

Sofia Grecco

List of Publications by Year in descending order

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Version: 2024-02-01

10
papers

179
citations

1478505

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h-index

1372567

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g-index

10
all docs

10
docs citations

10
times ranked

216
citing authors

#	ARTICLE	IF	CITATIONS
1	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
2	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
3	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
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	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
7	Origin, spreading and genetic variability of chicken anaemia virus. Avian Pathology, 2021, 50, 311-320.	2.0	7
8	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
9	Domestic dog origin of Carnivore Protoparvovirus 1 infection in a rescued free-ranging guinea pig (Cavia porcellus) from Brazil. PLoS ONE, 2022, 17, e0263563.	3.0	6
10	Genome Sequence of a Distinct Infectious Bursal Disease Virus. Genome Announcements, 2015, 3, .	0.8	5