

Michael Henry

List of Publications by Year in Descending Order

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Version: 2024-04-09

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

110 papers	2,506 citations	28 h-index	44 g-index
112 ext. papers	2,880 ext. citations	4.6 avg, IF	4.73 L-index

#	Paper	IF	Citations
110	Characterisation of the Tumour Proteome in Primary Extramedullary Multiple Myeloma Identifies Key Proteins Associated with Transendothelial Migration. <i>Blood</i> , 2021 , 138, 2665-2665	2.2	
109	Phosphoproteomic Analysis of Primary Myeloma Patient Samples Identifies Distinct Phosphorylation Signatures Correlating with Chemo-Sensitivity Profiles in an Ex Vivo Drug Sensitivity Testing Platform. <i>Blood</i> , 2021 , 138, 2666-2666	2.2	
108	Differential expression of miRNAs and functional role of mir-200a in high and low productivity CHO cells expressing an Fc fusion protein. <i>Biotechnology Letters</i> , 2021 , 43, 1551-1563	3	1
107	Mass Spectrometric Profiling of Extraocular Muscle and Proteomic Adaptations in the Model of Duchenne Muscular Dystrophy. <i>Life</i> , 2021 , 11,	3	3
106	Neonicotinoid residues in honey from urban and rural environments. <i>Environmental Science and Pollution Research</i> , 2021 , 28, 28179-28190	5.1	4
105	Copper toxicity of inflection point in human intestinal cell line Caco-2 dissected: influence of temporal expression patterns. <i>In Vitro Cellular and Developmental Biology - Animal</i> , 2021 , 57, 359-371	2.6	2
104	Proteomic profiling of the interface between the stomach wall and the pancreas in dystrophinopathy. <i>European Journal of Translational Myology</i> , 2021 , 31,	2.1	2
103	Mapping the molecular basis for growth related phenotypes in industrial producer CHO cell lines using differential proteomic analysis. <i>BMC Biotechnology</i> , 2021 , 21, 43	3.5	1
102	and modulation of NADPH oxidase activity and reactive oxygen species production in human neutrophils by Bantitrypsin. <i>ERJ Open Research</i> , 2021 , 7,	3.5	1
101	Global phosphoproteomic study of high/low specific productivity industrially relevant mAb producing recombinant CHO cell lines. <i>Current Research in Biotechnology</i> , 2021 , 3, 49-56	4.8	2
100	Dataset on the mass spectrometry-based proteomic profiling of the kidney from wild type and the dystrophic mouse model of X-linked muscular dystrophy. <i>Data in Brief</i> , 2020 , 28, 105067	1.2	1
99	LC-MS/MS-based quantitative proteomic and phosphoproteomic analysis of CHO-K1 cells adapted to growth in glutamine-free media. <i>Biotechnology Letters</i> , 2020 , 42, 2523-2536	3	4
98	LC-MS proteomic profiling of Caco-2 human intestinal cells exposed to the copper-chelating agent, triethylenetetramine: A preliminary study. <i>Biochemical and Biophysical Research Communications</i> , 2020 , 524, 847-852	3.4	
97	Characterisation and proteomic profiling of continuously exposed Cu-resistant variants of the Caco-2 cell line. <i>Toxicology in Vitro</i> , 2020 , 65, 104773	3.6	1
96	Protocol for the Bottom-Up Proteomic Analysis of Mouse Spleen. <i>STAR Protocols</i> , 2020 , 1, 100196	1.4	9
95	Proteomic and cell biological profiling of the renal phenotype of the mdx-4cv mouse model of Duchenne muscular dystrophy. <i>European Journal of Cell Biology</i> , 2020 , 99, 151059	6.1	11
94	Proteome-wide Changes in the mdx-4cv Spleen due to Pathophysiological Cross Talk with Dystrophin-Deficient Skeletal Muscle. <i>iScience</i> , 2020 , 23, 101500	6.1	10

