Vicki H Wysocki

List of Publications by Year in Descending Order

257 papers	11,607 citations	56 h-index	99 g-index
275 ext. papers	12,837 ext. citations	7.8 avg, IF	6.27 L-index

#	Paper	IF	Citations
257	Native Mass Spectrometry: Recent Progress and Remaining Challenges.. *Annual Review of Biophysics*, **2022**,	21.1	6
256	Optimization of proteomics sample preparation for identification of host and bacterial proteins in mouse feces.. *Analytical and Bioanalytical Chemistry*, **2022**, 414, 2317	4.4	1
255	Native Mass Spectrometry and Surface Induced Dissociation Provide Insight into the Post-Translational Modifications of Tetrameric AQP0 Isolated from Bovine Eye Lens.. *Analytical Chemistry*, **2022**,	7.8	2
254	Characterization of a Salmonella transcription factor-DNA complex and identification of the inducer by native mass spectrometry.. *Journal of Molecular Biology*, **2022**, 167480	6.5	0
253	Surface-Induced Dissociation for Protein Complex Characterization. *Methods in Molecular Biology*, **2022**, 211-237	1.4	1
252	Surface-induced Dissociation Mass Spectrometry as a Structural Biology Tool. *Chemical Reviews*, **2021**,	68.1	4
251	Implementing Digital-Waveform Technology for Extended / Range Operation on a Native Dual-Quadrupole FT-IM-Orbitrap Mass Spectrometer. *Journal of the American Society for Mass Spectrometry*, **2021**, 32, 2812-2820	3.5	3
250	Use of tandem affinity-buffer exchange chromatography online with native mass spectrometry for optimizing overexpression and purification of recombinant proteins. *Methods in Enzymology*, **2021**, 659, 37-70	1.7	1
249	Purification, reconstitution, and mass analysis of archaeal RNase P, a multisubunit ribonucleoprotein enzyme. *Methods in Enzymology*, **2021**, 659, 71-103	1.7	0
248	Surface-induced dissociation of protein complexes on a cyclic ion mobility spectrometer. *Analyst, The*, **2021**, 146, 6861-6873	5	3
247	Oligomeric complexes formed by Red□ single strand annealing protein in its different DNA bound states. *Nucleic Acids Research*, **2021**, 49, 3441-3460	20.1	3
246	Surface-Induced Dissociation of Protein Complexes Selected by Trapped Ion Mobility Spectrometry. *Analytical Chemistry*, **2021**, 93, 5513-5520	7.8	7
245	Tandem surface-induced dissociation of protein complexes on an ultrahigh resolution platform. *International Journal of Mass Spectrometry*, **2021**, 461,	1.9	3
244	Cellular mRNA triggers structural transformation of Ebola virus matrix protein VP40 to its essential regulatory form. *Cell Reports*, **2021**, 35, 108986	10.6	5
243	Transferrin receptor targeting by de novo sheet extension. *Proceedings of the National Academy of Sciences of the United States of America*, **2021**, 118,	11.5	6
242	Variable-Temperature Electrospray Ionization for Temperature-Dependent Folding/Refolding Reactions of Proteins and Ligand Binding. *Analytical Chemistry*, **2021**, 93, 6924-6931	7.8	9
241	Surface-Induced Dissociation of Anionic vs Cationic Native-Like Protein Complexes. *Journal of the American Chemical Society*, **2021**, 143, 7698-7706	16.4	6

240	Prediction of Protein Complex Structure Using Surface-Induced Dissociation and Cryo-Electron Microscopy. *Analytical Chemistry*, **2021**, 93, 7596-7605	7.8	4
239	STK11/LKB1 Loss of Function Is Associated with Global DNA Hypomethylation and -Adenosyl-Methionine Depletion in Human Lung Adenocarcinoma. *Cancer Research*, **2021**, 81, 4194-4204	10.1	1
238	Generation of ordered protein assemblies using rigid three-body fusion. *Proceedings of the National Academy of Sciences of the United States of America*, **2021**, 118,	11.5	6
