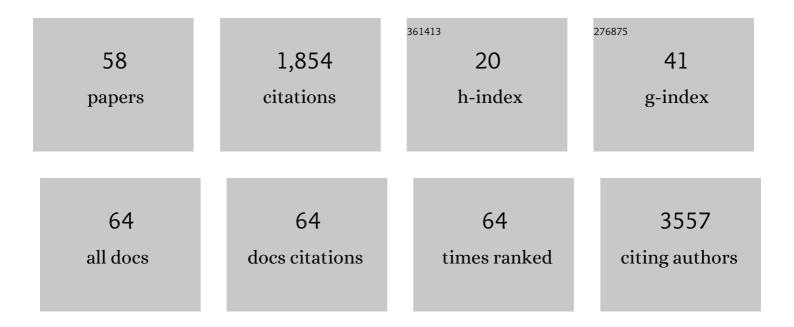
Wendy E Heywood

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Increased cerebrospinal fluid soluble TREM2 concentration in Alzheimer's disease. Molecular Neurodegeneration, 2016, 11, 3.	10.8	236
2	Ambroxol for the Treatment of Patients With Parkinson Disease With and Without Glucocerebrosidase Gene Mutations. JAMA Neurology, 2020, 77, 427.	9.0	213
3	Reproducibility of Molecular Phenotypes after Long-Term Differentiation toÂHuman iPSC-Derived Neurons: A Multi-Site Omics Study. Stem Cell Reports, 2018, 11, 897-911.	4.8	135
4	Identification of novel CSF biomarkers for neurodegeneration and their validation by a high-throughput multiplexed targeted proteomic assay. Molecular Neurodegeneration, 2015, 10, 64.	10.8	121
5	†The long tail of Covid-19' - The detection of a prolonged inflammatory response after a SARS-CoV-2 infection in asymptomatic and mildly affected patients. F1000Research, 2020, 9, 1349.	1.6	116
6	Development of a Novel Targeting System for Lethal Photosensitization of Antibiotic-Resistant Strains of Staphylococcus aureus. Antimicrobial Agents and Chemotherapy, 2005, 49, 3690-3696.	3.2	95
7	â€~The long tail of Covid-19' - The detection of a prolonged inflammatory response after a SARS-CoV-2 infection in asymptomatic and mildly affected patients. F1000Research, 2020, 9, 1349.	1.6	95
8	The Identification of New Biomarkers for Identifying and Monitoring Kidney Disease and Their Translation into a Rapid Mass Spectrometry-Based Test: Evidence of Presymptomatic Kidney Disease in Pediatric Fabry and Type-I Diabetic Patients. Journal of Proteome Research, 2013, 12, 2013-2021.	3.7	63
9	Regulation of post-Golgi LH3 trafficking is essential for collagen homeostasis. Nature Communications, 2016, 7, 12111.	12.8	54
10	Cytolethal distending toxin: creating a gap in the cell cycle. Journal of Medical Microbiology, 2005, 54, 207-216.	1.8	49
11	Programming of defective rat pancreatic \hat{l}^2 -cell function in offspring from mothers fed a low-protein diet during gestation and the suckling periods. Clinical Science, 2004, 107, 37-45.	4.3	48
12	New Role for LEKTI in Skin Barrier Formation: Label-Free Quantitative Proteomic Identification of Caspase 14 as a Novel Target for the Protease Inhibitor LEKTI. Journal of Proteome Research, 2010, 9, 4289-4294.	3.7	41
13	Mechanism of internalization of the cytolethal distending toxin of Actinobacillus actinomycetemcomitans. Microbiology (United Kingdom), 2005, 151, 1395-1402.	1.8	38
14	Proteomic Analysis of the Myocardium in Hypertrophic Obstructive Cardiomyopathy. Circulation Genomic and Precision Medicine, 2018, 11, e001974.	3.6	38
15	Proteomic Analysis of the Myocardium in Hypertrophic Obstructive Cardiomyopathy. Circulation Genomic and Precision Medicine, 2018, 11, .	3.6	34
16	Identification of a Multiplex Biomarker Panel for Hypertrophic Cardiomyopathy Using Quantitative Proteomics and Machine Learning. Molecular and Cellular Proteomics, 2020, 19, 114-127.	3.8	32
17	Rapid, proteomic urine assay for monitoring progressive organ disease in Fabry disease. Journal of Medical Genetics, 2020, 57, 38-47.	3.2	26
18	CSF pro-orexin and amyloid-β38 expression in Alzheimer's disease and frontotemporal dementia. Neurobiology of Aging, 2018, 72, 171-176.	3.1	25

