# Pier Giorgio Righetti

# List of Publications by Year in Descending Order

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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.

118 642 24,047 70 h-index g-index citations papers 6.76 672 25,426 4.1 avg, IF L-index ext. citations ext. papers

#	Paper	IF	Citations
642	Jack London and White Fang: a lost struggle. <i>Comptes Rendus Chimie</i> , <b>2022</b> , 25, 115-123	2.7	
641	Richard the Lionheart and the Ferocious Saladin Face to Face in Arsuf: A Proteomic Study. <i>Heritage</i> , <b>2021</b> , 4, 3382-3401	1.6	
640	Fiat Lux how Alessandro Volta illuminated his scripts. <i>Comptes Rendus Chimie</i> , <b>2021</b> , 24, 361-371	2.7	O
639	Radial distribution of figures in Leonardo and Renaissance paintings. <i>Digital Applications in Archaeology and Cultural Heritage</i> , <b>2021</b> , 20, e00178	2.1	
638	New baits for fishing in cultural heritage's Mare Magnum. <i>Journal of Proteomics</i> , <b>2021</b> , 235, 104113	3.9	4
637	Software <b>P</b> inxit⊞Hail Magister Leonardo!. <i>Heritage</i> , <b>2021</b> , 4, 917-936	1.6	
636	Meta-proteomic analysis of the Shandrin mammoth by EVA technology and high-resolution mass spectrometry: what is its gut microbiota telling us?. <i>Amino Acids</i> , <b>2021</b> , 53, 1507-1521	3.5	O
635	Combinatorial peptides: A library that continuously probes low-abundance proteins. <i>Electrophoresis</i> , <b>2021</b> ,	3.6	2
634	Proteomics and metabolomics composition of the ink of a letter in a fragment of a Dead Sea Scroll from Cave 11 (P1032-Fr0). <i>Journal of Proteomics</i> , <b>2021</b> , 249, 104370	3.9	1
633	"1984": What Orwell could not predict. Proteomic analysis of his scripts. <i>Electrophoresis</i> , <b>2020</b> , 41, 1931	-139340	2
632	Never boring: Non-invasive palaeoproteomics of mummified human skin. <i>Journal of Archaeological Science</i> , <b>2020</b> , 119, 105145	2.9	5
631	Detection of Plant Low-Abundance Proteins by Means of Combinatorial Peptide Ligand Library Methods. <i>Methods in Molecular Biology</i> , <b>2020</b> , 2139, 381-404	1.4	
630	A novel tool for assessing microbiomes in cultural heritage documents. <i>IOP Conference Series:</i> Materials Science and Engineering, <b>2020</b> , 949, 012116	0.4	
629	EVA Technology and Proteomics: A Two-Pronged Attack on Cultural Heritage. <i>Journal of Proteome Research</i> , <b>2020</b> , 19, 2914-2925	5.6	6
628	Associating 2-DE and CPLLs for low-abundance protein discovery: A winning strategy <b>2020</b> , 183-207		
627	Surface analysis of ancient parchments via the EVA film: The Aleppo Codex. <i>Analytical Biochemistry</i> , <b>2020</b> , 604, 113824	3.1	7
626	Stalin's "black dog": a postmortem diagnosis. Analytical and Bioanalytical Chemistry, 2020, 412, 7701-77	0.8.4	1