237	Analysis of Tagged Proteins Using Tandem Affinity-Buffer Exchange Chromatography Online with Native Mass Spectrometry. *Biochemistry*, **2021**, 60, 1876-1884	3.2	7
236	De novo design of transmembrane barrels. *Science*, **2021**, 371,	33.3	25
235	Protein cofactors and substrate influence Mg2+-dependent structural changes in the catalytic RNA of archaeal RNase P. *Nucleic Acids Research*, **2021**, 49, 9444-9458	20.1	1
234	Tunable Heteroassembly of a Plant Pseudoenzyme-Enzyme Complex. *ACS Chemical Biology*, **2021**, 16, 2315-2325	4.9	3
233	De novo design of tyrosine and serine kinase-driven protein switches. *Nature Structural and Molecular Biology*, **2021**, 28, 762-770	17.6	5
232	Optimization of proteomics sample preparation for forensic analysis of skin samples. *Journal of Proteomics*, **2021**, 249, 104360	3.9	1
231	Using SLIM-Based IMS-IMS Together with Cryogenic Infrared Spectroscopy for Glycan Analysis. *Analytical Chemistry*, **2020**, 92, 9079-9085	7.8	21
230	Population Distributions from Native Mass Spectrometry Titrations Reveal Nearest-Neighbor Cooperativity in the Ring-Shaped Oligomeric Protein TRAP. *Biochemistry*, **2020**, 59, 2518-2527	3.2	6
229	A Tilted Surface and Ion Carpet Array for SID. *Journal of the American Society for Mass Spectrometry*, **2020**, 31, 458-462	3.5	5
228	De novo design of protein logic gates. *Science*, **2020**, 368, 78-84	33.3	88
227	Quaternary Structure of the Tryptophan Synthase Subunit Homolog BX1 from. *Journal of the American Society for Mass Spectrometry*, **2020**, 31, 227-233	3.5	1
226	Characterization of [2Fe2S]-Cluster-Bridged Protein Complexes and Reaction Intermediates by use of Native Mass Spectrometric Methods. *Angewandte Chemie*, **2020**, 132, 6790-6794	3.6	
225	Collision Cross Sections of Charge-Reduced Proteins and Protein Complexes: A Database for Collision Cross Section Calibration. *Analytical Chemistry*, **2020**, 92, 4475-4483	7.8	18
224	Ion Mobility and Surface Collisions: Submicrometer Capillaries Can Produce Native-like Protein Complexes. *Analytical Chemistry*, **2020**, 92, 2460-2467	7.8	9
223	Rapid online buffer exchange for screening of proteins, protein complexes and cell lysates by native mass spectrometry. *Nature Protocols*, **2020**, 15, 1132-1157	18.8	46

222	Characterization of [2Fe-2S]-Cluster-Bridged Protein Complexes and Reaction Intermediates by use of Native Mass Spectrometric Methods. *Angewandte Chemie - International Edition*, **2020**, 59, 6724-6728	16.4	13
221	Comparative Structural Analysis of 20S Proteasome Ortholog Protein Complexes by Native Mass Spectrometry. *ACS Central Science*, **2020**, 6, 573-588	16.8	24
220	Chapter 11:Surface-induced Dissociation in Biomolecular Mass Spectrometry. *New Developments in Mass Spectrometry*, **2020**, 281-336	2.3	2
219	Cytoplasmic mRNA recapping has limited impact on proteome complexity. *Open Biology*, **2020**, 10, 200313	7	3
218	Solution structure of the nucleotide hydrolase BlsM: Implication of its substrate specificity. *Protein Science*, **2020**, 29, 1760-1773	6.3	2
217	Probing the structure of nanodiscs using surface-induced dissociation mass spectrometry. *Chemical Communications*, **2020**, 56, 15651-15654	5.8	10
216	Simple and Minimally Invasive SID Devices for Native Mass Spectrometry. *Analytical Chemistry*, **2020**, 92, 11195-11203	7.8	19
215	Coupling 193 nm Ultraviolet Photodissociation and Ion Mobility for Sequence Characterization of Conformationally-Selected Peptides. *Journal of the American Society for Mass Spectrometry*, **2020**, 31, 2313-2320	3.5	6