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19	A New Method for the Rapid Diagnosis of Protein N-linked Congenital Disorders of Glycosylation. Journal of Proteome Research, 2013, 12, 3471-3479.	3.7	24
20	An optimised method for the proteomic profiling of full thickness human skin. Biological Procedures Online, 2016, 18, 15.	2.9	23
21	Proteomic Discovery and Development of a Multiplexed Targeted MRM-LC-MS/MS Assay for Urine Biomarkers of Extracellular Matrix Disruption in Mucopolysaccharidoses I, II, and VI. Analytical Chemistry, 2015, 87, 12238-12244.	6.5	20
22	Comparison of proteomic profiles in the zebrafish retina during experimental degeneration and regeneration. Scientific Reports, 2017, 7, 44601.	3.3	20
23	Changes in regulation of human monocyte proteins in response to IgG from patients with antiphospholipid syndrome. Blood, 2014, 124, 3808-3816.	1.4	19
24	The development of a peptide SRM-based tandem mass spectrometry assay for prenatal screening of Down syndrome. Journal of Proteomics, 2012, 75, 3248-3257.	2.4	17
25	Comparative proteomic analysis of normal and gliotic PVR retina and contribution of Müller glia to this profile. Experimental Eye Research, 2018, 177, 197-207.	2.6	17
26	Applying modern Omic technologies to the Neuronal Ceroid Lipofuscinoses. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2020, 1866, 165498.	3.8	17
27	2D DIGE analysis of maternal plasma for potential biomarkers of Down Syndrome. Proteome Science, 2011, 9, 56.	1.7	16
28	Identification of new biomarkers for Down's syndrome in maternal plasma. Journal of Proteomics, 2012, 75, 2621-2628.	2.4	16
29	Multiplex High-Throughput Targeted Proteomic Assay To Identify Induced Pluripotent Stem Cells. Analytical Chemistry, 2017, 89, 2440-2448.	6.5	15
30	Progression in multiple sclerosis is associated with low endogenous <scp>NCAM</scp> . Journal of Neurochemistry, 2013, 125, 766-773.	3.9	14
31	A geroscience approach for Parkinson's disease: Conceptual framework and design of PROPAG-AGEING project. Mechanisms of Ageing and Development, 2021, 194, 111426.	4.6	14
32	Early downregulation of hsa-miR-144-3p in serum from drug-naÃ⁻ve Parkinson's disease patients. Scientific Reports, 2022, 12, 1330.	3.3	14
33	Current applications of biomarkers in cardiomyopathies. Expert Review of Cardiovascular Therapy, 2015, 13, 825-837.	1.5	13
34	Global glycosphingolipid analysis in urine and plasma of female Fabry disease patients. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2019, 1865, 2726-2735.	3.8	13
35	Urine proteomics analysis of patients with neuronal ceroid lipofuscinoses. IScience, 2021, 24, 102020.	4.1	12
36	The identification of a new role for LEKTI in the skin: The use of protein †bait' arrays to detect defective trafficking of dermcidin in the skin of patients with Netherton syndrome. Journal of Proteomics, 2012, 75, 3925-3937.	2.4	11

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37	Metabolite and lipoprotein profiles reveal sex-related oxidative stress imbalance in de novo drug-naive Parkinson's disease patients. Npj Parkinson's Disease, 2022, 8, 14.	5.3	11
38	Free urinary glycosylated hydroxylysine as an indicator of altered collagen degradation in the mucopolysaccharidoses. Journal of Inherited Metabolic Disease, 2020, 43, 309-317.	3.6	10
39	Investigation of pathology, expression and proteomic profiles in human <i>TREM2</i> variant postmortem brains with and without Alzheimer's disease. Brain Pathology, 2020, 30, 794-810.	4.1	10
