

Davy Guillarme

List of Publications by Year in Descending Order

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Version: 2024-04-10

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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.

245 papers	10,040 citations	57 h-index	84 g-index
260 ext. papers	11,219 ext. citations	4.7 avg, IF	6.74 L-index

#	Paper	IF	Citations
245	A New Practice to Monitor the Fabrication Process of Fab-Targeting Ligands from Bevacizumab by LC-MS: Preparation and Analytical Characterization. <i>Scientia Pharmaceutica</i> , 2022 , 90, 5	4.3	
244	Bispecific antibody characterization by a combination of intact and site-specific/chain-specific LC/MS techniques. <i>Talanta</i> , 2022 , 236, 122836	6.2	4
243	Automated ion exchange chromatography screening combined with in silico multifactorial simulation for efficient method development and purification of biopharmaceutical targets.. <i>Analytical and Bioanalytical Chemistry</i> , 2022 , 414, 3581	4.4	1
242	Monitoring multiple quality attributes of a complex Fc-fusion protein during cell culture production processes by mD-LC-MS peptide mapping.. <i>Talanta</i> , 2022 , 246, 123519	6.2	1
241	Quantitative Glycan Profiling of Therapeutic Monoclonal Antibodies Performed by Middle-Up Level HILIC-HRMS Analysis. <i>Pharmaceutics</i> , 2021 , 13,	6.4	2
240	New wide-pore superficially porous stationary phases with low hydrophobicity applied for the analysis of monoclonal antibodies. <i>Journal of Chromatography A</i> , 2021 , 1642, 462050	4.5	3
239	Expanding the range of sub/supercritical fluid chromatography: Advantageous use of methanesulfonic acid in water-rich modifiers for peptide analysis. <i>Journal of Chromatography A</i> , 2021 , 1642, 462048	4.5	12
238	Ion mobility-high resolution mass spectrometry in anti-doping analysis. Part I: Implementation of a screening method with the assessment of a library of substances prohibited in sports. <i>Analytica Chimica Acta</i> , 2021 , 1152, 338257	6.6	9
237	Alternative mobile phase additives for the characterization of protein biopharmaceuticals in liquid chromatography - Mass spectrometry. <i>Analytica Chimica Acta</i> , 2021 , 1156, 338347	6.6	5
236	Aptamer-based immunoaffinity LC-MS using an ultra-short column for rapid attomole level quantitation of intact mAbs. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2021 , 1173, 122694	3.2	3
235	State-of-the-Art Native Mass Spectrometry and Ion Mobility Methods to Monitor Homogeneous Site-Specific Antibody-Drug Conjugates Synthesis. <i>Pharmaceutics</i> , 2021 , 14,	5.2	8
234	Negative gradient slope methods to improve the separation of closely eluting proteins. <i>Journal of Chromatography A</i> , 2021 , 1635, 461743	4.5	7
233	Use of Ultra-short Columns for Therapeutic Protein Separations, Part 2: Designing the Optimal Column Dimension for Reversed-Phase Liquid Chromatography. <i>Analytical Chemistry</i> , 2021 , 93, 1285-1293 ^{7,8}	7.8	7
232	Use of Ultrashort Columns for Therapeutic Protein Separations. Part 1: Theoretical Considerations and Proof of Concept. <i>Analytical Chemistry</i> , 2021 , 93, 1277-1284	7.8	9
231	Ultra-high performance supercritical fluid chromatography coupled to tandem mass spectrometry for antidoping analyses: Assessment of the inter-laboratory reproducibility with urine samples. <i>Analytical Science Advances</i> , 2021 , 2, 68-75	1.1	2
230	Analytical challenges encountered and the potential of supercritical fluid chromatography: A perspective of five experts. <i>Analytical Science Advances</i> , 2021 , 2, 76-80	1.1	1
229	Therapeutic Fc-fusion proteins: Current analytical strategies. <i>Journal of Separation Science</i> , 2021 , 44, 35-62	3.4	26

