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Cover: Stripe-necked Mongoose Urva vi icolla in poster colours, adapted from photograph by Ashni Dhawale, by Pooja Ramdas Patil.

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Phylogenetic insights on the delineation of Mysore and Malabar subspecies of the Grey Slender Loris *Loris lydekkerianus* in southern India

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Abstract: Slender lorises are a threatened genus of small and nocturnal strepsirrhine primates confined to India and Sri Lanka. The Grey Slender Loris *Loris lydekkerianus* is divided into several subspecies based on morphological variation and geographical distribution but not supported by molecular data. We investigated the phylogenetic divergence of two subspecies of the Grey Slender Loris in southern India: the Mysore Slender Loris *Loris lydekkerianus* ssp. *lydekkerianus* and the Malabar Slender Loris *Loris lydekkerianus* ssp. *malabaricus*. We generated whole genome shotgun sequence data and assembled the whole mitochondrial genomes of representative individuals from their distribution in southern India and compared them with publicly available mitogenomes of other lorises. We found that the Mysore and Malabar Slender Loriss of 13 protein-coding and two ribosomal RNA genes in the mitochondrial genome showed that the Mysore and Malabar Slender Loris *Loris tardigradus*. Considering this relatively high sequence variation and evolutionary divergence together with their already established morphological differences and geographically distinct habitats, we propose to recognize the Mysore and Malabar Slender Lorises as two distinct species *Loris lydekkerianus* and *Loris malabaricus*.

Keywords: Grey Slender Loris, Malabar Slender Loris, molecular dating, Mysore Slender Loris, phylogenetics.

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Author contributions: G.U. and H.N.K conceived the idea for this study and collected the samples. V.T. and S.M. analyzed the data and drafted the manuscript. All authors contributed to the revision of the manuscript and agreed to the final version of the manuscript.

For Tamil abstract see end of this article.

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INTRODUCTION

Slender lorises (genus Loris) are one of the two genera of extremely specialized nocturnal primates that inhabit India (Nekaris 2014). They belong to the family Lorisidae, which also includes Slow lorises, Pottos, and Angwantibos. Slender lorises are confined to India and Sri Lanka, where they inhabit dry to moist and lowland to montane forests (Singh et al. 2021). Slender lorises are characterized by their small size, long limbs, vestigial tail, large eyes, and slow locomotion. They are adapted for arboreal life, using their opposable thumbs and toes to grasp branches and their binocular vision to judge distances, visual acuity, precise hand-eye coordination and social communication. They feed mainly on insects but also consume fruits, flowers, gums, and other plant materials (Nekaris & Rasmussen 2003; Radhakrishna & Kumara 2010). They have a variety of vocalizations that may help them avoid predators and communicate with conspecifics (Radhakrishna & Singh 2002). Slender lorises are divided into two species: the Grey Slender Loris Loris lydekkerianus found in southern India and Sri Lanka and the Red Slender Loris Loris tardigradus found only in Sri Lanka (Groves 2001). Both species show high phenotypic variation in fur color, body size, and cranial morphology, leading to the recognition of several subspecies, most of which are refuted by molecular studies (Nijman et al. 2020). The Mysore Slender Loris Loris lydekkerianus ssp. lydekkerianus (Image 1) and the Malabar Slender Loris Loris lydekkerianus ssp. malabaricus (Image 2), which live in the dry and wet forests of the Eastern and Western Ghats, respectively, are the two subspecies that have been recognized thus far in southern India (Kumara et al. 2013). There are several regions in their distribution where slender lorises face serious threats to their existence such as habitat loss due to deforestation and urbanization, electrocution on live wires, road accidents, pet trade, and illegal poaching for traditional medicine and black magic (Dittus et al. 2022; Gnanaolivu et al. 2022). The IUCN Red List of Threatened Species classifies the Mysore Slender Loris (Kumara et al. 2022a) and Malabar Slender Loris (Kumara et al. 2022b) as 'Near Threatened' and they are listed under Schedule I of the Indian Wild Life (Protection) Act, 1972. Recently, Tamil Nadu became the first Indian state to notify a sanctuary for slender lorises spanning 118.06 km², which is crucial for protecting their habitat and ensuring the survival of this unique primate species (Government of Tamil Nadu 2022).

The Mysore and Malabar subspecies of the Grey Slender Loris were delineated based on their

morphological differences and geographic distribution (Groves 2001; Kumara et al. 2013). The Mysore Slender Loris is relatively larger (ca. 260 g) than Malabar Slender Loris (ca. 180 g) (Kumara et al. 2006). The Mysore Slender Loris has a grayish-brown coat and a prominent white stripe on its forehead, whereas the Malabar Slender Loris has a reddish-brown coat and a less distinct forehead stripe (Groves 2001; Kumara et al. 2006). The relative distribution of the two subspecies as well as their comparative densities and the extent of overlap between their distributions have been very well established (Kumara et al. 2013). The Mysore Slender Loris is found in the Eastern Ghats and eastern foothills of the southern Western Ghats, while the Malabar Slender Loris is confined to the western slope of the entire Western Ghats (Kumara et al. 2013). The Mysore Slender Loris prefers dry deciduous forests with moderate canopy cover and high tree density, while the Malabar Slender Loris prefers moist evergreen forests with high canopy cover and low tree density (Kumara et al. 2013). Their distributions overlap along the southern ridges of the Western Ghats, where hybridization may occur. While the diet of Mysore Slender Loris mostly consists of insects, plant material, and gum, the feeding behavior of the Malabar Slender Loris is not well studied (Radhakrishna & Kumara 2010). The reproductive biology and social system of the Mysore subspecies is influenced by factors such as seasonality, food availability, predation risk, and population density. It also has a seasonal breeding cycle that coincides with periods of high food availability (Radhakrishna & Singh 2004). No such information on the reproductive biology of the Malabar subspecies is available. Behavioral studies on lorises have always been more challenging than on relatively large, diurnal, and group-living primates such as macaques and langurs because they are nocturnal, small in size, and mostly semi-gregarious. Given the distinct habitat preferences and morphology of these two subspecies, understanding their evolutionary history and genetic differences is vital to address their conservation status and management issues.

Therefore, the main objective of this study is to investigate the phylogenetic relationship and genetic divergence between the Mysore and Malabar Slender Lorises in southern India. To achieve this, we sequenced and assembled the whole mitochondrial sequences from three representative samples. We aligned these sequences with the publicly available sequences of other lorises and constructed phylogenetic trees. We estimated the sequence divergence and divergence time between the two subspecies using phylogenetic analysis. Phylogenetic insights on the delineation of Mysore and Malabar Loris in southern India



Image 1. Mysore Slender Loris Loris lydekkerianus ssp. lydekkerianus.



Image 2. Malabar Slender Loris Loris lydekkerianus ssp. malabaricus.

Our results support the morphological and geographical delineation of the Malabar and Mysore Slender Lorises and advocates for recognizing them as two distinct species. This study will contribute to the understanding of the biogeography and speciation processes of these threatened lorises and provide crucial insights for their conservation and management.

MATERIALS AND METHODS

Sample Collection, DNA extraction, and Sequencing

We followed the sample collection guidelines of the animal ethics committees of the CSIR-Centre for Cellular and Molecular Biology and Salim Ali Centre for Ornithology and Natural History. Necessary permissions for sample collection were obtained from the Central Zoo Authority of India, Ministry of Environment, Forests & Climate Change, Government of India, vide Ref. No. 9-2/2005-CZA(M) Vol III. Rescued lorises of known wild origin within the IUCN designated ranges (Figure 1) that were captive in Mysore and Hyderabad zoos were the sources of our samples. Blood samples were collected in EDTA vacutainers by qualified zoo veterinarians from three representative individuals of Loris lydekkerianus ssp. lydekkerianus (N = 2) and Loris lydekkerianus ssp. malabaricus (N = 1). We used the Qiagen DNeasy Blood and Tissue Kit to isolate the genomic DNA from the blood samples. We measured the quality and quantity of genomic DNA using Nanodrop and Qubit 4. We constructed whole genome libraries using the Truseq PCR-free library preparation kit according to Illumina's protocols. Briefly, 1 ug of genomic DNA was sheared to

approximately 350 bp using the Covaris ultrasonicator. The fragmented DNA was then end-repaired and bluntend ligated with sequencing adapters containing unique dual indices from IDT. The library was then size-selected using SPRI beads and verified on the Agilent fragment analyzer. The cleaned-up libraries were finally quantified in qPCR using the standards and Illumina adapterspecific primers from the Roche library quantification kit. Libraries having good concentration were pooled along with other samples and sequenced on the Illumina Novaseq 6000 platform for 300 cycles in paired-end mode.

Mitochondrial genome assembly

We demultiplexed the base call files to separate the three samples with the dual-indexed barcodes using the BCL2FASTQ tool from Illumina. Raw reads were quality-filtered with a phred quality score threshold of 15 using FASTP v0.20 (Chen et al. 2018). We subsampled 10 million quality filtered reads to de novo assemble the circular mitochondrial genomes of all three samples using GetOrganelle v1.7.1 (Jin et al. 2020). We then annotated all the mitogenomes using MITOS2 (Bernt et al. 2013) with the Refseq 89 Metazoa reference mitochondrial database and the vertebrate mitochondrial genetic code. All the coding and non-coding genes were extracted from the mitochondrial genomes using the annotations.

Sequence and Phylogenetic analyses

We aligned the full-length COX1 and CYTB genes of lorises using Clustal Omega with the "distmat" flag and calculated the pairwise distances between the sequences (Sievers & Higgins 2021). To build the phylogeny, along



Figure 1. IUCN Red List distribution range of the Grey Slender Loris subspecies in southern India and Sri Lanka.

with our samples we used the NCBI RefSeg mitochondrial sequences from strepsirrhines namely, Loris lydekkerianus, Loris tardigradus, Nycticebus coucang, Nycticebus bengalensis, Galago senegalensis, and Lemur catta. We aligned the 13 protein-coding genes and two non-coding ribosomal RNA genes individually from the assembled mitochondrial genomes and reference sequences using the MUSCLE algorithm in MEGA7 (Kumar et al. 2016) and checked for the presence of any sequencing errors or frameshifts for codon position. We then concatenated all the gene alignments using MEGA7 (Kumar et al. 2016) and identified the optimum nucleotide substitution model for each partition based on the corrected Akaike information criterion (AICc) values using PartitionFinder2 (Lanfear et al. 2017) (Supplementary file 1). We built the maximum likelihood (ML) tree based using IQ-TREE (Minh et al. 2020) with 1000 times bootstrapping. The ML tree was visualized in Evolview v3 (Subramanian et al. 2019).

We utilized BEAST2.5 (Bouckaert et al. 2014) to create a divergence time tree using the same concatenated alignment of 13 coding and two non-coding genes from the complete mitochondrial genomes. We used the same partitioning scheme and substitution models identified by PartitionFinder2. We then chose two fossil calibration points:

1) We calibrated the crown node of Galagos with 38 mya based on the age of the fossil *Saharagalago misrensis* (PaleoDB collection 67706) (Seiffert et al. 2003). We applied a normal distribution at 40 Mya (SD = 0.04; 95% range: 36–43)

2) We calibrated the crown node of Slow Lorises with 13.82 mya based on the age of the fossil *Nycticebus linglom* (PaleoDB collection 48126) (Harrison 2010). We applied a normal distribution at 14 Mya (SD = 0.05; 95% range: 9–17)

For all the partitions, we created a relaxed lognormal clock and employed a birth-death process using prior distributions. To get to the final tree, we ran for 40 million generation runs, sampling every 2,000th generation using TreeAnnotator (Helfrich et al. 2018) with a 10% burnin. We verified that all the ESS values were over 200 in Tracer 1.7 (Rambaut et al. 2018) and visualized the tree in FigTree v1.4.4 (Rambaut 2014).

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RESULTS

Phylogenetic analyses support the morphological and geographical delineation of Mysore & Malabar Slender Lorises

To investigate the genetic differences between the two subspecies of slender loris in southern India, we first assembled three new circular mitochondrial genomes with an average length of 16,771 bp from two samples of the Mysore Slender Loris Loris lydekkerianus ssp. lydekkerianus and one sample of Malabar Slender Loris Loris lydekkerianus ssp. malabaricus. We annotated the mitochondrial genomes along with published reference sequences and obtained the full-length sequences of 13 protein-coding genes and two ribosomal RNA genes. To check the variation in the nucleotide sequence within the Loris genus, we estimated the pairwise sequence similarity in the COX1 and CYTB regions spanning 2,682 bp between all six Loris samples (Table 1). We observed the highest average sequence variation of 2.82% (S.D. 0.16) between the four sequences of the Grey Slender Loris Loris lydekkerianus and the two sequences of the Red Slender Loris Loris tardigradus as they belong to two different species within the Loris genus. While there was no sequence variation found within the two sequences of Red Slender Loris, there was considerable variation within the four sequences of Grey Slender Loris contributed by the difference between the two subspecies. We found about 2.09% (S.D 0.0) variation in the COX1 and CYTB sequences of the Mysore and Malabar Slender Lorises.

We then used phylogenetic analyses to understand the evolutionary relationships between the two subspecies. Along with our three samples, we included reference mitochondrial sequences from two species of slender lorises (*L. lydekkerianus, L. tardigradus*) and two species of slow lorises (*Nycticebus bengalensis, N. coucang*) along with galago and lemur as outgroups (Figure 2). The phylogenetic tree recapitulates the broad evolutionary relationships of slender lorises with slow lorises and the outgroups. It reveals an interesting pattern within the clade of slender lorises where the Mysore Slender Loris L.I. ssp. lydekkerianus clusters with the reference sequence of Grey Slender Loris to form a monophyletic clade and the Malabar Slender Loris L.I. ssp. malabaricus forms a separate monophyletic clade with very strong statistical support. We noted that the Malabar Slender Loris appears more closely related to the Red Slender Loris L. tardigradus, albeit with a very small branch length (Figure 2). To estimate the divergence time between the two subspecies and other lorises, we constructed a fossil-calibrated Bayesian tree (Figure 3). Our results suggest that the split between the Grey Slender Loris L. lydekkerianus and Red Slender Loris L. tardigradus occurred approximately 1.087 million years ago (mya). This was immediately followed by diversification of the Mysore Slender Loris L.I. ssp. lydekkerianus and Malabar Slender Loris L.I. ssp. malabaricus at around 1.049 mya (Posterior probability = 1) (Figure 3).

DISCUSSION

Our results from the phylogenetic analyses based on the mitochondrial sequences show that the Mysore and Malabar Slender Lorises have significant genetic variation (2.09%) in the COX1 and CYTB genes and form distinct monophyletic clades in the phylogenetic tree that diverged a long time ago (1.049 mya), shortly after the divergence of Red Slender Loris from the Grey Slender Loris (1.087 mya). The observed sequence variation and divergence time between the Mysore and the Malabar Slender Lorises are surprisingly high, which is not very common between primate subspecies. Since they have been evolving independently for a

Sample	Loris tardigradus-1	Loris tardigradus-2	Loris lydekkerianus malabaricus	Loris lydekkerianus lydekkerianus-1	Loris lydekkerianus lydekkerianus-2	Loris lydekkerianus-Ref
Loris tardigradus-1	-	0	2.58	2.82	2.95	2.9
Loris tardigradus-2	0	-	2.58	2.82	2.97	2.9
Loris lydekkerianus malabaricus	2.58	2.58	-	2.09	2.09	2.09
Loris lydekkerianus lydekkerianus-1	2.82	2.82	2.09	-	0.16	0.08
Loris lydekkerianus lydekkerianus-2	2.97	2.97	2.09	0.16	-	0.16
Loris lydekkerianus- Ref	2.9	2.9	2.09	0.08	0.16	-

Table 1. Sequence distance matrix of the Loris genus based on full-length cytochrome b and cytochrome oxidase 1 genes.