# (2017-2020)

625	Low-abundance plant protein enrichment with peptide libraries to enlarge proteome coverage and related applications. <i>Plant Science</i> , <b>2020</b> , 290, 110302	5.3	4
624	What Sherlock sorely missed: the EVA technology for cultural heritage exploration. <i>Expert Review of Proteomics</i> , <b>2019</b> , 16, 533-542	4.2	6
623	De re metallica. Johannes Kepler and alchemy. <i>Talanta</i> , <b>2019</b> , 204, 82-88	6.2	5
622	Il n'y a pas d'amour heureux pour Casanova: Chemical- and bio-analysis of his Memoirs. <i>Electrophoresis</i> , <b>2019</b> , 40, 3050-3056	3.6	8
621	Leonardo's Donna Nuda unveiled. <i>Journal of Proteomics</i> , <b>2019</b> , 207, 103450	3.9	12
620	Progress in farm animal proteomics: The contribution of combinatorial peptide ligand libraries. <i>Journal of Proteomics</i> , <b>2019</b> , 197, 1-13	3.9	11
619	Proteomic fingerprinting of apple fruit, juice, and cider via combinatorial peptide ligand libraries and MS analysis. <i>Electrophoresis</i> , <b>2019</b> , 40, 266-271	3.6	6
618	Noninvasive wearable sensor for indirect glucometry. <i>Electrophoresis</i> , <b>2018</b> , 39, 2344-2350	3.6	3
617	Of mice and men: Traces of life in the death registries of the 1630 plague in Milano. <i>Journal of Proteomics</i> , <b>2018</b> , 180, 128-137	3.9	22
616	Towards the non-invasive proteomic analysis of cultural heritage objects. <i>Microchemical Journal</i> , <b>2018</b> , 139, 450-457	4.8	25
615	Anton Chekhov and Robert Koch Cheek to Cheek: A Proteomic Study. <i>Proteomics</i> , <b>2018</b> , 18, e1700447	4.8	15
614	Protein biomarkers for early detection of diseases: The decisive contribution of combinatorial peptide ligand libraries. <i>Journal of Proteomics</i> , <b>2018</b> , 188, 1-14	3.9	26
613	Novel low-abundance allergens from mango via combinatorial peptide libraries treatment: A proteomics study. <i>Food Chemistry</i> , <b>2018</b> , 269, 652-660	8.5	15
612	Method for Noninvasive Analysis of Proteins and Small Molecules from Ancient Objects. <i>Analytical Chemistry</i> , <b>2017</b> , 89, 3310-3317	7.8	33
611	A miniaturized sensor for detection of formaldehyde fumes. <i>Electrophoresis</i> , <b>2017</b> , 38, 2168-2174	3.6	12
610	Proteomic fingerprinting of mistletoe (Viscum album L.) via combinatorial peptide ligand libraries and mass spectrometry analysis. <i>Journal of Proteomics</i> , <b>2017</b> , 164, 52-58	3.9	8
609	Proteomics of Grapevines and Wines <b>2017</b> , 405-414		
608	Unearthing Bulgakov's trace proteome from the Master i Margarita manuscript. <i>Journal of Proteomics</i> , <b>2017</b> , 152, 102-108	3.9	27

607	Polyphemus, Odysseus and the ovine milk proteome. <i>Journal of Proteomics</i> , <b>2017</b> , 152, 58-74	3.9	10
606	Orange proteomic fingerprinting: From fruit to commercial juices. <i>Food Chemistry</i> , <b>2016</b> , 196, 739-49	8.5	26
605	Identification of plum and peach seed proteins by nLC-MS/MS via combinatorial peptide ligand libraries. <i>Journal of Proteomics</i> , <b>2016</b> , 148, 105-12	3.9	12
604	An in depth proteomic analysis based on ProteoMiner, affinity chromatography and nano-HPLC-MS/MS to explain the potential health benefits of bovine colostrum. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , <b>2016</b> , 121, 297-306	3.5	15
603	Global proteome analysis in plants by means of peptide libraries and applications. <i>Journal of Proteomics</i> , <b>2016</b> , 143, 3-14	3.9	24
602	Maestro, Marguerite, morphine: The last years in the life of Mikhail Bulgakov. <i>Journal of Proteomics</i> , <b>2016</b> , 131, 199-204	3.9	21
601	The secrets of Oriental panacea: Panax ginseng. <i>Journal of Proteomics</i> , <b>2016</b> , 130, 150-9	3.9	15
600	Zeus, Aesculapius, Amalthea and the proteome of goat milk. <i>Journal of Proteomics</i> , <b>2015</b> , 128, 69-82	3.9	21
599	Widening and diversifying the proteome capture by combinatorial peptide ligand libraries via Alcian Blue dye binding. <i>Analytical Chemistry</i> , <b>2015</b> , 87, 4814-20	7.8	15
598	From hundreds to thousands: Widening the normal human Urinome (1). <i>Journal of Proteomics</i> , <b>2015</b> , 112, 53-62	3.9	35
597	Combinatorial peptide ligand libraries as a "Trojan Horse" in deep discovery proteomics. <i>Analytical Chemistry</i> , <b>2015</b> , 87, 293-305	7.8	27
596	A sarabande of tropical fruit proteomics: Avocado, banana, and mango. <i>Proteomics</i> , <b>2015</b> , 15, 1639-45	4.8	15
595	Proteomics of fruits and beverages. Current Opinion in Food Science, 2015, 4, 76-85	9.8	13
594	Extensive heterogeneity of human urokinase, as detected by two-dimensional mapping. <i>Analytical Chemistry</i> , <b>2015</b> , 87, 1509-13	7.8	3
593	Sample treatment methods involving combinatorial peptide ligand libraries for improved proteomes analyses. <i>Methods in Molecular Biology</i> , <b>2015</b> , 1243, 55-82	1.4	8
592	Mixed-bed affinity chromatography: principles and methods. <i>Methods in Molecular Biology</i> , <b>2015</b> , 1286, 131-58	1.4	
591	It's time to pop a cork on champagne's proteome!. Journal of Proteomics, 2014, 105, 351-62	3.9	19
590	Plant proteomics methods to reach low-abundance proteins. <i>Methods in Molecular Biology</i> , <b>2014</b> , 1072, 111-29	1.4	7

