Sarah Cianferani

List of Publications by Year in descending order

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Version: 2024-02-01

147801 182427 3,724 143 31 51 citations h-index g-index papers 151 151 151 5163 docs citations times ranked citing authors all docs

#	Article	lF	Citations
1	Proteomic analysis of Rhodospirillum rubrum after carbon monoxide exposure reveals an important effect on metallic cofactor biosynthesis. Journal of Proteomics, 2022, 250, 104389.	2.4	3
2	Probing the mechanism of peroxiredoxin decamer interaction with its reductase sulfiredoxin from the single molecule to the solution scale. Nanoscale Horizons, 2022, , .	8.0	0
3	A Confinementâ€Driven Nucleation Mechanism of Metal Oxide Nanoparticles Obtained via Thermal Decomposition in Organic Media. Small, 2022, 18, e2200414.	10.0	5
4	Repeated Exposure of Macrophages to Synthetic Amorphous Silica Induces Adaptive Proteome Changes and a Moderate Cell Activation. Nanomaterials, 2022, 12, 1424.	4.1	3
5	Combination of IM-Based Approaches to Unravel the Coexistence of Two Conformers on a Therapeutic Multispecific mAb. Analytical Chemistry, 2022, 94, 7981-7989.	6.5	9
6	Does size matter? A proteomics-informed comparison of the effects of polystyrene beads of different sizes on macrophages. Environmental Science: Nano, 2022, 9, 2827-2840.	4.3	4
7	Upconversion in a d–f [RuYb ₃] Supramolecular Assembly. Journal of the American Chemical Society, 2022, 144, 13356-13365.	13.7	16
8	Structural and Biochemical Investigation of the Heterodimeric Murine tRNA-Guanine Transglycosylase. ACS Chemical Biology, 2022, 17, 2229-2247.	3.4	7
9	Optimized Sample Preparation and Data Processing of Data-Independent Acquisition Methods for the Robust Quantification of Trace-Level Host Cell Protein Impurities in Antibody Drug Products. Journal of Proteome Research, 2021, 20, 923-931.	3.7	17
10	Fast Afucosylation Profiling of Glycoengineered Antibody Subunits by Middle-Up Mass Spectrometry. Methods in Molecular Biology, 2021, 2271, 73-83.	0.9	2
11	Structural studies of RNase M5 reveal two-metal-ion supported two-step dsRNA cleavage for 5S rRNA maturation. RNA Biology, 2021, 18, 1-11.	3.1	1
12	Combining labelâ€free and labelâ€based accurate quantifications with SWATHâ€MS: Comparison with SRM and PRM for the evaluation of bovine muscle type effects. Proteomics, 2021, 21, e2000214.	2.2	5
13	Pushing the limits of native MS: Online SEC-native MS for structural biology applications. International Journal of Mass Spectrometry, 2021, 461, 116502.	1.5	22
14	A structural signature motif enlightens the origin and diversification of nuclear receptors. PLoS Genetics, 2021, 17, e1009492.	3.5	8
15	A Class of Valuable (Pro-)Activity-Based Protein Profiling Probes: Application to the Redox-Active Antiplasmodial Agent, Plasmodione. Jacs Au, 2021, 1, 669-689.	7.9	4
16	Temporal multiomic modeling reveals a B-cell receptor proliferative program in chronic lymphocytic leukemia. Leukemia, 2021, 35, 1463-1474.	7.2	6
17	Proteo3Dnet: a web server for the integration of structural information with interactomics data. Nucleic Acids Research, 2021, 49, W567-W572.	14.5	5
18	A proteomic view of cellular responses of macrophages to copper when added as ion or as copper-polyacrylate complex. Journal of Proteomics, 2021, 239, 104178.	2.4	1

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19	A proteomic-informed view of the changes induced by loss of cellular adherence: The example of mouse macrophages. PLoS ONE, 2021, 16, e0252450.	2.5	2
20	State-of-the-Art Native Mass Spectrometry and Ion Mobility Methods to Monitor Homogeneous Site-Specific Antibody-Drug Conjugates Synthesis. Pharmaceuticals, 2021, 14, 498.	3.8	16
