Jennifer S Brodbelt

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.

386 80 12,298 57 h-index g-index citations papers 6.6 6.95 13,658 422 L-index avg, IF ext. papers ext. citations

#	Paper	IF	Citations
386	Analysis of Viral Spike Protein N-Glycosylation Using Ultraviolet Photodissociation Mass Spectrometry <i>Analytical Chemistry</i> , 2022 ,	7.8	2
385	Influence of Primary Structure on Fragmentation of Native-Like Proteins by Ultraviolet Photodissociation. <i>Journal of the American Society for Mass Spectrometry</i> , 2021 , 32, 2860-2873	3.5	1
384	Structural Characterization of Carbonic Anhydrase-Arylsulfonamide Complexes Using Ultraviolet Photodissociation Mass Spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2021 , 32, 1370-1379	3.5	1
383	Evaluating Spatiotemporal Dynamics of Phosphorylation of RNA Polymerase II Carboxy-Terminal Domain by Ultraviolet Photodissociation Mass Spectrometry. <i>Journal of the American Chemical Society</i> , 2021 , 143, 8488-8498	16.4	2
382	Directed Evolution of an Improved Aminoacyl-tRNA Synthetase for Incorporation of L-3,4-Dihydroxyphenylalanine (L-DOPA). <i>Angewandte Chemie - International Edition</i> , 2021 , 60, 14811-14	816 ^{.4}	4
381	Directed Evolution of an Improved Aminoacyl-tRNA Synthetase for Incorporation of L-3,4-Dihydroxyphenylalanine (L-DOPA). <i>Angewandte Chemie</i> , 2021 , 133, 14937-14942	3.6	4
380	Absorption Mode Fourier Transform Ion Mobility Mass Spectrometry Multiplexing Combined with Half-Window Apodization Windows Improves Resolution and Shortens Acquisition Times. <i>Analytical Chemistry</i> , 2021 , 93, 9513-9520	7.8	3
379	Investigation of the Experimental Parameters of Ultraviolet Photodissociation for the Structural Characterization of Chondroitin Sulfate Glycosaminoglycan Isomers. <i>Journal of the American Society for Mass Spectrometry</i> , 2021 , 32, 1759-1770	3.5	2
378	Recent developments in the characterization of nucleic acids by liquid chromatography, capillary electrophoresis, ion mobility, and mass spectrometry (2010-2020). <i>Journal of Separation Science</i> , 2021 , 44, 340-372	3.4	7
377	What's all the phos about? Insights into the phosphorylation state of the RNA polymerase II C-terminal domain mass spectrometry. <i>RSC Chemical Biology</i> , 2021 , 2, 1084-1095	3	1
376	Characterization of the T4 gp32-ssDNA complex by native, cross-linking, and ultraviolet photodissociation mass spectrometry. <i>Chemical Science</i> , 2021 , 12, 13764-13776	9.4	O
375	Characterization of Isomers of Lipid A from PAO1 by Liquid Chromatography with Tandem Mass Spectrometry with Higher-Energy Collisional Dissociation and Ultraviolet Photodissociation. <i>Analytical Chemistry</i> , 2021 , 93, 4255-4262	7.8	3
374	Characterization of HLA-A*02:01 MHC Immunopeptide Antigens Enhanced by Ultraviolet Photodissociation Mass Spectrometry. <i>Analytical Chemistry</i> , 2021 , 93, 13134-13142	7.8	1
373	Relative Quantitation of Unsaturated Phosphatidylcholines Using 193 nm Ultraviolet Photodissociation Parallel Reaction Monitoring Mass Spectrometry. <i>Journal of the American Chemical Society</i> , 2021 , 143, 14622-14634	16.4	3
372	Uniting Native Capillary Electrophoresis and Multistage Ultraviolet Photodissociation Mass Spectrometry for Online Separation and Characterization of Ribosomal Proteins and Protein Complexes. <i>Analytical Chemistry</i> , 2020 , 92, 15202-15211	7.8	12
371	Double Bond Characterization of Free Fatty Acids Directly from Biological Tissues by Ultraviolet Photodissociation. <i>Analytical Chemistry</i> , 2020 , 92, 8386-8395	7.8	21
370	Unsaturation Elements and Other Modifications of Phospholipids in Bacteria: New Insight from Ultraviolet Photodissociation Mass Spectrometry. <i>Analytical Chemistry</i> , 2020 , 92, 9146-9155	7.8	11