214	HIV-1 Gag protein with or without p6 specifically dimerizes on the viral RNA packaging signal. *Journal of Biological Chemistry*, **2020**, 295, 14391-14401	5.4	12
213	Spectroscopic Evidence for Lactam Formation in Terminal Ornithine b and b Fragment Ions. *Journal of the American Society for Mass Spectrometry*, **2019**, 30, 1565-1577	3.5	2
212	Light Regulation of Enzyme Allostery through Photo-responsive Unnatural Amino Acids. *Cell Chemical Biology*, **2019**, 26, 1501-1514.e9	8.2	15
211	Surface-Induced Dissociation of Noncovalent Protein Complexes in an Extended Mass Range Orbitrap Mass Spectrometer. *Analytical Chemistry*, **2019**, 91, 3611-3618	7.8	48
210	Generation of a Stand-Alone Tryptophan Synthase β Subunit by Mimicking an Evolutionary Blueprint. *ChemBioChem*, **2019**, 20, 2747-2751	3.8	3
209	De novo design of tunable, pH-driven conformational changes. *Science*, **2019**, 364, 658-664	33.3	60
208	Relative interfacial cleavage energetics of protein complexes revealed by surface collisions. *Proceedings of the National Academy of Sciences of the United States of America*, **2019**, 116, 8143-8148	11.5	39
207	Predicting Protein Complex Structure from Surface-Induced Dissociation Mass Spectrometry Data. *ACS Central Science*, **2019**, 5, 1330-1341	16.8	19
206	Light-Regulation of Tryptophan Synthase by Combining Protein Design and Enzymology. *International Journal of Molecular Sciences*, **2019**, 20,	6.3	5
205	Oligomerization Affects the Ability of Human Cyclase-Associated Proteins 1 and 2 to Promote Actin Severing by Cofilins. *International Journal of Molecular Sciences*, **2019**, 20,	6.3	13

204	Design and Performance of a Second-Generation Surface-Induced Dissociation Cell for Fourier Transform Ion Cyclotron Resonance Mass Spectrometry of Native Protein Complexes. *Analytical Chemistry*, **2019**, 91, 14049-14057	7.8	13
203	Integrated Use of Biochemical, Native Mass Spectrometry, Computational, and Genome-Editing Methods to Elucidate the Mechanism of a Salmonella deglycase. *Journal of Molecular Biology*, **2019**, 431, 4497-4513	6.5	6
202	Programmable design of orthogonal protein heterodimers. *Nature*, **2019**, 565, 106-111	50.4	87
201	Surface-Induced Dissociation: An Effective Method for Characterization of Protein Quaternary Structure. *Analytical Chemistry*, **2019**, 91, 190-209	7.8	44
200	Stoichiometry of triple-sieve tRNA editing complex ensures fidelity of aminoacyl-tRNA formation. *Nucleic Acids Research*, **2019**, 47, 929-940	20.1	10
199	Salmonella-Mediated Inflammation Eliminates Competitors for Fructose-Asparagine in the Gut. *Infection and Immunity*, **2018**, 86,	3.7	9
198	How many human proteoforms are there?. *Nature Chemical Biology*, **2018**, 14, 206-214	11.7	324
197	Surface Induced Dissociation Coupled with High Resolution Mass Spectrometry Unveils Heterogeneity of a 211 kDa Multicopper Oxidase Protein Complex. *Journal of the American Society for Mass Spectrometry*, **2018**, 29, 723-733	3.5	13
196	Confirmation of intersubunit connectivity and topology of designed protein complexes by native MS. *Proceedings of the National Academy of Sciences of the United States of America*, **2018**, 115, 1268-1273	11.5	40
195	Identification of Bacterial Species That Can Utilize Fructose-Asparagine. *Applied and Environmental Microbiology*, **2018**, 84,	4.8	10
194	Investigation of sliding DNA clamp dynamics by single-molecule fluorescence, mass spectrometry and structure-based modeling. *Nucleic Acids Research*, **2018**, 46, 3103-3118	20.1	10
