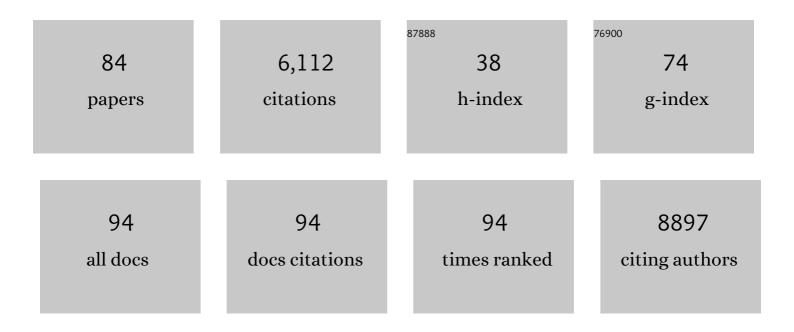
## Andrew I Webb

List of Publications by Year in descending order

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ANDREWIWERR

#	Article	IF	CITATIONS
1	The Pseudokinase MLKL Mediates Necroptosis via a Molecular Switch Mechanism. Immunity, 2013, 39, 443-453.	14.3	958
2	Linear ubiquitination prevents inflammation and regulates immune signalling. Nature, 2011, 471, 591-596.	27.8	805
3	Activation of the pseudokinase MLKL unleashes the four-helix bundle domain to induce membrane localization and necroptotic cell death. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 15072-15077.	7.1	484
4	CIS is a potent checkpoint in NK cell–mediated tumor immunity. Nature Immunology, 2016, 17, 816-824.	14.5	289
5	Familial autoinflammation with neutrophilic dermatosis reveals a regulatory mechanism of pyrin activation. Science Translational Medicine, 2016, 8, 332ra45.	12.4	241
6	A type III effector antagonizes death receptor signalling during bacterial gut infection. Nature, 2013, 501, 247-251.	27.8	238
7	Conformational switching of the pseudokinase domain promotes human MLKL tetramerization and cell death by necroptosis. Nature Communications, 2018, 9, 2422.	12.8	154
8	Lack of prominent peptide–major histocompatibility complex features limits repertoire diversity in virus-specific CD8+ T cell populations. Nature Immunology, 2005, 6, 382-389.	14.5	142
9	EspL is a bacterial cysteine protease effector that cleaves RHIM proteins to block necroptosis and inflammation. Nature Microbiology, 2017, 2, 16258.	13.3	141
10	A RIPK2 inhibitor delays NOD signalling events yet prevents inflammatory cytokine production. Nature Communications, 2015, 6, 6442.	12.8	112
11	A class of Î <sup>3</sup> δT cell receptors recognize the underside of the antigen-presenting molecule MR1. Science, 2019, 366, 1522-1527.	12.6	98
12	Necroptosis signalling is tuned by phosphorylation of MLKL residues outside the pseudokinase domain activation loop. Biochemical Journal, 2015, 471, 255-265.	3.7	91
13	Viral MLKL Homologs Subvert Necroptotic Cell Death by Sequestering Cellular RIPK3. Cell Reports, 2019, 28, 3309-3319.e5.	6.4	83
14	Enrichment of extracellular vesicles from human synovial fluid using size exclusion chromatography. Journal of Extracellular Vesicles, 2018, 7, 1490145.	12.2	78
15	Structure of Plasmodium falciparum Rh5–CyRPA–Ripr invasion complex. Nature, 2019, 565, 118-121.	27.8	74
16	Pathogenic Acinetobacter species have a functional type I secretion system and contact-dependent inhibition systems. Journal of Biological Chemistry, 2017, 292, 9075-9087.	3.4	73
17	Necroptotic signaling is primed in Mycobacterium tuberculosis-infected macrophages, but its pathophysiological consequence in disease is restricted. Cell Death and Differentiation, 2018, 25, 951-965.	11.2	72
18	Regulation of Starch Stores by a Ca2+-Dependent Protein Kinase Is Essential for Viable Cyst Development in Toxoplasma gondii. Cell Host and Microbe, 2015, 18, 670-681.	11.0	71