(2019-2020)

369	Comprehensive Middle-Down Mass Spectrometry Characterization of an Antibody-Drug Conjugate by Combined Ion Activation Methods. <i>Analytical Chemistry</i> , 2020 , 92, 9790-9798	7.8	7
368	Structural Motifs for CTD Kinase Specificity on RNA Polymerase II during Eukaryotic Transcription. <i>ACS Chemical Biology</i> , 2020 , 15, 2259-2272	4.9	4
367	Precision Mapping of O-Linked -Acetylglucosamine Sites in Proteins Using Ultraviolet Photodissociation Mass Spectrometry. <i>Journal of the American Chemical Society</i> , 2020 , 142, 11569-1157	7 ^{16.4}	20
366	Localization of Double Bonds in Bacterial Glycerophospholipids Using 193 nm Ultraviolet Photodissociation in the Negative Mode. <i>Analytical Chemistry</i> , 2020 , 92, 5986-5993	7.8	8
365	Ion Activation Methods for Peptides and Proteins. <i>Analytical Chemistry</i> , 2020 , 92, 227-251	7.8	42
364	Ultraviolet Photodissociation Mass Spectrometry for Analysis of Biological Molecules. <i>Chemical Reviews</i> , 2020 , 120, 3328-3380	68.1	71
363	Enhanced Characterization of Membrane Protein Complexes by Ultraviolet Photodissociation Mass Spectrometry. <i>Analytical Chemistry</i> , 2020 , 92, 899-907	7.8	8
362	Enhanced Sequence Coverage of Large Proteins by Combining Ultraviolet Photodissociation with Proton Transfer Reactions. <i>Analytical Chemistry</i> , 2020 , 92, 1041-1049	7.8	14
361	Defining principles that influence antimicrobial peptide activity against capsulated. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 27620-27626	11.5	16
360	Interlaboratory Study for Characterizing Monoclonal Antibodies by Top-Down and Middle-Down Mass Spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2020 , 31, 1783-1802	3.5	32
359	Mapping a Conformational Epitope of Hemagglutinin A Using Native Mass Spectrometry and Ultraviolet Photodissociation. <i>Analytical Chemistry</i> , 2020 , 92, 11869-11878	7.8	7
358	Elusive structural changes of New Delhi metallo-🏿 actamase revealed by ultraviolet photodissociation mass spectrometry. <i>Chemical Science</i> , 2020 , 11, 8999-9010	9.4	4
357	Structural basis for assembly of non-canonical small subunits into type I-C Cascade. <i>Nature Communications</i> , 2020 , 11, 5931	17.4	3
356	Structural Evaluation of Protein/Metal Complexes via Native Electrospray Ultraviolet Photodissociation Mass Spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2020 , 31, 1140-1150	3.5	8
355	Hybrid 193 nm Ultraviolet Photodissociation Mass Spectrometry Localizes Cardiolipin Unsaturations. <i>Analytical Chemistry</i> , 2019 , 91, 12509-12516	7.8	16
354	Mapping RNAPII CTD Phosphorylation Reveals That the Identity and Modification of Seventh Heptad Residues Direct Tyr1 Phosphorylation. <i>ACS Chemical Biology</i> , 2019 , 14, 2264-2275	4.9	5
353	Total Synthesis of an Immunogenic Trehalose Phospholipid from Salmonella Typhi and Elucidation of Its sn-Regiochemistry by Mass Spectrometry. <i>Organic Letters</i> , 2019 , 21, 5126-5131	6.2	1
352	A Lysine-Targeted Affinity Label for Serine-Lactamase Also Covalently Modifies New Delhi Metallo-Elactamase-1 (NDM-1). <i>Biochemistry</i> , 2019 , 58, 2834-2843	3.2	16