4.1 Lemur catta NC 059325.1 Roo taked Lam Galago sevegalensis NC_012761.1 Nycticebus coucang NC_002765.1 Nycticebus bengalensis NC_021958.1 Bengal Size Lone Loris lydekkerianus lydekkerianus-2. OR115513.1 Loris lydekkerlanus lydekkerlanus-1 OR115511.1 **Wykow Stender Lock** Loris lydekkerianus lydekkerianus-Ref NC_021955.1 100 Loris lydekkerianus malabaricus OR115512.1 14 oris landigrachus-2 AB371094,1 d SenderLone oris landigradua-1 NC_012763.1

Figure 2. Maximum likelihood tree based on 13 protein-coding and two ribosomal RNA genes from whole mitochondrial genomes. The bootstrap values are denoted at the nodes.



Figure 3. Fossil-calibrated Bayesian inference tree based on 13 protein-coding and two ribosomal RNA genes from whole mitochondrial genomes showing the divergence time estimates at the nodes in Mya. Bars indicate 95% CI.

long period comparable to the divergence time of their closest species (*Loris tardigradus*), the Mysore and Malabar Slender Lorises deserve independent recognition. Moreover, they occupy a geographically different landscape and unique habitat, where the Malabar Slender Loris occupies the wet zone of the Western Ghats, while the Mysore Slender Loris occupies the dry habitat of the eastern slope of the Western Ghats, dry forests of the Deccan plateau and Eastern Ghats (Kumara et al. 2006, 2009; 2013). They are also morphologically distinct, where the Malabar Slender Loris appears reddish in color and almost half the body size of the greyish colored Mysore Slender Loris (Kumara et al. 2006). Considering these significant differences in the morphological, geographical, and genetic factors, we propose to recognize the Mysore Slender Loris *Loris lydekkerianus* ssp. *lydekkerianus* and Malabar Slender Loris *Loris lydekkerianus* ssp. *malabaricus* as two distinct

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species, Loris lydekkerianus and Loris malabaricus, respectively.

The divergence time estimates are supported by a number of molecular markers in the whole mitochondrial genome and is consistent with previous studies on the evolutionary history of this genus (Finstermeier et al. 2013). Several environmental, climatic, and geographical factors might have influenced the divergence of the Mysore and Malabar Slender Lorises about one million years ago. The mid-Pleistocene transition (1.25–0.7 Mya) was a time of dramatic climatic change and glaciation that influenced the environments and biogeography of Earth (Herbert 2023). The glaciation and interglaciation cycles affected the sea level, precipitation, temperature, vegetation, and habitat availability. The environmental conditions in India specifically during the Pleistocene were diverse and dynamic, ranging from deserts, tropical forests to grasslands (Morley & Morley 2022). The variability of monsoon coupled with expansion and contraction of forests due to glacial-interglacial cycles could have influenced availability of resources, fragmentation of habitats, and changes in forest cover promoting genetic differentiation and divergence of the Mysore and Malabar Slender Lorises.

Understanding the genetic structure and variation of species is crucial for the scientific management of threatened species and their eventual recovery. The findings of this study have important implications for the conservation and management of the slender lorises in India. With a clearer understanding of the genetic differences between the Mysore and Malabar Slender Lorises, it will be possible to more accurately identify and classify individual animals, which will in turn facilitate the development of an effective conservation breeding program. Such a program can be particularly beneficial for species like the slender loris that are threatened by habitat loss and fragmentation, and whose populations have been declining in recent years (Kumara et al. 2006, 2016).

The main drawback of this study is the limited sample size which we duly acknowledge. It is to be noted that the construction of whole mitochondrial genomes from WGS data for accurate molecular dating often requires good-quality DNA from animals of known geographic origin which is very difficult to obtain, especially for the Malabar Slender Loris. More samples from the Malabar Slender Loris could better resolve the phylogenetic tree and nuclear markers could also be used to confirm our findings and validate the species delimitation. Furthermore, sampling the individuals from the range edges and the overlapping ranges in the southern ridge of Western Ghats would provide more statistical power to establish the monophyly and identify any hybridization. It would also be prudent to include samples of the Mysore subspecies from Sri Lanka in future studies to fully comprehend the diversity and understand the evolutionary history of the slender lorises throughout its geographical range. Comprehensive genome sequencing of all the subspecies of slender loris would also help to understand the genomic basis of morphological differences and their adaptations to respective niches.

In conclusion, this study provides the first molecular evidence for the genetic divergence and distinctiveness of the Mysore and Malabar Slender Lorises. The sequence analysis, phylogenetic analyses, and molecular dating suggest that the Mysore and Malabar Slender Lorises are genetically distinct and have been evolving independently for a significant period. The high level of genetic divergence between them highlights the importance of preserving their genetic diversity and underscores the need for more efforts to conserve them in the wild. By considering the significant differences in the morphological, geographical, and genetic factors, we recommend to elevate L.I. ssp. lydekkerianus and L.I. ssp. malabaricus to the species level. We propose to recognize them as two distinct species Loris lydekkerianus and Loris malabaricus as each of them represents a unique evolutionary lineage and deserves separate recognition and protection. We advocate for further studies to validate the species delimitation with larger sample sizes and recommend for separate conservation measures and management actions to preserve their unique genetic diversity in the wild and captivity.

Data Availability Statement

The three whole mitochondrial genome sequences generated in this study have been submitted to the NCBI database under the accessions OR115511, OR115512, and OR115513.

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Supplementary file 1. Partitioning scheme and nucleotide substitution model selection output from PartitionFinder2.

Best partitioning scheme

Scheme InL :-64454.10696411133 Scheme AICc : 129409.318896 Number of params : 246 Number of sites : 13594 Number of subsets : 33

Subset | Best Model | # sites | Partition names

1	GTR+I+X 622 NAD1_pos1, NAD4I_pos3, ATP6_pos1
2	TRN+I+X 533 NAD1_pos2, ATP6_pos2
3	HKY+X 223 ATP6_pos3
4	HKY+X 132 ATP8_pos1, ATP8_pos2
5	GTR+X 317 ATP8_pos3, COX3_pos3
6	HKY+G+X 368 COB_pos1
7	GTR+I+X 368 COB_pos2
8	TRN+I+X 368 COB_pos3
9	GTR+G+X 496 COX1_pos1
10	HKY+G+X 496 COX1_pos2
11	HKY+G+X 496 COX1_pos3
12	K80 222 COX2_pos1
13	TRN+X 222 COX2_pos2
14	TRN+I+X 424 NAD3_pos1, NAD4I_pos2, COX2_pos3
15	JC 251 COX3_pos1
16	HKY+X 251 COX3_pos2
17	HKY+X 310 NAD1_pos3
18	GTR+I+X 340 NAD2_pos1
19	HKY+I+X 340 NAD2_pos2
20	HKY+I+X 339 NAD2_pos3
21	TRN+I+X 111 NAD3_pos2
22	GTR+I+X 201 NAD4I_pos1, NAD3_pos3
23	HKY+X 457 NAD4_pos1
24	HKY+X 456 NAD4_pos2
25	HKY+I+X 456 NAD4_pos3
26	HKY+I+X 596 NAD5_pos1
27	GTR+I+X 596 NAD5_pos2
28	HKY+I+X 596 NAD5_pos3
29	GTR+G+X 181 NAD6_pos1
30	HKY+I+X 180 NAD6_pos2
31	GTR+G+X 180 NAD6_pos3
32	GTR+G+X 1454 rrns
33	GTR+I+X 1012 rrnl

Tamil: மெலிந்த தேவாங்குகள், இந்தியா மற்றும் இலங்கையில் மட்டும் வாழும் இரவில் நடமாடுகிற சிறிய வகை ஈரமூக்கு கொண்ட முதனிகளில் ஒன்றான பேரினமாகும். இந்தப் பேரினம் தற்பொழுது அழிவுநிலைக்கு அச்சுறுத்தப்பட்டுள்ளது.. சாம்பல் நிற மெலிந்த தேவாங்கு (Loris lydekkerianus), உருவவியல் மாறுபாடு மற்றும் புவியியல் பரவல் ஆகியவற்றின் அடிப்படையில் பல துணை இனங்களாகப் பிரிக்கப்பட்டுள்ள போதும், இந்தப் பிரிவினை மூலக்கூறு சான்றுகளால் ஆதரிக்கப்படவில்லை. தென்னிந்தியாவில் வாழும் இந்த சாம்பல் நிற மெலிந்த தேவாங்குகளின் இரண்டு துணை இனங்கள்: அதாவது மைதர் மெலிந்த தேவாங்கு (Loris lydekkerianus ssp. lydekkerianus) மற்றும் மலபார் மெலிந்த தேவாங்கு (Loris lydekkerianus ssp. malabaricus), பரிணாம வரலாற்றில் எப்பொழுது பிரிந்தன என்று ஆராய்ந்தோம். தென்னிந்தியாவில் வாழும் அவற்றின் புவியியல் தொகைகளில் இருந்து மாதிரிகள் எடுத்து, அவ்விலங்குகளின் முழு இழைமணி மரபணு தகவல்களை shotgun sequence எனப்படும் தகவல்களில் இருந்து ஒருங்கிணைத்து, பொது தரவுத்தளங்களில் கிடைக்கும் பிற தேவாங்குகளின் மரபணு தகவல்களுடன் ஒப்பிட்டோம். இதில் இருந்து மைதர் மற்றும் மலபார் மெலிந்த தேவாங்குகளை ஒப்பிடுகையில், அவற்றின் இடையே cox1 மற்றும் cyrB மரபணு பகுதிகள் 2.09 சதவிகிதம் வேறுபடுவதை நாங்கள் கண்டறிந்தோம். மேலும், இழைமணி மரபணுவில் காணப்படும் 13 புரதங்கள் மற்றும் 2 ரைபோசோமல் RNAக்களைக் குறியிடக்கூடிய மரபணு தகவல்களை பகுப்பாய்வு செய்கையில், மைதர் மற்றும் மலபார் மெலிந்த தேவாங்குகள், பரிணாம வரலாற்றில் சுமார் 10. 49 இலட்சம் ஆண்டுகளுக்கு முன்பு பிரிந்தன என விளங்கியது. இது சிவப்பு மெலிந்த தேவாங்கு (Loris tardigradus) பிரிந்த சற்று காலத்திற்குப் பின் நிகழ்ந்த நிகழ்வாகும். மரபணு தகவல்களில் உள்ள மாற்றங்கள் மற்றும் பரிணாம வேறுபாடுகளை கருத்தில் கொண்டு , அவற்றுடன் ஏற்கனவே நிறுவப்பட்ட உருவ வேறுபாடுகள் மற்றும் புவியியல் ரீதியாக தனித்துவமான வாழ்விடங்களை சேர்க்கையில், மைசூர் மற்றும் மலபார் மெலிந்த தேவாங்குகளளை Loris lydekkerianus மற்றும் Loris malabaricus என இரண்டு தனித்துவமான இனங்களாக அங்கீகரிக்க நாங்கள் முன்மொழிகிறோம்.



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New localities and sexual dichromatism in Blue-green eyed Spotted Cuscus Spilocuscus wilsoni Helgen et Flannery, 2004 (Mammalia: Marsupialia:

Phalangeridae) from Biak Island, Indonesia

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Abstract: Spotted cuscuses are medium-sized tree-dwelling mammals native to tropical forests of Australo-Papua that primarily feed on fruits and leaves. They belong to the phalangerid genus *Spilocuscus* (Gray, 1862). The difference in pelage color between male and female *Spilocuscus wilsoni* has not yet been well described morphologically. In the present study, we describe the coat color of four *S. wilsoni* individuals: a male adult, a sub-adult male, a sub-adult female, and a juvenile female. Dorsal, lateral, and ventral body section images were captured on camera, and body weight & length, tail & ear length were measured. The adult male *S. wilsoni* had brown spot and blotch patterns on the dorsal and lateral regions, and the ventral region was plain beige. The sub-adult male had distinct spot patterns without blotches on the dorsal and lateral regions, while the ventral region was plain with a cream base color. The sub-adult female had a mottled pattern that blended with the base color, making a silvery appearance. The female juvenile was spotless throughout, with a foundation hue ranging from creamy to somewhat yellow.

Keywords: Biak Island Spotted Cuscus, coat color, medium-sized tree-dwelling mammal, morphologically describe.

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Competing interests: The authors declare no competing interests.

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Author contributions: AMY: field work, conceptualization, and manuscript writing; KAW: conceptualization, manuscript writing, and review; TAW, HI, and BS: review.

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INTRODUCTION

Spotted cuscuses are medium-sized tree-dwelling marsupials that primarily feed on fruits and leaves and are native to tropical forests of Australo-Papua. They belong to the phalangerid genus *Spilocuscus* (Gray, 1862). Previously, Biak Island's Spotted Cuscus was considered belonging to the group *S. maculatus* (Flannery 1995b), but later it was separated as a new species *Spilocuscus wilsoni* Helgen et Flannery, 2004. Among all the species of Spilocuscus, this is the only one that possesses blue-green eyes. *S. wilsoni* is one of the smaller species from genus *Spilocuscus* that coexist with *S. maculatus* on Biak Island.

S. wilsoni is found exclusively on the oceanic islands of Biak-Supiori, located in the northern region of New Guinea. Biak-Supiori Island has an area of 2,497 km², located off the coast of Sahul, which has no connection with mainland New Guinea. The endemicity and restricted range of S. wilsoni make it 'Critically Endangered' on the IUCN Red List (Aplin & Helgen 2016). The description of S. wilsoni was based on two samples; a juvenile male (holotype) and an adult male (paratype) from the Rijksmuseum van Natuurlijke Historie, Leiden, Netherlands (RMNH) (now Naturalis) (Helgen & Flannery 2004). Furthermore, an immature individual of unspecified sex, residing as a domesticated animal within a family setting on Biak, was captured in photograph by Flannery 1992, and constitutes an additional paratype (Helgen & Flannery 2004).

The description of *S. wilsoni* is based on craniodental characters and coat color diagnosis of an adult male (paratype) on dry skin. The adult paratype has a pure white coat dorsally and ventrally, shared only with *S.m. maculatus* of northern New Guinea (Helgen & Flannery 2004). The immature holotype is known to be male, however, there is no information about coat color and body size (Helgen & Flannery 2004).

The pelage color description is essential in the identification of species and individuals. Although the identification of species involved adult individuals, immature individuals also need to be known because the pelage colors of mammals are not necessarily fixed throughout their lifetimes (Caro & Mallarino 2020). Baby marsupials, including cuscus, have pink skin and very little hair, and most weigh less than 0.01% of the mother's weight at birth (Hughes & Hall 1988). The difference in pelage color between male and female *S. wilsoni* has not yet been well described morphologically. We found that sub-adult *S. wilsoni* show sexual dicromatism (mottled in females versus spotted and pale color in males), a

limited phenomenon among mammals (Caro 2009) but not unusual among cuscuses (Flannery 1995a,b; Caro 2013).

In this study, we describe the coat color pattern of an adult male, sub-adult male, sub-adult female, and juvenile female of *S. wilsoni*. We also measured the body and marked the location where *S. wilsoni* was found for this distribution data.

MATERIAL AND METHODS

This study was conducted from July 2021 to October 2021. Four individuals of S. wilsoni from Biak represented the age categories of adult male, sub-adult male, subadult female, and juvenile female, one individual, respectively. Information on the origin of the cuscus habitat was obtained directly from a local resident for 1, 2, and 3 and from a keeper in the Biak Bird and Orchid Park for 4 (Figure 1). The subjects were recorded using Canon EOS 750D digital camera with Canon lens EF-S 18-55 mm/F3.5-5.6 (Canon, Tokyo, Japan) in the RAW format. Photographs of the dorsal, lateral, and ventral areas are made in the same frame with a color checker passport. S. wilsoni body measurements include weight (W), head-body length (HBL), tail length (TL), and ear length (EL). All length measurements are in centimeters and weight measurements are in grams. This study has received approval from the IPB Animal Ethics Commission (Number 207-2021 IPB).