93	Proteomic analysis of pancreatic ductal adenocarcinoma. <i>Expert Review of Proteomics</i> , 2020 , 17, 453-467	4.2	3
92	Activation of complement component 3 is associated with airways disease and pulmonary emphysema in alpha-1 antitrypsin deficiency. <i>Thorax</i> , 2020 , 75, 321-330	7.3	19
91	Proteomic Analysis of Cell Lines and Primary Tumors in Pancreatic Cancer Identifies Proteins Expressed Only In Vitro and Only In Vivo. <i>Pancreas</i> , 2020 , 49, 1109-1116	2.6	2
90	Antitrypsin therapy modulates the neutrophil membrane proteome and secretome. <i>European Respiratory Journal</i> , 2020 , 55,	13.6	7
89	Increased mAb production in amplified CHO cell lines is associated with increased interaction of CREB1 with transgene promoter. <i>Current Research in Biotechnology</i> , 2019 , 1, 49-57	4.8	4
88	Increased growth rate and productivity following stable depletion of miR-7 in a mAb producing CHO cell line causes an increase in proteins associated with the Akt pathway and ribosome biogenesis. <i>Journal of Proteomics</i> , 2019 , 195, 23-32	3.9	9
87	Improvements in single-use bioreactor film material composition leads to robust and reliable Chinese hamster ovary cell performance. <i>Biotechnology Progress</i> , 2019 , 35, e2824	2.8	5
86	Proteogenomic Annotation of Chinese Hamsters Reveals Extensive Novel Translation Events and Endogenous Retroviral Elements. <i>Journal of Proteome Research</i> , 2019 , 18, 2433-2445	5.6	12
85	Proteomic profiling of the mouse diaphragm and refined mass spectrometric analysis of the dystrophic phenotype. <i>Journal of Muscle Research and Cell Motility</i> , 2019 , 40, 9-28	3.5	17
84	Transcriptomic analysis of IgG4 Fc-fusion protein degradation in a panel of clonally-derived CHO cell lines using RNASeq. <i>Biotechnology and Bioengineering</i> , 2019 , 116, 1556-1562	4.9	6
83	Circulating Truncated Alpha-1 Antitrypsin Glycoprotein in Patient Plasma Retains Anti-Inflammatory Capacity. <i>Journal of Immunology</i> , 2019 , 202, 2240-2253	5.3	14
82	Proteomic analysis of the sarcolemma-enriched fraction from dystrophic mdx-4cv skeletal muscle. <i>Journal of Proteomics</i> , 2019 , 191, 212-227	3.9	28
81	Purification and Identification of Membrane Proteins from Urinary Extracellular Vesicles using Triton X-114 Phase Partitioning. <i>Journal of Proteome Research</i> , 2018 , 17, 86-96	5.6	10
80	The Expression Pattern of the Phosphoproteome Is Significantly Changed During the Growth Phases of Recombinant CHO Cell Culture. <i>Biotechnology Journal</i> , 2018 , 13, e1700221	5.6	15
79	Subproteomic profiling of sarcolemma from dystrophic skeletal muscle. <i>Data in Brief</i> , 2018 , 17, 980-993	1.2	4
78	Clonal variation in productivity and proteolytic clipping of an Fc-fusion protein in CHO cells: Proteomic analysis suggests a role for defective protein folding and the UPR. <i>Journal of Biotechnology</i> , 2018 , 281, 21-30	3.7	6
77	Glycosylation Repurposes Alpha-1 Antitrypsin for Resolution of Community-acquired Pneumonia. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2018 , 197, 1346-1349	10.2	26
76	A proteomic profiling dataset of recombinant Chinese hamster ovary cells showing enhanced cellular growth following miR-378 depletion. <i>Data in Brief</i> , 2018 , 21, 2679-2688	1.2	3