351	A Focus Honoring Carol Robinson's Election to the National Academy of Sciences. <i>Journal of the American Society for Mass Spectrometry</i> , 2019 , 30, 1-3	3.5	
350	Impact of charge state on 193 nm ultraviolet photodissociation of protein complexes. <i>Physical Chemistry Chemical Physics</i> , 2019 , 21, 9265-9276	3.6	18
349	Localization of Cyclopropane Modifications in Bacterial Lipids via 213 nm Ultraviolet Photodissociation Mass Spectrometry. <i>Analytical Chemistry</i> , 2019 , 91, 6820-6828	7.8	27
348	Characterization of Antigenic Oligosaccharides from Gram-Negative Bacteria via Activated Electron Photodetachment Mass Spectrometry. <i>Analytical Chemistry</i> , 2019 , 91, 4672-4679	7.8	5
347	Colistin heteroresistance in Enterobacter cloacae is regulated by PhoPQ-dependent 4-amino-4-deoxy-l-arabinose addition to lipid A. <i>Molecular Microbiology</i> , 2019 , 111, 1604-1616	4.1	24
346	Structural determinants for accurate dephosphorylation of RNA polymerase II by its cognate C-terminal domain (CTD) phosphatase during eukaryotic transcription. <i>Journal of Biological Chemistry</i> , 2019 , 294, 8592-8605	5.4	8
345	Structural Characterization of Glycosaminoglycan Carbohydrates Using Ultraviolet Photodissociation. <i>Analytical Chemistry</i> , 2019 , 91, 6019-6026	7.8	22
344	Top-Down Analysis of Proteins in Low Charge States. <i>Journal of the American Society for Mass Spectrometry</i> , 2019 , 30, 704-717	3.5	12
343	Investigation of GTP-dependent dimerization of G12X K-Ras variants using ultraviolet photodissociation mass spectrometry. <i>Chemical Science</i> , 2019 , 10, 8025-8034	9.4	16
342	Top-Down Characterization of Lipooligosaccharides from Antibiotic-Resistant Bacteria. <i>Analytical Chemistry</i> , 2019 , 91, 9608-9615	7.8	7
341	Ubiquitin receptors are required for substrate-mediated activation of the proteasome's unfolding ability. <i>Scientific Reports</i> , 2019 , 9, 14506	4.9	12
340	Tyr1 phosphorylation promotes phosphorylation of Ser2 on the C-terminal domain of eukaryotic RNA polymerase II by P-TEFb. <i>ELife</i> , 2019 , 8,	8.9	18
339	Modification of Ser7 of the RNA Polymerase II C-terminal domain (CTD) regulates CTD phosphorylation patterns and transcription. <i>FASEB Journal</i> , 2019 , 33, 458.6	0.9	1
338	Systematic bromodomain protein screens identify homologous recombination and R-loop suppression pathways involved in genome integrity. <i>Genes and Development</i> , 2019 , 33, 1751-1774	12.6	47
337	Footprints of Nanoscale DNA-Silver Cluster Chromophores Activated-Electron Photodetachment Mass Spectrometry. <i>ACS Nano</i> , 2019 , 13, 14070-14079	16.7	8
336	A Focus Honoring Carol Robinson Election to the National Academy of Sciences. <i>Journal of the American Society for Mass Spectrometry</i> , 2019 , 30, 1-3	3.5	2
335	Extending Proteome Coverage by Combining MS/MS Methods and a Modified Bioinformatics Platform Adapted for Database Searching of Positive and Negative Polarity 193 nm Ultraviolet Photodissociation Mass Spectra. <i>Journal of Proteome Research</i> , 2018 , 17, 1340-1347	5.6	7
334	Directed Evolution of Heterologous tRNAs Leads to Reduced Dependence on Post-transcriptional Modifications. <i>ACS Synthetic Biology</i> , 2018 , 7, 1315-1327	5.7	5

