

Szabolcs Fekete

List of Publications by Year in descending order

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124
papers

5,766
citations

57719

44
h-index

91828

69
g-index

125
all docs

125
docs citations

125
times ranked

3101
citing authors

#	ARTICLE	IF	CITATIONS
1	Fast liquid chromatography: The domination of core-shell and very fine particles. <i>Journal of Chromatography A</i> , 2012, 1228, 57-71.	1.8	232
2	Theory and practice of size exclusion chromatography for the analysis of protein aggregates. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2014, 101, 161-173.	1.4	226
3	Chromatographic, Electrophoretic, and Mass Spectrometric Methods for the Analytical Characterization of Protein Biopharmaceuticals. <i>Analytical Chemistry</i> , 2016, 88, 480-507.	3.2	205
4	Ion-exchange chromatography for the characterization of biopharmaceuticals. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2015, 113, 43-55.	1.4	186
5	Comparative study of new shell-type, sub-2 μ m fully porous and monolith stationary phases, focusing on mass-transfer resistance. <i>Journal of Chromatography A</i> , 2010, 1217, 3642-3653.	1.8	159
6	Current and future trends in UHPLC. <i>TrAC - Trends in Analytical Chemistry</i> , 2014, 63, 2-13.	5.8	140
7	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.	1.2	135
8	Method development for the separation of monoclonal antibody charge variants in cation exchange chromatography, Part I: Salt gradient approach. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2015, 102, 33-44.	1.4	133
9	New trends in reversed-phase liquid chromatographic separations of therapeutic peptides and proteins: Theory and applications. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2012, 69, 9-27.	1.4	120
10	Importance of instrumentation for fast liquid chromatography in pharmaceutical analysis. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2014, 87, 105-119.	1.4	113
11	Method development for the separation of monoclonal antibody charge variants in cation exchange chromatography, Part II: pH gradient approach. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2015, 102, 282-289.	1.4	113
12	Analytical strategies for the characterization of therapeutic monoclonal antibodies. <i>TrAC - Trends in Analytical Chemistry</i> , 2013, 42, 74-83.	5.8	104
13	Hydrophobic interaction chromatography for the characterization of monoclonal antibodies and related products. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2016, 130, 3-18.	1.4	104
14	Maximizing kinetic performance in supercritical fluid chromatography using state-of-the-art instruments. <i>Journal of Chromatography A</i> , 2013, 1314, 288-297.	1.8	94
15	The impact of extra-column band broadening on the chromatographic efficiency of 5cm long narrow-bore very efficient columns. <i>Journal of Chromatography A</i> , 2011, 1218, 5286-5291.	1.8	92
16	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, 8307-8315.	3.2	90
17	Efficiency of the new sub-2 μ m core-shell (Kinetex ® , C_8) column in practice, applied for small and large molecule separation. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2011, 54, 482-490.	1.4	87
18	Recent Advances in Chromatography for Pharmaceutical Analysis. <i>Analytical Chemistry</i> , 2019, 91, 210-239.	3.2	85

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19	Potential of hydrophilic interaction chromatography for the analytical characterization of protein biopharmaceuticals. <i>Journal of Chromatography A</i> , 2016, 1448, 81-92.	1.8	80
20	Therapeutic Fc fusion proteins: Current analytical strategies. <i>Journal of Separation Science</i> , 2021, 44, 35-62.	1.3	78
21	Kinetic evaluation of new generation of column packed with 1.3µm core-shell particles. <i>Journal of Chromatography A</i> , 2013, 1308, 104-113.	1.8	77
22	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.	3.2	77
23	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.	2.6	76
24	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.	1.2	73
25	Evaluation of a new wide pore core-shell material (Aeris, WIDEPORE) and comparison with other existing stationary phases for the analysis of intact proteins. <i>Journal of Chromatography A</i> , 2012, 1236, 177-188.	1.8	72
26	The effect of pressure and mobile phase velocity on the retention properties of small analytes and large biomolecules in ultra-high pressure liquid chromatography. <i>Journal of Chromatography A</i> , 2012, 1270, 127-138.	1.8	66
27	Shell and small particles; Evaluation of new column technology. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2009, 49, 64-71.	1.4	64
28	Pushing the performance limits of reversed-phase ultra high performance liquid chromatography with 1.3µm core-shell particles. <i>Journal of Chromatography A</i> , 2013, 1311, 90-97.	1.8	64
29	Evaluation of size exclusion chromatography columns packed with sub-3 µm particles for the analysis of biopharmaceutical proteins. <i>Journal of Chromatography A</i> , 2017, 1498, 80-89.	1.8	64
30	Impact of mobile phase temperature on recovery and stability of monoclonal antibodies using recent reversed-phase stationary phases. <i>Journal of Separation Science</i> , 2012, 35, 3113-3123.	1.3	62
31	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.	1.8	61
32	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.	1.4	61
33	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.	1.2	59
34	Current possibilities of liquid chromatography for the characterization of antibody-drug conjugates. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2018, 147, 493-505.	1.4	54
35	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.	1.2	51
36	Rapid high performance liquid chromatography method development with high prediction accuracy, using 5cm long narrow bore columns packed with sub-2µm particles and Design Space computer modeling. <i>Journal of Chromatography A</i> , 2009, 1216, 7816-7823.	1.8	49