21	High-Resolution IMS–MS to Assign Additional Disulfide Bridge Pairing in Complementarity-Determining Regions of an IgG4 Monoclonal Antibody. Journal of the American Society for Mass Spectrometry, 2021, 32, 2505-2512.	2.8	13
22	Iron Stearate Structures: An Original Tool for Nanoparticles Design. Inorganic Chemistry, 2021, 60, 12445-12456.	4.0	14
23	Rab7-harboring vesicles are carriers of the transferrin receptor through the biosynthetic secretory pathway. Science Advances, 2021, 7, .	10.3	11
24	Strongly Adhesive and Antimicrobial Peptide-Loaded, Alginate–Catechol-Based Gels for Application against Periimplantitis. Applied Sciences (Switzerland), 2021, 11, 10050.	2.5	2
25	Hands on Native Mass Spectrometry Analysis of Multi-protein Complexes. Methods in Molecular Biology, 2021, 2247, 173-191.	0.9	1
26	Bicyclo[6.1.0]nonyne carboxylic acid for the production of stable molecular probes. RSC Advances, 2021, 11, 36777-36780.	3.6	0
27	Toxoplasma gondii ROP16 kinase silences the cyclin B1 gene promoter by hijacking host cell UHRF1-dependent epigenetic pathways. Cellular and Molecular Life Sciences, 2020, 77, 2141-2156.	5.4	15
28	VHH characterization.Recombinant VHHs: Production, characterization and affinity. Analytical Biochemistry, 2020, 589, 113491.	2.4	9
29	Homogeneous antibody-drug conjugates: DAR 2 anti-HER2 obtained by conjugation on isolated light chain followed by mAb assembly. MAbs, 2020, 12, 1702262.	5.2	9
30	Automated linkage of proteins and payloads producing monodisperse conjugates. Chemical Science, 2020, 11, 1210-1215.	7.4	19
31	Structural basis for DNA recognition and allosteric control of the retinoic acid receptors RAR–RXR. Nucleic Acids Research, 2020, 48, 9969-9985.	14.5	17
32	Molecular determinants of MED1 interaction with the DNA bound VDR–RXR heterodimer. Nucleic Acids Research, 2020, 48, 11199-11213.	14.5	17
33	Structures of B.Âsubtilis Maturation RNases Captured on 50S Ribosome with Pre-rRNAs. Molecular Cell, 2020, 80, 227-236.e5.	9.7	12
34	How Reversible Are the Effects of Fumed Silica on Macrophages? A Proteomics-Informed View. Nanomaterials, 2020, 10, 1939.	4.1	7
35	Software Requirements for the Analysis and Interpretation of Native Ion Mobility Mass Spectrometry Data. Analytical Chemistry, 2020, 92, 10881-10890.	6.5	17
36	Computational Strategies and Challenges for Using Native Ion Mobility Mass Spectrometry in Biophysics and Structural Biology. Analytical Chemistry, 2020, 92, 10872-10880.	6.5	24

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37	Frontispiece: Investigating Ugi/Passerini Multicomponent Reactions for the Siteâ€Selective Conjugation of Native Trastuzumab. Chemistry - A European Journal, 2020, 26, .	3.3	O
38	The Importance of Charge in Perturbing the Aromatic Glue Stabilizing the Protein–Protein Interface of Homodimeric tRNA-Guanine Transglycosylase. ACS Chemical Biology, 2020, 15, 3021-3029.	3.4	3
39	Toward Automation of Collision-Induced Unfolding Experiments through Online Size Exclusion Chromatography Coupled to Native Mass Spectrometry. Analytical Chemistry, 2020, 92, 12900-12908.	6.5	18
40	On the use of DNA as a linker in antibody-drug conjugates: synthesis, stability and in vitro potency. Scientific Reports, 2020, 10, 7691.	3.3	20
41	Middle Level IM–MS and CIU Experiments for Improved Therapeutic Immunoglobulin Subclass Fingerprinting. Analytical Chemistry, 2020, 92, 8827-8835.	6.5	14
42	Glycan-Mediated Technology for Obtaining Homogeneous Site-Specific Conjugated Antibody–Drug Conjugates: Synthesis and Analytical Characterization by Using Complementary Middle-up LC/HRMS Analysis. Analytical Chemistry, 2020, 92, 8170-8177.	6.5	17