(2017-2018)

333	UV-POSIT: Web-Based Tools for Rapid and Facile Structural Interpretation of Ultraviolet Photodissociation (UVPD) Mass Spectra. <i>Journal of the American Society for Mass Spectrometry</i> , 2018 , 29, 1323-1326	3.5	15	
332	Top-Down Characterization of Heavily Modified Histones Using 193 nm Ultraviolet Photodissociation Mass Spectrometry. <i>Journal of Proteome Research</i> , 2018 , 17, 1138-1145	5.6	33	
331	Expanding the Scope of Cross-Link Identifications by Incorporating Collisional Activated Dissociation and Ultraviolet Photodissociation Methods. <i>Analytical Chemistry</i> , 2018 , 90, 6385-6389	7.8	5	
330	Determination of Collision Cross-Sections of Protein Ions in an Orbitrap Mass Analyzer. <i>Analytical Chemistry</i> , 2018 , 90, 5896-5902	7.8	16	
329	Probing the Interaction of Aspergillomarasmine A with Metallo-Elactamases NDM-1, VIM-2, and IMP-7. <i>ACS Infectious Diseases</i> , 2018 , 4, 135-145	5.5	37	
328	Extensive Characterization of Heavily Modified Histone Tails by 193 nm Ultraviolet Photodissociation Mass Spectrometry via a Middle-Down Strategy. <i>Analytical Chemistry</i> , 2018 , 90, 1042	5 ⁷ 1043	33 ¹⁶	
327	Multistage Ultraviolet Photodissociation Mass Spectrometry To Characterize Single Amino Acid Variants of Human Mitochondrial BCAT2. <i>Analytical Chemistry</i> , 2018 , 90, 9904-9911	7.8	19	
326	Desorption Electrospray Ionization Coupled with Ultraviolet Photodissociation for Characterization of Phospholipid Isomers in Tissue Sections. <i>Analytical Chemistry</i> , 2018 , 90, 10100-10104	7.8	47	
325	Custom selenoprotein production enabled by laboratory evolution of recoded bacterial strains. <i>Nature Biotechnology</i> , 2018 , 36, 624-631	44.5	21	
324	Characterization of Disulfide Linkages in Proteins by 193 nm Ultraviolet Photodissociation (UVPD) Mass Spectrometry. <i>Analytical Chemistry</i> , 2018 , 90, 8523-8530	7.8	24	
323	Mapping phosphate modifications of substituted lipid A via a targeted MS CID/UVPD strategy. <i>Analyst, The</i> , 2018 , 143, 3091-3099	5	8	
322	Tracking the Catalytic Cycle of Adenylate Kinase by Ultraviolet Photodissociation Mass Spectrometry. <i>Analytical Chemistry</i> , 2018 , 90, 839-846	7.8	23	
321	Structural and Functional Studies of a gem-Dimethylating Methyltransferase from a trans-Acyltransferase Assembly Line. <i>ACS Chemical Biology</i> , 2018 , 13, 3306-3314	4.9	5	
320	PepSAVI-MS reveals anticancer and antifungal cycloviolacins in Viola odorata. <i>Phytochemistry</i> , 2018 , 152, 61-70	4	27	
319	Desorption Electrospray Ionization Mass Spectrometry Imaging of Proteins Directly from Biological Tissue Sections. <i>Analytical Chemistry</i> , 2018 , 90, 7785-7789	7.8	74	
318	Implementation of Fragment Ion Protection (FIP) during Ultraviolet Photodissociation (UVPD) Mass Spectrometry. <i>Analytical Chemistry</i> , 2018 , 90, 8583-8591	7.8	7	
317	Towards mapping electrostatic interactions between Kdo-lipid A and cationic antimicrobial peptides via ultraviolet photodissociation mass spectrometry. <i>Analyst, The</i> , 2018 , 143, 3607-3618	5	2	
316	Structural Characterization of Phosphatidylcholines Using 193 nm Ultraviolet Photodissociation Mass Spectrometry. <i>Analytical Chemistry</i> , 2017 , 89, 1516-1522	7.8	79	