RESULTS

Adult male

The adult male appears to have a creamy base color and brown spotted and blotched morphs. The spotted and blotched morphs dominate the dorsum and flank, spreading from the head, back, limbs, and half of the tail. The blotch of the head is very dark, and the ears are covered with hair. The muzzle is hairless and darker in color compared to the chin area. The pelage on the foot is darker than the arms. Some ends of the hair strands look blackish and silvery in the dorsum and flank areas, while the ventral area does not. The ventral coat is creamy from the chin and belly to the limbs. This individual has blue-green eyes. The body measurements are as follows: W = 2,480 g; HBL = 46.5 cm; TL = 44 cm; and EL = 2.5 cm. This individual was found in the secondary forest around Warsa village, northern Biak (Image 1).

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Figure 1. Original habitat of Spilocuscus wilsoni.



Image 1. Adult male of Spilocuscus wilsoni. © Yohanita AM, 2021.

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Sub-adult male

The sub-adult male appears to have a more dominant creamy base color, and the spotted morph is brown. Spotted morphs spread from the head, back, limbs, and half of the tail but is less than that of adults. The blotch area of the head is brown, the area from the cheeks to the chin is creamy, and the ears are covered with hair. The pelage on the foot is dark. The ventral coat is creamy and a little orange in the chest area. This individual has blue-green eyes. The body measurements are as follows: W = 1,300 g; HBL = 36 cm; TL = 35 cm; and EL = 2 cm. This individual was found in the secondary forest around Makmakerbo Village, eastern Biak (Image 2).

Sub-adult female

The whole body of this sub-adult female is a mixture of creamy, light brown, and dark brown colors. The hair of the head area is a mixture of light brown on the face and dark brown on the head. The ears are covered with brown hair. The hair on the dorsum and flank areas has a mottled pattern of creamy and dark brown, while the tail area is light brown. The ends of the hair strands on the dorsum and flank areas appear silvery-buff hairs. The pelage on the foot is darker than the arms. The creamy-colored ventral area looks like a coat from head to legs. The belly part has an unopened sac. This individual has blue-green eyes. The body measurements are as follows: W = 1,100 g; HBL = 33 cm; TL = 30.7 cm; and EL = 1.5 cm. This individual was found in the secondary forest around Swandiwe Village, western Biak (Image 3).

Juvenile female

The whole body of this juvenile female looks creamy and unspotted (dorsum, flank, and belly visible). The hair on the head is thinner than that on the body. The muzzle is hairless and pink, and the ears are covered with light yellow hair. The dorsum and flank parts, including the legs, are creamy and look a little yellow in the upper back. Some ends of the hair strand on the dorsum and flank areas appear blackish and silvery. The belly part has an unopened sac. This individual has yellow-green



Image 2. Sub-adult male of Spilocuscus wilsoni. © Yohanita AM, 2021.

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Image 3. Sub-adult female of *Spilocuscus wilsoni*. © Yohanita AM, 2021.



Image 4. Juvenile female of Spilocuscus wilsoni. © Yohanita AM, 2021.

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Species	Pelage colo	luis color		
Species	Male	Female		
S. wilsoni	Adult males unspotted are yellowish white (Helgen 2007). Sub-adult has a more dominant creamy base color, and the spotted morph is brown. The adult male has a creamy base color and, more brown blotch that are dominant in the dorsum and flank (present study).	Sub-adult female is red-spotted (ZMB 91706) (Helgen 2007). The juvenile is creamy and unspotted in the entire body (dorsum, flank, and belly visible). Sub-adults have a mottled pattern that looks like a mixture of creamy, light brown, and dark brown colors (present study).	Blue-green (Helgen & Flannery 2004). Yellow-green (juvenile) and blue-green (sub-adult and adult).	
S. papuensis	The complete lower surface and base color of the dorsum are creamy in apperance, while the spots covering the back, head, and limbs are dark brown or blackish. Frequently, the upper surface of the body is washed with a yellowish hue and commonly exhibits lighter patches of yellow and brownish speckles, while the tail is typically characterized by gold or red-brown spotting. The markings are more prominent and blotchy in males (Helgen 2007).	The female has the same basic coloration and spot pattern all over the body as the male. However, the spots are smaller and more discrete (Helgen 2007).	Carmine-red (Jentink 1885); as either brown or hazel (Flannery 1994, 1995b; Gray, 1862).	
S. maculatus from northern and western New Guinea	Mature males typically have a yellowish-white or orange hue, often featuring substantial orange and white spots or blotches on their mid-back (Helgen 2007).	Mature females exhibit colors ranging from yellowish-white to orange, and some may have yellowish-white coats with orange markings (Helgen 2007).	Brown to hazel (Flannery, 1994, 1995a, 1995b; Gray, 1862).	
S. rufoniger	The dorsum of the animal has a creamy base-color with a superimposed pattern of intensely blackish (ranging from jet-black to maroon) spots or blotches on the mid-back and hind legs. The face, head, nape, shoulders, hands, feet, and sometimes the body of the limbs are covered in a vibrant red-orange or golden fur. The tail can be either golden or whitish, and the underside of the throat and chin is typically white fur that often extends as a crescent shape to the cheeks and ears, creating a striking contrast with the intense color of the head (Helgen 2007).	The female displays a coloration identical to that of the male, with the exception of the absence of black spots on the back. Instead, they have a black saddle marking that extends over the mid-back and hind limbs (Helgen 2007).	Brown to hazel (Flannery 1994, 1995a, 1995b; Gray 1862).	

eyes. The body measurements are as follows: W = 825 g; HBL = 30 cm; TL = 28.5 cm; and EL = 1.5 cm. This individual cuscus was found in the secondary forest around Warbekwan Village, northern Biak (Image 4).

DISCUSSION

Our observation of the coat colors of four individual S. wilsoni showed differences in pelage color patterns between males and females. The female had a mottled pattern throughout the dorsal and lateral to ventral edges and appeared to be wearing a coat. The male had a spotted and blotched pattern on the dorsal and lateral areas, while the ventral area was unspotted. We conclude that the sub-adult S. wilsoni shows sexual dichromatism (mottled in females versus spotted and pale color in males). Some cuscuses have spots or dorsal stripes; the spotted cuscuses S. maculatus and S. rufoniger show sexual dichromatism as females lack spots (Flannery 1995a; Helgen & Flannery 2004; Caro 2013), except for S. papuensis in which both males and females had spots (Table 1). The spotted cuscus has a unique color, especially in females, and it is recorded that four

species inhabit the mainland and islands of Papua. The female *S. maculatus* in the northern islands is plain yellowish-white, while in mainland Papua it is yellowish-white with orange markings from mid-back to the abdomen. Furthermore, the *S. rufoniger* female displays black saddle markings that cover both the mid-back and hind limbs (Helgen 2007).

The sub-adult and adult males in this study showed a brown spotted pattern. Nevertheless, the spots on the sub-adult individual are smaller and more discrete, so the beige base is more dominant. In adults, a blotch on the head extends to the upper back to the forelimbs, and a blotch on the lower back area to the tail and hind limbs; therefore, the brown blotch is more dominant. S. wilsoni juvenile female has a plain cream coloration all over the body, and it seems that pale, plain colors are common among juveniles of Spilocuscus. The colors of mammal pelage may not remain constant over their entire lifespan (Caro & Mallarino 2020). Certain pigs and peccaries experience age-related transformation; for example, they are born with spotted and striped coats that eventually become consistent as their young become mobile (Caro et al. 2018).

The immature S. wilsoni photographed by Flannery

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in 1992 was of unknown sex and was used as additional paratype information (Helgen & Flannery 2004). We saw similar color patterns between the photo and *S. wilsoni* in this study (Image 3), and we conclude that its morphology belongs to the sub-adult female individual. We also found differences in the iris color of juvenile (yellow-green) and sub-adult or adult individuals (blue-green), but this needs further investigation.

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Indonesian Abstract: Kuskus bertotol adalah penghuni pohon berukuran sedang berasal dari hutan tropis Australia-Papua yang memakan buah dan daun. Kuskus bertotol termasuk ke dalam famili Phalageridae dan genus Spilocuscus (Gray, 1862). Perbedaan warna rambut antara S. wilsoni jantan dan betina belum terdeskripsikan secara morfologi. Pada penelitian ini, kami mendeskripsikan warna rambut dan mengukur bagian tubuh eksternal dari empat individu S. wilsoni: jantan dewasa, jantan dewasa muda, betina dewasa muda, dan betina remaia. Pengambilan foto bagian tubuh area dorsal, lateral, dan ventral dilakukan menggunakan kamera. Pengukuran tubuh meliputi bobot, panjang tubuh, panjang telinga, dan panjang ekor. S. wilsoni jantan memiliki pola totol dan bercak berwarna coklat di area dorsal dan lateral, sementara area ventral berwarna krem tanpa totol. Individu jantan dewasa muda memiliki pola totol di area dorsal dan lateral, sementara area ventral berwarna krem tanpa totol. Selanjutnya, individu betina dewasa muda memiliki pola perpaduan bintik warna coklat dan krem yang tampak seperti lurik dengan ujung keperakan. Individu betina remaja, satu-satunya yang

tidak memiliki totol maupun bercak dan keseluruhan tubuh berwarna krem sampai kuning muda.

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Nest construction and repairing habits of Baya Weaver *Ploceus philippinus* (Aves: Passeriformes: Ploceidae) in the agricultural landscape of Villupuram District, Tamil Nadu, India

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Abstract: The intricate nesting habits of Baya Weaver Ploceus philippinus were studied on two Palmyrah Palm Borassus flabellifer trees in Chendur village, Villupuram district, Tamil Nadu between 20 March and 30 November 2020. Observations concentrated on sources of fibers, developmental stages of nests, re-construction & repairing of nests, deposition of clay in the nest walls, and various threats. A total of 98 nests of various developmental stages (wad stage-4, helmet stage-31, egg-chamber closed stage-5, and complete nests-58) were studied on these two nest colonies. The birds used leaf fibers of Indian date Palm Phoenix sylvestris and Sugarcane Saccharum officinarum as nest materials, and took 6-48 days for construction of a complete nest. 95% of helmet stage nests (n = 126) contained clay deposits. Analysis of plastered clay revealed it was alkaline with pH 9, and dry weight ranged from 5.1–5.8 g. Males re-constructed new nests from the remnant stalks attached to tips of palm fronds, and also made repairs on damaged nests. Anthropogenic factors, wind, rain, and avian predators, such as House Crow Corvus splendens, Long-billed Crow Corvus macrorhynchos, Rufous Treepie Dendocitta vagabunda, and Shikra Accipeter badius posed threats to Baya Weavers. A total of 42 nests, 11 broken eggs, and 14 dead chicks were found fallen under the two nest supporting trees. The detailed systematic survey covering entire district, rapid urbanization, and the anthropogenic pressures will help in drafting an action plan to conserve local populations of Baya Weaver.

Keywords: Clay deposit, nest fall, nest materials, nest re-construction, nest repair, threats.

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INTRODUCTION

Baya Weaver Ploceus philippinus (Linnaeus, 1766) is a social, polygamous, colonial nester. It occurs in the Indian subcontinent (Ali et al. 1956), Java, Malacca and Sumatra (Blyth 1845; Wood 1926), China, Indonesia, Laos, Myanmar, Singapore, Thailand, and Vietnam (BirdLife International 2016). The IUCN Red List of Threatened Species classifies Ploceus philippinus under 'Least Concern' (Birdlife International 2016). In India, the breeding season of Baya Weaver is from May-November (Ali & Ripley 1987; Rasmussen & Anderton 2005). While weavers select a variety of trees for nesting, they most prefer tall, unbranched trunks, and long-swaying foliage of palm trees to keep away predators and provide convenient leaf strips for building nests (Davis 1974). Males usually build partial helmet stage nests and complete them only after females select them and mate (Ali et al. 1956). Nesting birds prefer Cocos nucifera (Arecaceae) along the west coast of the Indian peninsula, B. flabellifer (Arecaceae) along the east coast, and Vachellia nilotica (Fabaceae) in the arid northwestern region (Sharma 1989). The breeding biology of this species was studied by Ali & Ambedkar (1956), Ambedkar (1964), and Mathew (1977). Several researchers have reported construction of abnormal nests (Ali & Ambedkar 1956; Ambedkar 1964; Crook 1964; Sharma 1989; Pandian 2018). Asokan et al. (2008) studied the timeline of nest construction. No other detailed studies have been carried out on the time taken for construction of various stages of nests, nest repairing, nest reconstruction, and physico-chemical analysis of clay deposited in the helmet stage nests of this species in Tamil Nadu. To fill this gap the present study was carried out.

The following questions were kept in mind: (1) How do weavers select substrata on the trees for nest construction? (2) What are the sources of nesting material? (3) What are the developmental stages of nests like wad, helmet, egg-chamber closed and complete nests, re-construction of nests and repairing of damaged nests? (4) How much time is taken to build various stages of nests? (5) What is the physico-chemical nature of clay deposits? And, (6) what are probable threats to weaver populations?

MATERIALS AND METHODS

Study Area

The present study was carried out in Chendur Village, Tindivanam Taluk, Villupuram District of northeastern Tamil Nadu. The district spreads over 3,715 km², with a human population of c. 2,090,000 (Figure 1). Agriculture is the primary occupation of the people. The major crops of the area are Paddy Oryza sativa, Jowar Sorghum bicolor, Pearl Millet Pennisetum glaucum, Finger Millet Eleusine coracana, Sugarcane Saccharum officinarum, Groundnut Arachis hypogaea, and Green Gram Vigna radiata. Three nest-supporting plant species, such as Palmyrah Palm, Indian Date Palm, and Coconut occur abundantly in the agricultural lands. Among them, only two individuals of B. flabellifer were chosen for study, considering the past history of Baya Weavers selecting these two trees for nest construction, proximity to road, and location of trees suitable for study by fixing camera. The maximum and minimum temperatures in the district are 36°C and 20°C, respectively. The average annual rainfall is 1,060 mm (Viluppuram 2021).

Methods

With help from field assistants/informants (4), I identified two B. flabellifer nesting trees in Chendur village having a history of Baya Weavers constructing nests since 2016. These two nesting trees were surveyed with the help of field assistants on two shifts, i.e., one person each from 0600 h to 1200 h and 1200 h to 1800 h on a daily basis between 20 March and 30 November 2020. All the nests found in these two trees were treated as nest colony-I and colony-II. The height of the trees, gbh, and distances from nesting trees with source of fibers, distance of source of wet clay, and cultivation of grains crop were measured using a 100 m measuring tape. The locations of nesting trees were determined using GPS. Using 10 x 42 field binoculars (Nikon-Monarch-7), the nests, males plucking nest fibers, developmental stages of nests, clay deposits on inner wall of helmet stage nests, and the total number of birds visiting nesting trees were observed. Clay deposits from helmet stage nests were collected separately from each side of inner walls of fallen nests and analyzed. The physico-chemical analyses of collected clay samples including temperature, salinity, dissolved oxygen, oxidation reduction potential, specific conductivity, electrical conductivity, total dissolved solids, and other metals & minerals were carried out by using YSI multiparameter (Model: 600XL-B-O, 650MDS, YSI Incorporated, Yellow Springs, Ohio 45387, USA) and Perkin Elmer Atomic Absorption Spectrophotometer

Pandian



Figure 1. Study area map: a—India map showing Tamil Nadu | b—Tamil Nadu map showing Villupuram district | c—Villupuram district map showing locations of two nest colonies in Chendur village.

