Maja Kosecka-Strojek

List of Publications by Year
in descending order

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26	266	1039406 9	1058022 14
papers	citations	h-index	g-index

27	27	27	402
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	May StaphylococcusÂlugdunensis Be an Etiological Factor of Chronic Maxillary Sinuses Infection?. International Journal of Molecular Sciences, 2022, 23, 6450.	1.8	2
2	Staphylococcal saoABC Operon Codes for a DNA-Binding Protein SaoC Implicated in the Response to Nutrient Deficit. International Journal of Molecular Sciences, 2022, 23, 6443.	1.8	1
3	First Case of Staphylococci Carrying Linezolid Resistance Genes from Laryngological Infections in Poland. Pathogens, 2021, 10, 335.	1.2	6
4	Epidemiology and Pathogenesis of Staphylococcus Bloodstream Infections in Humans: a Review. Polish Journal of Microbiology, 2021, 70, 13-23.	0.6	19
5	Effect of Staphylococcus aureus infection on the heat stress protein 70 (HSP70) level in chicken embryo tissues. Poultry Science, 2021, 100, 101119.	1.5	5
6	Human skin microbiota-friendly lysostaphin. International Journal of Biological Macromolecules, 2021, 183, 852-860.	3.6	7
7	Distribution and antibiotic-resistance of different Staphylococcus species identified by matrix assisted laser desorption ionization-time of flight mass spectrometry (MALDI-TOF MS) isolated from the oral cavity. Journal of Oral Microbiology, 2021, 13, 1983322.	1.2	13
8	Presence of egc-positive major clones ST 45, 30 and 22 among methicillin-resistant andÂmethicillin-susceptible oral Staphylococcus aureus strains. Scientific Reports, 2020, 10, 18889.	1.6	22
9	Identification of Clinically Relevant Streptococcus and Enterococcus Species Based on Biochemical Methods and 16S rRNA, sodA, tuf, rpoB, and recA Gene Sequencing. Pathogens, 2020, 9, 939.	1.2	4
10	Emergence of linezolid-resistant Staphylococcus epidermidis in the tertiary childrenâ€™s hospital in Cracow, Poland. European Journal of Clinical Microbiology and Infectious Diseases, 2020, 39, 1717-1725.	1.3	23
11	Coagulase-negative staphylococci (CoNS) as a significant etiological factor of laryngological infections: a review. Annals of Clinical Microbiology and Antimicrobials, 2020, 19, 26.	1.7	30
12	Preliminary evaluation of application of a 3-dimensional network structure of siloxanes Dergall preparation on chick embryo development and microbiological status of eggshells. Poultry Science, 2020, 99, 1581-1590.	1.5	2
13	Development and Validation of a Reference Data Set for Assigning Staphylococcus Species Based on Next-Generation Sequencing of the 16S-23S rRNA Region. Frontiers in Cellular and Infection Microbiology, 2019, 9, 278.	1.8	18
14	Development of a reference data set for assigning Streptococcus and Enterococcus species based on next generation sequencing of the 16Sâ€“23S rRNA region. Antimicrobial Resistance and Infection Control, 2019, 8, 178.	1.5	12
15	Thymol derivatives from the roots of Xerolekia speciosissima an endemic species of the pre-Alpine area. Phytochemistry Letters, 2019, 30, 235-237.	0.6	2
16	New Insight into Genotypic and Phenotypic Relatedness of Staphylococcus aureus Strains from Human Infections or Animal Reservoirs. Polish Journal of Microbiology, 2019, 68, 93-104.	0.6	3
17	Experimental Animal Models in Evaluation of Staphylococcal Pathogenicity. , 2018, , 265-279.		1
18	Staphylococcal Ecology and Epidemiology. , 2018, , 11-24.		8

#	Article	IF	Citations
19	Phage-Associated Virulence Determinants of Staphylococcus aureus. , 2018, , 173-183.		0
20	Characteristics of advanced methods used for typing bacterial isolates from mastitis with particular reference to Staphylococci. Polish Journal of Veterinary Sciences, 2018, 21, 229-239.	0.2	6
21	Daptomycin-resistant Staphylococcus pettenkoferi of human origin.. Acta Biochimica Polonica, 2016, 63, 297-301.	0.3	10
22	Identification of Secreted Exoproteome Fingerprints of Highly-Virulent and Non-Virulent Staphylococcus aureus Strains. Frontiers in Cellular and Infection Microbiology, 2016, 6, 51.	1.8	15
23	Multiple-locus variable-number tandem repeat fingerprinting as a method for rapid and cost-effective typing of animal-associated Staphylococcus aureus strains from lineages other than sequence type 398. Journal of Medical Microbiology, 2016, 65, 1494-1504.	0.7	7
24	Clonal Structure and Characterization of Staphylococcus aureus Strains from Invasive Infections in Paediatric Patients from South Poland: Association between Age, spa Types, Clonal Complexes, and Genetic Markers. PLoS ONE, 2016, 11, e0151937.	1.1	36
25	Species determination within Staphylococcus genus by extended PCR-restriction fragment length polymorphism of saoC gene. FEMS Microbiology Letters, 2015, 362, 1-11.	0.7	6
26	Clustering of Staphylococcus aureus bovine mastitis strains from regions of Central-Eastern Poland based on their biochemical and genetic characteristics. Polish Journal of Veterinary Sciences, 2015, 18, 333-342.	0.2	7