

Peter Schulz-Knappe

List of Publications by Year in descending order

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Version: 2024-02-01

51
papers

4,389
citations

230014

27
h-index

232693

48
g-index

52
all docs

52
docs citations

52
times ranked

4869
citing authors

#	ARTICLE	IF	CITATIONS
1	LEAP-1, a novel highly disulfide-bonded human peptide, exhibits antimicrobial activity. FEBS Letters, 2000, 480, 147-150.	1.3	1,108
2	HUPO Plasma Proteome Project specimen collection and handling: Towards the standardization of parameters for plasma proteome samples. Proteomics, 2005, 5, 3262-3277.	1.3	515
3	hBD-1: a novel β -defensin from human plasma. FEBS Letters, 1995, 368, 331-335.	1.3	502
4	Peptidomic analysis of human blood specimens: Comparison between plasma specimens and serum by differential peptide display. Proteomics, 2005, 5, 3414-3422.	1.3	265
5	Composition of the peptide fraction in human blood plasma: database of circulating human peptides. Biomedical Applications, 1999, 726, 25-35.	1.7	171
6	Peptidomics technologies for human body fluids. Trends in Biotechnology, 2001, 19, 55-60.	4.9	166
7	Peptidomics The Comprehensive Analysis of Peptides in Complex Biological Mixtures. Combinatorial Chemistry and High Throughput Screening, 2012, 4, 207-217.	0.6	166
8	Isolation and biochemical characterization of LEAP-2, a novel blood peptide expressed in the liver. Protein Science, 2003, 12, 143-152.	3.1	161
9	Peptidomics technologies for human body fluids. Trends in Biotechnology, 2001, 19, S55-S60.	4.9	156
10	Peptides in body fluids and tissues as markers of disease. Expert Review of Molecular Diagnostics, 2005, 5, 145-157.	1.5	89
11	The circulating bioactive form of human guanylin is a high molecular weight peptide (10.3 kDa). FEBS Letters, 1993, 318, 205-209.	1.3	85
12	A functional annotation of subproteomes in human plasma. Proteomics, 2005, 5, 3506-3519.	1.3	82
13	Historical perspective of peptidomics. EuPA Open Proteomics, 2014, 3, 171-182.	2.5	77
14	Peptide bank generated by large-scale preparation of circulating human peptides. Journal of Chromatography A, 1997, 776, 125-132.	1.8	72
15	Liquid chromatography and electrospray mass spectrometric mapping of peptides from human plasma filtrate. Journal of the American Society for Mass Spectrometry, 1999, 10, 45-54.	1.2	70
16	The Peptidomics Concept. Combinatorial Chemistry and High Throughput Screening, 2005, 8, 697-704.	0.6	62
17	GCAP-II: Isolation and characterization of the circulating form of human uroguanylin. FEBS Letters, 1995, 374, 34-38.	1.3	60
18	Human hemofiltrate as a source of circulating bioactive peptides: Determination of amino acids, peptides and proteins. Biomedical Chromatography, 1994, 8, 90-94.	0.8	50