Analyst 400. The recorded results were tabulated (SPSS software). Re-constructions of nests, repairing of damaged nests, fall of nests, eggs & chicks, and impact of avian predators were recorded and photographed. Photographs were taken using Nikon P1000 digital camera. Collected data were tabulated, analyzed as total number of fronds used by the birds, average number of nests per frond and shown as tables.

RESULTS

It was observed that no old or torn nests from previous years were found on these two male *B. fabellifer* trees when the study was commenced on 20 March 2020 (Image 1). Baya Weavers constructed a total of 98 nests (Wad stage—4, helmet stage—31, egg-chamber closed stage—5, and complete nests—58) on two male *B. fabellifer* trees. Birds failed to continue constructions on four wad stage nests, and 31 helmet stage nests and

five egg-chamber closed stage nests. The remaining 58 nests were complete nests with entrance tubes. The study revealed that the birds built an average of 1.17 nests per palm frond. Nest colony-I contained 62.24% nests (n = 61) and the remaining 37.76% nests (n = 37) were found in nest colony-II (Table 1).

Commencement of nest construction

All the males had commenced nest constructions on 02 April 2020. From 20 March to 01 April, no Baya Weaver was observed on these two nesting trees. On 02 April between 0600 h and 0830 h, 16 males with partial plumage first visited on these two palm trees probably searching for suitable substrata for construction of nests.

Selection of Palm fronds

Males visited distal ends of palm fronds randomly, except the lowermost dried and the uppermost partially opened young fronds. After selection of the distal ends of fronds, they bit the margins by using their beaks and made the margins serrate/rough and also made punctures on the leaf blades probably to make the knots strong. Out of 248 leaf tips studied, the margins of 232 leaf tips was serrate, and 16 leaf tips had serrate as well as punctures. This process of making frond margins serrate continued for five days, i.e., from 02-06 April and the males were observed on nesting trees between 0600 h and 0830 h. During these periods, no activities of nest construction were noted. After 0830 h, they left the nesting trees for foraging in the crop fields and perching on nearby tress/shrubs. They did not return to the nesting trees till the next morning.

Sources of fibers

Males plucked fibers from Indian date palm P. sylvestris trees (n = 6) and Sugarcane crops S. officinarum situated within c.120 m distance from the nest-supporting trees. Males moved to west to puck fibers from *P. sylvestris* and to all directions to get fibers from S. officinarum crops. The study revealed that the males had visited P. sylvestris trees daily from 0600 h to 1730 h, perched on rachis/leaflets, made incisions on the margins of leaflets near the bases and speedily tears off fine fibers toward the distal ends. The birds tore off fibers in this manner and carried to the nesting trees. Observations on 100 trips from the sources of fibers (P. sylvestris and S. officinarum) to nest-supporting trees revealed that the birds carried 2–5 fibers per trip. They selected young fronds for peeling fibers and avoided the old fronds on the bottom of the tree crown. Study on 10 fronds from four P. sylvestris trees where birds tore

off fibers revealed that the mean size of fronds was 106 cm (Standard Error of 4.73) and the birds had selected leaflets from the distal half of rachis, i.e., from middle to distal part of the rachis and never selected hardened leaflets found on the lower half of rachis, i.e., towards leaf bases. Similarly, males tore off fibers from young and green leaf blades of sugarcane crop and no incident of selection of fibers from dried and partly dried leaves were observed (Image 2).

Behaviour of stealing fibers

Thirty-two incidents of males stealing fibers from adjacent nests when the resident birds of the nests were away were observed. Such incidents of stealing fibers from nests of other birds within the colony were observed throughout the breeding season.

Stages of nest constructions

Four developmental stages of nest constructions such as wad stage, helmet stage, egg-chamber closed stage, and complete nest stage were taken into account and studied in detail (Image 5).

Wad stage

The males plait knots round the margins of leaf blades by using legs and beak called wad stage (Image 5a,b). The study on 98 wad stage nests revealed that the time taken for construction of wad stage varied and the males took minimum two hours to maximum of nine days for construction of wads. In an exceptional case, a male plaited knot continuously for nine days and the wad stage became an amorphous ball like structure. The males usually plaited knot on one leaf tip, but in many cases they used up to six leaf tips for plaiting a knot/wad stage (Image 3). The males built 22.45% wad stage nests (n = 22) in 1–2 days, 54.09% wad stage nests (n = 53) in 3–4 days, 19.38% wad stage nests (n = 19) in 5–6 days, and 4.08% wad stage nests (n = 4) in 7–9 days.

Helmet stage

The males took 1–15 days to construct helmet stage nests. Out of 98 wad stage, 94 were developed into helmet stage nests (birds abandoned 4 wad stage nests). When the females reached the nesting colony, the males perched on helmet stage nests and made loud noises by flapping their wings (Image 4 d). The males built 84% helmet stage nests (n = 79) in 1–5 days (including 27 nests were built within one day from dawn to dusk), 13.8% helmet stage nests (n = 13) in 6–10 days, and 2.1% helmet stage nests (n = 2) in 11–15 days.

Nest construction and repairing habits of *Ploceus philippinus* in Villupuram District, India

Nesting trees with GPS	Height (m)	GBH (cm)	Total no. of fronds found in the crown	No. of fronds without nests	No. of fronds used by birds for construction of nests	Total no. of nests (including all developmental stages) counted	Average no. of nests per frond
Borassus flabellifer (colony-I) (12.123446 N 79.591657 E)	7.5	78	54	30	24	61	0.88
Borassus flabellifer (colony-II) (12.113396 N 79.580264 E)	9.5	82	61	25	18	37	1.64
Total			115	55	42	98	1.17 (average)

Table 1. Details of fiest colonies of baya weaver of bolassus habennet fiees in chemuu village, villubulant ustrict



Image 1. Nest colonies on Borassus flabellifer trees in Chendur village: a-Nesting colony-I | b-Nesting colony -II. © M. Pandian.



Image 2. Baya Weaver: a—Male | b—Female | c & d—Male makes Borassus flabellifer frond margins rough around midrib | e & f—Male bird plucking fibers from leaflets of Phoenix sylvestris. © M. Pandian.

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Table 2. Details of number of days taken by the birds to construct complete nests.

Number of complete nests	Percentage (%)	Number of days taken
02	3.4	1–5
18	31	06–10
20	34.4	11–15
08	13.8	16–20
03	5.1	21–25
7	12	Above 26

Egg-chamber closed stage

eb.

Out of 94 helmet stage nests, 31 helmet stage nests were abandoned by the males and did not develop further probably due to non-selection of helmets by females or abandoning by resident males themselves. The remaining 63 helmet stage nests were developed further into egg-chamber closed stage nests. After construction of helmet stage nests, males wait for arrival of females for selection of their helmet stage nests followed by pairing. Hence, further development of nests depends on the chances of selection of helmet stage nests by females and time taken for such selection followed by pairings. My studies revealed that the number of days taken for development of helmet stage nests including the time taken for arrival of females, selection of helmets, and followed by pairing were found varied from one day to 29 days. Out of 63 helmet stage nests, 35 helmet stage nests were developed into eggchamber closed stage nests in 1-5 days, followed by 20 nests in 6–10 days, four nests in 11–15 days, three nests in 16–20 days and one nest took 29 days (Image 4e).

Complete nests

The birds abandoned five egg-chamber closed stage nests without any further development. Birds took 1–28 days to complete the construction of entrance tube. Out of 58 complete nests studied, in 69% nests (n=40), the entrance tubes were constructed in 1–5 days, while in 12 nests, it took 6–10 days, 11–15 days for four nests, and 21–28 days for two nests (Image 4f; Table 2). During the entire study period, neither courtships nor mating were observed on the helmet stage nests or source of fibers or on nest-supporting trees.

Deposition of clay in the nests

After completion of construction of helmet stage nests and before arrival of females to select such nests, the males plastered two sides of the inner walls of



Image 3. Nests attached on tip of palm frond: a—Nest attached with two leaf tips | b—Nest attached with four leaf tips. ${\ensuremath{\mathbb C}}$ M. Pandian.

helmet stage nests with wet clay. Observation on 132 helmet stage nests (94 first time built helmet stage nests and 38 re-built helmet stage nests) revealed that 95 % of nests (n = 126) contained clay deposits. Only a very small percentage (4.65%; n = 6) did not have clay. My studies revealed that the males did not take readily available wet clay from the paddy fields, situated c.300 m from the two nesting trees. Males waited for the frequent spell of rainfall during south west monsoon. Immediately after rainfall, the next day morning between 0600 h and 0730 h the males swarmed to the wet fallow land and mud road situated c.40 m distance from the nesting trees and scooped wet clay through their beaks in many trips and carried it to helmet stage nests. Continuous observations revealed that the males did not take clay directly from wet soil surfaces from all the sites. They selected sites where wet clay was exposed in tire tracks left by vehicles on mud roads and fallow land. The practice of males scooping clay after rainfall was observed from April-October 2020 in the morning between 0600 h and 0730 h. It was not possible to ascertain whether the birds added clay on the inner walls after closing of egg-chamber and construction of entrance tube. Dissection of two fallen nests (helmet stage—1 and egg-chamber closed—1) revealed that the

Tab	le 3	. Details	of properties	of clay (deposited	in the nests.
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Parameters	Soil sample collected from the site where Baya Weaver took soil	Egg-chamber closed stage (Left wall)	Egg-chamber closed stage (Right wall)	Helmet stage (Left wall)	Helmet stage (Right wall)
Weight (g)	25	5.1	5.2	5.4	5.8
Temperature (°C)	25.91	26.01	25.93	26.02	25.92
Specific Conductance (Ms/Cm)	0.048	0.048	0.035	0.051	0.024
Conductivity (Ms/Cm)	0.049	0.049	0.036	0.051	0.025
Resistivity (Ω Cm)	22193.3	19805.2	27440.7	19490.3	40131.3
Total Dissolved Solids (TDS %)	0.031	0.033	0.023	0.032	0.016
Salinity (Sal)	0.02	0.02	0.02	0.02	0.01
Dissolved Oxygen (DO %)	32.4	37.2	23.7	30.7	19.4
Dissolved Oxygen milligrams per litre (DO Mg/L)	2.72	2.94	1.94	2.42	1.89
Dissolved Oxygen charge (DO Ch)	15.5	16.5	13.5	15.5	12.4
Potential of Hydrogen (pH)	9.36	9.30	9.30	9.23	9.30
Potential of Hydrogen in Milli Volt (pHml/)	-179.3	-175.0	-175.9	-172.4	-176.4
Oxidation Reduction Potential (ORP)	-110.4	-103.5	-103.4	-99.7	-102.1

Table 4. Details of month-wise nest fall from two nesting trees.

Month	No. of nests felled down
May 2020	11
June 2020	13
July 2020	17
August 2020	1
Total	42

Table 5. Details of impact of avian predators on Baya Weaver colonies in Chendur village.

	Name of the predator	No. of sightings noted	No. of nests damaged	No. of Baya Weaver killed
1	Corvus splendens	72	3	0
2	Corvus macrorhynchos	27	0	0
3	Dendrocitta vagabunda	4	4	0
4	Accipiter badius	7	0	1
	Total	110	7	1

males smudged two patches of clay on each side of the nest. The surface of dried plaster of clay had many beak marks as scars (Image 5f). It was observed in one eggchamber closed stage nest that even after plastering of clay, another layer of fresh fibres was found on the patches of clay. It indicates that even after smudging clay, males further added fibres above the layer of clay (Image 6e). No female was seen scooping clay and carrying to nests (Image 5).

Physico-chemical analysis of clay deposits taken from one helmet and one egg-chamber closed stage nests revealed it was alkaline (pH 9). The dry weights of the clay ranged 5.1–5.8 g. The other parameters also showed no major variations. The physical and chemical properties of clay collected from both walls of two nests matched with the soil sample collected from the nearby ground where male birds scooped clay (Table 3).

Falling of nests

Of 98 nests constructed during the study period, 43% (n = 42) of various developmental stages (helmet stage—22, egg-chamber closed stage—03, and complete nests—17) fell from the nest-supporting trees due to biotic and abiotic factors. 31 nests fell after rainfall, and the remaining 11 fell when no rainfall occurred (Table 4).

Re-construction of nests

The males started to rebuild 38 nests from the remnants of wad fibers found attached to the tips of palm fronds. Twenty-three nests were developed into helmet stage nests but did not progress further. The remaining 15 nests were successfully developed into complete nests. When studying the number of days taken to re-built a complete nest revealed that the birds took 6–37 days to re-built complete nests. The study reveals that the birds had constructed 13 complete nests in 6–25 days and for another two nests took 26 days and

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Image 4. Various stages of nest development: a —Male individual brought fiber to tie a knot | b—Male bird perching on wad stage nest | c— Ring stage nest | d—Helmet stage nest | e—Egg-chamber closed stage nest | f—Complete nest with entrance tube. © M. Pandian.

37 days, respectively.

One re-built helmet stage nest was again felled down on 21 May and a male had started to construct another helmet stage nest at the same tip of frond on 24 May and completed the construction of helmet stage nest on 25 May. Later the helmet stage nest did not develop further. On 30 June, another re-built helmet stage nest was felled down and a male had again constructed helmet stage nest from the same tip of palm frond within two days i.e., on 01 and 02 July. Later in nine days, i.e., on 11 July and it was developed into a complete nest (Image 6).

Repairing of damaged nests

Incidents of partial damages to seven nests (eggchamber closed stage—4, and complete nests—3) by House Crow and Rufous Treepie were recorded in the study area. In all these nests the birds brought fresh plant fibers and plait on the edges of damaged walls. Then the birds had continued further construction activities and repaired all the nests. The repaired nests resembled two different colours, i.e., the older part resembled pale colour and the repaired portion resembled green colour due to the addition of fresh green fibers (Image 7).



Image 5. Plastering of wet clay on the inner walls of helmet stage nests: a & b—Males engaged in collection of clay from wet ground | c—Plastering of clay on the inner wall of helmet stage nest | d—Clay deposit found in dissected nest | e—Layer of fibres found above clay deposit | f—Beak marks on clay deposit. © M. Pandian.

Threats

Opportunistic sightings of predatory birds such as House Crow *C. splendens*, Long-billed Crow *C. macrorhynchos*, Rufous Treepie *D. vagabunda*, and Shikra *A. badius* were observed on these two nesting trees. On 24 July at 1240 h, one Shikra had chased the individuals of Baya Weaver from nesting colony-I and when seeing the predator, all the individuals of Baya Weaver fled from the nest colony. One male Baya became a prey to Shikra and later took the victim to a nearby shrub and ate it completely, except feathers (Image 8 j). On 11 June, one Rufous Treepie visited nesting colonies and made punctures on the egg-chambers. The predator had inserted its head into the egg-chamber but we were unable to ascertain whether it prey upon the eggs/chicks from the nests. Individuals of House Crow and Largebilled Crow were found perching on nesting trees and chased the individuals of Baya Weaver but preying adult birds/chicks were not observed during the study period. On 28 May, a House Crow plucked fibers from three nests (complete nests-3) and caused partial damages to the nests (Table 5).

On 30 September, one land holder had uprooted and removed *P. sylvestris* trees (n = 14) found on bunds of fallow lands situated 60 m from nesting trees while clearing the land for cultivation. The males had plucked fibers from these trees for construction of nests. A total of 42 fallen nests, 11 broken eggs, and 14 dead chicks

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Image 6. Re-construction of nest: a—Stalk of the fallen nest | b—Plaiting green fresh fibers on pale old stalk | c—Remnant stalk of old nest | d—Male plaiting green fresh fibers around old stalk | e—Re-construction of new helmet from old stalk | f—Re-constructed complete nest. © M. Pandian.

were observed under the nesting trees. In one instance, farmer burnt bushes around nesting tree which caused temporary driven of birds from nesting tree (Image 9).

