Sonja Hess

List of Publications by Year in descending order

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172457 110387 4,240 67 29 64 h-index citations g-index papers 67 67 67 8970 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	PD-L1 ⁺ neutrophils contribute to injury-induced infection susceptibility. Science Advances, 2021, 7, .	10.3	24
2	Resistance to Pyrrolobenzodiazepine Dimers Is Associated with SLFN11 Downregulation and Can Be Reversed through Inhibition of ATR. Molecular Cancer Therapeutics, 2021, 20, 541-552.	4.1	18
3	Characterization of Citrullination Sites in Neutrophils and Mast Cells Activated by Ionomycin via Integration of Mass Spectrometry and Machine Learning. Journal of Proteome Research, 2021, 20, 3150-3164.	3.7	9
4	Metabolomic Profiling of Human Urine Samples Using LC-TIMS-QTOF Mass Spectrometry. Journal of the American Society for Mass Spectrometry, 2021, 32, 2072-2080.	2.8	9
5	Staphylococcus aureus drives expansion of low-density neutrophils in diabetic mice. Journal of Clinical Investigation, 2019, 129, 2133-2144.	8.2	30
6	Bioorthogonal Noncanonical Amino Acid Tagging (BONCAT) Enables Time-Resolved Analysis of Protein Synthesis in Native Plant Tissue. Plant Physiology, 2017, 173, 1543-1553.	4.8	43
7	Middle-down electron capture dissociation and electron transfer dissociation for histone analysis. Journal of Analytical Science and Technology, 2015, 6, .	2.1	3
8	High resolution data-independent acquisition with electron transfer dissociation mass spectrometry: Multiplexed analysis of post-translationally modified proteins. International Journal of Mass Spectrometry, 2015, 390, 155-162.	1.5	2
9	Recent proteomic advances in developmental, regeneration, and cancer governing signaling pathways. Proteomics, 2015, 15, 1014-1025.	2.2	8
10	Recent advances in defining the ubiquitylome. Expert Review of Proteomics, 2014, 11, 477-490.	3.0	13
11	The topâ€down, middleâ€down, and bottomâ€up mass spectrometry approaches for characterization of histone variants and their postâ€translational modifications. Proteomics, 2014, 14, 489-497.	2.2	122
12	The emerging field of chemo―and pharmacoproteomics. Proteomics - Clinical Applications, 2013, 7, 171-180.	1.6	15
13	Cand1 Promotes Assembly of New SCF Complexes through Dynamic Exchange of F Box Proteins. Cell, 2013, 153, 206-215.	28.9	228
14	Evaluation and Optimization of Mass Spectrometric Settings during Data-dependent Acquisition Mode: Focus on LTQ-Orbitrap Mass Analyzers. Journal of Proteome Research, 2013, 12, 3071-3086.	3.7	148
15	Comprehensive Profiling of N-Linked Glycosylation Sites in HeLa Cells Using Hydrazide Enrichment. Journal of Proteome Research, 2013, 12, 248-259.	3.7	21
16	Data-Dependent Middle-Down Nano-Liquid Chromatography–Electron Capture Dissociation-Tandem Mass Spectrometry: An Application for the Analysis of Unfractionated Histones. Analytical Chemistry, 2013, 85, 3501-3507.	6. 5	35
17	Structural basis for high-affinity binding of LEDGF PWWP to mononucleosomes. Nucleic Acids Research, 2013, 41, 3924-3936.	14.5	182
18	Activation of p107 by Fibroblast Growth Factor, Which Is Essential for Chondrocyte Cell Cycle Exit, Is Mediated by the Protein Phosphatase $2A/B55\hat{l}\pm Holoenzyme$. Molecular and Cellular Biology, 2013, 33, 330-3342.	2.3	26