43	${\rm A\hat{l}^2(1\text{-}42)}$ tetramer and octamer structures reveal edge conductivity pores as a mechanism for membrane damage. Nature Communications, 2020, 11, 3014.	12.8	162
44	Ethynylation of Cysteine Residues: From Peptides to Proteins in Vitro and in Living Cells. Angewandte Chemie - International Edition, 2020, 59, 10961-10970.	13.8	46
45	The longer the worse: a combined proteomic and targeted study of the long-term <i>versus</i> short-term effects of silver nanoparticles on macrophages. Environmental Science: Nano, 2020, 7, 2032-2046.	4.3	11
46	Investigating Ugi/Passerini Multicomponent Reactions for the Siteâ€Selective Conjugation of Native Trastuzumab**. Chemistry - A European Journal, 2020, 26, 13797-13805.	3.3	17
47	Proline: an efficient and user-friendly software suite for large-scale proteomics. Bioinformatics, 2020, 36, 3148-3155.	4.1	155
48	(Thia)calixarenephosphonic Acids as Potent Inhibitors of the Nucleic Acid Chaperone Activity of the HIV-1 Nucleocapsid Protein with a New Binding Mode and Multitarget Antiviral Activity. ACS Infectious Diseases, 2020, 6, 687-702.	3.8	9
49	Determination of size variants by CE-SDS for approved therapeutic antibodies: Key implications of subclasses and light chain specificities. Journal of Pharmaceutical and Biomedical Analysis, 2020, 184, 113166.	2.8	30
50	Probing Protein Interaction Networks by Combining MS-Based Proteomics and Structural Data Integration. Journal of Proteome Research, 2020, 19, 2807-2820.	3.7	6
51	Insight on the Impact of the Reduction Step on the Siteâ€Directed Conjugation of an Antiâ€HER2 Cysteineâ€Engineered Antibody. ChemistrySelect, 2020, 5, 3187-3190.	1.5	0
52	Drug Loading and Distribution of ADCs After Reduction or IdeS Digestion and Reduction. Methods in Molecular Biology, 2020, 2078, 187-195.	0.9	2
53	Analysis of ADCs by Native Mass Spectrometry. Methods in Molecular Biology, 2020, 2078, 197-211.	0.9	2
54	VHH characterization. Comparison of recombinant with chemically synthesized antiâ∈HER2 VHH. Protein Science, 2019, 28, 1865-1879.	7.6	16

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55	A Case Study to Identify the Drug Conjugation Site of a Site-Specific Antibody-Drug-Conjugate Using Middle-Down Mass Spectrometry. Journal of the American Society for Mass Spectrometry, 2019, 30, 2419-2429.	2.8	23
56	An Investigation into the Stephens–Castro Synthesis of Dehydrotriaryl[12]annulenes: Factors Influencing the Cyclotrimerization. European Journal of Organic Chemistry, 2019, 2019, 6783-6795.	2.4	1
57	Cutting-edge multi-level analytical and structural characterization of antibody-drug conjugates: present and future. Expert Review of Proteomics, 2019, 16, 337-362.	3.0	47
58	A Proteomic View of Cellular Responses to Anticancer Quinoline-Copper Complexes. Proteomes, 2019, 7, 26.	3 . 5	12
59	A Combined N-terminomics and Shotgun Proteomics Approach to Investigate the Responses of Human Cells to Rapamycin and Zinc at the Mitochondrial Level. Molecular and Cellular Proteomics, 2019, 18, 1085-1095.	3.8	3
60	How reversible are the effects of silver nanoparticles on macrophages? A proteomic-instructed view. Environmental Science: Nano, 2019, 6, 3133-3157.	4.3	21
61	Synthesis and biological evaluation of 2.4 nm thiolate-protected gold nanoparticles conjugated to Cetuximab for targeting glioblastoma cancer cells via the EGFR. Nanotechnology, 2019, 30, 184005.	2.6	24
62	Recurrent activating mutations of PPAR \hat{I}^3 associated with luminal bladder tumors. Nature Communications, 2019, 10, 253.	12.8	44