Roosting

During the entire breeding period from April– November, no Baya Weaver was found night roosting on the nest-supporting trees. Between 1745 h and 1810 h all the birds used to fly away from the nest colonies and roost on the shrubs found 1–1.5 km from nest colonies and return to their nest colonies the next morning. Continuous observations revealed that some females entered their nests during the evening hours did not come out and stayed in the nests itself. These females might have incubated their eggs or nestlings.

DISCUSSION

Commencement of nest construction

The study revealed that starts of the breeding season of Baya Weaver vary from area to area in India. For example, date of commencement of nest construction was 22 May 1930 in Kolaba district of Maharashtra (Ali 1931), early June in Parbati Hill, Poona city (Ali et al. 1956), mid-June in Chorao Island, Goa (Borges et al. 2002), and mid-April in Rajampet Taluk of Cuddapah district, Andhra Pradesh (Mathew 1972). In the present



Image 7. Nest damage and repair: a—Male perching on damaged nest | b—View of damaged nest | c—Repairing of damaged nest | d—Further growth of nest. © M. Pandian.

study, breeding commences in the first week of April which corroborate the observation of Mathew (1972). It indicates that the breeding of Baya Weaver is related to South-West monsoon in Tamil Nadu.

Selection of Palm fronds

Baya Weavers construct nests from the distal ends of midribs of the coconut palm in South Goa (Borkar & Komarpant 2003). The birds wrap considerable amounts of fibers around a chosen frond and makes a strong base for further development of the nest (Wood 1926). Also in the present study nests were found attached at the distal ends of the midribs of *B. flabellifer* fronds. This indicates that the males selected the distal ends of palm fronds for construction of nests. No nest was found in the middle or basal parts of fronds in the two nest colonies.

Source of fibers

Except in northern India, the birds used leaf fibers of *C. nucifera* and *P. sylvestris* for nest construction in other parts of the country (Dewar 1909). Baya Weaver used pliant grass and fibers from palm fronds in the Northern Province of Sri Lanka (Wood 1926), and leaves of *Phoenix* sp., coarse grass and paddy in Kolaba district, Maharashtra (Ali 1931) as nest materials. The present findings of Baya Weaver using fibers from *P. sylvestris* for construction of nests matches with the observations of Dewar (1909), Wood (1926), and Ali (1931). Apart from that the birds also used sugarcane leaves as nest material in the study area.

Time taken for construction of nest

Asokan et al. (2008) stated that the birds took 18 days to construct one complete nest in Nagapattinam and Tiruvarur districts of Tamil Nadu. Achegawe et al. (2016) had also found similar results in Nanded region of Maharashtra. The present study revealed that the time taken for construction of a complete nest was not uniform for all the nests. The birds took 6–48 days for construction of complete nests. The study also revealed that the males had capable of constructing helmet stage nest in one day, i.e., from dawn to dusk. Hence, the present findings of number of days taken to construct complete nest in Villupuram district found differ with the observations of Asokan et al. (2008)

Plastering of inner nest walls with wet clay

The habits of smudging of clay in the nests are observed only in Asian weavers (*P. manyar, P. benghalensis,* and *P. philippinus*) and not found in African



Image 8. Various threats to Baya Weavers: a & b—Nests dangling from the stalk | c & d Fallen nests | d—Broken egg near fallen nest | e–g– Dead chicks under the nesting tree | i—Bushes burnt under nesting tree | j—Shikra eating a male Baya Weaver individual. © M. Pandian.

weavers and the quantity of mud varies from region to region in India (Crook 1963; Davis 1973). Plastering of inner walls of nest with wet clay is done when the nest construction reaches the helmet stage prior to pairing with females (Dewar 1909; Ali 1931; Ambedkar 1964; Borkar & Komarpant 2003).

Davis (1973) stated that about 18.33% nests did not show presence of mud blobs on the inner walls of nests in South Goa. He added that females were never found bringing mud. In the present study also, 4.54 % of helmet stage nests did not have clay deposits, and no female was observed carrying clay. This matches with the findings of Davis (1973). Ali (1931) had observed 0.5-1 oz of mud in the nest. Davis (1973) also found that the average dry weight of mud deposit per nest was 66.2 g. But in the present study, the dry weight of clay ranged from 5.1–5.8 g. Average weight of mud blobs on left side was greater in comparison with right side (Borkar & Komarpant 2003). The present study on two nests revealed that there was no major variation in the weight of clay deposited on the right and left side of the inner walls. In the present study, the clay collected from nests and in the original sites where birds took clay was found alkaline (pH 9). The exact causes of plastering of mud in the nest walls require further studies.

Fallen nests

The practice of male cutting down the nest of rival cock was common when the owner had gone to fetch building materials (Ali et al. 1956). Pandian (2021) had observed that male Baya Weaver cut down a complete nest occupied by White-rumped Munia *Lonchura striata* in Villupuram district. In the present study, 42.85% nests (n = 42) of various developmental stages were found fallen under the nest-supporting trees. Apart from rain and wind, these might have also occurred due to rival males as stated by Ali et al. (1956). Falling of such a great number of nests (42.85%) from two nest colonies in a single breeding season is of great concern and it needs further study.

Re-construction of nests

The study indicates that the birds are capable of constructing complete nests from the same stalks from where the earlier nests were detached. It was not possible to differentiate whether the same male commences construction of nest from the torn stalk or different male uses the stalk for further construction of nest. However the birds have the intelligent to re-construct their nests from the stalk of detached nests.

Repairing of damaged nests

Baya Weavers have the capacity to repair their damaged nests with fresh green strips of fibers, i.e., various types of artificial nest mutilations (Ali & Ambedkar 1957; Collias & Collias 1959, 1962). In the present study also the birds had repaired heavily damaged nests by using fresh fibers and hence it matches with the findings of Ali & Ambedkar (1957) and Collias & Collias (1959, 1962).

Threats

Ali (1931) had stated that agitated behaviour of Baya Weaver was observed when Crow Pheasants *Centropus sinensis* appeared in close proximity of the nesting tree. He also observed a Shikra making an unsuccessful stoop on nest colony. In the present study also, Baya Weaver had exhibited agitated behaviours when House Crows visited nesting trees and all birds fled away from tree crown when they saw Shikra and a Rufous Treepie near the nesting trees.

Ali et al. (1956) had observed that many completed nests were blown down due to recurring spells of bad weather during June–August in Bombay area and was major natural mortality factor of nest colonies. He also noted accidental drowning of chicks from nests. Similarly in the present study also, 31 nests were found fallen down immediately after rainfall. Out of 14 dead chicks, five were found under the nesting trees after rain and wind. Hence, the present observations match with the findings of Ali et al. (1956).

CONCLUSION

This is a systematic study on the nesting habits on Baya Weaver on two Palmyrah Palm trees in a confined geographical area of one village in Villupuram district. Increasing urbanization by conversion of cultivated lands into residential areas, expansion of roads, abandoning cultivations along with indiscriminate felling of principal nest-supporting palm plants, such as Palmyrah Palm B. flabellifer, Coconut C. nucifera, and Indian Date Palm P. sylvestris that are vital for Baya Weaver is a conservation issue in this landscape. Increasing practices of monoculture of Casuarina, Sugarcane, and flower crops, declining areas of cultivation of cereals and millets also causes shortage of grains to birds. Fall of viable nests due to various anthropogenic factors, winds, rain and avian predators cause severe stress on the breeding of Baya Weaver. Therefore, it is essential to conduct sustained surveys and monitor the nesting sites

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during the subsequent breeding seasons and efforts should be taken to create suitable nesting habitats by not destroying the nesting trees. Based on the above the following could be options for securing the bird's habitat from the area.

(a) Establish a special management plan for the area, considering the anthropogenic and natural stresses that the habitat is currently subjected to.

(b) Local community, particularly land holders, and agricultural workers should be sensitized to understand the need to preserve the precious nesting sites.

(c) The detailed systematic survey covering the entire Villupuram district will help in drafting an action plan to conserve the populations of Baya Weaver.

(d) The impact of abiotic factors such as rain and wind on the nest fall need further studies.

(e) The exact causes of males depositing wet clay on the inner walls of nests during helmet stage, fall of large number of nests, and mortality to chicks require further studies.

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A checklist of the avifauna of Samanatham tank, Madurai, Tamil Nadu, India

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Abstract: This avifaunal checklist from the Samanatham tank of Madurai District of Tamil Nadu is the first baseline data for this wetland. The study was done from August 2015 to March 2019 and from August 2020 to January 2022. A total of 150 species of birds comprising 17 orders and 52 families were documented. Order wise, Passeriformes 35% (n = 53), Pelecaniformes 15% (n = 22), and Charadriiformes 14% (n = 21) dominated the wetland bird community. The relative diversity (RDi) of families with the most species were Anatidae, Scolopacidae, and Accipitridae with 7.33%. Among the 150 species, 37 (25%) were winter visitors; and one passage migrant Rosy Starling Pastor roseus. The relative abundance indicated that 56% (84 species) were common, 28.6 % (43 species) were uncommon, and 15.4 % (23 species) were rare. The study recorded eight globally Red Listed 'Near Threatened' species-Oriental Darter Anhinga melanogaster, Painted Stork Mycteria leucocephala, Black-headed Ibis Threskiornis melanocephalus, Spot-billed Pelican Pelecanus philippensis, Black-tailed Godwit Limosa limosa, Bar-tailed Godwit Limosa lapponica, River Tern Sterna aurantia, & Osprey Pandion haliaetu-and two 'Vulnerable' species—Indian Spotted Eagle Clanga hastata & Great Spotted Eagle Clanga clanga.

Keywords: Baseline, bird sanctuaries, shorebirds, waterbirds, wetland, winter visitors.

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INTRODUCTION

Bird diversity is high among vertebrates due to their wide distribution. They are biodiversity indicators (Canterbury et al. 2000; Piersma & Lindström 2004) and documentation of their diversity, distribution, and taxonomy in India has been widely done (Ali & Ripley 1987; Kazmierczak 2006; Grimmett et al. 2011; Deomurari et al. 2023). India's share is 12.5% of the global avifaunal richness (Grimmett et al. 2011; Praveen & Jayapal 2023). As there are continuous landscape changes due to economic development in new areas (Jha et al. 2000; Purvis & Hector 2000), regional-level monitoring of birds is essential (Gadgil 1996). Avifaunal studies help to understand the long-term changes in the landscape and the vegetative structures (Rika & Santosa 2007; Rashiba et al. 2022) which aids in conservation planning (Paul & Cooper 2005). Baseline data from any site is essential for any long-term conservation efforts (Peterson et al. 2000; Llanos et al. 2011).

Avifaunal wetland studies in Tamil Nadu had been done in many wetlands like Pallikaranai (Raj et al. 2010), Karaivetti (Gokula 2010), Vaduvoor (Gokula & Raj 2011), and Karangad (Byju et al. 2023). In Madurai the avifaunal studies are scanty and literature search revealed few older studies (Nichols 1944a,b, 1945). Recent studies done on diversity and distribution were restricted to urban landscapes (Sathasivam 2015; Rajagopal et al. 2022) or thorn forest landscapes (Roopha et al. 2022). Wetlands worldwide are on the decline and India is no exception due to human interventions (Pringent et al. 2012; Sievers et al. 2018). The wetland diversity of Madurai is less documented, except for a few scanty works mentioned from the urban areas, no detailed longterm studies have been made on the diversity of birds in the wetland areas of Madurai. In this background, the current study highlights the status, composition, feeding guilds, and diversity of birds of Samanatham tank, one of the important wetlands of Madurai, Tamil Nadu.

MATERIALS AND METHODS

Study Area

Samanatham tank 9.866674°N, 78.14719°E is located in Thiruparankundram block in the Madurai District of Tamil Nadu, India. The tank and its surrounding areas have various types of vegetation, including shrubs, trees, and aquatic plants. The main habitat types observed in the tank include: 1. Open water habitat (WL = Wetland), 2. Agricultural land (AL), 3. Trees (Tr) bordering the wetland, 4. Grassland (GL) on the wetland area, and 5. Scrub habitat (OS = Open scrub type). Some of the common trees found in the area include Neem Azadirachta indica, Mango Mangifera indica, Banyan Ficus benghalensis, Acacia nilotica, and Tamarind Tamarindus indica. The tank and its surrounding areas also have invasive species like Prosopis juliflora. The surrounding area also supports a variety of flowering plants, including Jasmine, Hibiscus, Lotus, and many others. In addition, the tank is home to a variety of aquatic plants, such as water lilies, Water Hyacinth, and Cattails. The area supports a variety of animals including reptiles like snakes & turtles, amphibians like frogs & toads, and fishes like catfish, tilapia, & carp. Overall, the flora and fauna in and around Samanatham tank are an important aspect of the ecosystem of the region and contribute to its rich biodiversity.

Madurai, located in the southern part of India, has a tropical climate with hot and humid summers and mild winters. The temperature ranges from 20° C to around 38°C. The average annual rainfall is 800 mm. The highest amounts of rainfall are during October and November.

Bird survey method

In the Samanatham tank, bird monitoring was continued for seven years (2015–2022) to maintain the baseline data. The period of observation was carried out every month from August 2015 to March 2019 and from August 2020 to January 2022. The birds were observed during the peak hours of their activity from 0600-1000 h and 1600–1800 h. Later, bird surveys were conducted using block count and direct visual count methods (Howes & Bakewell 1989; Bibby et al. 2000). In this method, six scanning points were identified (Figure 1) and birds in the blocks were observed using field binoculars (10 x 52 Olympus; Celestron outland 10 x 42) and recorded with a camera (Nikon D750, Canon 100-400 mm f/5.6 lens). We began counting five minutes after the arrival at each scanning points for the waterbirds to acclimate to the human presence. The identification of birds was done using the following field guides (Ali 2002; Grimmett et al. 2011). The common name, scientific name, IUCN Red List status, and migratory status are followed using (Praveen & Jayapal 2023). The feeding guild data for each species were collected from the existing literature (Ali & Ripley 1987; Byju et al. 2023). The data recorded were later analyzed for relative abundance based on the frequency of sightings, as per MacKinnon & Phillips (1993): common (C) sighted from seven to nine times; un-common (UC) sighted from three to six times; rare (R) sighted once or twice. The relative diversity (RDi) of

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Figure 1. Map of the Samanatham tank, Madurai.

families was calculated using the following the formula by La Torre-Cuadros et al. (2007).

RDi = Total number of species

RESULTS AND DISCUSSION

A total of 150 species of birds belonging to 52 families under 17 orders were recorded from the Samanatham tank (Table 1; Images 1–18). The order Passeriformes dominated the study area with a maximum number of species, i.e., 53 species (35%), and with the lowest Strigiformes, Bucerotiformes, Falconiformes, Psittaciformes, sharing one species (0.66%) each (Figure 2). Similarly, family-wise Scolopacidae, Anatidae, and Accipitridae with relative abundance (RDi) of 7.33 % (11 species each), followed by Ardeidae (nine species) comprised the maximum number of birds species (Table 2).

As per the IUCN Red List (IUCN 2021), the wetland

supports, eight globally 'Near Threatened' (NT) species -Oriental Darter Anhinga melanogaster, Painted Stork Mycteria leucocephala, Black-headed Ibis Threskiornis melanocephalus, Spot-billed Pelican Pelecanus philippensis, Black-tailed Godwit Limosa limosa, Bartailed Godwit Limosa lapponica, River Tern Sterna aurantia, & Osprey Pandion haliaetus; two 'Vulnerable' (Vu) species – Indian Spotted Eagle Clanga hastata & Great Spotted Eagle Clanga clanga; and the remaining 140 species are under 'Least Concern' (LC).