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37	Analysis of recombinant monoclonal antibodies by RPLC: Toward a generic method development approach. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2012, 70, 158-168.	1.4	49
38	Critical evaluation of fast size exclusion chromatographic separations of protein aggregates, applying sub-2 μ m particles. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2013, 78-79, 141-149.	1.4	49
39	Ultra-high-performance liquid chromatography for the characterization of therapeutic proteins. <i>TrAC - Trends in Analytical Chemistry</i> , 2014, 63, 76-84.	5.8	49
40	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.	1.2	48
41	Facts and myths about columns packed with sub-3 μ m and sub-2 μ m particles. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2010, 51, 56-64.	1.4	47
42	Evaluation of recent very efficient wide-pore stationary phases for the reversed-phase separation of proteins. <i>Journal of Chromatography A</i> , 2012, 1252, 90-103.	1.8	47
43	Development of a rapid method for the determination and confirmation of nitroimidazoles in six matrices by fast liquid chromatography-tandem mass spectrometry. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2012, 64-65, 40-48.	1.4	47
44	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.	1.3	47
45	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.	1.4	46
46	Evaluation of stationary phases packed with superficially porous particles for the analysis of pharmaceutical compounds using supercritical fluid chromatography. <i>Journal of Chromatography A</i> , 2014, 1360, 275-287.	1.8	44
47	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.	1.4	44
48	Contribution of various types of liquid chromatography-mass spectrometry instruments to band broadening in fast analysis. <i>Journal of Chromatography A</i> , 2013, 1310, 45-55.	1.8	42
49	Adsorption and recovery issues of recombinant monoclonal antibodies in reversed-phase liquid chromatography. <i>Journal of Separation Science</i> , 2015, 38, 1-8.	1.3	42
50	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.	1.2	42
51	Comparison of liquid chromatography and supercritical fluid chromatography coupled to compact single quadrupole mass spectrometer for targeted in vitro metabolism assay. <i>Journal of Chromatography A</i> , 2014, 1371, 244-256.	1.8	40
52	Characterization of new types of stationary phases for fast liquid chromatographic applications. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2009, 50, 703-709.	1.4	39
53	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.	1.4	38
54	Reliability of simulated robustness testing in fast liquid chromatography, using state-of-the-art column technology, instrumentation and modelling software. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2014, 89, 67-75.	1.4	36

