

Petra S Langendijk-Genevaux

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/173875/petra-s-langendijk-genevaux-publications-by-year.pdf>

Version: 2024-04-25

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

22
papers

7,250
citations

18
h-index

25
g-index

25
ext. papers

7,965
ext. citations

13.7
avg, IF

6.49
L-index

#	Paper	IF	Citations
22	Phylogenetic Diversity of Lhr Proteins and Biochemical Activities of the Thermococcales aLhr2 DNA/RNA Helicase. <i>Biomolecules</i> , 2021 , 11,	5.9	1
21	RNA processing machineries in Archaea: the 5'3'5'4' exonuclease aRNase J of the ECASP family is engaged specifically with the helicase ASH-Ski2 and the 3'5'4' exonucleolytic RNA exosome machinery. <i>Nucleic Acids Research</i> , 2020 , 48, 3832-3847	20.1	7
20	Universal RNA-degrading enzymes in Archaea: Prevalence, activities and functions of ECASP ribonucleases. <i>Biochimie</i> , 2015 , 118, 278-85	4.6	17
19	Archaeal ECASP ribonucleases of the aCPSF1 family are orthologs of the eukaryal CPSF-73 factor. <i>Nucleic Acids Research</i> , 2013 , 41, 1091-103	20.1	35
18	PROSITE, a protein domain database for functional characterization and annotation. <i>Nucleic Acids Research</i> , 2010 , 38, D161-6	20.1	600
17	The Universal Protein Resource (UniProt) in 2010. <i>Nucleic Acids Research</i> , 2010 , 38, D142-8	20.1	1035
16	From protein sequences to 3D-structures and beyond: the example of the UniProt knowledgebase. <i>Cellular and Molecular Life Sciences</i> , 2010 , 67, 1049-64	10.3	25
15	The Universal Protein Resource (UniProt) 2009. <i>Nucleic Acids Research</i> , 2009 , 37, D169-74	20.1	509
14	The universal protein resource (UniProt). <i>Nucleic Acids Research</i> , 2008 , 36, D190-5	20.1	746
13	The 20 years of PROSITE. <i>Nucleic Acids Research</i> , 2008 , 36, D245-9	20.1	375
12	New developments in the InterPro database. <i>Nucleic Acids Research</i> , 2007 , 35, D224-8	20.1	397
11	The Universal Protein Resource (UniProt). <i>Nucleic Acids Research</i> , 2007 , 35, D193-7	20.1	437
10	ScanProsite: detection of PROSITE signature matches and ProRule-associated functional and structural residues in proteins. <i>Nucleic Acids Research</i> , 2006 , 34, W362-5	20.1	1055
9	The PROSITE database. <i>Nucleic Acids Research</i> , 2006 , 34, D227-30	20.1	649
8	ProRule: a new database containing functional and structural information on PROSITE profiles. <i>Bioinformatics</i> , 2005 , 21, 4060-6	7.2	54
7	In vivo analysis of the overlapping functions of DnaK and trigger factor. <i>EMBO Reports</i> , 2004 , 5, 195-200	6.5	140
6	Recent improvements to the PROSITE database. <i>Nucleic Acids Research</i> , 2004 , 32, D134-7	20.1	299

5	Sulfate-reducing bacteria in relation with other potential periodontal pathogens. <i>Journal of Clinical Periodontology</i> , 2001 , 28, 1151-7	7.7	21
4	Decrease of sulfate-reducing bacteria after initial periodontal treatment. <i>Journal of Dental Research</i> , 2001 , 80, 1637-42	8.1	8
3	Sulfate-reducing bacteria in association with human periodontitis. <i>Journal of Clinical Periodontology</i> , 2000 , 27, 943-50	7.7	41
2	Sulfate-reducing bacteria in periodontal pockets and in healthy oral sites. <i>Journal of Clinical Periodontology</i> , 1999 , 26, 596-9	7.7	28
1	Quantitative fluorescence in situ hybridization of Bifidobacterium spp. with genus-specific 16S rRNA-targeted probes and its application in fecal samples. <i>Applied and Environmental Microbiology</i> , 1995 , 61, 3069-75	4.8	770