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19	Mechanism of an ATP-independent Protein Disaggregase. Journal of Biological Chemistry, 2013, 288, 13420-13430.	3.4	17
20	Electron Capture Dissociation of Hydrogen-Deficient Peptide Radical Cations. Journal of the American Society for Mass Spectrometry, 2012, 23, 1729-1740.	2.8	6
21	Dopaminergic modulation of the hippocampal neuropil proteome identified by bioorthogonal noncanonical amino acid tagging (<scp>BONCAT</scp>). Proteomics, 2012, 12, 2464-2476.	2.2	58
22	Characterization of the <i>Mycobacterium tuberculosis</i> Proteome by Liquid Chromatography Mass Spectrometry-based Proteomics Techniques: A Comprehensive Resource for Tuberculosis Research. Journal of Proteome Research, 2012, 11, 119-130.	3.7	61
23	A Universal HPLC-MS Method to Determine the Stereochemistry of Common and Unusual Amino Acids. Methods in Molecular Biology, 2012, 828, 63-75.	0.9	5
24	Effect of mass spectrometric parameters on peptide and protein identification rates for shotgun proteomic experiments on an LTQâ€orbitrap mass analyzer. Proteomics, 2012, 12, 21-31.	2.2	50
25	Click Chemistry Facilitates Formation of Reporter Ions and Simplified Synthesis of Amine-Reactive Multiplexed Isobaric Tags for Protein Quantification. Journal of the American Chemical Society, 2012, 134, 2672-2680.	13.7	30
26	Designer Reagents for Mass Spectrometry-Based Proteomics: Clickable Cross-Linkers for Elucidation of Protein Structures and Interactions. Analytical Chemistry, 2012, 84, 2662-2669.	6.5	41
27	Broad activation of the ubiquitin–proteasome system by Parkin is critical for mitophagy. Human Molecular Genetics, 2011, 20, 1726-1737.	2.9	851
28	Peptide Fragmentation by Corona Discharge Induced Electrochemical Ionization. Journal of the American Society for Mass Spectrometry, 2010, 21, 2051-2061.	2.8	7
29	A combined topâ€down and bottomâ€up MS approach for the characterization of hemoglobin variants in Rhesus monkeys. Proteomics, 2010, 10, 3657-3668.	2.2	7
30	Mass Spectrometry in the Elucidation of the Glycoproteome of Bacterial Pathogens. Current Proteomics, 2010, 7, 57-81.	0.3	11
31	Preparation of stable isotope-labeled peripheral cannabinoid receptor CB2 by bacterial fermentation. Protein Expression and Purification, 2010, 70, 236-247.	1.3	30
32	An Allosteric Mechanism for Inhibiting HIV-1 Integrase with a Small Molecule. Molecular Pharmacology, 2009, 76, 824-832.	2.3	48
33	A corona discharge initiated electrochemical electrospray ionization technique. Journal of the American Society for Mass Spectrometry, 2009, 20, 1988-1996.	2.8	16
34	Analysis of a G protein-coupled receptor for neurotensin by liquid chromatography–electrospray ionization–mass spectrometry. Analytical Biochemistry, 2008, 376, 13-24.	2.4	12
35	Adipose proteome analysis: focus on mediators of insulin resistance. Expert Review of Proteomics, 2008, 5, 827-839.	3.0	25
36	The <i>sim</i> Operon Facilitates the Transport and Metabolism of Sucrose Isomers in <i>Lactobacillus casei</i> ATCC 334. Journal of Bacteriology, 2008, 190, 3362-3373.	2.2	11