75	A Comparative Quantitative LC-MS/MS Profiling Analysis of Human Pancreatic Adenocarcinoma, Adjacent-Normal Tissue, and Patient-Derived Tumour Xenografts. <i>Proteomes</i> , 2018 , 6,	4.6	15
74	Acute exposure to organic and inorganic sources of copper: Differential response in intestinal cell lines. <i>Food Science and Nutrition</i> , 2018 , 6, 2499-2514	3.2	4
73	Utilization of dried and long-term stored polyacrylamide gels for the advanced proteomic profiling of mitochondrial contact sites from rat liver. <i>Biology Methods and Protocols</i> , 2018 , 3, bpy008	2.4	4
72	Depletion of endogenous miRNA-378-3p increases peak cell density of CHO DP12 cells and is correlated with elevated levels of ubiquitin carboxyl-terminal hydrolase 14. <i>Journal of Biotechnology</i> , 2018 , 288, 30-40	3.7	9
71	Proteomic profiling of liver tissue from the - mouse model of Duchenne muscular dystrophy. <i>Clinical Proteomics</i> , 2018 , 15, 34	5	15
70	Proteomic profiling of mdx-4cv serum reveals highly elevated levels of the inflammation-induced plasma marker haptoglobin in muscular dystrophy. <i>International Journal of Molecular Medicine</i> , 2017 , 39, 1357-1370	4.4	26
69	Phosphopeptide Enrichment and LC-MS/MS Analysis to Study the Phosphoproteome of Recombinant Chinese Hamster Ovary Cells. <i>Methods in Molecular Biology</i> , 2017 , 1603, 195-208	1.4	3
68	Filter-Aided Sample Preparation (FASP) for Improved Proteome Analysis of Recombinant Chinese Hamster Ovary Cells. <i>Methods in Molecular Biology</i> , 2017 , 1603, 187-194	1.4	17
67	Differential Phosphoproteomic Analysis of Recombinant Chinese Hamster Ovary Cells Following Temperature Shift. <i>Journal of Proteome Research</i> , 2017 , 16, 2339-2358	5.6	14
66	Identification of the metabolic alterations associated with the multidrug resistant phenotype in cancer and their intercellular transfer mediated by extracellular vesicles. <i>Scientific Reports</i> , 2017 , 7, 44541 ⁹	4.9	47
65	Protein and chemotherapy profiling of extracellular vesicles harvested from therapeutic induced senescent triple negative breast cancer cells. <i>Oncogenesis</i> , 2017 , 6, e388	6.6	54
64	Neutrophil Membrane Cholesterol Content is a Key Factor in Cystic Fibrosis Lung Disease. <i>EBioMedicine</i> , 2017 , 23, 173-184	8.8	20
63	A novel inhibitory anti-invasive MAb isolated using phenotypic screening highlights AnxA6 as a functionally relevant target protein in pancreatic cancer. <i>British Journal of Cancer</i> , 2017 , 117, 1326-1335	8.7	12
62	Residual urinary extracellular vesicles in ultracentrifugation supernatants after hydrostatic filtration dialysis enrichment. <i>Journal of Extracellular Vesicles</i> , 2017 , 6, 1267896	16.4	20
61	Proteomic profiling of the dystrophin complex and membrane fraction from dystrophic mdx muscle reveals decreases in the cytolinker desmoglein and increases in the extracellular matrix stabilizers biglycan and fibronectin. <i>Journal of Muscle Research and Cell Motility</i> , 2017 , 38, 251-268	3.5	24
60	Novel panel of protein biomarkers to predict response to bortezomib-containing induction regimens in multiple myeloma patients. <i>BBA Clinical</i> , 2017 , 8, 28-34		12
59	Clinical Proteomics: Liquid Chromatography-Mass Spectrometry (LC-MS) Purification Systems. <i>Methods in Molecular Biology</i> , 2017 , 1485, 375-388	1.4	2
58	Parallel mRNA, proteomics and miRNA expression analysis in cell line models of the intestine. <i>World Journal of Gastroenterology</i> , 2017 , 23, 7369-7386	5.6	12

