

# Minia Antelo

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1371978/publications.pdf>

Version: 2024-02-01

11  
papers

246  
citations

1307594

7  
h-index

1281871

11  
g-index

11  
all docs

11  
docs citations

11  
times ranked

393  
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular reprogramming and phenotype switching in <i>Staphylococcus aureus</i> lead to high antibiotic persistence and affect therapy success. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	62
2	SppI Forms a Membrane Protein Complex with SppA and Inhibits Its Protease Activity in <i>Bacillus subtilis</i> . MSphere, 2020, 5, .	2.9	3
3	Functional association of the stress-responsive LiaH protein and the minimal TatAyCy protein translocase in <i>Bacillus subtilis</i> . Biochimica Et Biophysica Acta - Molecular Cell Research, 2020, 1867, 118719.	4.1	6
4	Membrane Modulation of Super-Secreting <i>Bacillus</i> Expressing the Major <i>Staphylococcus aureus</i> Antigen – A Mass-Spectrometry-Based Absolute Quantification Approach. Frontiers in Bioengineering and Biotechnology, 2020, 8, 143.	4.1	8
5	Ariadne's Thread in the Analytical Labyrinth of Membrane Proteins: Integration of Targeted and Shotgun Proteomics for Global Absolute Quantification of Membrane Proteins. Analytical Chemistry, 2019, 91, 11972-11980.	6.5	7
6	Identification and optimization of PrsA in <i>Bacillus subtilis</i> for improved yield of amylase. Microbial Cell Factories, 2019, 18, 158.	4.0	33
7	<i>Escherichia coli</i> Can Adapt Its Protein Translocation Machinery for Enhanced Periplasmic Recombinant Protein Production. Frontiers in Bioengineering and Biotechnology, 2019, 7, 465.	4.1	8
8	A retrospective cross-sectional quantitative molecular approach in biological samples from patients with syphilis. Microbial Pathogenesis, 2017, 104, 296-302.	2.9	10
9	Genome-scale analysis of the non-cultivable <i>Treponema pallidum</i> reveals extensive within-patient genetic variation. Nature Microbiology, 2017, 2, 16190.	13.3	81
10	In Silico Scrutiny of Genes Revealing Phylogenetic Congruence with Clinical Prevalence or Tropism Properties of <i>Chlamydia trachomatis</i> Strains. G3: Genes, Genomes, Genetics, 2015, 5, 9-19.	1.8	14
11	<i>Chlamydia trachomatis</i> In Vivo to In Vitro Transition Reveals Mechanisms of Phase Variation and Down-Regulation of Virulence Factors. PLoS ONE, 2015, 10, e0133420.	2.5	14