315	Hydrocarbon-stapled lipopeptides exhibit selective antimicrobial activity. <i>Biopolymers</i> , 2017 , 108, e230	0 6 .2	6
314	Comprehensive de Novo Peptide Sequencing from MS/MS Pairs Generated through Complementary Collision Induced Dissociation and 351 nm Ultraviolet Photodissociation. <i>Analytical Chemistry</i> , 2017 , 89, 3747-3753	7.8	11
313	Ultraviolet Photodissociation of Selected Gold Clusters: Ultraefficient Unstapling and Ligand Stripping of Au(pMBA) and Au(pMBA). <i>Journal of Physical Chemistry Letters</i> , 2017 , 8, 1283-1289	6.4	25
312	ZMYM3 regulates BRCA1 localization at damaged chromatin to promote DNA repair. <i>Genes and Development</i> , 2017 , 31, 260-274	12.6	47
311	High-Throughput Analysis of Intact Human Proteins Using UVPD and HCD on an Orbitrap Mass Spectrometer. <i>Journal of Proteome Research</i> , 2017 , 16, 2072-2079	5.6	57
310	Structural heterogeneity in the intrinsically disordered RNA polymerase II C-terminal domain. <i>Nature Communications</i> , 2017 , 8, 15231	17.4	37
309	Middle-Down 193-nm Ultraviolet Photodissociation for Unambiguous Antibody Identification and its Implications for Immunoproteomic Analysis. <i>Analytical Chemistry</i> , 2017 , 89, 6498-6504	7.8	13
308	Modulation of Protein Fragmentation Through Carbamylation of Primary Amines. <i>Journal of the American Society for Mass Spectrometry</i> , 2017 , 28, 1587-1599	3.5	15
307	Characterization of trimethoprim resistant dihydrofolate reductase mutants by mass spectrometry and inhibition by propargyl-linked antifolates. <i>Chemical Science</i> , 2017 , 8, 4062-4072	9.4	24
306	Improving Performance Metrics of Ultraviolet Photodissociation Mass Spectrometry by Selective Precursor Ejection. <i>Analytical Chemistry</i> , 2017 , 89, 837-846	7.8	6
305	Characterization of Lipid A Variants by Energy-Resolved Mass Spectrometry: Impact of Acyl Chains. Journal of the American Society for Mass Spectrometry, 2017 , 28, 1118-1126	3.5	13
304	Integrating Carbamylation and Ultraviolet Photodissociation Mass Spectrometry for Middle-Down Proteomics. <i>Analytical Chemistry</i> , 2017 , 89, 11772-11778	7.8	6
303	Pinpointing Double Bond and sn-Positions in Glycerophospholipids via Hybrid 193 nm Ultraviolet Photodissociation (UVPD) Mass Spectrometry. <i>Journal of the American Chemical Society</i> , 2017 , 139, 156	68 ¹⁶ 156	59 1 0 ⁰ 4
302	Characterization of hydrogen bonding motifs in proteins: hydrogen elimination monitoring by ultraviolet photodissociation mass spectrometry. <i>Physical Chemistry Chemical Physics</i> , 2017 , 19, 20057-	20674	9
301	SITS Derivatization of Peptides to Enhance 266Ihm Ultraviolet Photodissociation (UVPD). <i>Journal of the American Society for Mass Spectrometry</i> , 2017 , 28, 1462-1472	3.5	5
300	Mapping the Phosphorylation Pattern of Drosophila melanogaster RNA Polymerase II Carboxyl-Terminal Domain Using Ultraviolet Photodissociation Mass Spectrometry. <i>ACS Chemical Biology</i> , 2017 , 12, 153-162	4.9	18
299	193 nm Ultraviolet Photodissociation Mass Spectrometry for Phosphopeptide Characterization in the Positive and Negative Ion Modes. <i>Journal of Proteome Research</i> , 2016 , 15, 2739-48	5.6	34
298	Integrating Weak Anion Exchange and Ultraviolet Photodissociation Mass Spectrometry with Strategic Modulation of Peptide Basicity for the Enrichment of Sulfopeptides. <i>Analytical Chemistry</i> , 2016 , 88, 11037-11045	7.8	11