193	Measurement of Fructose-Asparagine Concentrations in Human and Animal Foods. *Journal of Agricultural and Food Chemistry*, **2018**, 66, 212-217	5.7	11
192	The Exon Junction Complex Undergoes a Compositional Switch that Alters mRNP Structure and Nonsense-Mediated mRNA Decay Activity. *Cell Reports*, **2018**, 25, 2431-2446.e7	10.6	33
191	Proteogenomic Analysis of Surgically Resected Lung Adenocarcinoma. *Journal of Thoracic Oncology* , **2018**, 13, 1519-1529	8.9	8
190	Localization of Protein Complex Bound Ligands by Surface-Induced Dissociation High-Resolution Mass Spectrometry. *Analytical Chemistry*, **2018**, 90, 12796-12801	7.8	21
189	RNA-binding proteins and heat-shock protein 90 are constituents of the cytoplasmic capping enzyme interactome. *Journal of Biological Chemistry*, **2018**, 293, 16596-16607	5.4	6
188	Identifying Unknown Enzyme-Substrate Pairs from the Cellular Milieu with Native Mass Spectrometry. *ChemBioChem*, **2017**, 18, 613-617	3.8	8
187	Eng1 and Exg8 Are the Major β-Glucanases Secreted by the Fungal Pathogen. *Journal of Biological Chemistry*, **2017**, 292, 4801-4810	5.4	26

186	Surface induced dissociation as a tool to study membrane protein complexes. *Chemical Communications*, **2017**, 53, 3106-3109	5.8	30
185	Infrared Multiple-Photon Dissociation Action Spectroscopy of the b Ion from PPG: Evidence of Third Residue Affecting b Fragment Structure. *Journal of the American Society for Mass Spectrometry*, **2017**, 28, 1482-1488	3.5	13
184	Chemical and pathogen-induced inflammation disrupt the murine intestinal microbiome. *Microbiome*, **2017**, 5, 47	16.6	64
183	Surface-Induced Dissociation of Protein Complexes in a Hybrid Fourier Transform Ion Cyclotron Resonance Mass Spectrometer. *Analytical Chemistry*, **2017**, 89, 895-901	7.8	21
182	Biogenic manganese oxide nanoparticle formation by a multimeric multicopper oxidase Mnx. *Nature Communications*, **2017**, 8, 746	17.4	46
181	Foldability of a Natural De Novo Evolved Protein. *Structure*, **2017**, 25, 1687-1696.e4	5.2	19
180	Human Argonaute3 has slicer activity. *Nucleic Acids Research*, **2017**, 45, 11867-11877	20.1	46
179	Evolutionary diversification of protein-protein interactions by interface add-ons. *Proceedings of the National Academy of Sciences of the United States of America*, **2017**, 114, E8333-E8342	11.5	13
178	A novel double kink-turn module in euryarchaeal RNase P RNAs. *Nucleic Acids Research*, **2017**, 45, 7432-7440	20.1	18
177	Use of chemical modification and mass spectrometry to identify substrate-contacting sites in proteinaceous RNase P, a tRNA processing enzyme. *Nucleic Acids Research*, **2016**, 44, 5344-55	20.1	13
176	Extended Gas-Phase Trapping Followed by Surface-Induced Dissociation of Noncovalent Protein Complexes. *Analytical Chemistry*, **2016**, 88, 1218-21	7.8	19
175	Possible isomers in ligand protected Ag11 cluster ions identified by ion mobility mass spectrometry and fragmented by surface induced dissociation. *Chemical Communications*, **2016**, 52, 3805-8	5.8	36
174	R vs. S fluoroproline ring substitution: trans/cis effects on the formation of b2 ions in gas-phase peptide fragmentation. *Physical Chemistry Chemical Physics*, **2016**, 18, 2202-9	3.6	6
173	A metabolic intermediate of the fructose-asparagine utilization pathway inhibits growth of a Salmonella fraB mutant. *Scientific Reports*, **2016**, 6, 28117	4.9	14
172	A Protein-derived Oxygen Is the Source of the Amide Oxygen of Nitrile Hydratases. *Journal of Biological Chemistry*, **2016**, 291, 7822-9	5.4	5