228	Fast Afucosylation Profiling of Glycoengineered Antibody Subunits by Middle-Up Mass Spectrometry. <i>Methods in Molecular Biology</i> , 2021 , 2271, 73-83	1.4	1
227	Algorithms to optimize multi-column chromatographic separations of proteins. <i>Journal of Chromatography A</i> , 2021 , 1637, 461838	4.5	1
226	Multi-dimensional LC-MS: the next generation characterization of antibody-based therapeutics by unified online bottom-up, middle-up and intact approaches. <i>Analyst, The</i> , 2021 , 146, 747-769	5	29
225	Characterization of Glycosylated Proteins at Subunit Level by HILIC/MS. <i>Methods in Molecular Biology</i> , 2021 , 2271, 85-95	1.4	1
224	Metamorphosis of supercritical fluid chromatography: A viable tool for the analysis of polar compounds?. <i>TrAC - Trends in Analytical Chemistry</i> , 2021 , 141, 116304	14.6	16
223	Ion mobility-high resolution mass spectrometry in doping control analysis. Part II: Comparison of acquisition modes with and without ion mobility. <i>Analytica Chimica Acta</i> , 2021 , 1175, 338739	6.6	5
222	Chromatographic Strategies for the Successful Characterization of Protein Biopharmaceuticals 2021 , 57-71		
221	Empirical correction of non-linear pH gradients and a tool for application to protein ion exchange chromatography. <i>Journal of Chromatography A</i> , 2021 , 1651, 462320	4.5	
220	Isolation and Identification of Isocoumarin Derivatives With Specific Inhibitory Activity Against Wnt Pathway and Metabolome Characterization of. <i>Frontiers in Chemistry</i> , 2021 , 9, 664489	5	2
219	Using 1.5mm internal diameter columns for optimal compatibility with current liquid chromatographic systems. <i>Journal of Chromatography A</i> , 2021 , 1650, 462258	4.5	3
218	Towards a simple on-line coupling of ion exchange chromatography and native mass spectrometry for the detailed characterization of monoclonal antibodies. <i>Journal of Chromatography A</i> , 2021 , 1655, 462499	4.5	9
217	The importance of being metal-free: The critical choice of column hardware for size exclusion chromatography coupled to high resolution mass spectrometry. <i>Analytica Chimica Acta</i> , 2021 , 1183, 338987	6.6	6
216	Ultra-short ion-exchange columns for fast charge variants analysis of therapeutic proteins. <i>Journal of Chromatography A</i> , 2021 , 1657, 462568	4.5	4
215	Inter-laboratory study to evaluate the performance of automated online characterization of antibody charge variants by multi-dimensional LC-MS/MS. <i>Talanta</i> , 2021 , 234, 122628	6.2	7
214	Glycan-Mediated Technology for Obtaining Homogeneous Site-Specific Conjugated Antibody-Drug Conjugates: Synthesis and Analytical Characterization by Using Complementary Middle-up LC/HRMS Analysis. <i>Analytical Chemistry</i> , 2020 , 92, 8170-8177	7.8	8
213	Impact of the column on effluent pH in cation exchange pH gradient chromatography, a practical study. <i>Journal of Chromatography A</i> , 2020 , 1626, 461350	4.5	3
212	Applicability of Supercritical fluid chromatography-Mass spectrometry to metabolomics. II-Assessment of a comprehensive library of metabolites and evaluation of biological matrices. <i>Journal of Chromatography A</i> , 2020 , 1620, 461021	4.5	26
211	Interlaboratory and Interplatform Study of Steroids Collision Cross Section by Traveling Wave Ion Mobility Spectrometry. <i>Analytical Chemistry</i> , 2020 , 92, 5013-5022	7.8	28