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55	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.	1.8	36
56	Systematic evaluation of mobile phase additives for the LC-MS characterization of therapeutic proteins. <i>Talanta</i> , 2015, 136, 60-67.	2.9	34
57	Characterization of cation exchanger stationary phases applied for the separations of therapeutic monoclonal antibodies. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2015, 111, 169-176.	1.4	34
58	Fast gradient screening of pharmaceuticals with 5 cm long, narrow bore reversed-phase columns packed with sub-3 μm core-shell and sub-2 μm totally porous particles. <i>Talanta</i> , 2011, 84, 416-423.	2.9	32
59	Possibilities of new generation columns packed with 1.3 μm core-shell particles in gradient elution mode. <i>Journal of Chromatography A</i> , 2013, 1320, 86-95.	1.8	32
60	Influence of pressure and temperature on molar volume and retention properties of peptides in ultra-high pressure liquid chromatography. <i>Journal of Chromatography A</i> , 2013, 1311, 65-71.	1.8	32
61	Robust UHPLC Separation Method Development for Multi-API Product Containing Amlodipine and Bisoprolol: The Impact of Column Selection. <i>Chromatographia</i> , 2014, 77, 1119-1127.	0.7	32
62	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.	1.4	32
63	Systematic comparison of a new generation of columns packed with sub-2 μm superficially porous particles. <i>Journal of Separation Science</i> , 2014, 37, 189-197.	1.3	31
64	Development of a fast workflow to screen the charge variants of therapeutic antibodies. <i>Journal of Chromatography A</i> , 2017, 1498, 147-154.	1.8	31
65	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.	1.4	31
66	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, 1032, 91-102.	1.2	30
67	Proof of Concept To Achieve Infinite Selectivity for the Chromatographic Separation of Therapeutic Proteins. <i>Analytical Chemistry</i> , 2019, 91, 12954-12961.	3.2	30
68	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.	1.4	30
69	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.	3.2	29
70	Estimation of pressure-, temperature- and frictional heating-related effects on proteins' retention under ultra-high-pressure liquid chromatographic conditions. <i>Journal of Chromatography A</i> , 2015, 1393, 73-80.	1.8	28
71	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.	1.4	28
72	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.	1.2	28

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73	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.	1.4	28
74	A generic workflow for the characterization of therapeutic monoclonal antibodiesâ€™ application to daratumumab. <i>Analytical and Bioanalytical Chemistry</i> , 2019, 411, 4615-4627.	1.9	28
75	Current and future trends in reversed-phase liquid chromatography-mass spectrometry of therapeutic proteins. <i>TrAC - Trends in Analytical Chemistry</i> , 2020, 130, 115962.	5.8	28
76	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.	1.8	28
77	Comparative study of recent wide-pore materials of different stationary phase morphology, applied for the reversed-phase analysis of recombinant monoclonal antibodies. <i>Analytical and Bioanalytical Chemistry</i> , 2013, 405, 3137-3151.	1.9	26
78	Use of Ultrashort Columns for Therapeutic Protein Separations. Part 1: Theoretical Considerations and Proof of Concept. <i>Analytical Chemistry</i> , 2021, 93, 1277-1284.	3.2	26
79	High resolution reversed phase analysis of recombinant monoclonal antibodies by ultra-high pressure liquid chromatography column coupling. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2013, 83, 273-278.	1.4	25
80	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.	1.4	25
81	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.	1.2	25
82	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.	1.4	25
83	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.	1.4	24
84	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.	1.8	24
85	Estimation of the effects of longitudinal temperature gradients caused by frictional heating on the solute retention using fully porous and superficially porous sub-2 $\frac{1}{4}$ μ m materials. <i>Journal of Chromatography A</i> , 2014, 1359, 124-130.	1.8	23
86	The importance of system band broadening in modern size exclusion chromatography. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2017, 135, 50-60.	1.4	23
87	Computer assisted liquid chromatographic method development for the separation of therapeutic proteins. <i>Analyst</i> , The, 2016, 141, 5488-5501.	1.7	22
88	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.	1.4	21
89	Size Exclusion and Ion Exchange Chromatographic Hardware Modified with a Hydrophilic Hybrid Surface. <i>Analytical Chemistry</i> , 2022, 94, 3360-3367.	3.2	19
90	ANALYSIS OF SULFONAMIDE RESIDUES IN REAL HONEY SAMPLES USING LIQUID CHROMATOGRAPHY WITH FLUORESCENCE AND TANDEM MASS SPECTROMETRY DETECTION. <i>Journal of Liquid Chromatography and Related Technologies</i> , 2013, 36, 1105-1125.	0.5	18