57	Proteomic strategies in the search for novel pancreatic cancer biomarkers and drug targets: recent advances and clinical impact. <i>Expert Review of Proteomics</i> , 2016 , 13, 383-94	4.2	7
56	Proteomic analysis of dystrophin deficiency and associated changes in the aged mdx-4cv heart model of dystrophinopathy-related cardiomyopathy. <i>Journal of Proteomics</i> , 2016 , 145, 24-36	3.9	34
55	The iron-responsive microsomal proteome of <i>Aspergillus fumigatus</i> . <i>Journal of Proteomics</i> , 2016 , 136, 99-111	3.9	19
54	Multidrug resistant tumour cells shed more microvesicle-like EVs and less exosomes than their drug-sensitive counterpart cells. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2016 , 1860, 618-27	4	38
53	Quantitative label-free mass spectrometry analysis of formalin-fixed, paraffin-embedded tissue representing the invasive cutaneous malignant melanoma proteome. <i>Oncology Letters</i> , 2016 , 12, 3296-3304	3.6	9
52	Proteomic differences in recombinant CHO cells producing two similar antibody fragments. <i>Biotechnology and Bioengineering</i> , 2016 , 113, 1902-12	4.9	22
51	Data supporting the shedding of larger extracellular vesicles by multidrug resistant tumour cells. <i>Data in Brief</i> , 2016 , 6, 1023-7	1.2	7
50	Process-relevant concentrations of the leachable bDtbPP impact negatively on CHO cell production characteristics. <i>Biotechnology Progress</i> , 2016 , 32, 1547-1558	2.8	24
49	Metabolomic and proteomic analysis of breast cancer patient samples suggests that glutamate and 12-HETE in combination with CA15-3 may be useful biomarkers reflecting tumour burden. <i>Metabolomics</i> , 2015 , 11, 620-635	4.7	14
48	Abnormal levels of heterogeneous nuclear ribonucleoprotein A2B1 (hnRNP A2B1) in tumour tissue and blood samples from patients diagnosed with lung cancer. <i>Molecular BioSystems</i> , 2015 , 11, 743-52		24
47	Label-free mass spectrometric analysis reveals complex changes in the brain proteome from the mdx-4cv mouse model of Duchenne muscular dystrophy. <i>Clinical Proteomics</i> , 2015 , 12, 27	5	18
46	Label-free LC-MS analysis of HER2+ breast cancer cell line response to HER2 inhibitor treatment. <i>DARU, Journal of Pharmaceutical Sciences</i> , 2015 , 23, 40	3.9	16
45	Label-free mass spectrometric analysis of the mdx-4cv diaphragm identifies the matricellular protein periostin as a potential factor involved in dystrophinopathy-related fibrosis. <i>Proteomics</i> , 2015 , 15, 2318-31	4.8	37
44	Re-programming CHO cell metabolism using miR-23 tips the balance towards a highly productive phenotype. <i>Biotechnology Journal</i> , 2015 , 10, 1029-40	5.6	37
43	Concurrent Label-Free Mass Spectrometric Analysis of Dystrophin Isoform Dp427 and the Myofibrosis Marker Collagen in Crude Extracts from Skeletal Muscles. <i>Proteomes</i> , 2015 , 3, 298-327	4.6	25
42	Simultaneous Pathoproteomic Evaluation of the Dystrophin-Glycoprotein Complex and Secondary Changes in the mdx-4cv Mouse Model of Duchenne Muscular Dystrophy. <i>Biology</i> , 2015 , 4, 397-423	4.9	27
41	Comparative Label-Free Mass Spectrometric Analysis of Mildly versus Severely Affected mdx Mouse Skeletal Muscles Identifies Annexin, Lamin, and Vimentin as Universal Dystrophic Markers. <i>Molecules</i> , 2015 , 20, 11317-44	4.8	32
40	Elevated levels of 14-3-3 proteins, serotonin, gamma enolase and pyruvate kinase identified in clinical samples from patients diagnosed with colorectal cancer. <i>Clinica Chimica Acta</i> , 2015 , 441, 133-41	6.2	26