(2016-2016)

297	Cysteine-Selective Peptide Identification: Selenium-Based Chromophore for Selective S-Se Bond Cleavage with 266 nm Ultraviolet Photodissociation. <i>Analytical Chemistry</i> , 2016 , 88, 7222-9	7.8	12	
296	Methylation follows condensation in the gephyronic acid modular polyketide synthase. <i>Chemical Communications</i> , 2016 , 52, 8822-5	5.8	16	
295	UVliPiD: A UVPD-Based Hierarchical Approach for De Novo Characterization of Lipid A Structures. <i>Analytical Chemistry</i> , 2016 , 88, 1812-20	7.8	24	
294	Shotgun Analysis of Rough-Type Lipopolysaccharides Using Ultraviolet Photodissociation Mass Spectrometry. <i>Analytical Chemistry</i> , 2016 , 88, 1044-51	7.8	64	
293	Ion Activation Methods for Peptides and Proteins. Analytical Chemistry, 2016, 88, 30-51	7.8	123	
292	Integration of Ultraviolet Photodissociation with Proton Transfer Reactions and Ion Parking for Analysis of Intact Proteins. <i>Analytical Chemistry</i> , 2016 , 88, 1008-16	7.8	26	
291	UVnovo: A de Novo Sequencing Algorithm Using Single Series of Fragment Ions via Chromophore Tagging and 351 nm Ultraviolet Photodissociation Mass Spectrometry. <i>Analytical Chemistry</i> , 2016 , 88, 3990-7	7.8	24	
290	Characterization of Therapeutic Monoclonal Antibodies at the Subunit-Level using Middle-Down 193 nm Ultraviolet Photodissociation. <i>Analytical Chemistry</i> , 2016 , 88, 4004-13	7.8	47	
289	Charge site assignment in native proteins by ultraviolet photodissociation (UVPD) mass spectrometry. <i>Analyst, The</i> , 2016 , 141, 166-76	5	37	
288	Ultraviolet Photodissociation of Native Proteins Following Proton Transfer Reactions in the Gas Phase. <i>Analytical Chemistry</i> , 2016 , 88, 12354-12362	7.8	10	
287	Characterization of the Cysteine Content in Proteins Utilizing Cysteine Selenylation with 266hm Ultraviolet Photodissociation (UVPD). <i>Journal of the American Society for Mass Spectrometry</i> , 2016 , 27, 1344-50	3.5	7	
286	Exploitation of the Ornithine Effect Enhances Characterization of Stapled and Cyclic Peptides. Journal of the American Society for Mass Spectrometry, 2016 , 27, 856-63	3.5	10	
285	Statistical Examination of the a and a + 1 Fragment Ions from 193hm Ultraviolet Photodissociation Reveals Local Hydrogen Bonding Interactions. <i>Journal of the American Society for Mass Spectrometry</i> , 2016 , 27, 1443-53	3.5	11	
284	Click and chemically triggered declick reactions through reversible amine and thiol coupling via a conjugate acceptor. <i>Nature Chemistry</i> , 2016 , 8, 968-973	17.6	66	
283	Ultraviolet Photodissociation Induced by Light-Emitting Diodes in a Planar Ion Trap. <i>Angewandte Chemie</i> , 2016 , 128, 12605-12609	3.6	1	
282	Ultraviolet Photodissociation Induced by Light-Emitting Diodes in a Planar Ion Trap. <i>Angewandte Chemie - International Edition</i> , 2016 , 55, 12417-21	16.4	3	
281	Impact of G12 Mutations on the Structure of K-Ras Probed by Ultraviolet Photodissociation Mass Spectrometry. <i>Journal of the American Chemical Society</i> , 2016 , 138, 13187-13196	16.4	26	
280	193 nm Ultraviolet Photodissociation Mass Spectrometry of Tetrameric Protein Complexes Provides Insight into Quaternary and Secondary Protein Topology. <i>Journal of the American Chemical Society</i> , 2016 , 138, 10849-59	16.4	50	

