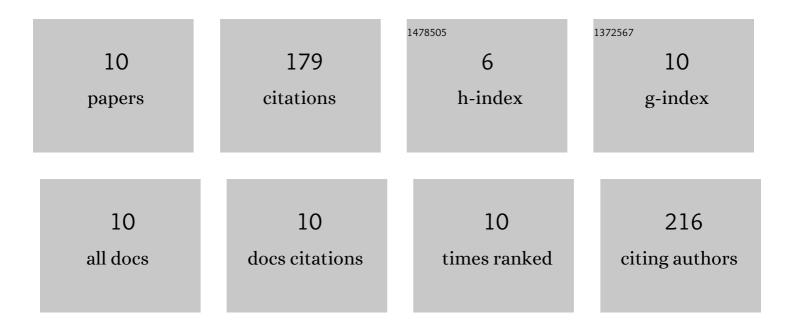
Sofia Grecco

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4354847/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Consecutive deletions in a unique Uruguayan SARS-CoV-2 lineage evidence the genetic variability potential of accessory genes. PLoS ONE, 2022, 17, e0263563.	2.5	7

2 Domestic dog origin of Carnivore Protoparvovirus 1 infection in a rescued free $\hat{a} \in r$ anging gui $\tilde{A} \pm a$ () Tj ETQq0 0 0 rgBT Overlock 10 Tf 50

3	Origin, spreading and genetic variability of chicken anaemia virus. Avian Pathology, 2021, 50, 311-320.	2.0	7
	ongin, spreading and genetic variability of chicken anachila viras. Avian Factology, 2021, 30, 311 320.		
4	Origin and global spreading of an ancestral lineage of the infectious bursal disease virus. Transboundary and Emerging Diseases, 2020, 67, 1198-1212.	3.0	11
5	First Molecular Identification of Canine Parvovirus Type 2 (CPV2) in Chile Reveals High Occurrence of CPV2c Antigenic Variant. Frontiers in Veterinary Science, 2020, 7, 194.	2.2	11
6	Inter- and intracontinental migrations and local differentiation have shaped the contemporary epidemiological landscape of canine parvovirus in South America. Virus Evolution, 2018, 4, vey011.	4.9	38
7	Development of an RT-qPCR assay for the specific detection of a distinct genetic lineage of the infectious bursal disease virus. Avian Pathology, 2017, 46, 150-156.	2.0	9
8	Development of RT-qPCR assays for the specific identification of two major genotypes of avian infectious bronchitis virus. Journal of Virological Methods, 2016, 235, 21-25.	2.1	12
9	Genome Sequence of a Distinct Infectious Bursal Disease Virus. Genome Announcements, 2015, 3, .	0.8	5
10	Phylogenetic and Genome-Wide Deep-Sequencing Analyses of Canine Parvovirus Reveal Co-Infection with Field Variants and Emergence of a Recent Recombinant Strain. PLoS ONE, 2014, 9, e111779.	2.5	73