An analysis of the feeding guilds of these birds revealed that 38% (57 species) were insectivorous, 33% (49 species) were carnivorous, 14% (21 species) were omnivorous, 7% (11 species) were granivorous, 5% (eight species each) were frugivorous, and 3% (four species) were nectarivorous (Figure 3). Further analysis of relative abundance based on the frequency of observation indicated that 56 % (84 species) were C (common), 28.6 % (43 species) were UC (uncommon), and 15.4 % (23 species) were Ra (rare). The analysis of data on residential status revealed that out of 150 species, 37 (25%) were winter visitors (WV) and one

checklist of avifauna of Samanatham tank, Madurai, India

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Table 1. Checklist of bird species in Samanatham tank, Madurai, Tamil Nadu, India.

	Order/Family/Common name	Scientific name	IUCN Red List status	Habitat	Resident status	Relative abundance	Feeding habits
	Anseriformes: Anatidae						
1	Knob-billed Duck	Sarkidiornis melanotos	LC	WL/AL	R/NB	UC	0
2	Common Teal	Anas crecca	LC	WL	WV	UC	0
3	Bar-headed Goose	Anser indicus	LC	WL	wv	UC	0
4	Eurasian Wigeon	Mareca penelope	LC	WL	WV	UC	0
5	Fulvous Whistling Duck	Dendrocygna bicolor	LC	WL	LM	С	0
6	Garganey	Spatula querquedula	LC	WL	WV	UC	0
7	Northern Shoveler	Spatula clypeata	LC	WL	WV	UC	0
8	Indian Spot-billed Duck	Anas poecilorhyncha	LC	WL	R	С	0
9	Lesser Whistling Duck	Dendrocygna javanica	LC	WL	R/NB	С	0
10	Northern Pintail	Anas acuta	LC	WL	WV	Ra	0
11	Ruddy Shelduck	Tadorna ferruginea	LC	WL	WV	Ra	0
	Phoenicopteriformes: Phoenicopteridae						
12	Greater Flamingo	Phoenicopterus roseus	LC	WL	LM	Ra	I
	Podicipedidae						
13	Little Grebe	Tachybaptus ruficollis	LC	WL	R	С	С
	Columbiformes: Columbidae						
14	Rock Pigeon	Columba livia	LC	AL/OS/GL	R	с	G
15	Spotted Dove	Spilopelia chinensis	LC	AL/OS/GL	R	с	G
16	Eurasian collared Dove	Streptopelia decaocto	LC	AL/OS/GL	R	С	G
17	Laughing Dove	Spilopelia senegalensis	LC	AL/OS/GL	R	с	G
	Caprimulgiformes: Apodidae						
18	Asian Palm Swift	Cypsiurus balasiensis	LC	Tr	R	с	1
19	Alpine Swift	Apus melba	LC	Tr	R	UC	I
	Cuculiformes: Cuculidae						
20	Asian Koel	Eudynamys scolopaceus	LC	OS	R	с	0
21	Common Hawk-Cuckoo	Hierococcyx varius	LC	OS	R	Ra	0
22	Greater Coucal	Centropus sinensis	LC	OS	R	с	С
23	Blue faced Malkoha	Phaenicophaeus viridirostris	LC	OS	R	С	I
24	Pied Crested Cuckoo	Clamator jacobinus	LC	OS	R/NB	С	I
	Gruiformes: Rallidae	1					
25	Baillon's Crake	Zapornia pusilla	LC	WL	R	UC	С
26	Eurasian Coot	Fulica atra	LC	WL	R	С	С
27	Eurasian Moorhen	Gallinula chloropus	LC	WL	R	с	С
28	Gray-headed Swamphen	Porphyrio poliocephalus	LC	WL	R	С	С
29	White-breasted Waterhen	Amaurornis phoenicurus	LC	IWB	R	С	С
	Galliformes: Phasianidae	1					
30	Grey Francolin	Ortygornis pondicerianus	LC	GL/OS	R	С	G
31	Indian Peafowl	Pavo cristatus	LC	AL/OS	R	С	0
	Pelecaniformes: Ciconiidae	1		1		1	
32	Asian Openbill Stork	Anastomus oscitans	LC	WL	R	С	С
33	Painted Stork	Mycteria leucocephala	NT	WL	R	с	с
34	Asian Wooly-necked Stork	Ciconia episcopus	LC	WL	wv	Ra	С
	Pelecanidae						
35	Spot-billed Pelican	Pelecanus philippensis	NT	WL	R	с	С
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	Order/Family/Common name	Scientific name	IUCN Red List status	Habitat	Resident status	Relative abundance	Feeding habits
	Ardeidae						
36	Black-crowned Night Heron	Nycticorax nycticorax	LC	WL	R	С	С
37	Cattle Egret	Bubulcus ibis	LC	WL	R	С	С
38	Purple Heron	Ardea purpurea	LC	WL	R	С	С
39	Grey Heron	Ardea cinerea	LC	WL	R	С	С
40	Indian Pond-Heron	Ardeola grayii	LC	WL	R	С	С
41	Intermediate Egret	Ardea intermedia	LC	WL	R	С	С
42	Great Egret	Ardea alba	LC	WL	R	С	С
43	Little Egret	Egretta garzetta	LC	WL	R	С	С
44	Striated Heron	Butorides striata	LC	WL	R	С	С
	Threskiornithidae						
45	Black-headed Ibis	Threskiornis melanocephalus	NT	WL	R	С	С
46	Eurasian Spoonbill	Platalea leucorodia	LC	WL	R	С	С
47	Glossy Ibis	Plegadis falcinellus	LC	WL	R	С	С
48	Red-naped Ibis	Pseudibis papillosa	LC	WL/AL	R	UC	С
	Phalacrocoracidae						
49	Great Cormorant	Phalacrocorax carbo	LC	WL	R	UC	С
50	Indian Cormorant	Phalacrocorax fuscicollis	LC	WL	R	с	С
51	Little Cormorant	Microcarbo niger	LC	WL	R	С	С
	Anhingidae						
52	Oriental Darter	Anhinga melanogaster	NT	WL	R	UC	С
	Charadriiformes: Recurvirostridae	2					
53	Black Winged Stilt	Himantopus himantopus	LC	WL	R	С	1
	Charadriidae						
54	Kentish Plover	Charadrius alexandrinus	LC	WL	LM	UC	1
55	Little Ringed Plover	Charadrius dubius	LC	WL	WV	UC	I
56	Red-wattled Lapwing	Vanellus indicus	LC	WL	R	с	0
57	Yellow-wattled Lapwing	Vanellus malabaricus	LC	WL/AL	R	Ra	0
	Jacanidae						
58	Pheasant-tailed Jacana	Hydrophasianus chirurgus	LC	WL	R	С	I
	Scolopacidae						
59	Black-tailed Godwit	Limosa limosa	NT	WL	WV	UC	I
60	Bar-tailed Godwit	Limosa lapponica	NT	WL	WV	UC	I
61	Ruff	Calidris pugnax	LC	WL	WV	UC	0
62	Temminck's Stint	Calidris temminckii	LC	WL	WV	UC	I
63	Little Stint	Calidris minuta	LC	WL	WV	UC	I
64	Common Sandpiper	Actitis hypoleucos	LC	WL	WV	UC	I
65	Green Sandpiper	Tringa ochropus	LC	WL	WV	UC	I
66	Marsh Sandpiper	Tringa stagnatilis	LC	WL	WV	UC	I
67	Wood Sandpiper	Tringa glareola	LC	WL	WV	UC	I
68	Pin-tailed Snipe	Gallinago stenura	LC	WL	WV	UC	I
69	Common Greenshank	Tringa nebularia	LC	WL	WV	UC	I
70	Small Pratincole	Glareola lactea	LC	WL	WV	UC	I
	Laridae						
71	Gull-billed Tern	Gelochelidon nilotica	LC	WL	WV	Ra	С
72	Whiskered Tern	Chlidonias hybrida	LC	WL	WV	Ra	С

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	Order/Family/Common name	Scientific name	IUCN Red List status	Habitat	Resident status	Relative abundance	Feeding habits
73	River Tern	Sterna aurantia	NT	WL	WV	Ra	С
	Accipitriformes: Pandionidae						
74	Osprey	Pandion haliaetus	NT	WL/Tr	WV	Ra	с
	Accipitridae						
75	Booted Eagle	Hieraaetus pennatus	LC	Tr	wv	UC	с
76	Black Kite	Milvus migrans	LC	Tr	R	С	С
77	Black-winged Kite	Elanus caeruleus	LC	Tr	R	с	с
78	Greater Spotted Eagle	Clanga clanga	VU	Tr	WV	UC	с
79	Indian Spotted Eagle	Clanga hastata	VU	Tr	WV	UC	с
80	Shikra	Accipiter badius	LC	Tr	R	с	с
81	Short-toed Snake Eagle	Circaetus gallicus	LC	Tr	LM	Ra	с
82	Eurasian Marsh- Harrier	Circus aeruginosus	LC	Tr	WV	Ra	с
83	White-eyed Buzzard	Butastur teesa	LC	GL	R	Ra	с
84	Oriental Honey-Buzzard	Pernis ptilorhynchus	LC	Tr	LM	UC	с
85	Common Buzzard	Buteo buteo	LC	Tr	WV	Ra	с
	Strigiformes: Strigidae				-	-	
86	Spotted Owlet	Athene brama	LC	OS	R	С	с
	Bucerotiformes: Upupidae						
87	Common Hoopoe	Upupa epops	LC	OS/GL	R	С	I
	Piciformes: Picidae						
88	Black-rumped Flameback	Dinopium benghalense	LC	Tr	R	С	I
	Megalaimidae						
89	Coppersmith Barbet	Psilopogon haemacephalus	LC	Tr	R	С	F
	Coraciiformes:Meropidae						
90	Blue-tailed Bee-eater	Merops philippinus	LC	OS	R	С	I
91	Green Bee-eater	Merops orientalis	LC	OS	R	UC	I
	Coraciidae						
92	Indian Roller	Coracias benghalensis	LC	OS/GL	R	с	I
	Alcedinidae						
93	Pied Kingfisher	Ceryle rudis	LC	WL/OS	R	с	с
94	Common Kingfisher	Alcedo atthis	LC	WL/OS	R	С	С
95	White-throated Kingfisher	Halcyon smyrnensis	LC	WL/OS	R	с	с
	Falconiformes: Falconidae						
96	Red-necked Falcon	Falco chicquera	LC	OS/GL	WV	Ra	с
	Psittaciformes: Psittacidae						
97	Rose-ringed Parakeet	Psittacula krameri	LC	Tr	R	С	F
	Passeriformes: Oriolidae						
98	Indian Golden Oriole	Oriolus kundoo	LC	Tr	R	С	0
	Pittadae						
99	Indian Pitta	Pitta brachyura	LC	OS/GL	WV	Ra	I
	Artamidae						
100	Ashy Woodswallow	Artamus fuscus	LC	OS	R	С	I
	Dicruridae						
101	Black Drongo	Dicrurus macrocercus	LC	OS	R	С	1
102	Ashy Drongo	Dicrurus leucophaeus	LC	OS	R	UC	1

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	Order/Family/Common name	Scientific name	IUCN Red List status	Habitat	Resident status	Relative abundance	Feeding habits
	Laniidae						
103	Brown Shrike	Lanius cristatus	LC	OS	WV	UC	I
104	Long-tailed Shrike	Lanius schach	LC	OS	R	С	С
	Vangidae						
105	Common Woodshrike	Tephrodornis pondicerianus	LC	OS	R	Ra	I
	Corvidae						
106	House Crow	Corvus splendens	LC	OS/Tr	R	С	0
107	Rufous Treepie	Dendrocitta vagabunda	LC	OS	R	С	0
108	Large-billed Crow	Corvus macrorhynchos	LC	OS/Tr	R	с	0
	Monarchidae						
109	Indian Paradise-Flycatcher	Terpsiphone paradisi	LC	OS/GL	R/NB	UC	1
	Nectariniidae						
110	Loten's Sunbird	Cinnyris lotenius	LC	OS	R	UC	N
111	Purple-rumped Sunbird	Leptocoma zevlonica	LC	OS	R	с	N
112	Purple Sunbird	Cinnvris asiaticus	LC	OS	R	c	N
	Ploceidae	,					
113	Bava Weaver	Ploceus nhilinninus	10	05	R	C	G
114	Streaked Weaver	Ploceus manyar		05	R	C C	6
	Estrildidae			03			3
115	Tricolored Munia	Lonchurg malacca		05	P	C	6
115		Euodico malabarica		05	D D		6
117	Scolu broasted Munic			03	n n	C	G
11/				05	ĸ	C	6
440	Passeridae					-	
118	House Sparrow	Passer domesticus		AL/GL	ĸ	Ĺ	G
	Motacillidae						
119	Citrine Wagtail	Motacilla citreola	LC	GL	WV	UC	
120	Paddyfield Pipit	Anthus rufulus	LC	GL/AL	R	C	1
121	Grey Wagtail	Motacilla cinerea	LC	GL	R/NB	UC	1
122	White-browed Wagtail	Motacilla maderaspatensis	LC	WL	R	С	1
123	Western Yellow Wagtail	Motacilla flava	LC	OS	WV	Ra	I
	Alaudidae						
124	Ashy-crowned Sparrow-Lark	Eremopterix griseus	LC	AL/OS/GL	R	С	1
125	Jerdon's Bushlark	Mirafra affinis	LC	AL/OS/GL	R	С	1
126	Oriental Skylark	Alauda gulgula	LC	AL/OS/GL	R	С	1
	Cisticolidae						
127	Common Tailorbird	Orthotomus sutorius	LC	AL/OS/GL	R	С	I
128	Plain Prinia	Prinia inornata	LC	AL/OS/GL	R	С	1
129	Zitting Cisticola	Cisticola juncidis	LC	AL/OS/GL	R	С	I
130	Ashy Prinia	Prinia socialis	LC	AL/OS/GL	R	С	1
131	Jungle Prinia	Prinia sylvatica	LC	OS	R	С	1
	Leiothrichidae						
132	Yellow-billed Babbler	Argya affinis	LC	OS	R	С	I
	Acrocephalidae						
133	Blyth's Reed Warbler	Acrocephalus dumetorum	LC	OS	R/NB	UC	I
134	Booted Warbler	Iduna caligata	LC	OS	R/NB	Ra	I
135	Clamorous Reed-Warbler	Acrocephalus stentoreus	LC	OS	R	С	I

	Order/Family/Common name	Scientific name	IUCN Red List status	Habitat	Resident status	Relative abundance	Feeding habits
136	Thick-billed Warbler	Arundinax aedon	LC	OS	R/NB	Ra	I
137	Sykes's Warbler	Iduna rama	LC	OS	R/NB	Ra	I
	Hirundinidae						
138	Barn Swallow	Hirundo rustica	LC	AL/WL	WV	Ra	1
139	Red-rumped Swallow	Cecropis daurica	LC	AL/WL	R	UC	I
140	Dusky crag Martin	Ptyonoprogne concolor	LC	WL/AL	R	UC	1
	Pycnonotidae						
141	Red-vented Bulbul	Pycnonotus cafer	LC	OS	R	С	F
142	White-browed Bulbul	Pycnonotus luteolus	LC	OS	R	Ra	F
	Sturnidae						
143	Brahminy Starling	Sturnia pagodarum	LC	OS	R	UC	F
144	Common Myna	Acridotheres tristis	LC	OS	R	С	F
145	Chestnut-tailed Starling	Sturnia malabarica	LC	OS	WV	UC	F
146	Rosy Starling	Pastor roseus	LC	OS	PM	UC	F
	Muscicapidae						
147	Indian Robin	Copsychus fulicatus	LC	OS/AL	R	С	I
148	Oriental Magpie-Robin	Copsychus saularis	LC	OS/AL	R	С	I
149	Pied Bushchat	Saxicola caprata	LC	OS	R	С	1
	Dicaeidae						
150	Pale-billed Flowerpecker	Dicaeum erthrorhynchos	LC	OS	R	UC	N

IUCN Red list status: LC—Least Concern | NT—Near Threatened | Vu—Vulnerable. Resident status: WV—Winter Visitor | LM—Local Migrant | R—Resident | R/NB—Resident/Non-Breeding. Relative abundance: C—Common | UC—Uncommon | Ra—Rare. Habitats: W—Wetland | AL—Agricultural Land | Tr—Tree | GL—Grass Land | OS—Open Scrub. Feeding status: I—Insectivore | G—Granivore | C—Carnivore | O—Omnivore | N—Nectarivore | F—Frugivore.



