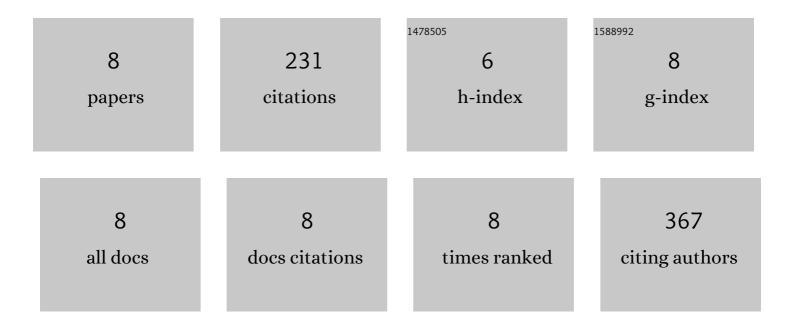
Klaus-Peter Pleissner

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Iterative data analysis is the key for exhaustive analysis of peptide mass fingerprints from proteins separated by two-dimensional electrophoresis. Journal of the American Society for Mass Spectrometry, 2003, 14, 943-956.	2.8	73
2	Identification of candidate antigens for serologic detection ofHelicobacter pylori-infected patients with gastric carcinoma. International Journal of Cancer, 2004, 108, 456-463.	5.1	51
3	Webâ€accessible proteome databases for microbial research. Proteomics, 2004, 4, 1305-1313.	2.2	41
4	Helicobacter pylori proteomics by 2-DE/MS, 1-DE-LC/MS and functional data mining. Proteomics, 2010, 10, 182-193.	2.2	35
5	A European Pathogenic Microorganism Proteome Database: Construction and Maintenance. Comparative and Functional Genomics, 2002, 3, 97-100.	2.0	12
6	Assessment of protein spot components applying correspondence analysis for peptide mass fingerprint data. Proteomics, 2004, 4, 2982-2986.	2.2	9
7	Short Communication– Presentation of differentially regulated proteins within a web-accessible proteome database system of microorganisms. Proteomics, 2004, 4, 2987-2990.	2.2	5
8	Assembling proteomics data as a prerequisite for the analysis of large scale experiments. Chemistry Central Journal, 2009, 3, 2.	2.6	5