

Human Serum Proteins Preseparated by Electrophoresis Tandem Mass Spectrometry

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Citation Report

#	ARTICLE	IF	CITATIONS
2	Current Awareness on Comparative and Functional Genomics. Comparative and Functional Genomics, 2004, 5, 658-669.	2.0	0
3	A strategy for high-resolution protein identification in surface-enhanced laser desorption/ionization mass spectrometry: Calgranulin A and chaperonin 10 as protein markers for endometrial carcinoma. Proteomics, 2005, 5, 1953-1966.	1.3	33
4	Comparison of alternative analytical techniques for the characterisation of the human serum proteome in HUPO Plasma Proteome Project. Proteomics, 2005, 5, 3423-3441.	1.3	104
5	A novel four-dimensional strategy combining protein and peptide separation methods enables detection of low-abundance proteins in human plasma and serum proteomes. Proteomics, 2005, 5, 3329-3342.	1.3	128
6	Efficient prefractionation of low-abundance proteins in human plasma and construction of a two-dimensional map. Proteomics, 2005, 5, 3386-3396.	1.3	121
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8	An evaluation, comparison, and accurate benchmarking of several publicly available MS/MS search algorithms: Sensitivity and specificity analysis. Proteomics, 2005, 5, 3475-3490.	1.3	332
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22	Efficient prefractionation of low-abundance proteins in human plasma and construction of a two-dimensional map. , 2006, , 201-219.		0

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