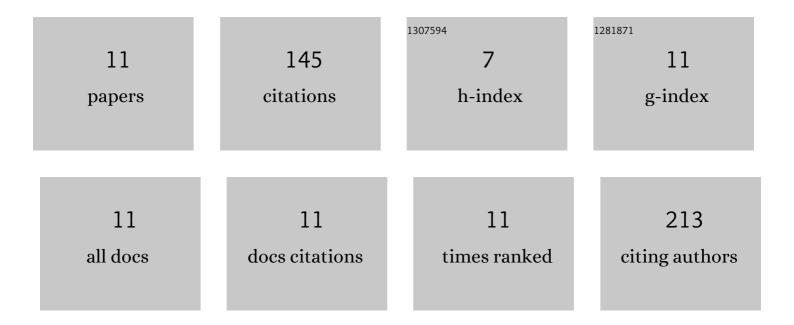
## David Pizarro

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6676363/publications.pdf Version: 2024-02-01



Πλυίο Ρισλάβο

#	Article	IF	CITATIONS
1	RNA SARS-CoV-2 Persistence in the Lung of Severe COVID-19 Patients: A Case Series of Autopsies. Frontiers in Microbiology, 2022, 13, 824967.	3.5	18
2	Differences in the Molecular Profile between Primary Breast Carcinomas and Their Cutaneous Metastases. Cancers, 2022, 14, 1151.	3.7	5
3	Neuropathological findings in fatal COVID-19 and their associated neurological clinical manifestations. Pathology, 2022, 54, 738-745.	0.6	12
4	Molecular Heterogeneity of High Grade Colorectal Adenocarcinoma. Cancers, 2021, 13, 233.	3.7	4
5	Epithelial Mesenchymal Transition and Immune Response in Metaplastic Breast Carcinoma. International Journal of Molecular Sciences, 2021, 22, 7398.	4.1	13
6	Clinical, Pathological, and Molecular Features of Breast Carcinoma Cutaneous Metastasis. Cancers, 2021, 13, 5416.	3.7	7
7	Genome-Wide Analysis of Biosynthetic Gene Cluster Reveals Correlated Gene Loss with Absence of Usnic Acid in Lichen-Forming Fungi. Genome Biology and Evolution, 2020, 12, 1858-1868.	2.5	28
8	Using target enrichment sequencing to study the higher-level phylogeny of the largest lichen-forming fungi family: Parmeliaceae (Ascomycota). IMA Fungus, 2020, 11, 27.	3.8	7
9	Whole-Genome Sequence Data Uncover Widespread Heterothallism in the Largest Group of Lichen-Forming Fungi. Genome Biology and Evolution, 2019, 11, 721-730.	2.5	15
10	Draft genome sequences of five Calonectria species from Eucalyptus plantations in China, Celoporthe dispersa, Sporothrix phasma and Alectoria sarmentosa. IMA Fungus, 2019, 10, 22.	3.8	17
11	Phylogenomic analysis of 2556 single-copy protein-coding genes resolves most evolutionary relationships for the major clades in the most diverse group of lichen-forming fungi. Fungal	12.3	19