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19	RIPLET, and not TRIM25, is required for endogenous RIGâ€lâ€dependent antiviral responses. Immunology and Cell Biology, 2019, 97, 840-852.	2.3	70
20	Protein kinase A negatively regulates Ca2+ signalling in Toxoplasma gondii. PLoS Biology, 2018, 16, e2005642.	5.6	65
21	Therapeutic blockade of activin-A improves NK cell function and antitumor immunity. Science Signaling, 2019, 12, .	3.6	64
22	Identification of MLKL membrane translocation as a checkpoint in necroptotic cell death using Monobodies. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 8468-8475.	7.1	64
23	Functional and Structural Characteristics of NY-ESO-1-related HLA A2-restricted Epitopes and the Design of a Novel Immunogenic Analogue. Journal of Biological Chemistry, 2004, 279, 23438-23446.	3.4	61
24	Suppressor of cytokine signaling (SOCS)5 ameliorates influenza infection via inhibition of EGFR signaling. ELife, 2017, 6, .	6.0	61
25	Plasmepsin V cleaves malaria effector proteins in a distinct endoplasmic reticulum translocation interactome for export to the erythrocyte. Nature Microbiology, 2018, 3, 1010-1022.	13.3	59
26	Targeting histone acetylation dynamics and oncogenic transcription by catalytic P300/CBP inhibition. Molecular Cell, 2021, 81, 2183-2200.e13.	9.7	59
27	Constraints within major histocompatibility complex class I restricted peptides: Presentation and consequences for T-cell recognition. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 5534-5539.	7.1	58
28	Lectin Switching During Dengue Virus Infection. Journal of Infectious Diseases, 2011, 203, 1775-1783.	4.0	58
29	T Cell Determinants Incorporating β-Amino Acid Residues Are Protease Resistant and Remain Immunogenic In Vivo. Journal of Immunology, 2005, 175, 3810-3818.	0.8	56
30	Conformational interconversion of MLKL and disengagement from RIPK3 precede cell death by necroptosis. Nature Communications, 2021, 12, 2211.	12.8	56
31	Person-Specific Biomolecular Coronas Modulate Nanoparticle Interactions with Immune Cells in Human Blood. ACS Nano, 2020, 14, 15723-15737.	14.6	55
32	The bacterial arginine glycosyltransferase effector NleB preferentially modifies Fas-associated death domain protein (FADD). Journal of Biological Chemistry, 2017, 292, 17337-17350.	3.4	53
33	The Cellular Redox Environment Alters Antigen Presentation. Journal of Biological Chemistry, 2014, 289, 27979-27991.	3.4	52
34	Epitope-specific TCRÎ <sup>2</sup> repertoire diversity imparts no functional advantage on the CD8 <sup>+</sup> T cell response to cognate viral peptides. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 2034-2039.	7.1	50
35	Synthesis of Biotinylated Episilvestrol: Highly Selective Targeting of the Translation Factors elF4Al/II. Organic Letters, 2013, 15, 1406-1409.	4.6	49
36	BAK core dimers bind lipids and can be bridged by them. Nature Structural and Molecular Biology, 2020, 27, 1024-1031.	8.2	49