210	Current and future trends in reversed-phase liquid chromatography-mass spectrometry of therapeutic proteins. <i>TrAC - Trends in Analytical Chemistry</i> , 2020 , 130, 115962	14.6	13
209	Non-invasive targeted iontophoretic delivery of cetuximab to skin. <i>Expert Opinion on Drug Delivery</i> , 2020 , 17, 589-602	8	8
208	Development of a 3D-LC/MS Workflow for Fast, Automated, and Effective Characterization of Glycosylation Patterns of Biotherapeutic Products. <i>Analytical Chemistry</i> , 2020 , 92, 4357-4363	7.8	20
207	Coupling non-denaturing chromatography to mass spectrometry for the characterization of monoclonal antibodies and related products. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2020 , 185, 113207	3.5	22
206	Determination of size variants by CE-SDS for approved therapeutic antibodies: Key implications of subclasses and light chain specificities. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2020 , 184, 113166	3.5	15
205	Automated middle-up approach for the characterization of biotherapeutic products by combining on-line hinge-specific digestion with RPLC-HRMS analysis. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2020 , 182, 113130	3.5	9
204	Improving selectivity and performing online on-column fractioning in liquid chromatography for the separation of therapeutic biopharmaceutical products. <i>Journal of Chromatography A</i> , 2020 , 1618, 460901	4.5	8
203	Fast and Automated Characterization of Monoclonal Antibody Minor Variants from Cell Cultures by Combined Protein-A and Multidimensional LC/MS Methodologies. <i>Analytical Chemistry</i> , 2020 , 92, 8506-8513	7.8	19
202	Supercritical fluid chromatography-mass spectrometry in routine anti-doping analyses: Estimation of retention time variability under reproducible conditions. <i>Journal of Chromatography A</i> , 2020 , 1616, 460780	4.5	8
201	From proof of concept to the routine use of an automated and robust multi-dimensional liquid chromatography mass spectrometry workflow applied for the charge variant characterization of therapeutic antibodies. <i>Journal of Chromatography A</i> , 2020 , 1615, 460740	4.5	21
200	Evaluation of Different Tandem MS Acquisition Modes to Support Metabolite Annotation in Human Plasma Using Ultra High-Performance Liquid Chromatography High-Resolution Mass Spectrometry for Untargeted Metabolomics. <i>Metabolites</i> , 2020 , 10,	5.6	4
199	Supercritical fluid chromatography - Mass spectrometry in metabolomics: Past, present, and future perspectives. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2020 , 1161, 122444	3.2	22
198	Investigating the use of unconventional temperatures in supercritical fluid chromatography. <i>Analytica Chimica Acta</i> , 2020 , 1134, 84-95	6.6	7
197	Development of an innovative salt-mediated pH gradient cation exchange chromatography method for the characterization of therapeutic antibodies. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2020 , 1160, 122379	3.2	6
196	Targeted Bottom-up Characterization of Recombinant Monoclonal Antibodies by Multidimensional LC/MS. <i>Analytical Chemistry</i> , 2020 , 92, 13420-13426	7.8	8
195	Evaluation of additives on reversed-phase chromatography of monoclonal antibodies using a 1000 Å stationary phase. <i>Journal of Chromatography A</i> , 2020 , 1610, 460562	4.5	10
194	Glycosylation of biosimilars: Recent advances in analytical characterization and clinical implications. <i>Analytica Chimica Acta</i> , 2019 , 1089, 1-18	6.6	37
193	Proof of Concept To Achieve Infinite Selectivity for the Chromatographic Separation of Therapeutic Proteins. <i>Analytical Chemistry</i> , 2019 , 91, 12954-12961	7.8	17