171	The Pseudomonas aeruginosa AmrZ C-terminal domain mediates tetramerization and is required for its activator and repressor functions. *Environmental Microbiology Reports*, **2016**, 8, 85-90	3.7	11
170	A dimer interface mutation in glyceraldehyde-3-phosphate dehydrogenase regulates its binding to AU-rich RNA. *Journal of Biological Chemistry*, **2015**, 290, 1770-85	5.4	34
169	Label-free detection and identification of protein ligands captured by receptors in a polymerized planar lipid bilayer using MALDI-TOF MS. *Analytical and Bioanalytical Chemistry*, **2015**, 407, 2777-89	4.4	3

168	Surface induced dissociation yields substructure of 20S proteasome complexes. *International Journal of Mass Spectrometry*, **2015**, 377, 201-204	1.9	12
167	Top-down-assisted bottom-up method for homologous protein sequencing: hemoglobin from 33 bird species. *Journal of the American Society for Mass Spectrometry*, **2015**, 26, 1875-84	3.5	5
166	Illustration of SID-IM-SID (surface-induced dissociation-ion mobility-SID) mass spectrometry: homo and hetero model protein complexes. *Analyst, The*, **2015**, 140, 7012-9	5	16
165	Surface-Induced Dissociation Mass Spectra as a Tool for Distinguishing Different Structural Forms of Gas-Phase Multimeric Protein Complexes. *Analytical Chemistry*, **2015**, 87, 11879-86	7.8	45
164	Laser desorption ionization of small molecules assisted by tungsten oxide and rhenium oxide particles. *Journal of Mass Spectrometry*, **2015**, 50, 891-8	2.2	15
163	Resolution of Stepwise Cooperativities of Copper Binding by the Homotetrameric Copper-Sensitive Operon Repressor (CsoR): Impact on Structure and Stability. *Angewandte Chemie*, **2015**, 127, 12986-12990	3.6	
162	Resolution of Stepwise Cooperativities of Copper Binding by the Homotetrameric Copper-Sensitive Operon Repressor (CsoR): Impact on Structure and Stability. *Angewandte Chemie - International Edition*, **2015**, 54, 12795-9	16.4	10
161	Surface-Induced Dissociation of Homotetramers with D2 Symmetry Yields their Assembly Pathways and Characterizes the Effect of Ligand Binding. *Chemistry and Biology*, **2015**, 22, 583-92		49
160	Low energy CID and action IRMPD provide insights into a minor subpopulation of the gas-phase conformers of triply charged bradykinin. *International Journal of Mass Spectrometry*, **2015**, 391, 2-10	1.9	4
159	Refining the Structural Model of a Heterohexameric Protein Complex: Surface Induced Dissociation and Ion Mobility Provide Key Connectivity and Topology Information. *ACS Central Science*, **2015**, 1, 477-487	16.8	49
158	Mass spectrometry: Bound in flight. *Nature Chemistry*, **2015**, 7, 189-90	17.6	1
157	Mutant poisoning demonstrates a nonsequential mechanism for digestion of double-stranded DNA by exonuclease trimers. *Biochemistry*, **2015**, 54, 942-51	3.2	3
156	Probing the run-on oligomer of activated SgrAI bound to DNA. *PLoS ONE*, **2015**, 10, e0124783	3.7	10
155	Shotgun Proteomic Analysis of Protein Expression in Mosquito Ovaries Post Blood Meal. *FASEB Journal*, **2015**, 29, 567.13	0.9	
154	Mass Spectrometry in Structural Biology: Surface-induced Dissociation/Ion Mobility of Protein Complexes. *FASEB Journal*, **2015**, 29, 360.3	0.9	1
153	Surface induced dissociation: dissecting noncovalent protein complexes in the gas phase. *Accounts of Chemical Research*, **2014**, 47, 1010-8	24.3	99
152	Human defensins facilitate local unfolding of thermodynamically unstable regions of bacterial protein toxins. *Immunity*, **2014**, 41, 709-21	32.3	46
151	Uncovering the stoichiometry of Pyrococcus furiosus RNase P, a multi-subunit catalytic ribonucleoprotein complex, by surface-induced dissociation and ion mobility mass spectrometry. *Angewandte Chemie - International Edition*, **2014**, 53, 11483-7	16.4	28

