

# Anke Burmester

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/733084/publications.pdf>

Version: 2024-02-01

9  
papers

677  
citations

1163117  
8  
h-index

1474206  
9  
g-index

9  
all docs

9  
docs citations

9  
times ranked

540  
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparative and functional genomics provide insights into the pathogenicity of dermatophytic fungi. <i>Genome Biology</i> , 2011, 12, R7.	9.6	181
2	The current Indian epidemic of superficial dermatophytosis due to <i>Trichophyton mentagrophytes</i> – A molecular study. <i>Mycoses</i> , 2019, 62, 336-356.	4.0	164
3	Alarming India-wide phenomenon of antifungal resistance in dermatophytes: A multicentre study. <i>Mycoses</i> , 2020, 63, 717-728.	4.0	122
4	Spread of Terbinafine-Resistant <i>Trichophyton mentagrophytes</i> Type VIII (India) in Germany – “The Tip of the Iceberg”. <i>Journal of Fungi</i> (Basel, Switzerland), 2020, 6, 207.	3.5	73
5	A clarion call for preventing taxonomical errors of dermatophytes using the example of the novel <i>Trichophyton mentagrophytes</i> genotype VIII uniformly isolated in the Indian epidemic of superficial dermatophytosis. <i>Mycoses</i> , 2019, 62, 6-10.	4.0	62
6	Point mutations in the squalene epoxidase gene of Indian ITS genotype VIII <i>T. mentagrophytes</i> identified after DNA isolation from infected scales. <i>Medical Mycology Case Reports</i> , 2019, 26, 23-24.	1.3	26
7	Indian <i>Trichophyton mentagrophytes</i> squalene epoxidase <i>erg1</i> double mutants show high proportion of combined fluconazole and terbinafine resistance. <i>Mycoses</i> , 2020, 63, 1175-1180.	4.0	23
8	Point mutations in the squalene epoxidase <i>erg1</i> and sterol 14 $\alpha$ -demethylase <i>erg11</i> gene of <i>T. indotineae</i> isolates indicate that the resistant mutant strains evolved independently. <i>Mycoses</i> , 2022, 65, 97-102.	4.0	20
9	Efficacy of antifungal agents against fungal spores: An in vitro study using microplate laser nephelometry and an artificially infected 3D skin model. <i>MicrobiologyOpen</i> , 2022, 11, e1257.	3.0	6