

The speciation of the proteome

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Two-dimensional electrophoresis-based characterization of post-translational modifications of mammalian 20S proteasome complexes. <i>Proteomics</i> , 2008, 8, 5025-5037.	1.3	56
2	Proteome changes in human bronchoalveolar cells following styrene exposure indicate involvement of oxidative stress in the molecular response mechanism. <i>Proteomics</i> , 2009, 9, 4920-4933.	1.3	19
3	Finding one's way in proteomics: a protein species nomenclature. <i>Chemistry Central Journal</i> , 2009, 3, 11.	2.6	229
4	Assembling proteomics data as a prerequisite for the analysis of large scale experiments. <i>Chemistry Central Journal</i> , 2009, 3, 2.	2.6	5
5	Comparison of two label-free global quantitation methods, APEX and 2D gel electrophoresis, applied to the <i>Shigella dysenteriae</i> proteome. <i>Proteome Science</i> , 2009, 7, 22.	0.7	28
6	Proteomic identification of secreted proteins of <i>Propionibacterium acnes</i> . <i>BMC Microbiology</i> , 2010, 10, 230.	1.3	142
7	<i>Helicobacter pylori</i> proteomics by 2-DE/MS, 1-DE-LC/MS and functional data mining. <i>Proteomics</i> , 2010, 10, 182-193.	1.3	35
8	Viral myocarditis induced by Coxsackievirus B3 in <i>A.BY/Sn</i> mice: Analysis of changes in the myocardial proteome. <i>Proteomics</i> , 2010, 10, 1802-1818.	1.3	15
9	An Empirical Strategy for Characterizing Bacterial Proteomes across Species in the Absence of Genomic Sequences. <i>PLoS ONE</i> , 2010, 5, e13968.	1.1	16
10	Immunology and Cell Biology of Parasitic Diseases. <i>Journal of Biomedicine and Biotechnology</i> , 2010, 1-5.	3.0	4
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17	Functional analysis of proteins and protein species using shotgun proteomics and linear mathematics. <i>Amino Acids</i> , 2011, 41, 329-341.	1.2	10
18	Adaptation of proteomic techniques for the identification and characterization of protein species from murine heart. <i>Amino Acids</i> , 2011, 41, 401-414.	1.2	12

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21	Superoxide dismutase type 1 in monocytes of chronic kidney disease patients. <i>Amino Acids</i> , 2011, 41, 427-438.	1.2	15
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24	Lights and shadows of proteomic technologies for the study of protein species including isoforms, splicing variants and protein post-translational modifications. <i>Proteomics</i> , 2011, 11, 590-603.	1.3	19
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49	Amino acids: Chemistry, functionality and selected non-enzymatic post-translational modifications. <i>Journal of Proteomics</i> , 2012, 75, 2275-2296.	1.2	131
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111	Comparative physiology and proteomic analysis of two wheat genotypes contrasting in drought tolerance. <i>Journal of Proteomics</i> , 2015, 114, 1-15.	1.2	99
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