150	Gas-phase helical peptides mimic solution-phase behavior. *Journal of the American Chemical Society*, **2014**, 136, 14173-83	16.4	11
149	Histone H1 phosphorylation in breast cancer. *Journal of Proteome Research*, **2014**, 13, 2453-67	5.6	30
148	Investigations of the mechanism of the "proline effect" in tandem mass spectrometry experiments: the "pipecolic acid effect". *Journal of the American Society for Mass Spectrometry*, **2014**, 25, 1705-15	3.5	21
147	Paper spray ionization of noncovalent protein complexes. *Analytical Chemistry*, **2014**, 86, 1342-6	7.8	58
146	Interfacial residues promote an optimal alignment of the catalytic center in human soluble guanylate cyclase: heterodimerization is required but not sufficient for activity. *Biochemistry*, **2014**, 53, 2153-65	3.2	34
145	Surface induced dissociation yields quaternary substructure of refractory noncovalent phosphorylase B and glutamate dehydrogenase complexes. *Journal of the American Society for Mass Spectrometry*, **2014**, 25, 368-79	3.5	28
144	Uncovering the Stoichiometry of Pyrococcus furiosus RNase P, a Multi-Subunit Catalytic Ribonucleoprotein Complex, by Surface-Induced Dissociation and Ion Mobility Mass Spectrometry. *Angewandte Chemie*, **2014**, 126, 11667-11671	3.6	1
143	Gene regulation by substoichiometric heterocomplex formation of undecameric TRAP and trimeric anti-TRAP. *Proceedings of the National Academy of Sciences of the United States of America*, **2014**, 111, 3442-7	11.5	12
142	Ligand binding and unfolding of tryptophan synthase revealed by ion mobility-tandem mass spectrometry employing collision and surface induced dissociation. *International Journal for Ion Mobility Spectrometry*, **2013**, 16, 133-143	1.5	11
141	High-resolution identification of human adiponectin oligomers and regulation by pioglitazone in type 2 diabetic patients. *Analytical Biochemistry*, **2013**, 437, 150-60	3.1	8
140	Influence of N-terminal residue composition on the structure of proline-containing b2+ ions. *Journal of Physical Chemistry A*, **2013**, 117, 1291-8	2.8	22
139	Impact of charge state on gas-phase behaviors of noncovalent protein complexes in collision induced dissociation and surface induced dissociation. *Analyst, The*, **2013**, 138, 1353-62	5	62
138	A polymetamorphic protein. *Protein Science*, **2013**, 22, 641-9	6.3	6
137	Molecular model of a soluble guanylyl cyclase fragment determined by small-angle X-ray scattering and chemical cross-linking. *Biochemistry*, **2013**, 52, 1568-82	3.2	51
136	An unusual dimeric small heat shock protein provides insight into the mechanism of this class of chaperones. *Journal of Molecular Biology*, **2013**, 425, 1683-96	6.5	48
135	Structural analysis of activated SgrAI-DNA oligomers using ion mobility mass spectrometry. *Biochemistry*, **2013**, 52, 4373-81	3.2	18
134	Low mass MS/MS fragments of protonated amino acids used for distinction of their 13C-isotopomers in metabolic studies. *Journal of the American Society for Mass Spectrometry*, **2013**, 24, 622-31	3.5	8
133	Dissecting the large noncovalent protein complex GroEL with surface-induced dissociation and ion mobility-mass spectrometry. *Analytical Chemistry*, **2013**, 85, 8262-7	7.8	74

132	Survival of host blood proteins in Ixodes scapularis (Acari: Ixodidae) ticks: a time course study. *Journal of Medical Entomology*, **2013**, 50, 1282-90	2.2	12
131	Influence of a Gamma Amino Acid on the Structures and Reactivity of Peptide a(3) Ions. *International Journal of Mass Spectrometry*, **2012**, 316-318, 259-267	1.9	4