192	Practical considerations on the particle size and permeability of ion-exchange columns applied to biopharmaceutical separations. <i>Journal of Chromatography A</i> , 2019 , 1604, 460487	4.5	1
191	Cutting-edge multi-level analytical and structural characterization of antibody-drug conjugates: present and future. <i>Expert Review of Proteomics</i> , 2019 , 16, 337-362	4.2	30
190	The Emergence of Universal Chromatographic Methods in the Research and Development of New Drug Substances. <i>Accounts of Chemical Research</i> , 2019 , 52, 1990-2002	24.3	31
189	Tuning selectivity in cation-exchange chromatography applied for monoclonal antibody separations, part 2: Evaluation of recent stationary phases. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2019 , 172, 320-328	3.5	11
188	Utility of dry load injection for an efficient natural products isolation at the semi-preparative chromatographic scale. <i>Journal of Chromatography A</i> , 2019 , 1598, 85-91	4.5	20
187	Tuning selectivity in cation-exchange chromatography applied for monoclonal antibody separations, part 1: Alternative mobile phases and fine tuning of the separation. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2019 , 168, 138-147	3.5	22
186	Supercritical fluid chromatography [Mass spectrometry: Recent evolution and current trends. <i>TrAC - Trends in Analytical Chemistry</i> , 2019 , 118, 731-738	14.6	43
185	Impact of particle size gradients on the apparent efficiency of chromatographic columns. <i>Journal of Chromatography A</i> , 2019 , 1603, 208-215	4.5	6
184	Streamlined Characterization of an Antibody-Drug Conjugate by Two-Dimensional and Four-Dimensional Liquid Chromatography/Mass Spectrometry. <i>Analytical Chemistry</i> , 2019 , 91, 14896-14903	7.8	22
183	Analytical strategies for the determination of amino acids: Past, present and future trends. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2019 , 1132, 121819	3.2	35
182	Apparent efficiency of serially coupled columns in gradient elution liquid chromatography: Extension to the combination of any column formats. <i>Journal of Chromatography A</i> , 2019 , 1588, 159-162	4.5	5
181	Is hydrophobic interaction chromatography the most suitable technique to characterize site-specific antibody-drug conjugates?. <i>Journal of Chromatography A</i> , 2019 , 1586, 149-153	4.5	14
180	Recent Advances in Chromatography for Pharmaceutical Analysis. <i>Analytical Chemistry</i> , 2019 , 91, 210-239	7.8	47
179	Computer-assisted UHPLC-MS method development and optimization for the determination of 24 antineoplastic drugs used in hospital pharmacy. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2019 , 164, 395-401	3.5	24
178	A generic workflow for the characterization of therapeutic monoclonal antibodies-application to daratumumab. <i>Analytical and Bioanalytical Chemistry</i> , 2019 , 411, 4615-4627	4.4	18
177	A scoring approach for multi-platform acquisition in metabolomics. <i>Journal of Chromatography A</i> , 2019 , 1592, 47-54	4.5	25
176	Orthogonal Middle-up Approaches for Characterization of the Glycan Heterogeneity of Etanercept by Hydrophilic Interaction Chromatography Coupled to High-Resolution Mass Spectrometry. <i>Analytical Chemistry</i> , 2019 , 91, 873-880	7.8	21
175	Characterization of an antibody-drug conjugate by hydrophilic interaction chromatography coupled to mass spectrometry. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2018 , 1080, 37-41	3.2	32

