

Roger E Moore

List of Publications by Year in descending order

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Version: 2024-02-01

10
papers

876
citations

1163117

8
h-index

1372567

10
g-index

11
all docs

11
docs citations

11
times ranked

1159
citing authors

#	ARTICLE	IF	CITATIONS
1	Qscore: An algorithm for evaluating SEQUEST database search results. <i>Journal of the American Society for Mass Spectrometry</i> , 2002, 13, 378-386.	2.8	352
2	Composition and Functional Characterization of the Yeast Spliceosomal Penta-snRNP. <i>Molecular Cell</i> , 2002, 9, 31-44.	9.7	225
3	Proteomic analysis of in vivo-assembled pre-mRNA splicing complexes expands the catalog of participating factors. <i>Nucleic Acids Research</i> , 2007, 35, 3928-3944.	14.5	102
4	A Microscale Electrospray Interface Incorporating a Monolithic, Poly(styrene- <i>co</i> -divinylbenzene) Support for On-Line Liquid Chromatography/Tandem Mass Spectrometry Analysis of Peptides and Proteins. <i>Analytical Chemistry</i> , 1998, 70, 4879-4884.	6.5	81
5	Method for screening peptide fragment ion mass spectra prior to database searching. <i>Journal of the American Society for Mass Spectrometry</i> , 2000, 11, 422-426.	2.8	53
6	Osteoclasts Secrete the Chemotactic Cytokine Mim-1. <i>Biochemical and Biophysical Research Communications</i> , 2001, 281, 180-185.	2.1	31
7	Proteomic Analysis of Surface and Endosomal Membrane Proteins from the Avian LMH Epithelial Cell Line. <i>Journal of Proteome Research</i> , 2011, 10, 3973-3982.	3.7	12
8	MHC class I target recognition, immunophenotypes and proteomic profiles of natural killer cells within the spleens of day-14 chick embryos. <i>Developmental and Comparative Immunology</i> , 2012, 37, 446-456.	2.3	11
9	Protein Identification Using a Quadrupole Ion Trap Mass Spectrometer and SEQUEST Database Matching. <i>Current Protocols in Protein Science</i> , 2000, 22, Unit 16.10.	2.8	8
10	Introducing Samples Directly into Electrospray Ionization Mass Spectrometers Using Microscale Capillary Liquid Chromatography. <i>Current Protocols in Protein Science</i> , 2000, 22, Unit 16.9.	2.8	1