Table 1. Checklist of bird species in Samanatham tank, Madurai, Tamil Nadu, India.

Table 2. Relative diversity (RDi) of various avian families at Samanatham tank, Madurai.

	Family	No of species	RDi %
1	Anatidae	11	7.33
2	Phoenicopteridae	1	0.66
3	Podicipedidae	1	0.66
4	Columbidae	4	2.66
5	Apodidae	2	1.33
6	Cuculidae	5	3.33
7	Rallidae	5	3.33
8	Phasianidae	2	1.33
9	Ciconiidae	3	2.00
10	Pelecanidae	1	0.66
11	Ardeidae	9	6.00
12	Threskiornithidae	4	2.66
13	Phalacrocoracidae	3	2.00
14	Anhingidae	1	0.66
15	Recurvirostridae	1	0.66
16	Charadriidae	4	2.66
17	Jacanidae	1	0.66
18	Scolopacidae	11	7.33
19	Laridae	3	2.00
20	Pandionidae	1	0.66
21	Accipitridae	11	7.33
22	Strigidae	1	0.66
23	Upupidae	1	0.66
24	Picidae	1	0.66
25	Megalaimidae	1	0.66
26	Meropidae	1	0.66
27	Coraciidae	1	0.66

	Family	No of species	RDi %
28	Alcedinidae	3	2.00
29	Falconidae	1	0.66
30	Psittacidae	1	0.66
31	Oriolidae	1	0.66
32	Pittidae	1	0.66
33	Artamidae	1	0.66
34	Dicruridae	1	0.66
35	Laniidae	2	1.33
36	Vangidae	1	0.66
37	Corvidae	3	2.00
38	Monarchidae	1	0.66
39	Nectariniidae	3	2.00
40	Ploceidae	2	1.33
41	Estrildidae	3	2.00
42	Passeridae	1	0.66
43	Motacillidae	5	3.33
44	Alaudidae	3	2.00
45	Cisticolidae	5	3.33
46	Leiotrichidae	1	0.66
47	Acrocephalidae	5	3.33
48	Hirundinidae	3	2.00
49	Pycnonotidae	3	1.65
50	Sturnidae	4	2.66
51	Muscicapidae	3	1.65
52	Dicaeidae	1	0.66
	Total	150	



Figure 3. Feeding guilds of birds at Samanatham tank, Madurai.

passage migrant (PM) Rosy Starling Pastor roseus.

The wetland being used by the long-distance migrant shorebirds including the Bar-tailed Godwit and the Black-tailed Godwit, the two Near Threatened species in reasonable numbers makes this an important wintering area for those species. Similar works were reported recently from the Changaram wetlands of Kerala highlighting the need of conserving hitherto undocumented new areas of shorebird wintering sites (Anand et al. 2023). The consistent occurrence of the Near Threatened River Tern represents another ecologically significant species within this geographic region. It is also interesting to note that another two Near Threatened species namely Spot-billed Pelican and Oriental Darter were breeding in the Samanatham tank. Tamil Nadu is home to 535 bird species (Praveen et al.



Image 1–8. 1—Black-crowned Night Heron Nycticorax nycticorax | 2—Black-tailed Godwit Limosa limosa | 3—Black headed Ibis Threskiornis melanocephalus & Glossy Ibis Plegadis falcinellus | 4—Eurasian Spoonbill Platalea leucorodia | 5— Greater Flamingos Phoenicopterus roseus, Northern Shoveler Spatula clypeata, & Garganey Spatula querquedula | 6—Bar-headed Goose Anser indicus & Black-winged Stint Himantopus Himantopus | 7—Flocks of duck species with Greater Flamingos | 8—Spot-billed Pelican Pelecanus philippensis nesting. © N. Raveendran.

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Image 9–16. 9—Painted Stork *Mycteria leucocephala* | 10—Osprey *Pandion haliaetus* | 11—Oriental Darter *Anhinga melanogaster* | 12—Grey Heron *Ardea cinerea* | 13—*Prosopis juliflora* serves as nesting trees for cormorants and egrets | 14—*Prosopis juliflora* trees used as roosting spot for egrets | 15—Brown Shrike *Lanius cristatus* | 16—Greater Spotted Eagle *Clanga clanga*. © N. Raveendran.

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Image 17–18. 17—Indian Spotted Eagle Clanga hastata | 18—Samanatham tank view. © N. Raveendran.

2018) and we recorded 150 (28%) species during the current study at this single water tank in Madurai.

Our study provides important baseline information and the presence of many important waterbird species; this will help in the long-term monitoring of birds in the tank besides acting as an essential document in planning conservation efforts for the wetland. This tank is an artificial waterbody that supports a variety of plant and animal life. Samanatham tank plays a role in water management and flood control for the surrounding area. Hence, this tank should be recognized as a valuable ecosystem that should be elevated to protection status and conserved for future generations.

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Composition of avian communities in Ranjit Sagar Conservation Reserve, Punjab, India

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Abstract: The Ranjit Sagar wetland, a critical aquatic ecosystem supporting diverse flora and fauna, has been the subject of investigation from January 2013 to January 2015. The study revealed the presence of 167 avian species across 19 orders and 52 families. Notably, areas with minimal disturbance within the wetland serve as key niches for a substantial avian population, encompassing 47.31% migrant and 52.69% resident species. Significantly, 10 Near Threatened species, alongside the Vulnerable Common Pochard *Aythya ferina* and the Endangered Egyptian Vulture *Neophron percnopterus*, underscore the wetland's conservation value. Dominated by order Passeriformes (55 species), followed by Charadriiformes (29 species), and Anseriformes (20 species), the avian community exhibits varied feeding guilds, with insectivores (37.13%), omnivores (18.56%), aquatic animal feeders (16.16%), carnivores (14.97%), frugivores (5.39%), granivores (4.79%), and herbivores (3%) comprising its composition. This study elucidates the intricate avian ecosystem's dynamics in Ranjit Sagar wetland, emphasizing its conservation significance and ecological structure, thereby contributing valuable insights for preservation and future research.

Keywords: Aquatic ecosystems, avifauna, conservation, habitats, IUCN Red List, mining, niche, vegetation, water birds, wetlands.

Punjabi: ਪੰਜਾਬ ਦੇ ਰੁੱਖਾਂ ਅਤੇ ਜੀਵਾਂ ਦੇ ਨਿਰੀਖਣ ਲਈ ਰਣਜੀਤ ਸਾਗਰ ਬੀਲ ਨੂੰ ਚੁਣਿਆ ਗਿਆ ਜਿਸ ਦੌਰਾਨ ਜਨਵਰੀ 2013 ਤੋਂ ਜਨਵਰੀ 2015 ਤੱਕ ਦੇ ਅਧਿਅਨ ਵਿੱਚ ਪਾਇਆ ਗਿਆ ਕਿ ਇਸ ਥਾਂ ਉੱਤੇ 167 ਪੰਛੀ ਹਨ ਜੋ ਕਿ 19 ਆਰਡਰ, 52 ਫੇਮਿਲੀਆਂ ਨਾਲ ਸਬੰਧ ਰੱਖਦੇ ਹਨ। ਇਸ ਥਾਂ ਤੇ ਮਨੁੱਖੀ ਦਖਲਅੰਦਾਜੀ ਨਾ-ਮਾਤਰ ਹੋਣ ਕਰਕੇ ਇਹ ਇਲਾਕਾ ਪੰਛੀਆਂ ਲਈ ਪਸੰਦੀਦਾ ਥਾਂ ਹੈ ਜਿਸ ਕਰਕੇ ਇੱਥੇ 47.31 ਫੀਸਦੀ ਪ੍ਰਵਾਸੀ ਪੰਛੀ ਆਉਂਦੇ ਹਨ ਅਤੇ 52.69 ਫੀਸਦੀ ਪੰਛੀ ਇੱਥੋਂ ਦੇ ਹੀ ਵਸਨੀਕ ਹਨ। ਇੱਥੇ 10 ਦੇ ਆਸਪ।ਸ ਪ੍ਰਜਾਤੀਆਂ ਲੁਪਤ ਹੋਣ ਦੀ ਕਗਾਰ ਤੇ ਹਨ ਜਿਨ੍ਹਾਂ ਵਿੱਚੋਂ ਕਮਜੋਰ/vulnerable ਪ੍ਰਜਾਤੀ ਕਾਮਨ ਪੱਛੀਆਂ ਦੀ ਬੀਨਾ ਲਈ ਪਸੰਦੀਦਾ ਥਾਂ ਹੈ ਜਿਸ ਕਰਕੇ ਇੱਥੇ 47.31 ਫੀਸਦੀ ਪ੍ਰਵਾਸੀ ਪੰਛੀ ਆਉਂਦੇ ਹਨ ਅਤੇ 52.69 ਫੀਸਦੀ ਪੰਛੀ ਇੱਥੋਂ ਦੇ ਹੀ ਵਸਨੀਕ ਹਨ। ਇੱਥੇ 10 ਦੇ ਆਸਪ।ਸ ਪ੍ਰਜਾਤੀਆਂ ਲੁਪਤ ਹੋਣ ਦੀ ਕਗਾਰ ਤੇ ਹਨ ਜਿਨ੍ਹਾਂ ਵਿੱਚੋਂ ਕਮਜੋਰ/vulnerable ਪ੍ਰਜਾਤੀ ਕਾਮਨ ਪੋਰਾਰਡ (ਅਥਿਇਆ ਫੇਰਿਨਾ) ਸ਼ਾਮਿਲ ਹੈ ਅਤੇ ਖਤਰੇ ਦੀ ਕਗਾਰ/ Endangered ਤੇ ਖੜੀ ਚਿੰਦੀ ਗਿੰਦ (ਨਿਓਫਿਰੋਨ ਪਰਕਨੋਪਟੇਰਸ) ਸ਼ਾਮਿਲ ਹੈ। ਇਸ ਬੀਲ ਵਿੱਚ ਆਰਡਰ ਪੇਜਰੀਫਾਰਮਿਸ (55 ਪ੍ਰਜਾਤੀਆਂ) ਦੇ ਪੰਛੀਆਂ ਦੀ ਬਹੁਤਾਤ ਹੈ ਅਤੇ ਇਸ ਤੋਂ ਬਾਅਦ ਕਰਾਡਰੀਫਾਰਮਿਸ (29 ਪ੍ਰਜਾਤੀਆਂ) ਅਤੇ 20 ਪ੍ਰਜਾਤੀਆਂ ਅਨਜੇਰੀਫਾਰਮਿਸ ਆਰਡਰ ਦੀਆਂ ਹਨ। ਇਹ ਪੰਛੀ ਵੱਖ ਵੱਖ ਤਰ੍ਹਾਂ ਦੇ ਭੋਜਨ ਉੱਤੇ ਨਿਰਭਰ ਕਰਦੇ ਹਨ ਜਿਵੇਂ ਕਿ 37.13% ਕੀਤੇ-ਖਾਣ ਵਾਲੇ, 18.56% ਸਰਵਹਾਰੇ, 16.16% ਪਾਣੀ ਵਾਲੇ ਜੀਵ ਖਾਂਦੇ ਹਨ, 14.97% ਮਾਸਾਹਾਰੀ, 5.39% ਫਲ-ਖਾਣੇ, 4.79% ਦਾਣੇ ਖਾਣ ਵਾਲੇ ਅਤੇ 3% ਸ਼ਾਕਾਹਰੀ ਹਨ। ਇਸ ਅਧਿਅਨ ਤੋਂ ਰਣਜੀਤ ਸਾਗਰ ਬੀਲ ਦੇ ਗਤੀਸ਼ੀਲ/ਜੀਵੰਤ ਸੁਭਾਆ ਦਾ ਪਤਾ ਲਗਦਾ ਹੈ ਜਿਸ ਕਰਕੇ ਇਸ ਬੀਲ ਦੇ ਸਾਂਭ-ਸੰਭਾਲ ਦਾ ਅਪਣਾ ਇੱਕ ਮਹੱਤਵ ਹੈ।

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Author contributions: OSB—PhD supervisor, edited MS., SKS— Field survey and M.S writing, JS—Final manuscript preparation, editing and communication with journal.

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INTRODUCTION

Wetlands serve as a crucial transitional zone bridging terrestrial and aquatic ecosystems, characterized by a shallow water table or the presence of shallow water covering the land's surface. They play a pivotal role in providing habitat for diverse biota and offer a range of ecological services (Coppes et al. 2018; Kakati et al. 2021; Wu et al. 2021). India, in particular, boasts an impressive avian diversity, with 1,353 documented bird species, of which approximately 310 are known to be dependent on wetlands (Kumar et al. 2005; Praveen & Jayapal 2023). Wetlands hold special significance for birds, serving as essential sites for roosting, foraging, drinking, resting, shelter, and social interactions (Singh & Banyal 2013; Kumar & Sharma 2019; Yao et al. 2020; Casazza et al. 2021; Joshi et al. 2021; Teng et al. 2021).

Birds are reliable indicators of water quality and wetland health (Yao et al. 2020; Anand et al. 2023). In wetland ecosystems, birds can be broadly classified into two categories: wetland specialists, which exclusively nest, feed, and roost in wetlands and are entirely reliant on aquatic habitats, and generalists, which frequent wetlands but also inhabit other environments. The development of an avifaunal baseline checklist specific to this habitat plays a pivotal role in shaping site and species-specific conservation strategies, applicable to both protected and unprotected areas, as supported by previous research (Bibby 1998; Anand et al. 2023; Byju et al. 2023a,b). Birds offer a valuable avenue for assessing biodiversity, often indicative of a healthy ecology through their diverse populations. Enumeration of avian diversity also contributes to our understanding of the status of endangered or threatened species. Furthermore, it holds educational value, fostering greater regional species and ecosystem appreciation, while encouraging community participation in conservation initiatives. Multiple researchers have conducted extensive studies on bird diversity in Punjab's wetlands, yielding noteworthy findings (Ali et al. 1981; Robson 1996; Kazmierczak et al. 1998; Robson 1999; Sawant & Sudhagar 2013; Prasad 2008a,b; Singh & Brraich 2021). This comprehensive study serves to elucidate the importance of wetlands and their relevance in shaping effective conservation strategies.

MATERIALS AND METHODS

Study area

The Ranjit Sagar wetland came into existence in 2004 with the construction of a barrage on the river Ravi of the Indus River system a freshwater wetland (32.442° N and 75.725° E, at the altitudes of 540 m) located near Pathankot City, Punjab (Figure 1). This wetland falls into three states, i.e., Punjab, Himachal Pradesh, and Jammu & Kashmir, and is spread over an area of 87.60 km² with a catchment area spread over 6,086 km². In and around this wetland, several habitats, including marshy parts, shallow areas, riparian vegetation, deep areas, islands, large trees (for the roosting of birds), fallow fields near the banks (for the breeding purposes) have developed over the time, generally preferred by migratory birds. It also provides food for different birds in the form of fishes, amphibians, reptiles, insects, zooplankton, phytoplankton, and weeds.