589	From hundreds to thousands: Widening the normal human Urinome. <i>Data in Brief</i> , <b>2014</b> , 1, 25-8	1.2	11
588	According to the CPLL proteome sheriffs, not all aperitifs are created equal!. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , <b>2014</b> , 1844, 1493-9	4	5
587	Making Progress in Plant Proteomics for Improved Food Safety. <i>Comprehensive Analytical Chemistry</i> , <b>2014</b> , 64, 131-155	1.9	2
586	The "Dark Side" of Food Stuff Proteomics: The CPLL-Marshals Investigate. <i>Foods</i> , <b>2014</b> , 3, 217-237	4.9	4
585	The Monkey King: a personal view of the long journey towards a proteomic Nirvana. <i>Journal of Proteomics</i> , <b>2014</b> , 107, 39-49	3.9	6
584	The need for agriculture phenotyping: "moving from genotype to phenotype". <i>Journal of Proteomics</i> , <b>2013</b> , 93, 20-39	3.9	17
583	Biomedical Involvements of Low-Abundance Proteins <b>2013</b> , 197-231		
582	Combinatorial peptide libraries to overcome the classical affinity-enrichment methods in proteomics. <i>Amino Acids</i> , <b>2013</b> , 45, 219-29	3.5	17
581	Plant Proteomics and Food and Beverage Analysis via CPLL Capture <b>2013</b> , 159-196		
580	Analytical approaches for the characterization and identification of olive (Olea europaea) oil proteins. <i>Journal of Agricultural and Food Chemistry</i> , <b>2013</b> , 61, 10384-91	5.7	6
580 579		5·7 3.6	4
Ĭ	proteins. Journal of Agricultural and Food Chemistry, <b>2013</b> , 61, 10384-91		
579	proteins. Journal of Agricultural and Food Chemistry, 2013, 61, 10384-91  Bioanalysis: Heri, hodie, cras. Electrophoresis, 2013, 34, 1442-51  The peel and pulp of mango fruit: a proteomic samba. Biochimica Et Biophysica Acta - Proteins and	3.6	4
579 578	proteins. Journal of Agricultural and Food Chemistry, 2013, 61, 10384-91  Bioanalysis: Heri, hodie, cras. Electrophoresis, 2013, 34, 1442-51  The peel and pulp of mango fruit: a proteomic samba. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2013, 1834, 2539-45  Lemon peel and Limoncello liqueur: a proteomic duet. Biochimica Et Biophysica Acta - Proteins and	3.6	4 25
579 578 577	Bioanalysis: Heri, hodie, cras. <i>Electrophoresis</i> , <b>2013</b> , 34, 1442-51  The peel and pulp of mango fruit: a proteomic samba. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , <b>2013</b> , 1834, 2539-45  Lemon peel and Limoncello liqueur: a proteomic duet. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , <b>2013</b> , 1834, 1484-91  Optimized sample treatment protocol by solid-phase peptide libraries to enrich for protein traces.	3.6 4	4 25 14
579 578 577 576	Bioanalysis: Heri, hodie, cras. <i>Electrophoresis</i> , <b>2013</b> , 34, 1442-51  The peel and pulp of mango fruit: a proteomic samba. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , <b>2013</b> , 1834, 2539-45  Lemon peel and Limoncello liqueur: a proteomic duet. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , <b>2013</b> , 1834, 1484-91  Optimized sample treatment protocol by solid-phase peptide libraries to enrich for protein traces. <i>Amino Acids</i> , <b>2013</b> , 45, 1431-42	3.6 4	4 25 14
579 578 577 576 575	Bioanalysis: Heri, hodie, cras. <i>Electrophoresis</i> , <b>2013</b> , 34, 1442-51  The peel and pulp of mango fruit: a proteomic samba. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , <b>2013</b> , 1834, 2539-45  Lemon peel and Limoncello liqueur: a proteomic duet. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , <b>2013</b> , 1834, 1484-91  Optimized sample treatment protocol by solid-phase peptide libraries to enrich for protein traces. <i>Amino Acids</i> , <b>2013</b> , 45, 1431-42  Current Gel Electrophoresis Approaches 'to Low-Abundance Protein 'Marker Discovery <b>2013</b> , 175-190  In-depth proteomic analysis of banana (Musa spp.) fruit with combinatorial peptide ligand libraries.	3.6 4 4 3.5	4 25 14 9