130	PCR-electrospray ionization mass spectrometry: the potential to change infectious disease diagnostics in clinical and public health laboratories. *Journal of Molecular Diagnostics*, **2012**, 14, 295-304	5.1	71
129	The influence glutamic acid in protonated b -lb formation from VGEIG and related analogs. *International Journal of Mass Spectrometry*, **2012**, 325-327, 139-149	1.9	5
128	N-terminal region of CusB is sufficient for metal binding and metal transfer with the metallochaperone CusF. *Biochemistry*, **2012**, 51, 6767-75	3.2	30
127	Surface-induced dissociation of ion mobility-separated noncovalent complexes in a quadrupole/time-of-flight mass spectrometer. *Analytical Chemistry*, **2012**, 84, 6016-23	7.8	58
126	Study of the fragmentation of arginine isobutyl ester applied to arginine quantification in Aedes aegypti mosquito excreta. *Journal of Mass Spectrometry*, **2012**, 47, 1364-71	2.2	11
125	Development of a host blood meal database: de novo sequencing of hemoglobin from nine small mammals using mass spectrometry. *Biological Chemistry*, **2012**, 393, 195-201	4.5	16
124	Methods of Mass Spectrometry in Homeland Security Applications **2012**, 417-439		2
123	Structural influences on preferential oxazolone versus diketopiperazine b(2+) ion formation for histidine analogue-containing peptides. *Journal of Physical Chemistry A*, **2012**, 116, 4296-304	2.8	30
122	Protein Subunits Released by Surface Collisions of Noncovalent Complexes: Nativelike Compact Structures Revealed by Ion Mobility Mass Spectrometry. *Angewandte Chemie*, **2012**, 124, 4412-4415	3.6	4
121	Protein subunits released by surface collisions of noncovalent complexes: nativelike compact structures revealed by ion mobility mass spectrometry. *Angewandte Chemie - International Edition*, **2012**, 51, 4336-9	16.4	82
120	SQID-XLink: implementation of an intensity-incorporated algorithm for cross-linked peptide identification. *Bioinformatics*, **2012**, 28, 2548-50	7.2	10
119	ETD fragmentation features improve algorithm. *Expert Review of Proteomics*, **2012**, 9, 241-3	4.2	
118	Molecular structure and function of the novel BrnT/BrnA toxin-antitoxin system of Brucella abortus. *Journal of Biological Chemistry*, **2012**, 287, 12098-110	5.4	42
117	Activation by Oligomerization of an Allosteric Sequence Specific Endonuclease. *FASEB Journal*, **2012**, 26, lb91	0.9	
116	Characterization of a novel Aedes aegypti ferritin subunit identified utilizing proteomic techniques. *FASEB Journal*, **2012**, 26, 985.4	0.9	
115	Comparative analysis of PCR-electrospray ionization/mass spectrometry (MS) and MALDI-TOF/MS for the identification of bacteria and yeast from positive blood culture bottles. *Clinical Chemistry*, **2011**, 57, 1057-67	5.5	88

114	The role of proton bridges in selective cleavage of Ser-, Thr-, Cys-, Met-, Asp-, and Asn-containing peptides. *International Journal of Mass Spectrometry*, **2011**, 300, 108-117	1.9	8
113	Use of PCR coupled with electrospray ionization mass spectrometry for rapid identification of bacterial and yeast bloodstream pathogens from blood culture bottles. *Journal of Clinical Microbiology*, **2011**, 49, 345-53	9.7	87
112	Determinants of gas-phase disassembly behavior in homodimeric protein complexes with related yet divergent structures. *Analytical Chemistry*, **2011**, 83, 3881-9	7.8	25
111	Statistical analysis of electron transfer dissociation pairwise fragmentation patterns. *Analytical Chemistry*, **2011**, 83, 9540-5	7.8	31
