

# Shubhagata Das

## List of Publications by Year in descending order

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35  
papers

345  
citations

1039880

9  
h-index

839398

18  
g-index

36  
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36  
docs citations

36  
times ranked

316  
citing authors

#	ARTICLE	IF	CITATIONS
1	Prevalence and Antimicrobial Resistance Profile of <i>E. coli</i> and <i>Salmonella</i> spp. from Liver and Heart of Chickens. <i>Turkish Journal of Agriculture: Food Science and Technology</i> , 2022, 10, 1191-1196.	0.1	0
2	Structural Perspectives of Beak and Feather Disease Virus and Porcine Circovirus Proteins. <i>Viral Immunology</i> , 2021, 34, 49-59.	0.6	9
3	Lumpy skin disease outbreak in cattle population of Chattogram, Bangladesh. <i>Veterinary Medicine and Science</i> , 2021, 7, 1616-1624.	0.6	34
4	Retrospective Genotyping and Whole Genome Sequencing of a Canine Parvovirus Outbreak in Bangladesh. <i>Pathogens</i> , 2021, 10, 1373.	1.2	0
5	Molecular characterisation of a novel pathogenic avipoxvirus from the Australian magpie ( <i>Gymnorhina tibicen</i> ). <i>Virology</i> , 2020, 540, 1-16.	1.1	24
6	Diverse <i>Trichomonas</i> lineages in Australasian pigeons and doves support a columbid origin for the genus <i>Trichomonas</i> . <i>Molecular Phylogenetics and Evolution</i> , 2020, 143, 106674.	1.2	12
7	REPEAT SPILLOVER OF BEAK AND FEATHER DISEASE VIRUS INTO AN ENDANGERED PARROT HIGHLIGHTS THE RISK ASSOCIATED WITH ENDEMIC PATHOGEN LOSS IN ENDANGERED SPECIES. <i>Journal of Wildlife Diseases</i> , 2020, 56, 896-906.	0.3	9
8	Pigeon circoviruses from feral pigeons in Australia demonstrate extensive recombination and genetic admixture with other circoviruses. <i>Avian Pathology</i> , 2019, 48, 512-520.	0.8	12
9	Assessing circovirus gene flow in multiple spill-over events. <i>Virus Genes</i> , 2019, 55, 802-814.	0.7	5
10	The first complete mitogenome of Australia's largest raptor, the wedge-tailed eagle ( <i>Aquila</i> ) Tj ETQq0 0 0 rgBT <sub>2</sub> /Overlock 10 Tf 50	0.2	1
11	The first complete mitogenome of Indian ringneck ( <i>Psittacula krameri</i> ) demonstrates close phylogenetic relationship with <i>Electus</i> parrot. <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 3579-3581.	0.2	10
12	The first complete mitogenome of red-bellied parrot ( <i>Poicephalus rufiventris</i> ) resolves phylogenetic status within <i>Psittacidae</i> . <i>Mitochondrial DNA Part B: Resources</i> , 2018, 3, 195-197.	0.2	1
13	Genome sequence of an Australian strain of <i>canid alphaherpesvirus 1</i> . <i>Australian Veterinary Journal</i> , 2018, 96, 24-27.	0.5	25
14	Laboratory Diagnosis of Respiratory Tract Infections in Children – the State of the Art. <i>Frontiers in Microbiology</i> , 2018, 9, 2478.	1.5	39
15	The first complete mitogenome of Indian star tortoise ( <i>Geochelone elegans</i> ). <i>Mitochondrial DNA Part B: Resources</i> , 2018, 3, 1112-1113.	0.2	1
16	Pathogenic potentials and shedding probability of <i>Salmonella enterica</i> serotype Kentucky in experimentally infected backyard chicken. <i>Journal of Advanced Veterinary and Animal Research</i> , 2018, 5, 196.	0.5	2
17	A novel pathogenic aviadenovirus from red-bellied parrots ( <i>Poicephalus rufiventris</i> ) unveils deep recombination events among avian host lineages. <i>Virology</i> , 2017, 502, 188-197.	1.1	34
18	Genomic characterization of two novel pathogenic avipoxviruses isolated from pacific shearwaters ( <i>Ardenna</i> spp.). <i>BMC Genomics</i> , 2017, 18, 298.	1.2	51

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19	Structural insights into GDP-mediated regulation of a bacterial acyl-CoA thioesterase. <i>Journal of Biological Chemistry</i> , 2017, 292, 20461-20471.	1.6	9
20	Complete mitochondrial genome sequence of an Australian little penguin ( <i>Eudyptula minor</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 702 T	0.2	0
21	Beak and feather disease virus carriage by <i>Knemidocoptes pilae</i> in a sulphur-crested cockatoo ( <i>Cacatua galerita</i> ). <i>Australian Veterinary Journal</i> , 2017, 95, 486-489.	0.5	5
22	Characterization of Beak and Feather Disease Virus Genomes from Wild Musk Lorikeets ( <i>Glossopsitta</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 T	0.8	1
23	Whole-Genome Sequence of a Beak and Feather Disease Virus Isolate from a Fledgling Red-Capped Parrot ( <i>Purpureicephalus spurius</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	1
24	Molecular Characterization of a Beak and Feather Disease Virus Genome from a Purple Crowned Lorikeet ( <i>Glossopsitta porphyrocephala</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	0
25	Evolution of circoviruses in lorikeets lags behind its hosts. <i>Molecular Phylogenetics and Evolution</i> , 2016, 100, 281-291.	1.2	22
26	A comparison of PCR assays for beak and feather disease virus and high resolution melt (HRM) curve analysis of replicase associated protein and capsid genes. <i>Journal of Virological Methods</i> , 2016, 237, 47-57.	1.0	9
27	The complete mitochondrial genome sequence of an Endangered powerful owl ( <i>Ninox strenua</i> ). <i>Mitochondrial DNA Part B: Resources</i> , 2016, 1, 722-723.	0.2	5
28	Identification of Beak and Feather Disease Virus in an Unusual Novel Host ( <i>Merops ornatus</i> ) Using Nested PCR. <i>Genome Announcements</i> , 2016, 4, .	0.8	0
29	Molecular Detection of Infectious Bronchitis Virus Isolated from Commercial Breeder Farms in Chittagong District, Bangladesh. <i>Advances in Animal and Veterinary Sciences</i> , 2016, 4, 370-375.	0.1	2
30	Characterization of the Whole-Genome Sequence of a Beak and Feather Disease Virus Isolate from a Mallee Ringneck Parrot ( <i>Barnardius zonarius barnardi</i> ). <i>Genome Announcements</i> , 2014, 2, .	0.8	6
31	Molecular Characterization of Genome Sequences of Beak and Feather Disease Virus from the Australian Twenty-Eight Parrot ( <i>Barnardius zonarius semitorquatus</i> ). <i>Genome Announcements</i> , 2014, 2, .	0.8	8
32	Acanthomatous ameloblastoma in a female Spitz dog with rare atypical histomorphology: A case study. <i>Veterinary World</i> , 2013, 6, 219.	0.7	2
33	Pyometra in a German Shepherd Dog: A Clinical Case Report. <i>Asian Journal of Animal and Veterinary Advances</i> , 2012, 7, 446-451.	0.3	0
34	Spirocerosis in stray dogs of Chittagong metropolitan area of Bangladesh: an epidemiological and pathological investigation. <i>Veterinary World</i> , 2011, , 485.	0.7	6
35	A postmortem experience of Indian rock python ( <i>Python molurus molurus</i> ) that swallowed a whole barking deer. <i>Veterinary World</i> , 0, , 163-165.	0.7	0