

# Marcelina Osińska

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7520256/publications.pdf>

Version: 2024-02-01

15  
papers

197  
citations

1162367

8  
h-index

1058022

14  
g-index

17  
all docs

17  
docs citations

17  
times ranked

163  
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparative characteristics of sequence types, genotypes and virulence of multidrug-resistant <i>E. faecium</i> isolated from various hosts in eastern Poland. Spread of clonal complex 17 in humans and animals. <i>Research in Microbiology</i> , 2022, , 103925.	1.0	3
2	Airborne dermatophyte propagules concentration in cowsheds as an underestimated reservoir of potential zoonoses. <i>Journal of Applied Microbiology</i> , 2022, , .	1.4	1
3	Detection and identification of dermatophytes based on currently available methods – a comparative study. <i>Journal of Applied Microbiology</i> , 2021, 130, 278-291.	1.4	20
4	Comparative study of multidrug-resistant <i>Enterococcus faecium</i> obtained from different hosts. <i>Journal of Medical Microbiology</i> , 2021, 70, .	0.7	3
5	Are dogs and cats a reservoir of resistant and virulent <i>Enterococcus faecalis</i> strains and a potential threat to public health?. <i>Journal of Applied Microbiology</i> , 2021, 131, 2061-2071.	1.4	6
6	Complementary effect of mechanism of multidrug resistance in <i>Trichophyton mentagrophytes</i> isolated from human dermatophytoses of animal origin. <i>Mycoses</i> , 2021, 64, 537-549.	1.8	15
7	Comparison of in vitro activities of 11 antifungal agents against <i>Trichophyton verrucosum</i> isolates associated with a variety hosts and geographical origin. <i>Mycoses</i> , 2020, 63, 294-301.	1.8	10
8	A significant number of multi-drug resistant <i>Enterococcus faecalis</i> in wildlife animals; long-term consequences and new or known reservoirs of resistance?. <i>Science of the Total Environment</i> , 2020, 705, 135830.	3.9	21
9	Population differentiation, antifungal susceptibility, and host range of <i>Trichophyton mentagrophytes</i> isolates causing recalcitrant infections in humans and animals. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2020, 39, 2099-2113.	1.3	20
10	Assessment of the subtilisin gene profile in <i>Trichophyton verrucosum</i> isolated from human and animal dermatophytoses in two-stage multiplex PCR. <i>Journal of Applied Microbiology</i> , 2020, 131, 300-306.	1.4	6
11	Intrinsic resistance to terbinafine among human and animal isolates of <i>Trichophyton mentagrophytes</i> related to amino acid substitution in the squalene epoxidase. <i>Infection</i> , 2020, 48, 889-897.	2.3	39
12	Dermatophytosis with concurrent <i>Trichophyton verrucosum</i> and <i>T. benhamiae</i> in calves after long-term transport. <i>Veterinary Dermatology</i> , 2020, 31, 414.	0.4	6
13	Application of genotyping methods in the investigation of sources of dermatophytosis associated with vaccination in cattle. <i>Annals of Applied Biology</i> , 2020, 177, 325-332.	1.3	6
14	In search of the source of dermatophytosis: Epidemiological analysis of <i>Trichophyton verrucosum</i> infection in llamas and the breeder (case report). <i>Zoonoses and Public Health</i> , 2019, 66, 982-989.	0.9	28
15	THE PREVALENCE OF SYMPTOMATIC DERMATOPHYTOSES IN DOGS AND CATS AND THE PATHOMECHANISM OF DERMATOPHYTE INFECTIONS. <i>Postepy Mikrobiologii</i> , 2019, 58, 165-176.	0.1	13