63	Reduction–rebridging strategy for the preparation of ADPN-based antibody–drug conjugates. MedChemComm, 2018, 9, 827-830.	3.4	24
64	Hyphenation of size exclusion chromatography to native ion mobility mass spectrometry for the analytical characterization of therapeutic antibodies and related products. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2018, 1086, 176-183.	2.3	69
65	Arginine-selective bioconjugation with 4-azidophenyl glyoxal: application to the single and dual functionalisation of native antibodies. Organic and Biomolecular Chemistry, 2018, 16, 1305-1311.	2.8	30
66	An Online Four-Dimensional HIC×SEC-IM×MS Methodology for Proof-of-Concept Characterization of Antibody Drug Conjugates. Analytical Chemistry, 2018, 90, 1578-1586.	6.5	75
67	Adding a new separation dimension to MS and LC–MS: What is the utility of ion mobility spectrometry?. Journal of Separation Science, 2018, 41, 20-67.	2.5	140
68	Peptide deformylases from Vibrio parahaemolyticus phage and bacteria display similar deformylase activity and inhibitor binding clefts. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2018, 1866, 348-355.	2.3	4
69	Dual Data-Independent Acquisition Approach Combining Global HCP Profiling and Absolute Quantification of Key Impurities during Bioprocess Development. Analytical Chemistry, 2018, 90, 1241-1247.	6.5	26
70	Structure-Based Identification of HIV-1 Nucleocapsid Protein Inhibitors Active against Wild-Type and Drug-Resistant HIV-1 Strains. ACS Chemical Biology, 2018, 13, 253-266.	3.4	13
71	Combinatorial regulation of hepatic cytoplasmic signaling and nuclear transcriptional events by the OGT/REV-ERBα complex. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E11033-E11042.	7.1	35
72	A quantitative proteomic analysis of cofilin phosphorylation in myeloid cells and its modulation using the LIM kinase inhibitor Pyr1. PLoS ONE, 2018, 13, e0208979.	2.5	11

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73	Morphine Binds Creatine Kinase B and Inhibits Its Activity. Frontiers in Cellular Neuroscience, 2018, 12, 464.	3.7	7
74	A Novel Online Four-Dimensional SEC×SEC-IM×MS Methodology for Characterization of Monoclonal Antibody Size Variants. Analytical Chemistry, 2018, 90, 13929-13937.	6.5	49
75	The H ₂ O ₂ -Resistant Fe–S Redox Switch MitoNEET Acts as a pH Sensor To Repair Stress-Damaged Fe–S Protein. Biochemistry, 2018, 57, 5616-5628.	2.5	16
76	The RPAP3-Cterminal domain identifies R2TP-like quaternary chaperones. Nature Communications, 2018, 9, 2093.	12.8	59
77	Extended investigation of tube-gel sample preparation: a versatile and simple choice for high throughput quantitative proteomics. Scientific Reports, 2018, 8, 8260.	3.3	3
78	Formation of Mono- and Polynuclear Luminescent Lanthanide Complexes based on the Coordination of Preorganized Phosphonated Pyridines. Inorganic Chemistry, 2018, 57, 6095-6106.	4.0	21
79	Native Mass Spectrometry, Ion Mobility, and Collision-Induced Unfolding for Conformational Characterization of IgG4 Monoclonal Antibodies. Analytical Chemistry, 2018, 90, 8865-8872.	6.5	51
80	Stable isotopeâ€labelled morphine to study ⟨i⟩in vivo⟨ i⟩ central and peripheral morphine glucuronidation and brain transport in tolerant mice. British Journal of Pharmacology, 2018, 175, 3844-3856.	5 . 4	10
81	Austausch der Proteinkontaktflähen in der homodimeren tRNAâ€Guaninâ€Transglycosylase: ein Weg der funktionellen Regulation. Angewandte Chemie, 2018, 130, 10242-10247.	2.0	2
82	Multi-OMICS analyses unveil <i>STAT1</i> as a potential modifier gene in mevalonate kinase deficiency. Annals of the Rheumatic Diseases, 2018, 77, 1675-1687.	0.9	19
83	The ESCRT-III Protein CHMP1A Mediates Secretion of Sonic Hedgehog on a Distinctive Subtype of Extracellular Vesicles. Cell Reports, 2018, 24, 973-986.e8.	6.4	79