174	Natural compounds analysis using liquid and supercritical fluid chromatography hyphenated to mass spectrometry: Evaluation of a new design of atmospheric pressure ionization source. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2018 , 1083, 1-11	3.2	16
173	Extending the limits of size exclusion chromatography: Simultaneous separation of free payloads and related species from antibody drug conjugates and their aggregates. <i>Journal of Chromatography A</i> , 2018 , 1539, 19-29	4.5	15
172	Systematic evaluation of matrix effects in supercritical fluid chromatography versus liquid chromatography coupled to mass spectrometry for biological samples. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2018 , 1079, 51-61	3.2	31
171	Development of Comprehensive Online Two-Dimensional Liquid Chromatography/Mass Spectrometry Using Hydrophilic Interaction and Reversed-Phase Separations for Rapid and Deep Profiling of Therapeutic Antibodies. <i>Analytical Chemistry</i> , 2018 , 90, 5923-5929	7.8	58
170	Implementation of a generic liquid chromatographic method development workflow: Application to the analysis of phytocannabinoids and Cannabis sativa extracts. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2018 , 155, 116-124	3.5	23
169	Hyphenation of size exclusion chromatography to native ion mobility mass spectrometry for the analytical characterization of therapeutic antibodies and related products. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2018 , 1086, 176-183	3.2	52
168	On-tubing fluorescence measurements of the band broadening of contemporary injectors in ultra-high performance liquid chromatography. <i>Journal of Chromatography A</i> , 2018 , 1535, 44-54	4.5	10
167	An Online Four-Dimensional HICSEC-IMMS Methodology for Proof-of-Concept Characterization of Antibody Drug Conjugates. <i>Analytical Chemistry</i> , 2018 , 90, 1578-1586	7.8	57
166	Utility of a high coverage phenyl-bonding and wide-pore superficially porous particle for the analysis of monoclonal antibodies and related products. <i>Journal of Chromatography A</i> , 2018 , 1549, 63-76	4.5	29
165	What are the current solutions for interfacing supercritical fluid chromatography and mass spectrometry?. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2018 , 1083, 160-170	3.2	52
164	Current possibilities of liquid chromatography for the characterization of antibody-drug conjugates. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2018 , 147, 493-505	3.5	39
163	Influence of connection tubing in modern size exclusion chromatography and its impact on the characterization of mAbs. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2018 , 149, 22-32	3.5	3
162	Monoclonal antibody N-glycosylation profiling using capillary electrophoresis - Mass spectrometry: Assessment and method validation. <i>Talanta</i> , 2018 , 178, 530-537	6.2	42
161	Adding a new separation dimension to MS and LC-MS: What is the utility of ion mobility spectrometry?. <i>Journal of Separation Science</i> , 2018 , 41, 20-67	3.4	94
160	Unraveling the mysteries of modern size exclusion chromatography - the way to achieve confident characterization of therapeutic proteins. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2018 , 1092, 368-378	3.2	24
159	Characterizing various monoclonal antibodies with milder reversed phase chromatography conditions. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2018 , 1096, 1-10	3.2	18
158	An attempt to characterize the human Chorionic Gonadotropin protein by reversed phase liquid chromatography coupled with high-resolution mass spectrometry at the intact level. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2018 , 161, 35-44	3.5	12
157	Protocols for the analytical characterization of therapeutic monoclonal antibodies. III - Denaturing chromatographic techniques hyphenated to mass spectrometry. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2018 , 1096, 95-106	3.2	22

156	Apparent efficiency of serially coupled columns in isocratic and gradient elution modes. <i>Journal of Chromatography A</i> , 2018 , 1571, 121-131	4.5	12
155	Applicability of supercritical fluid chromatography - mass spectrometry to metabolomics. I - Optimization of separation conditions for the simultaneous analysis of hydrophilic and lipophilic substances. <i>Journal of Chromatography A</i> , 2018 , 1562, 96-107	4.5	55
154	New developments and possibilities of wide-pore superficially porous particle technology applied for the liquid chromatographic analysis of therapeutic proteins. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2018 , 158, 225-235	3.5	21
153	Development of a LC-MS/MS method for the determination of isomeric glutamyl peptides in food ingredients. <i>Journal of Separation Science</i> , 2018 , 41, 847-855	3.4	9
152	Improved separation by at-column dilution in preparative hydrophilic interaction chromatography. <i>Journal of Chromatography A</i> , 2018 , 1532, 136-143	4.5	2
151	5. What is the potential of SFC-MS for doping control analysis? 2018 , 111-128		
150	A Novel Online Four-Dimensional SECSEC-IMMS Methodology for Characterization of Monoclonal Antibody Size Variants. <i>Analytical Chemistry</i> , 2018 , 90, 13929-13937	7.8	37
149	First inter-laboratory study of a Supercritical Fluid Chromatography method for the determination of pharmaceutical impurities. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2018 , 161, 414-424	3.5	43
148	High-resolution separation of monoclonal antibodies mixtures and their charge variants by an alternative and generic CZE method. <i>Electrophoresis</i> , 2018 , 39, 2083-2090	3.6	18
147	Hydrophilic Interaction Chromatography Hyphenated with Mass Spectrometry: A Powerful Analytical Tool for the Comparison of Originator and Biosimilar Therapeutic Monoclonal Antibodies at the Middle-up Level of Analysis. <i>Analytical Chemistry</i> , 2017 , 89, 2086-2092	7.8	62
146	Separation of antibody drug conjugate species by RPLC: A generic method development approach. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2017 , 137, 60-69	3.5	20
145	Development of a fast workflow to screen the charge variants of therapeutic antibodies. <i>Journal of Chromatography A</i> , 2017 , 1498, 147-154	4.5	24
144	Achievable separation performance and analysis time in current liquid chromatographic practice for monoclonal antibody separations. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2017 , 141, 59-69	3.5	16
143	Protocols for the analytical characterization of therapeutic monoclonal antibodies. I - Non-denaturing chromatographic techniques. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2017 , 1058, 73-84	3.2	31
142	Optimized selection of liquid chromatography conditions for wide range analysis of natural compounds. <i>Journal of Chromatography A</i> , 2017 , 1504, 91-104	4.5	20
141	Quantitative determination of salbutamol sulfate impurities using achiral supercritical fluid chromatography. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2017 , 134, 170-180	3.5	39
140	Analysis of recombinant monoclonal antibodies in hydrophilic interaction chromatography: A generic method development approach. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2017 , 145, 24-32	3.5	21
139	Antineoplastic drugs and their analysis: a state of the art review. <i>Analyst, The</i> , 2017 , 142, 2273-2321	5	23