571	A decade of plant proteomics and mass spectrometry: translation of technical advancements to food security and safety issues. <i>Mass Spectrometry Reviews</i> , <b>2013</b> , 32, 335-65	11	59
57°	Proteomic analysis of Lycium barbarum (Goji) fruit via combinatorial peptide ligand libraries. <i>Electrophoresis</i> , <b>2013</b> , 34, 1729-1736	3.6	9
569	Combinatorial ligand libraries as a two-dimensional method for proteome analysis. <i>Journal of Chromatography A</i> , <b>2013</b> , 1297, 106-12	4.5	17
568	Artichoke and Cynar liqueur: two (not quite) entangled proteomes. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , <b>2013</b> , 1834, 119-26	4	15
567	Chromatographic and Electrophoretic Prefractionation Tools in Proteome Analysis <b>2013</b> , 13-40		
566	Introducing Low-Abundance Species in Proteome Analysis <b>2013</b> , 1-11		1
565	Low-Abundance Protein Access by Combinatorial Peptide Libraries 2013, 79-157		
564	Other Applications of Combinatorial Peptide Libraries <b>2013</b> , 233-261		
563	Current Low-Abundance Protein Access 2013, 41-77		1
562	Detailed Methodologies and Protocols <b>2013</b> , 263-319		2
562 561	Detailed Methodologies and Protocols <b>2013</b> , 263-319  Identification of olive (Olea europaea) seed and pulp proteins by nLC-MS/MS via combinatorial peptide ligand libraries. <i>Journal of Proteomics</i> , <b>2012</b> , 75, 2396-403	3.9	32
	Identification of olive (Olea europaea) seed and pulp proteins by nLC-MS/MS via combinatorial	3.9	
561	Identification of olive (Olea europaea) seed and pulp proteins by nLC-MS/MS via combinatorial peptide ligand libraries. <i>Journal of Proteomics</i> , <b>2012</b> , 75, 2396-403  Allergomic study of cypress pollen via combinatorial peptide ligand libraries. <i>Journal of Proteomics</i> ,		32
561 560	Identification of olive (Olea europaea) seed and pulp proteins by nLC-MS/MS via combinatorial peptide ligand libraries. <i>Journal of Proteomics</i> , <b>2012</b> , 75, 2396-403  Allergomic study of cypress pollen via combinatorial peptide ligand libraries. <i>Journal of Proteomics</i> , <b>2012</b> , 77, 101-10  "Cheek-to-cheek" urinary proteome profiling via combinatorial peptide ligand libraries: A novel,	3.9	32
561 560 559	Identification of olive (Olea europaea) seed and pulp proteins by nLC-MS/MS via combinatorial peptide ligand libraries. <i>Journal of Proteomics</i> , <b>2012</b> , 75, 2396-403  Allergomic study of cypress pollen via combinatorial peptide ligand libraries. <i>Journal of Proteomics</i> , <b>2012</b> , 77, 101-10  "Cheek-to-cheek" urinary proteome profiling via combinatorial peptide ligand libraries: A novel, unexpected elution system. <i>Journal of Proteomics</i> , <b>2012</b> , 75, 796-805  Resurrexit, sicut dixit, alleluia. Snake venomics from a 26-year old polyacrylamide focusing gel.	3.9	3 <sup>2</sup> 3 <sup>2</sup> 26
561 560 559 558	Identification of olive (Olea europaea) seed and pulp proteins by nLC-MS/MS via combinatorial peptide ligand libraries. <i>Journal of Proteomics</i> , <b>2012</b> , 75, 2396-403  Allergomic study of cypress pollen via combinatorial peptide ligand libraries. <i>Journal of Proteomics</i> , <b>2012</b> , 77, 101-10  "Cheek-to-cheek" urinary proteome profiling via combinatorial peptide ligand libraries: A novel, unexpected elution system. <i>Journal of Proteomics</i> , <b>2012</b> , 75, 796-805  Resurrexit, sicut dixit, alleluia. Snake venomics from a 26-year old polyacrylamide focusing gel. <i>Journal of Proteomics</i> , <b>2012</b> , 75, 1074-8	3·9 3·9	32 32 26
561 560 559 558 557	Identification of olive (Olea europaea) seed and pulp proteins by nLC-MS/MS via combinatorial peptide ligand libraries. <i>Journal of Proteomics</i> , <b>2012</b> , 75, 2396-403  Allergomic study of cypress pollen via combinatorial peptide ligand libraries. <i>Journal of Proteomics</i> , <b>2012</b> , 77, 101-10  "Cheek-to-cheek" urinary proteome profiling via combinatorial peptide ligand libraries: A novel, unexpected elution system. <i>Journal of Proteomics</i> , <b>2012</b> , 75, 796-805  Resurrexit, sicut dixit, alleluia. Snake venomics from a 26-year old polyacrylamide focusing gel. <i>Journal of Proteomics</i> , <b>2012</b> , 75, 1074-8  Harry Belafonte and the secret proteome of coconut milk. <i>Journal of Proteomics</i> , <b>2012</b> , 75, 914-20	3.9 3.9 3.9	32 32 26 6