279	Modulation of Phosphopeptide Fragmentation via Dual Spray Ion/Ion Reactions Using a Sulfonate-Incorporating Reagent. <i>Analytical Chemistry</i> , 2016 , 88, 8158-65	7.8	6
278	Characterization of Intra- and Intermolecular Protein Crosslinking by Top Down Ultraviolet Photodissociation Mass Spectrometry. <i>ChemistrySelect</i> , 2016 , 1, 590-593	1.8	7
277	Screen identifies bromodomain protein ZMYND8 in chromatin recognition of transcription-associated DNA damage that promotes homologous recombination. <i>Genes and Development</i> , 2015 , 29, 197-211	12.6	153
276	Selective cleavage upon ETD of peptides containing disulfide or nitrogenlitrogen bonds. <i>International Journal of Mass Spectrometry</i> , 2015 , 378, 127-133	1.9	4
275	The rice immune receptor XA21 recognizes a tyrosine-sulfated protein from a Gram-negative bacterium. <i>Science Advances</i> , 2015 , 1, e1500245	14.3	151
274	Structural Characterization of Dihydrofolate Reductase Complexes by Top-Down Ultraviolet Photodissociation Mass Spectrometry. <i>Journal of the American Chemical Society</i> , 2015 , 137, 9128-35	16.4	54
273	Impact of Protease on Ultraviolet Photodissociation Mass Spectrometry for Bottom-up Proteomics. Journal of Proteome Research, 2015 , 14, 2626-32	5.6	21
272	Comparison of Ultraviolet Photodissociation and Collision Induced Dissociation of Adrenocorticotropic Hormone Peptides. <i>Journal of the American Society for Mass Spectrometry</i> , 2015 , 26, 1570-9	3.5	12
271	Reinforcing Lipid A Acylation on the Cell Surface of Acinetobacter baumannii Promotes Cationic Antimicrobial Peptide Resistance and Desiccation Survival. <i>MBio</i> , 2015 , 6, e00478-15	7.8	93
270	Comparison of Glycopeptide Fragmentation by Collision Induced Dissociation and Ultraviolet Photodissociation. <i>International Journal of Mass Spectrometry</i> , 2015 , 377, 385-392	1.9	30
269	Ultraviolet photodissociation of protonated, fixed charge, and charge-reduced peptides. <i>International Journal of Mass Spectrometry</i> , 2015 , 390, 81-90	1.9	10
268	Structural characterization of holo- and apo-myoglobin in the gas phase by ultraviolet photodissociation mass spectrometry. <i>Chemical Science</i> , 2015 , 6, 1324-1333	9.4	60
267	High-throughput bioconjugation for enhanced 193 nm photodissociation via droplet phase initiated ion/ion chemistry using a front-end dual spray reactor. <i>Analytical Chemistry</i> , 2015 , 87, 9396-402	7.8	13
266	Extracellular zinc induces phosphoethanolamine addition to Pseudomonas aeruginosa lipid A via the ColRS two-component system. <i>Molecular Microbiology</i> , 2015 , 97, 166-78	4.1	51
265	A New Covalent Inhibitor of Class C Lactamases Reveals Extended Active Site Specificity. <i>Biochemistry</i> , 2015 , 54, 7375-84	3.2	7
264	Evolving tRNA(Sec) for efficient canonical incorporation of selenocysteine. <i>Journal of the American Chemical Society</i> , 2015 , 137, 46-9	16.4	43
263	Top-down 193-nm ultraviolet photodissociation mass spectrometry for simultaneous determination of polyubiquitin chain length and topology. <i>Analytical Chemistry</i> , 2015 , 87, 1812-20	7.8	33
262	The first pilot project of the consortium for top-down proteomics: a status report. <i>Proteomics</i> , 2014 , 14, 1130-40	4.8	73

