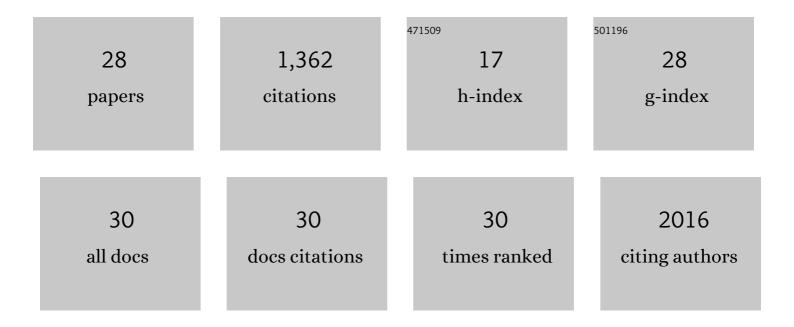
Susanne Engelmann

List of Publications by Year in descending order

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SUSANNE ENCELMANN

#	Article	IF	CITATIONS
1	Determining the bacterial cell biology of Planctomycetes. Nature Communications, 2017, 8, 14853.	12.8	175
2	A Proteomic View of an Important Human Pathogen – Towards the Quantification of the Entire Staphylococcus aureus Proteome. PLoS ONE, 2009, 4, e8176.	2.5	139
3	Proteomics uncovers extreme heterogeneity in the <i>Staphylococcus aureus</i> exoproteome due to genomic plasticity and variant gene regulation. Proteomics, 2010, 10, 1634-1644.	2.2	129
4	Impaired oxidative stress resistance ofBacillus subtilis sigBmutants and the role ofkatAandkatE. FEMS Microbiology Letters, 1996, 145, 63-69.	1.8	120
5	Staphylococcal serine protease–like proteins are pacemakers of allergic airway reactions to Staphylococcus aureus. Journal of Allergy and Clinical Immunology, 2017, 139, 492-500.e8.	2.9	118
6	Proteomic analysis of antioxidant strategies of <i>Staphylococcus aureus</i> : Diverse responses to different oxidants. Proteomics, 2008, 8, 3139-3153.	2.2	109
7	Global Analysis of the Staphylococcus aureus Response to Mupirocin. Antimicrobial Agents and Chemotherapy, 2012, 56, 787-804.	3.2	88
8	Oxidative stress triggers thiol oxidation in the glyceraldehyde-3-phosphate dehydrogenase of Staphylococcus aureus. Molecular Microbiology, 2004, 52, 133-140.	2.5	87
9	Secrets of the secretome in Staphylococcus aureus. International Journal of Medical Microbiology, 2014, 304, 133-141.	3.6	49
10	Specific serum IgG at diagnosis of Staphylococcus aureus bloodstream invasion is correlated with disease progression. Journal of Proteomics, 2015, 128, 1-7.	2.4	49
11	Costs of life - Dynamics of the protein inventory of Staphylococcus aureus during anaerobiosis. Scientific Reports, 2016, 6, 28172.	3.3	38
12	Aureolib — A Proteome Signature Library: Towards an Understanding of Staphylococcus aureus Pathophysiology. PLoS ONE, 2013, 8, e70669.	2.5	28
13	A nonâ€coding RNA from the intercellular adhesion (<i>ica</i>) locus of <i>Staphylococcus epidermidis</i> controls polysaccharide intercellular adhesion (PIA)â€mediated biofilm formation. Molecular Microbiology, 2019, 111, 1571-1591.	2.5	25
14	A Systematic Proteomic Analysis of Listeria monocytogenes House-keeping Protein Secretion Systems. Molecular and Cellular Proteomics, 2014, 13, 3063-3081.	3.8	23
15	Towards the characterization of the hidden world of small proteins in Staphylococcus aureus, a proteogenomics approach. PLoS Genetics, 2021, 17, e1009585.	3.5	23
16	Microarrayâ€based identification of human antibodies against <i>Staphylococcus aureus</i> antigens. Proteomics - Clinical Applications, 2015, 9, 1003-1011.	1.6	21
17	Human antibody responses against non-covalently cell wall-bound Staphylococcus aureus proteins. Scientific Reports, 2018, 8, 3234.	3.3	21
18	Adaptation of Staphylococcus aureus to Airway Environments in Patients With Cystic Fibrosis by Upregulation of Superoxide Dismutase M and Iron-Scavenging Proteins. Journal of Infectious Diseases, 2018, 217, 1453-1461.	4.0	20

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#	Article	IF	CITATIONS
19	Extracellular milieu grossly alters pathogen-specific immune response of mammary epithelial cells. BMC Veterinary Research, 2015, 11, 172.	1.9	18
20	smORFer: a modular algorithm to detect small ORFs in prokaryotes. Nucleic Acids Research, 2021, 49, e89-e89.	14.5	16
21	Global antibody response to Staphylococcus aureus live-cell vaccination. Scientific Reports, 2016, 6, 24754.	3.3	15
22	Hepatic Transcriptome Analysis Identifies Divergent Pathogen-Specific Targeting-Strategies to Modulate the Innate Immune System in Response to Intramammary Infection. Frontiers in Immunology, 2020, 11, 715.	4.8	15
23	Adaptive immune response to lipoproteins of Staphylococcus aureus in healthy subjects. Proteomics, 2016, 16, 2667-2677.	2.2	13
24	The hidden lipoproteome of Staphylococcus aureus. International Journal of Medical Microbiology, 2018, 308, 569-581.	3.6	7
25	Dinoroseobacter shibae Outer Membrane Vesicles Are Enriched for the Chromosome Dimer Resolution Site <i>dif</i> . MSystems, 2021, 6, .	3.8	7
26	Development and evaluation of a milk protein transcript depletion method for differential transcriptome analysis in mammary gland tissue. BMC Genomics, 2019, 20, 400.	2.8	4
27	Adaptation of Dinoroseobacter shibae to oxidative stress and the specific role of RirA. PLoS ONE, 2021, 16, e0248865.	2.5	2
28	Proteomic Signatures in Staphylococcus aureus. Methods in Molecular Biology, 2018, 1841, 113-130.	0.9	1