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91	Evolution and Current Trends in Liquid and Supercritical Fluid Chromatography. <i>Current Chromatography</i> , 2014, 1, 15-40.	0.1	18
92	Reliability of computer-assisted method transfer between several column dimensions packed with 1.3-5.14m core-shell particles and between various instruments. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2014, 94, 188-195.	1.4	18
93	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.	1.4	18
94	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.	1.8	18
95	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.	1.4	17
96	Evaluation of a new wide-pore superficially porous material with carbon core and nanodiamond-polymer shell for the separation of proteins. <i>Journal of Chromatography A</i> , 2015, 1414, 51-59.	1.8	16
97	Evaluation of new superficially porous particles with carbon core and nanodiamond-polymer shell for proteins characterization. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2015, 104, 130-136.	1.4	16
98	Negative gradient slope methods to improve the separation of closely eluting proteins. <i>Journal of Chromatography A</i> , 2021, 1635, 461743.	1.8	16
99	Apparent efficiency of serially coupled columns in isocratic and gradient elution modes. <i>Journal of Chromatography A</i> , 2018, 1571, 121-131.	1.8	15
100	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.	1.2	13
101	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.	1.8	13
102	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.	3.2	13
103	Ultra-short ion-exchange columns for fast charge variants analysis of therapeutic proteins. <i>Journal of Chromatography A</i> , 2021, 1657, 462568.	1.8	13
104	Pressure-Enhanced Liquid Chromatography, a Proof of Concept: Tuning Selectivity with Pressure Changes and Gradients. <i>Analytical Chemistry</i> , 2022, 94, 7877-7884.	3.2	13
105	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.	2.6	12
106	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.	1.8	11
107	Using 1.5mm internal diameter columns for optimal compatibility with current liquid chromatographic systems. <i>Journal of Chromatography A</i> , 2021, 1650, 462258.	1.8	11
108	Comparison of various silica-based monoliths for the analysis of large biomolecules. <i>Journal of Separation Science</i> , 2013, 36, 2231-2243.	1.3	10

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109	Impact of particle size gradients on the apparent efficiency of chromatographic columns. Journal of Chromatography A, 2019, 1603, 208-215.	1.8	10
110	Prototype sphere-on-sphere silica particles for the separation of large biomolecules. Journal of Chromatography A, 2016, 1431, 94-102.	1.8	9
111	Updating the European Pharmacopoeia impurity profiling method for terazosin and suggesting alternative columns. Journal of Pharmaceutical and Biomedical Analysis, 2020, 187, 113371.	1.4	9
112	Enhancing the Quality of Separation in One-Dimensional Peptide Mapping Using Mathematical Transformation. Chromatographia, 2012, 75, 305-312.	0.7	8
113	Importance of vial shape and type on the reproducibility of size exclusion chromatography measurement of monoclonal antibodies. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2016, 1032, 131-138.	1.2	8
114	New wide-pore superficially porous stationary phases with low hydrophobicity applied for the analysis of monoclonal antibodies. Journal of Chromatography A, 2021, 1642, 462050.	1.8	8
115	Salt gradient and ion-pair mediated anion exchange of intact messenger ribonucleic acids. Journal of Chromatography Open, 2022, 2, 100031.	0.8	8
116	Direct coupling of size exclusion chromatography and mass spectrometry for the characterization of complex monoclonal antibody products. Journal of Separation Science, 2022, 45, 1997-2007.	1.3	8
117	Investigating the secondary interactions of packing materials for size-exclusion chromatography of therapeutic proteins. Journal of Chromatography A, 2022, 1676, 463262.	1.8	8
118	Aptamer-based immunoaffinity LC-MS using an ultra-short column for rapid attomole level quantitation of intact mAbs. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2021, 1173, 122694.	1.2	7
119	Influence of connection tubing in modern size exclusion chromatography and its impact on the characterization of mAbs. Journal of Pharmaceutical and Biomedical Analysis, 2018, 149, 22-32.	1.4	5
120	Practical considerations on the particle size and permeability of ion-exchange columns applied to biopharmaceutical separations. Journal of Chromatography A, 2019, 1604, 460487.	1.8	5
121	Apparent efficiency of serially coupled columns in gradient elution liquid chromatography: Extension to the combination of any column formats. Journal of Chromatography A, 2019, 1588, 159-162.	1.8	5
122	Development of a Fast and Robust UHPLC Method for Apixaban In-Process Control Analysis. Molecules, 2021, 26, 3505.	1.7	5
123	Algorithms to optimize multi-column chromatographic separations of proteins. Journal of Chromatography A, 2021, 1637, 461838.	1.8	1
124	Empirical correction of non-linear pH gradients and a tool for application to protein ion exchange chromatography. Journal of Chromatography A, 2021, 1651, 462320.	1.8	1