 5174-5186.	0.8	183
56	Human Monocytes Undergo Functional Re-programming during Sepsis Mediated by Hypoxia-Inducible Factor-1 $\alpha$ . <i>Immunity</i> , 2015, 42, 484-498.	14.3	340
57	CD103+ Dendritic Cells Control Th17 Cell Function in the Lung. <i>Cell Reports</i> , 2015, 12, 1789-1801.	6.4	89
58	Genome-wide analysis of the genetic regulation of gene expression in human neutrophils. <i>Nature Communications</i> , 2015, 6, 7971.	12.8	23
59	Gene Essentiality Is a Quantitative Property Linked to Cellular Evolvability. <i>Cell</i> , 2015, 163, 1388-1399.	28.9	146
60	RNA sensing by conventional dendritic cells is central to the development of lupus nephritis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E6195-204.	7.1	49
61	Dengue Serotype Cross-Reactive, Anti-E Protein Antibodies Confound Specific Immune Memory for 1 Year after Infection. <i>Frontiers in Immunology</i> , 2014, 5, 388.	4.8	18
62	Human Dermal CD14 + Cells Are a Transient Population of Monocyte-Derived Macrophages. <i>Immunity</i> , 2014, 41, 465-477.	14.3	256
63	Selective Susceptibility of Human Skin Antigen Presenting Cells to Productive Dengue Virus Infection. <i>PLoS Pathogens</i> , 2014, 10, e1004548.	4.7	80
64	Calcium and Calcineurin-NFAT Signaling Regulate Granulocyte-Monocyte Progenitor Cell Cycle via Flt3-L. <i>Stem Cells</i> , 2014, 32, 3232-3244.	3.2	20
65	Metformin as adjunct antituberculosis therapy. <i>Science Translational Medicine</i> , 2014, 6, 263ra159.	12.4	404
66	Molecular Profiling Reveals a Tumor-Promoting Phenotype of Monocytes and Macrophages in Human Cancer Progression. <i>Immunity</i> , 2014, 41, 815-829.	14.3	240
67	High-dimensional analysis of the murine myeloid cell system. <i>Nature Immunology</i> , 2014, 15, 1181-1189.	14.5	349
68	Involvement of GABA Transporters in Atropine-Treated Myopic Retina As Revealed by iTRAQ Quantitative Proteomics. <i>Journal of Proteome Research</i> , 2014, 13, 4647-4658.	3.7	56
69	Enhancers Are Major Targets for Murine Leukemia Virus Vector Integration. <i>Journal of Virology</i> , 2014, 88, 4504-4513.	3.4	88
70	Human Regulatory B Cells Combine Phenotypic and Genetic Hallmarks with a Distinct Differentiation Fate. <i>Journal of Immunology</i> , 2014, 193, 2258-2266.	0.8	40
71	Ubiquitin-conjugating enzyme Ubc13 controls breast cancer metastasis through a TAK1-p38 MAP kinase cascade. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 13870-13875.	7.1	99
72	IRF4 Transcription Factor-Dependent CD11b+ Dendritic Cells in Human and Mouse Control Mucosal IL-17 Cytokine Responses. <i>Immunity</i> , 2013, 38, 970-983.	14.3	703

#	ARTICLE	IF	CITATIONS
73	The distinctive germinal center phase of IgE+ B lymphocytes limits their contribution to the classical memory response. <i>Journal of Experimental Medicine</i> , 2013, 210, 2755-2771.	8.5	139
74	Rational Design of a Live Attenuated Dengue Vaccine: 2- <i>O</i> -Methyltransferase Mutants Are Highly Attenuated and Immunogenic in Mice and Macaques. <i>PLoS Pathogens</i> , 2013, 9, e1003521.	4.7	98
75	An Unbiased Approach to Identifying Tau Kinases That Phosphorylate Tau at Sites Associated with Alzheimer Disease. <i>Journal of Biological Chemistry</i> , 2013, 288, 23331-23347.	3.4	99
76	Enhanced Neutralizing Antibody Titers and Th1 Polarization from a Novel <i>Escherichia coli</i> Derived Pandemic Influenza Vaccine. <i>PLoS ONE</i> , 2013, 8, e76571.	2.5	25
77	Protumoral role of monocytes in human B-cell precursor acute lymphoblastic leukemia: involvement of the chemokine CXCL10. <i>Blood</i> , 2012, 119, 227-237.	1.4	59
78	Human Tissues Contain CD141hi Cross-Presenting Dendritic Cells with Functional Homology to Mouse CD103+ Nonlymphoid Dendritic Cells. <i>Immunity</i> , 2012, 37, 60-73.	14.3	643
79	Optimal cellular preservation for high dimensional flow cytometric analysis of multicentre trials. <i>Journal of Immunological Methods</i> , 2012, 385, 79-89.	1.4	38
80	Cistrome: an integrative platform for transcriptional regulation studies. <i>Genome Biology</i> , 2011, 12, R83.	9.6	598
81	Sequence determinants of innate immune activation by short interfering RNAs. <i>BMC Immunology</i> , 2009, 10, 40.	2.2	57
82	Interfering ribonucleic acids that suppress expression of multiple unrelated genes. <i>BMC Biotechnology</i> , 2009, 9, 57.	3.3	3
83	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. <i>Virus Genes</i> , 2003, 26, 317-327.	1.6	11
84	The Relationships between West Nile and Kunjin Viruses. <i>Emerging Infectious Diseases</i> , 2001, 7, 697-705.	4.3	126
85	The Relationships between West Nile and Kunjin Viruses. <i>Emerging Infectious Diseases</i> , 2001, 7, 697-705.	4.3	90
86	Two contiguous outbreaks of dengue type 2 in north Queensland. <i>Medical Journal of Australia</i> , 1998, 168, 221-225.	1.7	66
87	Genetic Stability Among Temporally and Geographically Diverse Isolates of Barmah Forest Virus. <i>American Journal of Tropical Medicine and Hygiene</i> , 1997, 57, 230-234.	1.4	38
88	An outbreak of Japanese encephalitis in the Torres Strait, Australia, 1995. <i>Medical Journal of Australia</i> , 1996, 165, 256-260.	1.7	298
89	Molecular Characterization of the Japanese Encephalitis Serocomplex of the Flavivirus Genus. <i>Virology</i> , 1996, 218, 417-421.	2.4	126
90	Molecular Epidemiology and Evolution of Mosquito-Borne Flaviviruses and Alphaviruses Enzootic in Australia. , 1996, , 153-165.		0

#	ARTICLE	IF	CITATIONS
91	RNA-Seq Signatures Normalized by mRNA Abundance Allow Absolute Deconvolution of Human Immune Cells. SSRN Electronic Journal, 0, , .	0.4	3