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37	ROCK-mediated selective activation of PERK signalling causes fibroblast reprogramming and tumour progression through a CRELD2-dependent mechanism. Nature Cell Biology, 2020, 22, 882-895.	10.3	47
38	Universal Solid-Phase Protein Preparation (USP <sup>3</sup> ) for Bottom-up and Top-down Proteomics. Journal of Proteome Research, 2019, 18, 2915-2924.	3.7	43
39	Suppressor of Cytokine Signaling (SOCS) 5 Utilises Distinct Domains for Regulation of JAK1 and Interaction with the Adaptor Protein Shc-1. PLoS ONE, 2013, 8, e70536.	2.5	42
40	The ubiquitylation of IL-1β limits its cleavage by caspase-1 and targets it for proteasomal degradation. Nature Communications, 2021, 12, 2713.	12.8	40
41	Oligomerizationâ€driven MLKL ubiquitylation antagonizes necroptosis. EMBO Journal, 2021, 40, e103718.	7.8	39
42	Discovery and Validation of Novel Protein Biomarkers in Ovarian Cancer Patient Urine. Proteomics - Clinical Applications, 2018, 12, e1700135.	1.6	37
43	How IGF-II Binds to the Human Type 1 Insulin-like Growth Factor Receptor. Structure, 2020, 28, 786-798.e6.	3.3	36
44	Progranulin does not inhibit TNF and lymphotoxinâ€Î± signalling through TNF receptor 1. Immunology and Cell Biology, 2013, 91, 661-664.	2.3	35
45	Ensemble Properties of Bax Determine Its Function. Structure, 2018, 26, 1346-1359.e5.	3.3	34
46	The Structure of H-2Kb and Kbm8 Complexed to a Herpes Simplex Virus Determinant: Evidence for a Conformational Switch That Governs T Cell Repertoire Selection and Viral Resistance. Journal of Immunology, 2004, 173, 402-409.	0.8	31
47	BAX Activation: Mutations Near Its Proposed Non-canonical BH3 Binding Site Reveal Allosteric Changes Controlling Mitochondrial Association. Cell Reports, 2019, 27, 359-373.e6.	6.4	31
48	A small molecule interacts with VDAC2 to block mouse BAK-driven apoptosis. Nature Chemical Biology, 2019, 15, 1057-1066.	8.0	30
49	Serum microRNA is a biomarker for post-operative monitoring in glioma. Journal of Neuro-Oncology, 2020, 149, 391-400.	2.9	27
50	Structural and Biological Basis of CTL Escape in Coronavirus-Infected Mice. Journal of Immunology, 2008, 180, 3926-3937.	0.8	23
51	A Mechanism for Actin Filament Severing by Malaria Parasite Actin Depolymerizing Factor 1 via a Low Affinity Binding Interface. Journal of Biological Chemistry, 2014, 289, 4043-4054.	3.4	22
52	Menstrual fluid factors facilitate tissue repair: identification and functional action in endometrial and skin repair. FASEB Journal, 2019, 33, 584-605.	0.5	22
53	Targeting of RNA Polymerase II by a nuclear <i>Legionella pneumophila</i> Dot/Icm effector SnpL. Cellular Microbiology, 2018, 20, e12852.	2.1	21
54	Proteomic analysis of extracellular vesicles reveals an immunogenic cargo in rheumatoid arthritis synovial fluid. Clinical and Translational Immunology, 2020, 9, e1185.	3.8	21