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37	Protein disulfide isomerase acts as an injury response signal that enhances fibrin generation via tissue factor activation. Journal of Clinical Investigation, 2008, 118, 1110-22.	8.2	251
38	Mycobacterium tuberculosis produces pili during human infection. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 5145-5150.	7.1	164
39	High Resolution Footprinting of the Hepatitis C Virus Polymerase NS5B in Complex with RNA. Journal of Biological Chemistry, 2007, 282, 16907-16916.	3.4	20
40	Cellular Expression and Crystal Structure of the Murine Cytomegalovirus Major Histocompatibility Complex Class I-like Glycoprotein, m153. Journal of Biological Chemistry, 2007, 282, 35247-35258.	3.4	22
41	Amended Description of the Genes for Synthesis of Actinomyces naeslundii T14V Type 1 Fimbriae and Associated Adhesin. Infection and Immunity, 2007, 75, 4181-4185.	2.2	14
42	Endogenous retroviral insertion in Cryge in the mouse No3 cataract mutant. Genomics, 2007, 89, 512-520.	2.9	8
43	Solid-Phase Synthesis of Thermolytic DNA Oligonucleotides Functionalized with a Single 4-Hydroxy-1-butyl or 4-Phosphato-/Thiophosphato-1-butyl Thiophosphate Protecting Groupâ€. Journal of Organic Chemistry, 2007, 72, 805-815.	3.2	22
44	Genetic Requirements for Growth of Escherichia coli K12 on Methyl-α-D-glucopyranoside and the Five α-D-Glucosyl-D-fructose Isomers of Sucrose. Journal of Biological Chemistry, 2006, 281, 17900-17908.	3.4	19
45	Modulation of Replication Protein A Function by Its Hyperphosphorylation-induced Conformational Change Involving DNA Binding Domain B. Journal of Biological Chemistry, 2005, 280, 32775-32783.	3.4	55
46	Mass Spectrometric Analysis of the HIV-1 Integrase-Pyridoxal 5′-Phosphate Complex Reveals a New Binding Site for a Nucleotide Inhibitor. Journal of Biological Chemistry, 2005, 280, 7949-7955.	3.4	27
47	Thermolytic CpG-containing DNA oligonucleotides as potential immunotherapeutic prodrugs. Nucleic Acids Research, 2005, 33, 3550-3560.	14.5	30
48	Quantitative Proteomic Analysis of the Secretory Proteins from Rat Adipose Cells Using a 2D Liquid Chromatographyâ^MS/MS Approach. Journal of Proteome Research, 2005, 4, 570-577.	3.7	115
49	Genes malh and pagl of Clostridium acetobutylicum ATCC 824 Encode NAD+- and Mn2+-dependent Phospho-α-glucosidase(s). Journal of Biological Chemistry, 2004, 279, 1553-1561.	3.4	26
50	Identification of an inhibitor-binding site to HIV-1 integrase with affinity acetylation and mass spectrometry. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 6894-6899.	7.1	42
51	Chirality determination of unusual amino acids using precolumn derivatization and liquid chromatography–electrospray ionization mass spectrometry. Journal of Chromatography A, 2004, 1035, 211-219.	3.7	74
52	Neamphamide A, a New HIV-Inhibitory Depsipeptide from the Papua New Guinea Marine Sponge Neamphius huxleyi. Journal of Natural Products, 2004, 67, 1407-1411.	3.0	120
53	Isolation of Homodolastatin 16, a New Cyclic Depsipeptide from a Kenyan Collection of Lyngbyamajuscula. Journal of Natural Products, 2003, 66, 712-715.	3.0	81
54	Lignans Isolated from Valerian:  Identification and Characterization of a New Olivil Derivative with Partial Agonistic Activity at A1 Adenosine Receptors. Journal of Natural Products, 2002, 65, 1479-1485.	3.0	143

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55	The molecular structure of spider dragline silk: Folding and orientation of the protein backbone. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 10266-10271.	7.1	465
56	Identification and Characterization of Hydrophobic Escherichia coli Virulence Proteins by Liquid Chromatography–Electrospray Ionization Mass Spectrometry. Analytical Biochemistry, 2002, 302, 123-130.	2.4	13
57	Structure of cyclized green fluorescent protein. Acta Crystallographica Section D: Biological Crystallography, 2002, 58, 1400-1406.	2.5	7
58	Acid hydrolysis of silk fibroins and determination of the enrichment of isotopically labeled amino acids using precolumn derivatization and high-performance liquid chromatography–electrospray ionization–mass spectrometry. Analytical Biochemistry, 2002, 311, 19-26.	2.4	35
59	Plant annexins form calcium-independent oligomers in solution. Protein Science, 2002, 11, 2033-2040.	7.6	19
60	Recent advances in adenosine receptor antagonist research. Expert Opinion on Therapeutic Patents, 2001, 11, 1533-1561.	5.0	46
61	Solid-state NMR data support a helix-loop-helix structural model for the N-terminal half of HIV-1 rev in fibrillar form. Journal of Molecular Biology, 2001, 313, 845-859.	4.2	47
62	Effect of an adenosine A1 receptor agonist and a novel pyrimidoindole on membrane properties and neurotransmitter release in rat cortical and hippocampal neurons. Neurochemistry International, 2001, 38, 391-398.	3.8	8
63	Rapid and sensitive LC separation of new impurities in trimethoprim. Journal of Pharmaceutical and Biomedical Analysis, 2001, 25, 531-538.	2.8	9
64	Synthesis and immunological activity of water-soluble Thalidomide prodrugs. Bioorganic and Medicinal Chemistry, 2001, 9, 1279-1291.	3.0	38
65	Profiling indomethacin impurities using high-performance liquid chromatography and nuclear magnetic resonance. European Journal of Pharmaceutical Sciences, 2001, 14, 301-311.	4.0	22
66	In vivo mechanism-based inactivation of S-adenosylmethionine decarboxylases from Escherichia coli, Salmonella typhimurium, and Saccharomycescerevisiae. Proceedings of the National Academy of Sciences of the United States of America, 2001, 98, 10578-10583.	7.1	27
67	7-Deazaadenines Bearing Polar Substituents:  Structureâ^'Activity Relationships of New A1 and A3 Adenosine Receptor Antagonists. Journal of Medicinal Chemistry, 2000, 43, 4636-4646.	6.4	49