#	ARTICLE	IF	CITATIONS
19	Prerequisites for Peptidomic Analysis of Blood Samples: I. Evaluation of Blood Specimen Qualities and Determination of Technical Performance Characteristics. <i>Combinatorial Chemistry and High Throughput Screening</i> , 2005, 8, 725-733.	0.6	42
20	Expression Profiling of Breast Cancer Cells by Differential Peptide Display. <i>Breast Cancer Research and Treatment</i> , 2003, 79, 83-93.	1.1	37
21	Peptidomic analysis of breast cancer reveals a putative surrogate marker for estrogen receptor-negative carcinomas. <i>Laboratory Investigation</i> , 2006, 86, 246-253.	1.7	37
22	Serum autoantibodies for discovery of prostate cancer specific biomarkers. <i>Prostate</i> , 2012, 72, 427-436.	1.2	33
23	Matrix-assisted laser desorption/ionisation mass spectrometry guided purification of human guanylin from blood ultrafiltrate. <i>Journal of Chromatography A</i> , 1997, 776, 139-145.	1.8	32
24	Effects of various phosphodiesterase-inhibitors, forskolin, and sodium nitroprusside on porcine detrusor smooth muscle tonic responses to muscarinergic stimulation and cyclic nucleotide levels in vitro. , 1996, 15, 59-70.		31
25	Porcine detrusor cyclic nucleotide phosphodiesterase isoenzymes: Characterization and functional effects of various phosphodiesterase inhibitors in vitro. <i>Urology</i> , 1995, 45, 893-901.	0.5	29
26	Correlation-associated peptide networks of human cerebrospinal fluid. <i>Proteomics</i> , 2005, 5, 2789-2798.	1.3	29
27	High-resolution peptide mapping of cerebrospinal fluid: a novel concept for diagnosis and research in central nervous system diseases. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2002, 782, 353-361.	1.2	28
28	In vivo profiling of DPP4 inhibitors reveals alterations in collagen metabolism and accumulation of an amyloid peptide in rat plasma. <i>Biochemical Pharmacology</i> , 2009, 77, 228-237.	2.0	27
29	Posttranslationally Processed Forms of the Human Chemokine HCC-1. <i>Biochemistry</i> , 2000, 39, 10799-10805.	1.2	24
30	Structural and Functional Characterization of Vitronectin-Derived RGD-Containing Peptides from Human Hemofiltrate. <i>FEBS Journal</i> , 1996, 241, 557-563.	0.2	22
31	Screening for disulfide-rich peptides in biological sources by carboxyamidomethylation in combination with differential matrix-assisted laser desorption/ionization time-of-flight mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2001, 15, 1586-1592.	0.7	16
32	The RA-MAP Consortium: a working model for academia industry collaboration. <i>Nature Reviews Rheumatology</i> , 2018, 14, 53-60.	3.5	15
33	Comparison of pre-processing methods for multiplex bead-based immunoassays. <i>BMC Genomics</i> , 2016, 17, 601.	1.2	13
34	Sequential high-content profiling of the IgG-autoantibody repertoire reveals novel antigens in rheumatoid arthritis. <i>Arthritis Research and Therapy</i> , 2016, 18, 235.	1.6	13
35	Serum Autoantibodies in Chronic Prostate Inflammation in Prostate Cancer Patients. <i>PLoS ONE</i> , 2016, 11, e0147739.	1.1	13
36	Characterization of natural posttranslationally processed peptides from human blood: A new tool in the systematic investigation of native peptides. , 1993, , 553-557.		11

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37	Identification of Peptide Tumor Markers in a Tumor Graft Model in Immunodeficient Mice. Combinatorial Chemistry and High Throughput Screening, 2005, 8, 783-788.	0.6	10
38	Metrological sharp shooting for plasma proteins and peptides: The need for reference materials for accurate measurements in clinical proteomics and <i>in vitro</i> diagnostics to generate reliable results. Proteomics - Clinical Applications, 2007, 1, 1016-1035.	0.8	10
39	High diagnostic accuracy of histone H4-IgG autoantibodies in systemic lupus erythematosus. Rheumatology, 2018, 57, 533-537.	0.9	10
40	Safe cosmetics without animal testing? Contributions of the EU Project Sens-it-iv. Journal Fur Verbraucherschutz Und Lebensmittelsicherheit, 2009, 4, 41-48.	0.5	8
41	Label-free microarray-based detection of autoantibodies in human serum. Journal of Immunological Methods, 2018, 459, 44-49.	0.6	7
42	Comprehensive Longitudinal Surveillance of the IgG Autoantibody Repertoire in Established Systemic Lupus Erythematosus. Arthritis and Rheumatology, 2019, 71, 736-743.	2.9	7
43	Proteomic analysis to define predictors of treatment response to adalimumab or methotrexate in rheumatoid arthritis patients. Pharmacogenomics Journal, 2020, 20, 516-523.	0.9	6
44	High-performance liquid chromatographic determination of sulfated peptides in human hemofiltrate using a radioactivity monitor. Journal of Chromatography A, 1995, 691, 255-261.	1.8	5
45	Specific determination of tyrosine-phosphorylated proteins and peptides by differential iodination. Journal of Chromatography A, 1996, 743, 273-282.	1.8	4
46	RA-MAP, molecular immunological landscapes in early rheumatoid arthritis and healthy vaccine recipients. Scientific Data, 2022, 9, 196.	2.4	4
47	Peptidomic analysis of human peripheral monocytes persistently infected by Chlamydia trachomatis. Medical Microbiology and Immunology, 2007, 196, 103-114.	2.6	3
48	Profiling of IgG antibodies targeting unmodified and corresponding citrullinated autoantigens in a multicenter national cohort of early arthritis in Germany. Arthritis Research and Therapy, 2020, 22, 167.	1.6	3
49	Chapter 7 Clinical peptidomics: peptide-biomarker discovery in blood. Comprehensive Analytical Chemistry, 2005, 46, 385-409.	0.7	1
50	HUPO Plasma Proteome Project specimen collection and handling: Towards the standardization of parameters for plasma proteome samples. , 2006, , 63-89.		0
51	Peptidomics. , 0, , 91-104.		0