553	Assessment of the floral origin of honey via proteomic tools. <i>Journal of Proteomics</i> , <b>2012</b> , 75, 3688-93	3.9	53
552	Mark Twain: how to fathom the depth of your pet proteome. <i>Journal of Proteomics</i> , <b>2012</b> , 75, 4783-91	3.9	9
551	Breakfast at Tiffany's? Only with a low-abundance proteomic signature!. Electrophoresis, 2012, 33, 2228	-38	15
550	Identification of avocado (Persea americana) pulp proteins by nano-LC-MS/MS via combinatorial peptide ligand libraries. <i>Electrophoresis</i> , <b>2012</b> , 33, 2799-805	3.6	33
549	Polar electrophoresis: shape of two-dimensional maps is as important as size. <i>PLoS ONE</i> , <b>2012</b> , 7, e3091	13.7	4
548	"The quest for biomarkers": are we on the right technical track?. <i>Proteomics - Clinical Applications</i> , <b>2012</b> , 6, 22-41	3.1	20
547	Exploration of the sea urchin coelomic fluid via combinatorial peptide ligand libraries. <i>Biological Bulletin</i> , <b>2012</b> , 222, 93-104	1.5	8
546	Mixed Beds. Beyond the Frontiers of Classical Chromatography for Proteins. <i>Advances in Chromatography</i> , <b>2012</b> , 50, 1-46		1
545	Conventional isoelectric focusing. In gel slabs and capillaries and immobilized pH gradients. <i>Methods of Biochemical Analysis</i> , <b>2011</b> , 54, 379-409		4
544	Mehercules, adhuc Bacchus! The debate on wine proteomics continues. <i>Journal of Proteome Research</i> , <b>2011</b> , 10, 3789-801	5.6	32
543	Poppea's bath liquor: the secret proteome of she-donkey's milk. <i>Journal of Proteomics</i> , <b>2011</b> , 74, 2083-9	<b>9</b> .9	39
542	Cibacron Blue and proteomics: the mystery of the platoon missing in action. <i>Journal of Proteomics</i> , <b>2011</b> , 74, 2856-65	3.9	8
541	Horam nonam exclamavit: sitio. The trace proteome of your daily vinegar. <i>Journal of Proteomics</i> , <b>2011</b> , 75, 718-24	3.9	13
540	Facing challenges in Proteomics today and in the coming decade: Report of Roundtable Discussions at the 4th EuPA Scientific Meeting, Portugal, Estoril 2010. <i>Journal of Proteomics</i> , <b>2011</b> , 75, 4-17	3.9	8
539	Recent advances in electrophoretic techniques for the characterization of protein biomolecules: a poker of aces. <i>Journal of Chromatography A</i> , <b>2011</b> , 1218, 8727-37	4.5	21
538	Investigation of the applicability of Zernike moments to the classification of SDS 2D-PAGE maps. <i>Analytical and Bioanalytical Chemistry</i> , <b>2011</b> , 400, 1419-31	4.4	7
537	Mixed-bed chromatography as a way to resolve peculiar protein fractionation situations. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , <b>2011</b> , 879, 827-35	3.2	3
536	Combinatorial peptide ligand libraries: the conquest of the 'hidden proteome' advances at great strides. <i>Electrophoresis</i> , <b>2011</b> , 32, 960-6	3.6	38