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55	Idiopathic infertility in women is associated with distinct changes in proliferative phase uterine fluid proteinsâ€. Biology of Reproduction, 2018, 98, 752-764.	2.7	20
56	Dynamic reconfiguration of proâ€apoptotic BAK on membranes. EMBO Journal, 2021, 40, e107237.	7.8	20
57	Phosphorylation by Aurora B kinase regulates caspase-2 activity and function. Cell Death and Differentiation, 2021, 28, 349-366.	11.2	18
58	Role of salt bridges in the dimer interface of 14-3-3ζ in dimer dynamics, N-terminal α-helical order, and molecular chaperone activity. Journal of Biological Chemistry, 2018, 293, 89-99.	3.4	17
59	Prevention of Cytotoxic T Cell Escape Using a Heteroclitic Subdominant Viral T Cell Determinant. PLoS Pathogens, 2008, 4, e1000186.	4.7	14
60	Sperm proteins and cancerâ€ŧestis antigens are released by the seminiferous tubules in mice and men. FASEB Journal, 2021, 35, e21397.	0.5	14
61	Arrdc4â€dependent extracellular vesicle biogenesis is required for sperm maturation. Journal of Extracellular Vesicles, 2021, 10, e12113.	12.2	14
62	High Yield Production of a Soluble Human Interleukin-3 Variant from E. coli with Wild-Type Bioactivity and Improved Radiolabeling Properties. PLoS ONE, 2013, 8, e74376.	2.5	13
63	Proprotein convertase 5/6 cleaves platelet-derived growth factor A in the human endometrium in preparation for embryo implantation. Molecular Human Reproduction, 2015, 21, 262-270.	2.8	13
64	Holographic Interface for three-dimensional Visualization of MRI on HoloLens: A Prototype Platform for MRI Guided Neurosurgeries. , 2017, , .		13
65	Crystal structure of the hinge domain of Smchd1 reveals its dimerization mode and nucleic acid–binding residues. Science Signaling, 2020, 13, .	3.6	12
66	The search for RNA-binding proteins: a technical and interdisciplinary challenge. Biochemical Society Transactions, 2021, 49, 393-403.	3.4	10
67	A regulatory region on <scp>RIPK</scp> 2 is required for <scp>XIAP</scp> binding and <scp>NOD</scp> signaling activity. EMBO Reports, 2020, 21, e50400.	4.5	9
68	Cp1/cathepsin L is required for autolysosomal clearance in <i>Drosophila</i> . Autophagy, 2021, 17, 2734-2749.	9.1	9
69	Tankyrase-mediated ADP-ribosylation is a regulator of TNF-induced death. Science Advances, 2022, 8, eabh2332.	10.3	9
70	Loss of Bcl-G, a Bcl-2 family member, augments the development of inflammation-associated colorectal cancer. Cell Death and Differentiation, 2020, 27, 742-757.	11.2	8
71	Kâ€29 linked ubiquitination of Arrdc4 regulates its function in extracellular vesicle biogenesis. Journal of Extracellular Vesicles, 2022, 11, e12188.	12.2	8
72	Plasma membrane profiling during enterohemorrhagic E. coli infection reveals that the metalloprotease StcE cleaves CD55 from host epithelial surfaces. Journal of Biological Chemistry, 2018, 293, 17188-17199.	3.4	7

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73	Granulovirus PK-1 kinase activity relies on a side-to-side dimerization mode centered on the regulatory αC helix. Nature Communications, 2021, 12, 1002.	12.8	7
74	Multimodal regulation of encystation in Giardia duodenalis revealed by deep proteomics. International Journal for Parasitology, 2021, 51, 809-824.	3.1	7
75	<scp>SPATA</scp> 2 – Keeping the <scp>TNF</scp> signal short and sweet. EMBO Journal, 2016, 35, 1848-1850.	7.8	6
76	Clinical MDR1 inhibitors enhance Smac-mimetic bioavailability to kill murine LSCs and improve survival in AML models. Blood Advances, 2020, 4, 5062-5077.	5.2	6
77	Quantitative proteomic analysis of EZH2 inhibition in acute myeloid leukemia reveals the targets and pathways that precede the induction of cell death. Proteomics - Clinical Applications, 2017, 11, 1700013.	1.6	5
78	Mass Dynamics 1.0: A Streamlined, Web-Based Environment for Analyzing, Sharing, and Integrating Label-Free Data. Journal of Proteome Research, 2021, 20, 5180-5188.	3.7	4
79	SFPQ-ABL1 and BCR-ABL1 use different signaling networks to drive B-cell acute lymphoblastic leukemia. Blood Advances, 2022, 6, 2373-2387.	5.2	4
80	Generation of novel Id2 and E2-2, E2A and HEB antibodies reveals novel Id2 binding partners and species-specific expression of E-proteins in NK cells. Molecular Immunology, 2019, 115, 56-63.	2.2	3
81	Simplifying MS1 and MS2 spectra to achieve lower mass error, more dynamic range, and higher peptide identification confidence on the Bruker timsTOF Pro. PLoS ONE, 2022, 17, e0271025.	2.5	2
82	Optimisation of peptide-based cytotoxic T-cell determinants using. International Journal of Peptide Research and Therapeutics, 2003, 10, 561-569.	0.1	1
83	Proteomic Profiling of Cell Death: Stable Isotope Labeling and Mass Spectrometry Analysis. Methods in Molecular Biology, 2016, 1419, 277-286.	0.9	0

84 Budget Management. , 2016, , 339-357.

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