39	Glycosylation patterns of kidney proteins differ in rat diabetic nephropathy. <i>Kidney International</i> , 2015 , 87, 963-74	9.9	21
38	Transferrin-bound proteins as potential biomarkers for advanced breast cancer patients. <i>BBA Clinical</i> , 2014 , 2, 24-30		15
37	Identification and functional validation of RAD23B as a potential protein in human breast cancer progression. <i>Journal of Proteome Research</i> , 2014 , 13, 3212-22	5.6	25
36	7B7: a novel antibody directed against the Ku70/Ku80 heterodimer blocks invasion in pancreatic and lung cancer cells. <i>Tumor Biology</i> , 2014 , 35, 6983-97	2.9	10
35	Increased outer arm and core fucose residues on the N-glycans of mutated alpha-1 antitrypsin protein from alpha-1 antitrypsin deficient individuals. <i>Journal of Proteome Research</i> , 2014 , 13, 596-605	5.6	20
34	A neutrophil intrinsic impairment affecting Rab27a and degranulation in cystic fibrosis is corrected by CFTR potentiator therapy. <i>Blood</i> , 2014 , 124, 999-1009	2.2	105
33	Intricate effects of primary motor neuronopathy on contractile proteins and metabolic muscle enzymes as revealed by label-free mass spectrometry. <i>Bioscience Reports</i> , 2014 , 34,	4.1	9
32	PP2A inhibition overcomes acquired resistance to HER2 targeted therapy. <i>Molecular Cancer</i> , 2014 , 13, 157	42.1	34
31	Proteomic profiling of cardiomyopathic tissue from the aged mdx model of Duchenne muscular dystrophy reveals a drastic decrease in laminin, nidogen and annexin. <i>Proteomics</i> , 2013 , 13, 2312-23	4.8	37
30	Recruitment of host translation initiation factor eIF4G by the Vaccinia Virus ssDNA-binding protein I3. <i>Virology</i> , 2012 , 425, 11-22	3.6	20
29	Integrated miRNA, mRNA and protein expression analysis reveals the role of post-transcriptional regulation in controlling CHO cell growth rate. <i>BMC Genomics</i> , 2012 , 13, 656	4.5	68
28	Impact of miR-7 over-expression on the proteome of Chinese hamster ovary cells. <i>Journal of Biotechnology</i> , 2012 , 160, 251-62	3.7	51
27	A novel neutrophil derived inflammatory biomarker of pulmonary exacerbation in cystic fibrosis. <i>Journal of Cystic Fibrosis</i> , 2012 , 11, 100-7	4.1	12
26	Target Prediction Algorithms and Bioinformatics Resources for miRNA Studies 2012 , 29-48		1
25	The use of LC-MS to identify differentially expressed proteins in docetaxel-resistant prostate cancer cell lines. <i>Proteomics</i> , 2012 , 12, 2115-26	4.8	13
24	Utilization and evaluation of CHO-specific sequence databases for mass spectrometry based proteomics. <i>Biotechnology and Bioengineering</i> , 2012 , 109, 1386-94	4.9	42
23	Differential expression of fourteen proteins between uveal melanoma from patients who subsequently developed distant metastases versus those who did Not 2012 , 53, 4634-43		43
22	Proteomic Profiling of Temperature-Shifted CHO Cells to Identify Genes that Impact on Both Cell Growth and Recombinant Protein Productivity in Suspension Culture 2012 , 95-107		

