

# Petra S Langendijk-Genevaux

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/173875/publications.pdf>

Version: 2024-02-01

23  
papers

8,585  
citations

393982

19  
h-index

676716

22  
g-index

25  
all docs

25  
docs citations

25  
times ranked

14070  
citing authors

#	ARTICLE	IF	CITATIONS
1	ScanProsite: detection of PROSITE signature matches and ProRule-associated functional and structural residues in proteins. <i>Nucleic Acids Research</i> , 2006, 34, W362-W365.	6.5	1,428
2	The Universal Protein Resource (UniProt) in 2010. <i>Nucleic Acids Research</i> , 2010, 38, D142-D148.	6.5	1,131
3	Quantitative fluorescence in situ hybridization of <i>Bifidobacterium</i> spp. with genus-specific 16S rRNA-targeted probes and its application in fecal samples. <i>Applied and Environmental Microbiology</i> , 1995, 61, 3069-3075.	1.4	882
4	The Universal Protein Resource (UniProt). <i>Nucleic Acids Research</i> , 2007, 36, D190-D195.	6.5	852
5	The PROSITE database. <i>Nucleic Acids Research</i> , 2006, 34, D227-D230.	6.5	800
6	PROSITE, a protein domain database for functional characterization and annotation. <i>Nucleic Acids Research</i> , 2010, 38, D161-D166.	6.5	744
7	The Universal Protein Resource (UniProt) 2009. <i>Nucleic Acids Research</i> , 2009, 37, D169-D174.	6.5	548
8	The Universal Protein Resource (UniProt). <i>Nucleic Acids Research</i> , 2007, 35, D193-D197.	6.5	488
9	New developments in the InterPro database. <i>Nucleic Acids Research</i> , 2007, 35, D224-D228.	6.5	444
10	The 20 years of PROSITE. <i>Nucleic Acids Research</i> , 2007, 36, D245-D249.	6.5	441
11	Recent improvements to the PROSITE database. <i>Nucleic Acids Research</i> , 2004, 32, 134D-137.	6.5	350
12	In vivo analysis of the overlapping functions of DnaK and trigger factor. <i>EMBO Reports</i> , 2004, 5, 195-200.	2.0	163
13	ProRule: a new database containing functional and structural information on PROSITE profiles. <i>Bioinformatics</i> , 2005, 21, 4060-4066.	1.8	73
14	Sulfate-reducing bacteria in association with human periodontitis. <i>Journal of Clinical Periodontology</i> , 2000, 27, 943-950.	2.3	52
15	Archaeal $\hat{2}$ -CASP ribonucleases of the aCPSF1 family are orthologs of the eukaryal CPSF-73 factor. <i>Nucleic Acids Research</i> , 2013, 41, 1091-1103.	6.5	42
16	Sulfate-reducing bacteria in periodontal pockets and in healthy oral sites. <i>Journal of Clinical Periodontology</i> , 1999, 26, 596-599.	2.3	36
17	From protein sequences to 3D-structures and beyond: the example of the UniProt Knowledgebase. <i>Cellular and Molecular Life Sciences</i> , 2010, 67, 1049-1064.	2.4	33
18	Universal RNA-degrading enzymes in Archaea: Prevalence, activities and functions of $\hat{2}$ -CASP ribonucleases. <i>Biochimie</i> , 2015, 118, 278-285.	1.3	25

#	ARTICLE	IF	CITATIONS
19	Sulfate-reducing bacteria in relation with other potential periodontal pathogens. <i>Journal of Clinical Periodontology</i> , 2001, 28, 1151-1157.	2.3	24
20	RNA processing machineries in Archaea: the 5'→3' exoribonuclease aRNase J of the Î²-CASP family is engaged specifically with the helicase ASH-Ski2 and the 3'→5' exoribonucleolytic RNA exosome machinery. <i>Nucleic Acids Research</i> , 2020, 48, 3832-3847.	6.5	14
21	Decrease of Sulfate-reducing Bacteria after Initial Periodontal Treatment. <i>Journal of Dental Research</i> , 2001, 80, 1637-1642.	2.5	9
22	Phylogenetic Diversity of Lhr Proteins and Biochemical Activities of the Thermococcales aLhr2 DNA/RNA Helicase. <i>Biomolecules</i> , 2021, 11, 950.	1.8	4
23	Functionally and structurally relevant residues in PROSITE motif descriptors. , 2005, , .		0