

# Klaus-Peter Pleissner

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11911164/publications.pdf>

Version: 2024-02-01

8  
papers

231  
citations

1478505  
6  
h-index

1588992  
8  
g-index

8  
all docs

8  
docs citations

8  
times ranked

367  
citing authors

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