535	"Proteomineering" serum biomarkers. A study in scarlet. <i>Electrophoresis</i> , <b>2011</b> , 32, 976-80	3.6	9
534	Plasma proteomics for biomarker discovery: a study in blue. <i>Electrophoresis</i> , <b>2011</b> , 32, 3638-44	3.6	11
533	Going nuts for nuts? The trace proteome of a Cola drink, as detected via combinatorial peptide ligand libraries. <i>Journal of Proteome Research</i> , <b>2011</b> , 10, 2684-6	5.6	20
532	Popeye strikes again: The deep proteome of spinach leaves. <i>Journal of Proteomics</i> , <b>2011</b> , 74, 127-36	3.9	42
531	"Proteomineering" or not? The debate on biomarker discovery in sera continues. <i>Journal of Proteomics</i> , <b>2011</b> , 74, 589-94	3.9	31
530	In-depth proteomic analysis of non-alcoholic beverages with peptide ligand libraries. I: Almond milk and orgeat syrup. <i>Journal of Proteomics</i> , <b>2011</b> , 74, 1080-90	3.9	36
529	Capturing and amplifying impurities from recombinant therapeutic proteins via combinatorial peptide libraries: a proteomic approach. <i>Current Pharmaceutical Biotechnology</i> , <b>2011</b> , 12, 1537-47	2.6	8
528	Revisiting Jurassic Park: The Isolation of Proteins from Amber Encapsulated Organisms Millions of Years Old <b>2011</b> , 925-938		2
527	Specific and Surrogate Cerebrospinal Fluid Markers in Creutzfeldt Dakob Disease. <i>Advances in Neurobiology</i> , <b>2011</b> , 455-467	2.1	2
526	Wider Protein Detection from Biological Extracts by the Reduction of the Dynamic Concentration Range <b>2010</b> , 175-204		
525	In-depth exploration of cerebrospinal fluid by combining peptide ligand library treatment and label-free protein quantification. <i>Molecular and Cellular Proteomics</i> , <b>2010</b> , 9, 1006-21	7.6	106
524	The proteome buccaneers: how to unearth your treasure chest via combinatorial peptide ligand libraries. <i>Expert Review of Proteomics</i> , <b>2010</b> , 7, 373-85	4.2	59
523	Les Mattres de l'Orge: the proteome content of your beer mug. <i>Journal of Proteome Research</i> , <b>2010</b> , 9, 5262-9	5.6	63
522	The red blood cell proteome and interactome: an update. <i>Journal of Proteome Research</i> , <b>2010</b> , 9, 144-6	<b>3</b> 5.6	152
521	In depth exploration of the hemolymph of Limulus polyphemus via combinatorial peptide ligand libraries. <i>Journal of Proteome Research</i> , <b>2010</b> , 9, 3260-9	5.6	10
520	SDS-PAGE and two-dimensional maps in a radial gel format. <i>Electrophoresis</i> , <b>2010</b> , 31, 465-70	3.6	7
519	Third generation of focusing: gel matrices with immobilized cation gradients. <i>Electrophoresis</i> , <b>2010</b> , 31, 1747-53	3.6	1
518	Analysis of trace degradation products (decarboxylated diastereoisomers) of S-adenosylmethionine by electrophoresis in capillaries with cationic coatings (N-methylpolyvinylpyridinium or divalent barium). <i>Electrophoresis</i> , <b>2010</b> , 31, 3592-6	3.6	2

