

# Christine Hoogland

## List of Publications by Year in descending order

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39  
papers

5,658  
citations

304743  
22  
h-index

315739  
38  
g-index

43  
all docs

43  
docs citations

43  
times ranked

8545  
citing authors

#	ARTICLE	IF	CITATIONS
1	Integration of Two In-depth Quantitative Proteomics Approaches Determines the Kallikrein-related Peptidase 7 (KLK7) Degradome in Ovarian Cancer Cell Secretome. <i>Molecular and Cellular Proteomics</i> , 2019, 18, 818a-836.	3.8	16
2	Site-specific glycosylation of the Newcastle disease virus haemagglutinin-neuraminidase. <i>Glycoconjugate Journal</i> , 2017, 34, 181-197.	2.7	6
3	Mass-spectrometry data for <i>Rhizoctonia solani</i> proteins produced during infection of wheat and vegetative growth. <i>Data in Brief</i> , 2016, 8, 267-271.	1.0	5
4	Proteomic Analysis of <i>Rhizoctonia solani</i> Identifies Infection-specific, Redox Associated Proteins and Insight into Adaptation to Different Plant Hosts. <i>Molecular and Cellular Proteomics</i> , 2016, 15, 1188-1203.	3.8	37
5	EasyProt – An easy-to-use graphical platform for proteomics data analysis. <i>Journal of Proteomics</i> , 2013, 79, 146-160.	2.4	57
6	Management and dissemination of MS proteomic data with PROTEOMICSdb: Example of a quantitative comparison between methods of protein extraction. <i>Proteomics</i> , 2013, 13, 1457-1466.	2.2	25
7	Human Hemolysate Glycated Proteome. <i>Analytical Chemistry</i> , 2011, 83, 5673-5680.	6.5	15
8	Quantitative Analysis of Human Cerebrospinal Fluid Proteins Using a Combination of Cysteine Tagging and Amine-Reactive Isobaric Labeling. <i>Journal of Proteome Research</i> , 2011, 10, 249-258.	3.7	37
9	Combining low- and high-energy tandem mass spectra for optimized peptide quantification with isobaric tags. <i>Journal of Proteomics</i> , 2010, 73, 769-777.	2.4	99
10	The Gel Electrophoresis Markup Language (GelML) from the Proteomics Standards Initiative. <i>Proteomics</i> , 2010, 10, 3073-3081.	2.2	19
11	Guidelines for reporting the use of gel image informatics in proteomics. <i>Nature Biotechnology</i> , 2010, 28, 655-656.	17.5	22
12	A suite of tools to analyse and publish 2D-DE data. <i>Proteomics</i> , 2008, 8, 4907-4909.	2.2	3
13	The World-2DPAGE Constellation to promote and publish gel-based proteomics data through the ExPASy server. <i>Journal of Proteomics</i> , 2008, 71, 245-248.	2.4	47
14	MIAPEGelDB, a web-based submission tool and public repository for MIAPE gel electrophoresis documents. <i>Journal of Proteomics</i> , 2008, 71, 249-251.	2.4	19
15	Guidelines for reporting the use of gel electrophoresis in proteomics. <i>Nature Biotechnology</i> , 2008, 26, 863-864.	17.5	61
16	Using bioinformatic resources in the proteomic analysis of biological fluids. <i>Proteomics - Clinical Applications</i> , 2007, 1, 900-915.	1.6	2
17	2D PAGE Databases for Proteins in Human Body Fluids. , 2007, , 137-146.		1
18	SPS' Digest: The Swiss Proteomics Society selection of proteomics articles. <i>Proteomics</i> , 2005, 5, 3045-3047.	2.2	0

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19	Correlation of proteomic and transcriptomic profiles of <i>Staphylococcus aureus</i> during the post-exponential phase of growth. <i>Journal of Microbiological Methods</i> , 2005, 60, 247-257.	1.6	59
20	SWISS-2DPAGE, ten years later. <i>Proteomics</i> , 2004, 4, 2352-2356.	2.2	104
21	The molecular scanner: concept and developments. <i>Current Opinion in Biotechnology</i> , 2004, 15, 17-23.	6.6	28
22	The Make 2D-DB II package: Conversion of federated two-dimensional gel electrophoresis databases into a relational format and interconnection of distributed databases. <i>Proteomics</i> , 2003, 3, 1441-1444.	2.2	34
23	ExPASy: the proteomics server for in-depth protein knowledge and analysis. <i>Nucleic Acids Research</i> , 2003, 31, 3784-3788.	14.5	4,128
24	Hydrogen/deuterium exchange for higher specificity of protein identification by peptide mass fingerprinting. <i>Rapid Communications in Mass Spectrometry</i> , 2002, 16, 616-626.	1.5	18
25	The mouse SWISS-2D PAGE database: a tool for proteomics study of diabetes and obesity. <i>Proteomics</i> , 2001, 1, 136-163.	2.2	155
26	The establishment of a human liver nuclei two-dimensional electrophoresis reference map. <i>Electrophoresis</i> , 2000, 21, 3483-3487.	2.4	40
27	Constructing a 2-D Database for the World Wide Web. , 1999, 112, 411-416.		6
28	The SWISS-2DPAGE database: what has changed during the last year. <i>Nucleic Acids Research</i> , 1999, 27, 289-291.	14.5	18
29	Improving protein identification from peptide mass fingerprinting through a parameterized multi-level scoring algorithm and an optimized peak detection. <i>Electrophoresis</i> , 1999, 20, 3535-3550.	2.4	140
30	Two-dimensional electrophoresis resources available from ExPASy. <i>Electrophoresis</i> , 1999, 20, 3568-3571.	2.4	47
31	A Molecular Scanner To Automate Proteomic Research and To Display Proteome Images. <i>Analytical Chemistry</i> , 1999, 71, 4981-4988.	6.5	127
32	'98Escherichia coli SWISS-2DPAGE database update. <i>Electrophoresis</i> , 1998, 19, 1960-1971.	2.4	90
33	Current status of the SWISS-2DPAGE database. <i>Nucleic Acids Research</i> , 1998, 26, 332-333.	14.5	23
34	Title is missing!. <i>Genetica</i> , 1997, 100, 161-166.	1.1	31
35	Make2ddb: A simple package to set up a two-dimensional electrophoresis database for the World Wide Web. <i>Electrophoresis</i> , 1997, 18, 2755-2758.	2.4	24
36	Maintenance of transposable element copy number in natural populations of <i>Drosophila melanogaster</i> and <i>D. simulans</i> . <i>Contemporary Issues in Genetics and Evolution</i> , 1997, , 161-166.	0.9	1

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37	Chromosomal Distribution of Transposable Elements in <i>Drosophila melanogaster</i> Test of the Ectopic Recombination Model for Maintenance of Insertion Site Number. <i>Genetics</i> , 1996, 144, 197-204.	2.9	46
38	Chromosomal distribution and population dynamics of the 412 retrotransposon in a natural population of <i>Drosophila melanogaster</i> . <i>Chromosoma</i> , 1995, 103, 693-699.	2.2	1
39	Computational Methods and Bioinformatic Tools. , 0, , 769-904.		0