(2014-2014)

261	Direct identification of tyrosine sulfation by using ultraviolet photodissociation mass spectrometry. Journal of the American Society for Mass Spectrometry, 2014 , 25, 1461-71	3.5	38
260	Directed evolution of genetic parts and circuits by compartmentalized partnered replication. <i>Nature Biotechnology</i> , 2014 , 32, 97-101	44.5	103
259	Photodissociation mass spectrometry: new tools for characterization of biological molecules. <i>Chemical Society Reviews</i> , 2014 , 43, 2757-83	58.5	221
258	Hybridizing ultraviolet photodissociation with electron transfer dissociation for intact protein characterization. <i>Analytical Chemistry</i> , 2014 , 86, 10970-7	7.8	34
257	Top-Down Strategies for the Structural Elucidation of Intact Gram-negative Bacterial Endotoxins. <i>Chemical Science</i> , 2014 , 5, 4291-4301	9.4	19
256	Ultraviolet photodissociation for characterization of whole proteins on a chromatographic time scale. <i>Analytical Chemistry</i> , 2014 , 86, 2185-92	7.8	69
255	Antimicrobial peptide resistance of Vibrio cholerae results from an LPS modification pathway related to nonribosomal peptide synthetases. <i>ACS Chemical Biology</i> , 2014 , 9, 2382-92	4.9	41
254	193 nm ultraviolet photodissociation mass spectrometry for the structural elucidation of lipid A compounds in complex mixtures. <i>Analytical Chemistry</i> , 2014 , 86, 2138-45	7.8	45
253	Characterization of native protein complexes using ultraviolet photodissociation mass spectrometry. <i>Journal of the American Chemical Society</i> , 2014 , 136, 12920-8	16.4	83
252	Streamlining bottom-up protein identification based on selective ultraviolet photodissociation (UVPD) of chromophore-tagged histidine- and tyrosine-containing peptides. <i>Analytical Chemistry</i> , 2014 , 86, 6237-44	7.8	14
251	Probing the unfolding of myoglobin and domain C of PARP-1 with covalent labeling and top-down ultraviolet photodissociation mass spectrometry. <i>Analytical Chemistry</i> , 2014 , 86, 2534-42	7.8	25
250	Mechanically Modulating the Photophysical Properties of Fluorescent Protein Biocomposites for Ratio- and Intensiometric Sensors. <i>Angewandte Chemie</i> , 2014 , 126, 5188-5192	3.6	4
249	The molecular mechanism of eukaryotic elongation factor 2 kinase activation. <i>Journal of Biological Chemistry</i> , 2014 , 289, 23901-16	5.4	24
248	Improvement of shotgun proteomics in the negative mode by carbamylation of peptides and ultraviolet photodissociation mass spectrometry. <i>Analytical Chemistry</i> , 2014 , 86, 12285-90	7.8	23
247	Covalent inhibition of New Delhi metallo-Elactamase-1 (NDM-1) by cefaclor. <i>ChemBioChem</i> , 2014 , 15, 2541-8	3.8	34
246	Characterization of green fluorescent proteins by 193 nm ultraviolet photodissociation mass spectrometry. <i>Proteomics</i> , 2014 , 14, 1165-73	4.8	23
245	Identification of a broad family of lipid A late acyltransferases with non-canonical substrate specificity. <i>Molecular Microbiology</i> , 2014 , 91, 887-99	4.1	16
244	Characterization of Pseudomonas aeruginosa LpxT reveals dual positional lipid A kinase activity and co-ordinated control of outer membrane modification. <i>Molecular Microbiology</i> , 2014 , 94, 728-41	4.1	28

243	Differentiation and distributions of DNA/cisplatin crosslinks by liquid chromatography-electrospray ionization-infrared multiphoton dissociation mass spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2014 , 25, 71-9	3.5	8
242	Ligand Binding to Nucleic Acids. <i>Physical Chemistry in Action</i> , 2014 , 225-252		2
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