84	Deep Structural Analysis of RPAP3 and PIH1D1, Two Components of the HSP90 Co-chaperone R2TP Complex. Structure, 2018, 26, 1196-1209.e8.	3.3	36
85	Cytosolic Diffusion and Peptide-Assisted Nuclear Shuttling of Peptide-Substituted Circa 102 Gold Atom Nanoclusters in Living Cells. ACS Applied Nano Materials, 2018, 1, 4236-4246.	5.0	10
86	The nuclear bile acid receptor FXR is a PKA- and FOXA2-sensitive activator of fasting hepatic gluconeogenesis. Journal of Hepatology, 2018, 69, 1099-1109.	3.7	40
87	Evolutionary insights into Trm112-methyltransferase holoenzymes involved in translation between archaea and eukaryotes. Nucleic Acids Research, 2018, 46, 8483-8499.	14.5	37
88	Homodimer Architecture of QTRT2, the Noncatalytic Subunit of the Eukaryotic tRNA-Guanine Transglycosylase. Biochemistry, 2018, 57, 3953-3965.	2.5	8
89	Swapping Interface Contacts in the Homodimeric tRNAâ€Guanine Transglycosylase: An Option for Functional Regulation. Angewandte Chemie - International Edition, 2018, 57, 10085-10090.	13.8	10
90	Activation mode of the eukaryotic m ² G ₁₀ tRNA methyltransferase Trm11 by its partner protein Trm112. Nucleic Acids Research, 2017, 45, gkw1271.	14.5	23

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91	Generic Hybrid Ligand Binding Assay Liquid Chromatography High-Resolution Mass Spectrometry-Based Workflow for Multiplexed Human Immunoglobulin G1 Quantification at the Intact Protein Level: Application to Preclinical Pharmacokinetic Studies. Analytical Chemistry, 2017, 89, 2628-2635.	6.5	39
92	Development of a fast workflow to screen the charge variants of therapeutic antibodies. Journal of Chromatography A, 2017, 1498, 147-154.	3.7	31
93	Insights from native mass spectrometry approaches for top- and middle- level characterization of site-specific antibody-drug conjugates. MAbs, 2017, 9, 801-811.	5.2	55
94	Acyl Fluorides: Fast, Efficient, and Versatile Lysine-Based Protein Conjugation via Plug-and-Play Strategy. Bioconjugate Chemistry, 2017, 28, 1452-1457.	3.6	31
95	Differential proteomics highlights macrophage-specific responses to amorphous silica nanoparticles. Nanoscale, 2017, 9, 9641-9658.	5.6	31
96	A Direct Oneâ€Pot Synthesis of Asymmetric Dehydrobenzopyrido[12]annulenes and Their Physicochemical Properties. European Journal of Organic Chemistry, 2017, 2017, 4625-4632.	2.4	1
97	The intricate network between the p34 and p44 subunits is central to the activity of the transcription/DNA repair factor TFIIH. Nucleic Acids Research, 2017, 45, 10872-10883.	14.5	21
98	Epitope characterization of anti-JAM-A antibodies using orthogonal mass spectrometry and surface plasmon resonance approaches. MAbs, 2017, 9, 1317-1326.	5.2	11
99	Development and evaluation of \hat{l}^2 -galactosidase-sensitive antibody-drug conjugates. European Journal of Medicinal Chemistry, 2017, 142, 376-382.	5.5	38
100	EB1-binding–myomegalin protein complex promotes centrosomal microtubules functions. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E10687-E10696.	7.1	28
101	Structural studies of protein arginine methyltransferase 2 reveal its interactions with potential substrates and inhibitors. FEBS Journal, 2017, 284, 77-96.	4.7	25
102	Structural characterization of the yeast CF IA complex through a combination of mass spectrometry approaches. International Journal of Mass Spectrometry, 2017, 420, 57-66.	1.5	5
103	Distinct roles of Pcf11 zinc-binding domains in pre-mRNA 3′-end processing. Nucleic Acids Research, 2017, 45, 10115-10131.	14.5	11