21	Clinical proteomics: liquid chromatography-mass spectrometry purification systems. <i>Methods in Molecular Biology</i> , 2011 , 681, 473-83	1.4	
20	Bleomycin treatment of A549 human lung cancer cells results in association of MGr1-Ag and caveolin-1 in lipid rafts. <i>International Journal of Biochemistry and Cell Biology</i> , 2011 , 43, 98-105	5.6	9
19	Sustained productivity in recombinant Chinese hamster ovary (CHO) cell lines: proteome analysis of the molecular basis for a process-related phenotype. <i>BMC Biotechnology</i> , 2011 , 11, 78	3.5	53
18	2-D DIGE analysis of the mitochondrial proteome from human skeletal muscle reveals time course-dependent remodelling in response to 14 consecutive days of endurance exercise training. <i>Proteomics</i> , 2011 , 11, 1413-28	4.8	56
17	2D-DIGE analysis of phospho-enriched fractions from dasatinib-treated melanoma cell lines. <i>Journal of Proteomics</i> , 2011 , 74, 490-501	3.9	15
16	Proteomic Characterization of An Isogenic Multiple Myeloma Cell Line Model of Bortezomib Resistance. <i>Blood</i> , 2011 , 118, 1820-1820	2.2	
15	Recent advances in clinical proteomics using mass spectrometry. <i>Bioanalysis</i> , 2010 , 2, 1609-15	2.1	15
14	Microarray and proteomics expression profiling identifies several candidates, including the valosin-containing protein (VCP), involved in regulating high cellular growth rate in production CHO cell lines. <i>Biotechnology and Bioengineering</i> , 2010 , 106, 42-56	4.9	60
13	El Antitrypsin regulates human neutrophil chemotaxis induced by soluble immune complexes and IL-8. <i>Journal of Clinical Investigation</i> , 2010 , 120, 4236-50	15.9	191
12	Proteomic analysis of multidrug-resistance mechanisms in adriamycin-resistant variants of DLKP, a squamous lung cancer cell line. <i>Proteomics</i> , 2009 , 9, 1556-66	4.8	45
11	MiRNA-29a regulates the expression of numerous proteins and reduces the invasiveness and proliferation of human carcinoma cell lines. <i>European Journal of Cancer</i> , 2009 , 45, 3104-18	7.5	99
10	Identification of pancreatic cancer invasion-related proteins by proteomic analysis. <i>Proteome Science</i> , 2009 , 7, 3	2.6	54
9	Differential protein expression following low temperature culture of suspension CHO-K1 cells. <i>BMC Biotechnology</i> , 2008 , 8, 42	3.5	54
8	Proteomic profiling of CHO cells with enhanced rhBMP-2 productivity following co-expression of PACEsol. <i>Proteomics</i> , 2008 , 8, 2611-24	4.8	43
7	Proteomic analysis of conditioned media from glucose responsive and glucose non-responsive phenotypes reveals a panel of secreted proteins associated with beta cell dysfunction. <i>Electrophoresis</i> , 2008 , 29, 4141-9	3.6	11
6	Analysis of the saliva proteome from patients with head and neck squamous cell carcinoma reveals differences in abundance levels of proteins associated with tumour progression and metastasis. <i>Journal of Proteomics</i> , 2008 , 71, 168-75	3.9	93
5	Aldehyde dehydrogenase 1A1 and gelsolin identified as novel invasion-modulating factors in conditioned medium of pancreatic cancer cells. <i>Journal of Proteomics</i> , 2008 , 71, 561-71	3.9	15
4	Proteomic investigation of taxol and taxotere resistance and invasiveness in a squamous lung carcinoma cell line. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2008 , 1784, 1184-91	4	21

3	2-D difference gel electrophoresis of the lung squamous cell carcinoma versus normal sera demonstrates consistent alterations in the levels of ten specific proteins. <i>Electrophoresis</i> , 2007 , 28, 4302-10	3.6	65
2	Proteomic analysis of isolated membrane fractions from superinvasive cancer cells. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2007 , 1774, 93-101	4	67
1	Proteomic screening of glucose-responsive and glucose non-responsive MIN-6 beta cells reveals differential expression of proteins involved in protein folding, secretion and oxidative stress. <i>Proteomics</i> , 2006 , 6, 6578-87	4.8	46