40	The presubiculum is preserved from neurodegenerative changes in Alzheimer's disease. Acta Neuropathologica Communications, 2018, 6, 62.	5.2	9
41	Global serum glycoform profiling for the investigation of dystroglycanopathies & Congenital Disorders of Glycosylation. Molecular Genetics and Metabolism Reports, 2016, 7, 55-62.	1.1	8
42	An InÂVitro Whole-Organ Liver Engineering for Testing of Genetic Therapies. IScience, 2020, 23, 101808.	4.1	8
43	Proteomic profiling reveals sub proteomes of the human placenta. Placenta, 2017, 59, 69-72.	1.5	7
44	Growth associated protein (GAP-43): Cloning and the development of a sensitive ELISA for neurological disorders. Journal of Neuroimmunology, 2014, 276, 18-23.	2.3	6
45	Niemann–Pick type C disease as proofâ€ofâ€concept for intelligent biomarker panel selection in neurometabolic disorders. Developmental Medicine and Child Neurology, 0, , .	2.1	6
46	Cerebrospinal fluid neurofilament light levels in CLN2 disease patients treated with enzyme replacement therapy normalise after two years on treatment. F1000Research, 2021, 10, 614.	1.6	4
47	Tissue Proteome of 2-Hydroxyacyl-CoA Lyase Deficient Mice Reveals Peroxisome Proliferation and Activation of ω-Oxidation. International Journal of Molecular Sciences, 2022, 23, 987.	4.1	4
48	An Optimized Method for the Proteomic Analysis of Low Volumes of Cell Culture Media and the Secretome: The Application and the Demonstration of Altered Protein Expression in iPSC-Derived Neuronal Cell Lines from Parkinson's Disease Patients. Journal of Proteome Research, 2019, 18, 1198-1207.	3.7	2
49	Heterogeneity of prodromal Parkinson symptoms in siblings of Parkinson disease patients. Npj Parkinson's Disease, 2021, 7, 78.	5.3	2
50	Proteomic signatures for perioperative oxygen delivery in skin after major elective surgery: mechanistic sub-study of a randomised controlled trial. British Journal of Anaesthesia, 2021, 127, 511-520.	3.4	2
51	A High Throughput, Multiplexed and Targeted Proteomic CSF Assay to Quantify Neurodegenerative Biomarkers and Apolipoprotein E Isoforms Status. Journal of Visualized Experiments, 2016, , .	0.3	2
52	Cerebrospinal fluid neurofilament light chain levels in CLN2 disease patients treated with enzyme replacement therapy normalise after two years on treatment. F1000Research, 0, 10, 614.	1.6	2
53	Preparation of iPSCs for Targeted Proteomic Analysis. Methods in Molecular Biology, 2019, 1994, 131-139.	0.9	1
54	The development of a rapid, multiplexed UPLC–MS/MS assay for quantitation of lyso-Gb3 and Gb3 in dried blood spots. Molecular Genetics and Metabolism, 2015, 114, S107.	1.1	0

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55	A rapid high throughput proteomic method based on profiling of proteolytic free peptides to assess post-delivery degradation of placental tissue. Placenta, 2016, 44, 109-111.	1.5	Ο
56	An unforgettable change of perspective. British Journal of Nursing, 2016, 25, 509-509.	0.7	0
57	A Selected Reaction Monitoring Protocol for the Measurement of sTREM2 in Cerebrospinal Fluid. Neuromethods, 2018, , 169-177.	0.3	0
58	Mass Spectrometry Measurement of Albumin–Alpha Fetoprotein Ratio as an Indicator of iPSC-Derived Hepatocyte Differentiation. Methods in Molecular Biology, 2019, 1994, 149-156.	0.9	0