104	Importance of the Sequence-Directed DNA Shape for Specific Binding Site Recognition by the Estrogen-Related Receptor. Frontiers in Endocrinology, 2017, 8, 140.	3.5	12
105	Soaking suggests "alternative facts― Only co-crystallization discloses major ligand-induced interface rearrangements of a homodimeric tRNA-binding protein indicating a novel mode-of-inhibition. PLoS ONE, 2017, 12, e0175723.	2.5	30
106	A ruthenium anticancer compound interacts with histones and impacts differently on epigenetic and death pathways compared to cisplatin. Oncotarget, 2017, 8, 2568-2584.	1.8	44
107	Probing Supramolecular Interactions between a Crown Ether Appended Zinc Phthalocyanine and an Ammonium Group Appended to a C ₆₀ Derivative. Chemistry - A European Journal, 2016, 22, 2051-2059.	3.3	13
108	Step by Step Assembly of Polynuclear Lanthanide Complexes with a Phosphonated Bipyridine Ligand. Inorganic Chemistry, 2016, 55, 12962-12974.	4.0	15

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109	Functional and Structural Insights of the Zinc-Finger HIT protein family members Involved in Box C/D snoRNP Biogenesis. Journal of Molecular Biology, 2016, 428, 2488-2506.	4.2	20
110	SMART Digestâ,,¢ compared with pellet digestion for analysis of human immunoglobulin G1 in rat serum by liquid chromatography tandem mass spectrometry. Analytical Biochemistry, 2016, 501, 23-25.	2.4	8
111	Insights from native mass spectrometry and ion mobility-mass spectrometry for antibody and antibody-based product characterization. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2016, 1032, 79-90.	2.3	48
112	Quantitative analysis of hlgG1 in monkey serum by LC–MS/MS using mass spectrometric immunoassay. Bioanalysis, 2016, 8, 1035-1049.	1.5	3
113	The flexibility of a generic LC–MS/MS method for the quantitative analysis of therapeutic proteins based on human immunoglobulin G and related constructs in animal studies. Journal of Pharmaceutical and Biomedical Analysis, 2016, 131, 214-222.	2.8	11
114	Culture medium associated changes in the core proteome of macrophages and in their responses to copper oxide nanoparticles. Proteomics, 2016, 16, 2864-2877.	2.2	2
115	Palladiumâ€Catalyzed Chemoselective and Biocompatible Functionalization of Cysteineâ€Containing Molecules at Room Temperature. Chemistry - A European Journal, 2016, 22, 11365-11370.	3.3	51
116	Looking for Missing Proteins in the Proteome of Human Spermatozoa: An Update. Journal of Proteome Research, 2016, 15, 3998-4019.	3.7	66
117	Benchmarking sample preparation/digestion protocols reveals tube-gel being a fast and repeatable method for quantitative proteomics. Proteomics, 2016, 16, 2953-2961.	2.2	15
118	The use of generic surrogate peptides for the quantitative analysis of human immunoglobulin G1 in pre-clinical species with high-resolution mass spectrometry. Analytical and Bioanalytical Chemistry, 2016, 408, 1687-1699.	3.7	20
119	Spiked proteomic standard dataset for testing label-free quantitative software and statistical methods. Data in Brief, 2016, 6, 286-294.	1.0	30
120	A combined proteomic and targeted analysis unravels new toxic mechanisms for zinc oxide nanoparticles in macrophages. Journal of Proteomics, 2016, 134, 174-185.	2.4	41
121	Cutting-edge mass spectrometry methods for the multi-level structural characterization of antibody-drug conjugates. Expert Review of Proteomics, 2016, 13, 157-183.	3.0	91
122	Benchmarking quantitative label-free LC–MS data processing workflows using a complex spiked proteomic standard dataset. Journal of Proteomics, 2016, 132, 51-62.	2.4	68
123	MetAP1 and MetAP2 drive cell selectivity for a potent anti-cancer agent in synergy, by controlling glutathione redox state. Oncotarget, 2016, 7, 63306-63323.	1.8	32