110	Interactions between CusF and CusB identified by NMR spectroscopy and chemical cross-linking coupled to mass spectrometry. *Biochemistry*, **2011**, 50, 2559-66	3.2	41
109	Revealing the quaternary structure of a heterogeneous noncovalent protein complex through surface-induced dissociation. *Analytical Chemistry*, **2011**, 83, 2862-5	7.8	68
108	SQID: an intensity-incorporated protein identification algorithm for tandem mass spectrometry. *Journal of Proteome Research*, **2011**, 10, 1593-602	5.6	45
107	Proteomics analyses of the opportunistic pathogen Burkholderia vietnamiensis using protein fractionations and mass spectrometry. *Journal of Biomedicine and Biotechnology*, **2011**, 2011, 701928		9
106	NF45 and NF90 regulate HS4-dependent interleukin-13 transcription in T cells. *Journal of Biological Chemistry*, **2010**, 285, 8256-67	5.4	46
105	Mechanistic differences between two conserved classes of small heat shock proteins found in the plant cytosol. *Journal of Biological Chemistry*, **2010**, 285, 11489-97	5.4	70
104	Understanding and exploiting Peptide fragment ion intensities using experimental and informatic approaches. *Methods in Molecular Biology*, **2010**, 604, 73-94	1.4	13
103	Differential ammonia metabolism in Aedes aegypti fat body and midgut tissues. *Journal of Insect Physiology*, **2010**, 56, 1040-9	2.4	44
102	Analysis of deprotonated acids with silicon nanoparticle-assisted laser desorption/ ionization mass spectrometry. *Journal of Mass Spectrometry*, **2010**, 45, 1394-401	2.2	11
101	Separation and identification of structural isomers by quadrupole collision-induced dissociation-hydrogen/deuterium exchange-infrared multiphoton dissociation (QCID-HDX-IRMPD). *Journal of the American Society for Mass Spectrometry*, **2010**, 21, 1329-38	3.5	27
100	Differentiation and quantification of C1 and C2 (13)C-labeled glucose by tandem mass spectrometry. *Analytical Biochemistry*, **2010**, 404, 40-4	3.1	11
99	Rapid molecular genotyping and clonal complex assignment of Staphylococcus aureus isolates by PCR coupled to electrospray ionization-mass spectrometry. *Journal of Clinical Microbiology*, **2009**, 47, 1733-41	9.7	55
98	Pathogen profiling: rapid molecular characterization of Staphylococcus aureus by PCR/electrospray ionization-mass spectrometry and correlation with phenotype. *Journal of Clinical Microbiology*, **2009**, 47, 3129-37	9.7	54
97	Internal energy deposition with silicon nanoparticle-assisted laser desorption/ionization (SPALDI) mass spectrometry. *International Journal of Mass Spectrometry*, **2009**, 283, 200-205	1.9	32

96	Kinetics of surface-induced dissociation of N(CH3)4(+) and N(CD3)4(+) using silicon nanoparticle assisted laser desorption/ionization and laser desorption/ionization. *Journal of the American Society for Mass Spectrometry*, **2009**, 20, 957-64	3.5	5
95	Influence of cluster size and ion activation method on the dissociation of cesium iodide clusters. *International Journal of Mass Spectrometry*, **2009**, 287, 105-113	1.9	12
94	Evidence of diketopiperazine and oxazolone structures for HA b2+ ion. *Journal of the American Chemical Society*, **2009**, 131, 17528-9	16.4	87
93	Noncovalent protein tetramers and pentamers with "n" charges yield monomers with n/4 and n/5 charges. *Analytical Chemistry*, **2009**, 81, 1347-56	7.8	64
92	Disulfide-dependent self-assembly of adiponectin octadecamers from trimers and presence of stable octadecameric adiponectin lacking disulfide bonds in vitro. *Biochemistry*, **2009**, 48, 12345-57	3.2	30
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