138	Protocols for the analytical characterization of therapeutic monoclonal antibodies. II - Enzymatic and chemical sample preparation. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2017 , 1060, 325-335	3.2	45
137	The importance of system band broadening in modern size exclusion chromatography. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2017 , 135, 50-60	3.5	19
136	Optimization of non-linear gradient in hydrophobic interaction chromatography for the analytical characterization of antibody-drug conjugates. <i>Journal of Chromatography A</i> , 2017 , 1481, 82-91	4.5	19
135	Determination of isoelectric points and relative charge variants of 23 therapeutic monoclonal antibodies. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2017 , 1065-1066, 119-128	3.2	85
134	Characterization of 30 therapeutic antibodies and related products by size exclusion chromatography: Feasibility assessment for future mass spectrometry hyphenation. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2017 , 1065-1066, 35-43	3.2	49
133	A workflow for column interchangeability in liquid chromatography using modeling software and quality-by-design principles. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2017 , 146, 220-225	3.5	11
132	A systematic investigation of sample diluents in modern supercritical fluid chromatography. <i>Journal of Chromatography A</i> , 2017 , 1511, 122-131	4.5	54
131	Comprehensive study on the effects of sodium and potassium additives in size exclusion chromatographic separations of protein biopharmaceuticals. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2017 , 144, 242-251	3.5	17
130	Evaluation of size exclusion chromatography columns packed with sub-3 μ m particles for the analysis of biopharmaceutical proteins. <i>Journal of Chromatography A</i> , 2017 , 1498, 80-89	4.5	44
129	Theory and Practice of UHPLC and UHPLC/MS 2017 , 1-38		
128	Systematic evaluation of matrix effects in hydrophilic interaction chromatography versus reversed phase liquid chromatography coupled to mass spectrometry. <i>Journal of Chromatography A</i> , 2016 , 1439, 42-53	4.5	24
127	Computer assisted liquid chromatographic method development for the separation of therapeutic proteins. <i>Analyst, The</i> , 2016 , 141, 5488-501	5	17
126	Analysis of antibody-drug conjugates by comprehensive on-line two-dimensional hydrophobic interaction chromatography x reversed phase liquid chromatography hyphenated to high resolution mass spectrometry. II- Identification of sub-units for the characterization of even and odd load drug species. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2016 , 1052, 91-102	3.2	24
125	Preparative Scale MS-Guided Isolation of Bioactive Compounds Using High-Resolution Flash Chromatography: Antifungals from <i>Chiloscyphus polyanthos</i> as a Case Study. <i>Planta Medica</i> , 2016 , 82, 1051-7	3.1	10
124	Prototype sphere-on-sphere silica particles for the separation of large biomolecules. <i>Journal of Chromatography A</i> , 2016 , 1431, 94-102	4.5	9
123	Practical method development for the separation of monoclonal antibodies and antibody-drug-conjugate species in hydrophobic interaction chromatography, part 2: Optimization of the phase system. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2016 , 121, 161-173	3.5	36
122	Separation of substrates and closely related glucuronide metabolites using various chromatographic modes. <i>Journal of Chromatography A</i> , 2016 , 1435, 54-65	4.5	16
121	Chromatographic, Electrophoretic, and Mass Spectrometric Methods for the Analytical Characterization of Protein Biopharmaceuticals. <i>Analytical Chemistry</i> , 2016 , 88, 480-507	7.8	162