124	Semi-synthesis of a HGF/SF kringle one (K1) domain scaffold generates a potent in vivo MET receptor agonist. Chemical Science, 2015, 6, 2110-2121.	7.4	26
125	Combining native MS approaches to decipher archaeal box H/ACA ribonucleoprotein particle structure and activity. Proteomics, 2015, 15, 2851-2861.	2.2	9
126	Towards integrative structural mass spectrometry: Benefits from hybrid approaches. Methods, 2015, 89, 4-12.	3.8	24

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127	CBTF: New Amine-to-Thiol Coupling Reagent for Preparation of Antibody Conjugates with Increased Plasma Stability. Bioconjugate Chemistry, 2015, 26, 197-200.	3.6	53
128	Cuttingâ€edge mass spectrometry characterization of originator, biosimilar and biobetter antibodies. Journal of Mass Spectrometry, 2015, 50, 285-297.	1.6	109
129	Native mass spectrometry and ion mobility characterization of trastuzumab emtansine, a lysineâ€linked antibody drug conjugate. Protein Science, 2015, 24, 1210-1223.	7.6	113
130	What Glues a Homodimer Together: Systematic Analysis of the Stabilizing Effect of an Aromatic Hot Spot in the Protein–Protein Interface of the tRNA-Modifying Enzyme Tgt. ACS Chemical Biology, 2015, 10, 1897-1907.	3.4	19
131	Characterization of the N-Terminal Heterogeneities of Monoclonal Antibodies Using In-Gel Charge Derivatization of α-Amines and LC-MS/MS. Analytical Chemistry, 2015, 87, 3784-3790.	6.5	28
132	Intermolecular recognition of the non-coding RNA 7SK and HEXIM protein in perspective. Biochimie, 2015, 117, 63-71.	2.6	16
133	Ion mobility coupled to native mass spectrometry as a relevant tool to investigate extremely small ligand-induced conformational changes. Analyst, The, 2015, 140, 7234-7245.	3.5	32
134	Structure/Function Analysis of Protein–Protein Interactions Developed by the Yeast Pih1 Platform Protein and Its Partners in Box C/D snoRNP Assembly. Journal of Molecular Biology, 2015, 427, 2816-2839.	4.2	22
135	Computational and Mass-Spectrometry-Based Workflow for the Discovery and Validation of Missing Human Proteins: Application to Chromosomes 2 and 14. Journal of Proteome Research, 2015, 14, 3621-3634.	3.7	35
136	MAPN: First-in-Class Reagent for Kinetically Resolved Thiol-to-Thiol Conjugation. Bioconjugate Chemistry, 2015, 26, 1863-1867.	3.6	11
137	Multicentric Analyses of the CD148, CD180, and CD200 Combination for the Diagnosis of Mature B-Cell Neoplasm Using Flow Cytometry. Blood, 2015, 126, 2662-2662.	1.4	4
138	Comparative Expression Study of the Endo–G Protein Coupled Receptor (GPCR) Repertoire in Human Glioblastoma Cancer Stem-like Cells, U87-MG Cells and Non Malignant Cells of Neural Origin Unveils New Potential Therapeutic Targets. PLoS ONE, 2014, 9, e91519.	2.5	28
139	Innovative Native MS Methodologies for Antibody Drug Conjugate Characterization: High Resolution Native MS and IM-MS for Average DAR and DAR Distribution Assessment. Analytical Chemistry, 2014, 86, 10674-10683.	6.5	147
140	Interest of the CD148, CD180 and CD200 Combination in Flow Cytometry Analyses for Mature B-Cell Neoplasms Diagnosis. Blood, 2014, 124, 5407-5407.	1.4	0
141	Analytical characterization of biosimilar antibodies and Fc-fusion proteins. TrAC - Trends in Analytical Chemistry, 2013, 48, 81-95.	11.4	104
142	Time Resolved Native Ion-Mobility Mass Spectrometry to Monitor Dynamics of IgG4 Fab Arm Exchange and "Bispecific―Monoclonal Antibody Formation. Analytical Chemistry, 2013, 85, 9785-9792.	6.5	62
143	Strong Cell Surface Expression of the Toll-Like Receptor Homolog CD180 Identifies Circulating Cells of Marginal Zone Lymphoma From Other B-Cell Malignancies. Blood, 2012, 120, 1542-1542.	1.4	0