517	Improved instrumentation for large-size two-dimensional protein maps. <i>Electrophoresis</i> , <b>2010</b> , 31, 3863-	<b>-6</b> .6	5
516	Interaction among proteins and peptide libraries in proteome analysis: pH involvement for a larger capture of species. <i>Journal of Proteomics</i> , <b>2010</b> , 73, 733-42	3.9	62
515	Exploring the venom proteome of the African puff adder, Bitis arietans, using a combinatorial peptide ligand library approach at different pHs. <i>Journal of Proteomics</i> , <b>2010</b> , 73, 932-42	3.9	35
514	The egg white and yolk interactomes as gleaned from extensive proteomic data. <i>Journal of Proteomics</i> , <b>2010</b> , 73, 1028-42	3.9	28
513	In-depth exploration of Hevea brasiliensis latex proteome and "hidden allergens" via combinatorial peptide ligand libraries. <i>Journal of Proteomics</i> , <b>2010</b> , 73, 1368-80	3.9	64
512	Proteomics of wine additives: mining for the invisible via combinatorial peptide ligand libraries. Journal of Proteomics, <b>2010</b> , 73, 1732-9	3.9	65
511	Noah's nectar: the proteome content of a glass of red wine. <i>Journal of Proteomics</i> , <b>2010</b> , 73, 2370-7	3.9	59
510	Plucking, pillaging and plundering proteomes with combinatorial peptide ligand libraries. <i>Journal of Chromatography A</i> , <b>2010</b> , 1217, 893-900	4.5	36
509	Congenital dyserythropoietic anemia type II (CDAII) is caused by mutations in the SEC23B gene. <i>Human Mutation</i> , <b>2009</b> , 30, 1292-8	4.7	132
508	Immobilized pH gradients. <i>Electrophoresis</i> , <b>2009</b> , 30 Suppl 1, S112-21	3.6	15
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	An N-methylpolyvinylpyridinium cationic polymer for capillary coating in electrophoresis of		
507	An N-methylpolyvinylpyridinium cationic polymer for capillary coating in electrophoresis of proteins and peptides. <i>Electrophoresis</i> , <b>2009</b> , 30, 2313-20  Combinatorial peptide ligand libraries for urine proteome analysis: investigation of different	3.6	16
507	An N-methylpolyvinylpyridinium cationic polymer for capillary coating in electrophoresis of proteins and peptides. <i>Electrophoresis</i> , <b>2009</b> , 30, 2313-20  Combinatorial peptide ligand libraries for urine proteome analysis: investigation of different elution systems. <i>Electrophoresis</i> , <b>2009</b> , 30, 2405-11  Steady-state electrophoresis of RNA against a gradient of cationic charges in a polyacrylamide	3.6 3.6	16 91
507 506 505	An N-methylpolyvinylpyridinium cationic polymer for capillary coating in electrophoresis of proteins and peptides. <i>Electrophoresis</i> , <b>2009</b> , 30, 2313-20  Combinatorial peptide ligand libraries for urine proteome analysis: investigation of different elution systems. <i>Electrophoresis</i> , <b>2009</b> , 30, 2405-11  Steady-state electrophoresis of RNA against a gradient of cationic charges in a polyacrylamide matrix. <i>Electrophoresis</i> , <b>2009</b> , 30, 3696-700  Synergistic effect of trichostatin A and 5-aza-2'-deoxycytidine on growth inhibition of pancreatic	3.6 3.6 3.6	16 91 1
507 506 505	An N-methylpolyvinylpyridinium cationic polymer for capillary coating in electrophoresis of proteins and peptides. <i>Electrophoresis</i> , <b>2009</b> , 30, 2313-20  Combinatorial peptide ligand libraries for urine proteome analysis: investigation of different elution systems. <i>Electrophoresis</i> , <b>2009</b> , 30, 2405-11  Steady-state electrophoresis of RNA against a gradient of cationic charges in a polyacrylamide matrix. <i>Electrophoresis</i> , <b>2009</b> , 30, 3696-700  Synergistic effect of trichostatin A and 5-aza-2'-deoxycytidine on growth inhibition of pancreatic endocrine tumour cell lines: a proteomic study. <i>Proteomics</i> , <b>2009</b> , 9, 1952-66  The art of observing rare protein species in proteomes with peptide ligand libraries. <i>Proteomics</i> ,	3.6 3.6 3.6 4.8	16 91 1 34
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#### (1984-1985)

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2	Gel-Based Proteomics9-31		О

Steady-state electrolysis of a solution of nonamphotheric compounds138-142