120	Fast and sensitive supercritical fluid chromatography - tandem mass spectrometry multi-class screening method for the determination of doping agents in urine. <i>Analytica Chimica Acta</i> , 2016 , 915, 102-10	6.6	50
119	Evaluation of innovative stationary phase ligand chemistries and analytical conditions for the analysis of basic drugs by supercritical fluid chromatography. <i>Journal of Chromatography A</i> , 2016 , 1438, 244-53	4.5	25
118	Prediction of retention time in reversed-phase liquid chromatography as a tool for steroid identification. <i>Analytica Chimica Acta</i> , 2016 , 916, 8-16	6.6	44
117	Practical method development for the separation of monoclonal antibodies and antibody-drug-conjugate species in hydrophobic interaction chromatography, part 1: optimization of the mobile phase. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2016 , 118, 393-403	3.5	50
116	Analytical Strategies for Doping Control Purposes: Needs, Challenges, and Perspectives. <i>Analytical Chemistry</i> , 2016 , 88, 508-23	7.8	31
115	Analysis of antibody-drug conjugates by comprehensive on-line two-dimensional hydrophobic interaction chromatography x reversed phase liquid chromatography hyphenated to high resolution mass spectrometry. I - Optimization of separation conditions. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2016 , 1032, 103-111	3.2	41
114	Evaluation of thermally pretreated silica stationary phases under hydrophilic interaction chromatography conditions. <i>Journal of Separation Science</i> , 2016 , 39, 1611-8	3.4	
113	Comparison of originator and biosimilar therapeutic monoclonal antibodies using comprehensive two-dimensional liquid chromatography coupled with time-of-flight mass spectrometry. <i>MAbs</i> , 2016 , 8, 1224-1234	6.6	61
112	Liquid chromatography and supercritical fluid chromatography as alternative techniques to gas chromatography for the rapid screening of anabolic agents in urine. <i>Journal of Chromatography A</i> , 2016 , 1451, 145-155	4.5	50
111	Potential of hydrophilic interaction chromatography for the analytical characterization of protein biopharmaceuticals. <i>Journal of Chromatography A</i> , 2016 , 1448, 81-92	4.5	66
110	Ultra-high performance supercritical fluid chromatography coupled with quadrupole-time-of-flight mass spectrometry as a performing tool for bioactive analysis. <i>Journal of Chromatography A</i> , 2016 , 1450, 101-11	4.5	49
109	Hydrophobic interaction chromatography for the characterization of monoclonal antibodies and related products. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2016 , 130, 3-18	3.5	76
108	Importance of vial shape and type on the reproducibility of size exclusion chromatography measurement of monoclonal antibodies. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2016 , 1032, 131-138	3.2	7
107	Impact of organic modifier and temperature on protein denaturation in hydrophobic interaction chromatography. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2016 , 131, 124-132	3.5	26
106	Systematic evaluation of mobile phase additives for the LC-MS characterization of therapeutic proteins. <i>Talanta</i> , 2015 , 136, 60-7	6.2	29
105	Comparison of the most recent chromatographic approaches applied for fast and high resolution separations: Theory and practice. <i>Journal of Chromatography A</i> , 2015 , 1408, 1-14	4.5	57
104	Direct identification of rituximab main isoforms and subunit analysis by online selective comprehensive two-dimensional liquid chromatography-mass spectrometry. <i>Analytical Chemistry</i> , 2015 , 87, 8307-15	7.8	77
103	Strategies for formulating and delivering poorly water-soluble drugs. <i>Journal of Drug Delivery Science and Technology</i> , 2015 , 30, 342-351	4.5	84

102	Estimation of pressure-, temperature- and frictional heating-related effects on proteinsTretention under ultra-high-pressure liquid chromatographic conditions. <i>Journal of Chromatography A</i> , 2015 , 1393, 73-80	4.5	25
101	Characterization of cation exchanger stationary phases applied for the separations of therapeutic monoclonal antibodies. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2015 , 111, 169-76	3.5	27
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