Methods

Field surveys were conducted from January 2013 to January 2015, utilizing visual counting methods following Gaston's (1975) methodology for bird observation. To ensure comprehensive coverage of the wetland area, about 12 vantage points were established. Data collection was conducted year-round, with observations made on five days each month, both in the morning (0700–1000 h) and evening (1600–1830 h). Species identification was based on references from bird identification guides, including works by Ali & Ripley (1983), Grimmett & Inskipp (2010), and Grimmett et al. (2012). Bird species were categorized into three groups: annual, winter, and summer birds, following Mavi & Tiwana's (1993) classification system. The checklist was meticulously compiled using standardized common and scientific names, adhering to guidelines by Clements et al. (2022) and Praveen & Jayapal (2023).

RESULTS AND DISCUSSION

In the present study, which focused on avian diversity at Ranjit Sagar Wetland, a total of 167 species were observed. The majority, accounting for 157 species, fall within the IUCN's Least Concern category, while the remaining 10 species are classified as threatened. Notable among these are the Vulnerable Common Pochard Aythya ferina and the Endangered Egyptian Vulture Neophron percnopterus. Additionally, species such as the Himalayan Vulture Gyps himalayensis,



Figure 1. Ranjit Sagar Wetland on the map.

Ferruginous Duck Aythya nyroca, Black-tailed Godwit Limosa limosa, River Lapwing Vanellus duvaucelii, Eurasian Curlew Numenius arquata, Painted Stork Mycteria leucocephala, Black-headed Ibis Threskiornis melanocephalus, Alexandrine Parakeet Psittacula eupatria, and Oriental Darter Anhinga melanogaster are categorized as Near Threatened. Vulnerable River Tern Sterna aurantia, Common Pochard Aythya ferina, and the Endangered Egyptian Vulture Neophron percnopterus complete the list of threatened species.

Concerning dietary preferences, the observed species display diverse feeding habits. The majority, comprising 37.13% (62 species), are insectivorous, followed by 18.56% (31 species) categorized as omnivorous, 16.16% (27 species) primarily feeding on aquatic animals, and 14.97% (25 species) displaying carnivorous tendencies. Additionally, 5.39% of species are frugivorous (9 species), while 4.79% are granivorous (8 species).

Herbivorous birds constitute only 3% (5 species) of the observed population, as detailed in Table 1.

The Ranjit Sagar Wetland, characterized by its extensive open spaces in comparison to other wetlands in the region, provides an ideal habitat for waterfowls and aquatic birds. Furthermore, the presence of 44 fish species within this wetland area may attract piscivorous birds classified as Aquatic animal feeders. Adjacent agricultural lands offer ample opportunities to attract herbivorous migratory birds, including the Bar-headed Goose Anser indicus and Greylag Goose Anser anser. Nevertheless, the study underscores the negative impact of anthropogenic activities such as mining, poaching, and sewage discharge on wetland health. To mitigate these concerns and enhance the future attraction of migratory birds, it is recommended to implement conservation efforts that include conservation efforts, including habitat restoration, bans on sand mining and

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Table 1. A checklist of the avian species of Ranjit Sagar wetland, Punjab.

	Taxon	Family	Season	Feeding habit	Residential status	IUCN Status
Order	Anseriformes					
1	Bar-headed Goose Anser indicus (Latham, 1790)	Anatidae	w	н	Migrant	Least Concern
2	Knob-billed Duck Sarkidiornis melanotos (Pennant, 1769)	Anatidae	А	0	Resident	Least Concern
3	Common Pochard Aythya ferina (Linnaeus, 1758)	Anatidae	w	0	Migrant	Vulnerable
4	Cotton Pygmy Goose <i>Nettapus coromandelianus</i> (Gmelin, J.F., 1789)	Anatidae	W	0	Migrant	Least Concern
5	Common Teal Anas crecca (Linnaeus, 1758)	Anatidae	w	0	Migrant	Least Concern
6	Eurasian Wigeon Mareca penelope (Linnaeus, 1758)	Anatidae	w	н	Migrant	Least Concern
7	Gadwall Mareca strepera (Linnaeus, 1758)	Anatidae	w	н	Migrant	Least Concern
8	Garganey Spatula querquedula (Linnaeus, 1758)	Anatidae	w	0	Migrant	Least Concern
9	Greylag Goose Anser anser (Linnaeus, 1758)	Anatidae	w	н	Migrant	Least Concern
10	Indian Spot-billed Duck Anas poecilorhyncha (Forster, 1781)	Anatidae	А	н	Resident	Least Concern
11	Lesser Whistling Duck <i>Dendrocygna javanica</i> (Horsfeld, 1821)	Anatidae	S	0	Migrant	Least Concern
12	Northern Shoveler Spatula clypeata (Linnaeus, 1758)	Anatidae	w	0	Migrant	Least Concern
13	Northern Pintail Anas acuta (Linnaeus, 1758)	Anatidae	w	0	Migrant	Least Concern
14	Red-crested Pochard Netta rufina (Pallas, 1773)	Anatidae	w	0	Migrant	Least Concern
15	Ruddy Shelduck Tadorna ferruginea (Pallas, 1764)	Anatidae	w	0	Migrant	Least Concern
16	Common Shelduck Tadorna tadorna (Linnaeus, 1758)	Anatidae	w	о	Migrant	Least Concern
17	Ferruginous Duck Aythya nyroca (Guldenstadt, 1770)	Anatidae	w	0	Migrant	Near Threatened
18	Tufted Duck Aythya fuligula (Linnaeus, 1758)	Anatidae	w	0	Migrant	Least Concern
Order	Order: Galliformes					
19	Grey Francolin Ortygornis pondicerianus (Gmelin, 1789)	Phasinidae	А	0	Resident	Least Concern
20	Indian Peafowl Pavo cristatus (Linnaeus, 1758)	Phasinidae	А	0	Resident	Least Concern
Order	Phoenicopteriformes					·
21	Greater Flamingo Phoenicopterus roseus (Pallas, 1811)	Phoenicopteridae	w	AqA	Migrant	Least Concern
	Order: Podicipediformes					
22	Little Grebe Tachybaptus ruficollis (Pallas, 1764)	Podicipedidae	А	AqA	Resident	Least Concern
23	Great Crested Grebe Podiceps cristatus (Linnaeus, 1758)	Podicipedidae	w	AqA	Migrant	Least Concern
24	Black-necked Grebe Podiceps nigricollis (Brehm, 1831)	Podicipedidae	w	AqA	Migrant	Least Concern
Order	Columbiformes					
25	Rock Pigeon Columba livia (Gmelin, 1789)	Columbidae	А	G	Resident	Least Concern
26	Eurasian Collared Dove <i>Streptopelia decaocto</i> (Frivaldszky, 1838)	Columbidae	А	G	Resident	Least Concern
27	Laughing Dove Spilopelia senegalensis (Linnaeus, 1766)	Columbidae	А	G	Resident	Least Concern
28	Red Collared Dove <i>Streptopelia tranquebarica</i> (Hermann, 1804)	Columbidae	А	G	Resident	Least Concern
29	Spotted Dove Spilopelia chinensis (Scopoli, 1786)	Columbidae	А	G	Resident	Least Concern
30	Yellow-footed Green Pigeon Treron phoenicopterus (Latham, 1790)	Columbidae	А	F	Resident	Least Concern
Order	Cuculiformes					
31	Asian Koel Eudynamys scolopaceus (Linnaeus, 1758)	Cuculidae	S	I	Migrant	Least Concern
32	Greater Coucal Centropus sinensis (Stephens, 1815)	Cuculidae	А	С	Resident	Least Concern
33	Pied Cuckoo Clamator jacobinus(Boddaert, 1783)	Cuculidae	А	I	Resident	Least Concern
Order	Caprimulgiformes					
34	Alpine Swift Tachymarptis melba (Linnaeus, 1758)	Apodidae	А	I	Resident	Least Concern

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	Taxon	Family	Season	Feeding habit	Residential status	IUCN Status
Order	Gruiformes					
35	Eurasian Coot Fulica atra (Linnaeus, 1758)	Rallidae	w	0	Migrant	Least Concern
36	Common Moorhen Gallinula chloropus (Linnaeus, 1758)	Rallidae	А	0	Resident	Least Concern
37	Grey-headed Swamphen Porphyrio poliocephalus (Latham, 1801)	Rallidae	А	0	Resident	Least Concern
38	White-Breasted Waterhen <i>Amaurornis phoenicurus</i> (Pennant, 1769)	Rallidae	A	0	Resident	Least Concern
39	Common Crane Grus grus (Linnaeus, 1758)	Gruidae	w	0	Migrant	Least Concern
40	Demoiselle Crane Grus virgo (Linnaeus, 1758)	Gruidae	w	0	Migrant	Least Concern
Order	Charadriiformes					
41	Black-winged Stilt Himantopus himantopus (Linnaeus, 1758)	Recurvirostridae	А	I	Resident	Least Concern
42	Pied Avocet Recurvirostra avosetta Linnaeus, 1758	Recurvirostridae	w	С	Migrant	Least Concern
43	Little Ringed Plover Charadrius dubius Scopoli, 1786	Charadriidae	w	1	Migrant	Least Concern
44	Kentish Plover Charadrius alexandrinus Linnaeus, 1758	Charadriidae	w	1	Migrant	Least Concern
45	Red-wattled Lapwing Vanellus indicus (Boddaert, 1783)	Charadriidae	А	1	Resident	Least Concern
46	River Lapwing Vanellus duvaucelii (Lesson, 1826)	Charadriidae	w	AqA	Migrant	Near Threatened
47	White-tailed Lapwing Vanellus leucurus (Lichtenstein, MHC, 1823)	Charadriidae	w	I	Migrant	Least Concern
48	Yellow-wattled Lapwing Vanellus malabaricus (Boddaert, 1783)	Charadriidae	w	I	Migrant	Least Concern
49	Greater Painted-snipe <i>Rostratula benghalensis</i> (Linnaeus, 1758)	Rostratulidae	w	0	Migrant	Least Concern
50	Pheasant-tailed Jacana Hydrophasianus chirurgus (Scopoli, 1786)	Jacanidae	S	I	Migrant	Least Concern
51	Black-tailed Godwit Limosa limosa (Linnaeus, 1758)	Scolopacidae	w	0	Migrant	Near Threatened
52	Common Redshank Tringa totanus(Linnaeus, 1758)	Scolopacidae	w	I	Migrant	Least Concern
53	Common Sandpiper Actitis hypoleucos Linnaeus, 1758	Scolopacidae	w	I	Migrant	Least Concern
54	Common Snipe Gallinago gallinago (Linnaeus, 1758)	Scolopacidae	w	I	Migrant	Least Concern
55	Dunlin Calidris alpina (Linnaeus, 1758)	Scolopacidae	w	AqA	Migrant	Least Concern
56	Green Sandpiper Tringa ochropus (Linnaeus, 1758)	Scolopacidae	w	I	Migrant	Least Concern
57	Common Greenshank Tringa nebularia (Gunnerus, 1767)	Scolopacidae	w	AqA	Migrant	Least Concern
58	Marsh Sandpiper Tringa stagnatilis (Bechstein, 1803)	Scolopacidae	w	AqA	Migrant	Least Concern
59	Ruff Calidris pugnax (Linnaeus, 1758)	Scolopacidae	w	AqA	Migrant	Least Concern
60	Spotted Redshank Tringa erythropus (Pallas, 1764)	Scolopacidae	w	1	Migrant	Least Concern
61	Wood Sandpiper Tringa glareola Linnaeus, 1758	Scolopacidae	w	I	Migrant	Least Concern
62	Eurasian Curlew Numenius arquata (Linnaeus, 1758)	\$colopacidae	w	AqA	Migrant	Near threatened
63	Black-headed Gull Chroicocephalus ridibundus (Linnaeus, 1766)	Laridae	w	AqA	Migrant	Least Concern
64	Brown-headed Gull Chroicocephalus brunnicephalus (Jerdon, 1840)	Laridae	w	AqA	Migrant	Least Concern
65	Pallas's Gull Ichthyaetus ichthyaetus (Pallas, 1773)	Laridae	w	С	Migrant	Least Concern
66	Lesser Black-backed Gull Larus fuscus (Linnaeus, 1758)	Laridae	w	С	Migrant	Least Concern
67	River Tern Sterna aurantia (Gray, 1831)	Laridae	A	AqA	Resident	Near Threatened
68	Little Tern Sternula albifrons (Pallas, 1764)	Laridae	w	AqA	Migrant	Least Concern
69	Whiskered Tern Chlidonias hybrida (Pallas, 1811)	Laridae	w	AqA	Migrant	Least Concern
Order	Suliformes	1		1		1
70	Oriental Darter Anhinga melanogaster (Pennant, 1769)	Anhingidae	А	AqA	Resident	Near Threatened
71	Great Cormorant Phalacrocorax carbo (Linnaeus, 1758)	Phalacrocoracidae	А	AqA	Resident	Least Concern
72	Little Cormorant Microcarbo niger (Vieillot, 1817)	Phalacrocoracidae	А	AqA	Resident	Least Concern

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	Taxon	Family	Season	Feeding habit	Residential status	IUCN Status
73	Indian Cormorant Phalacrocorax fuscicollis Stephens, 1826	Phalacrocoracidae	w	AqA	Migrant	Least Concern
Order	: Pelecaniformes	1				
74	Cattle Egret Bubulcus ibis (Linnaeus, 1758)	Ardeidae	А	I	Resident	Least Concern
75	Grey Heron Ardea cinerea (Linnaeus, 1758)	Ardeidae	А	с	Resident	Least Concern
76	Indian Pond Heron Ardeola grayii (Sykes, 1832)	Ardeidae	А	с	Resident	Least Concern
77	Intermediate Egret Ardea intermedia (Wagler, 1829)	Ardeidae	А	С	Resident	Least Concern
78	Great Egret Ardea alba (Linnaeus, 1758)	Ardeidae	А	AqA	Resident	Least Concern
79	Little Egret Egretta garzetta (Linnaeus, 1766)	Ardeidae	А	AqA	Resident	Least Concern
80	Black-crowned Night Heron Nycticorax nycticorax (Linnaeus, 1758)	Ardeidae	A	AqA	Resident	Least Concern
81	Purple Heron Ardea purpurea (Linnaeus, 1766)	Ardeidae	A	AqA	Resident	Least Concern
82	Yellow Bittern Ixobrychus sinensis (Gmelin, 1789)	Ardeidae	S	AqA	Migrant	Least Concern
83	Striated Heron Butorides striata (Linnaeus, 1758)	Ardeidae	A	AqA	Resident	Least Concern
84	Red-naped Ibis Pseudibis papillosa (Temminck, 1824)	Threskiornithidae	A	С	Resident	Least Concern
85	Black-headed Ibis Threskiornis melanocephalus (Latham, 1790)	Threskiornithidae	А	с	Resident	Near Threatened
86	Glossy Ibis Plegadis falcinellus (Linnaeus, 1766)	Threskiornithidae	w	с	Migrant	Least Concern
87	Eurasian Spoonbill Platalea leucorodia Linnaeus, 1758	Threskiornithidae	w	С	Migrant	Least Concern
Order	: Accipitriformes					
88	Osprey Pandion haliaetus (Linnaeus, 1758)	Pandionidae	А	с	Migrant	Least Concern
89	Black Kite Milvus migrans (Boddaert, 1783)	Accipitrida <u>e</u>	А	с	Resident	Least Concern
90	Black-winged Kite Elanus caeruleus (Desfontaines, 1789)	Accipitridae	А	С	Resident	Least Concern
91	Brahminy Kite Haliastur indus (Boddaert, 1783)	Accipitridae	w	С	Migrant	Least Concern
92	Shikra Accipiter badius (Gmelin, 1788)	Accipitridae	А	С	Resident	Least Concern
93	Common Buzzard Buteo buteo (Linnaeus, 1758)	Accipitridae	w	С	Migrant	Least Concern
94	White-eyed buzzard Butastur teesa (Franklin, 1831)	Accipitridae	А	С	Resident	Least Concern
95	Himalayan Vulture Gyps himalayensis (Hume, 1869)	Accipitridae	w	С	Migrant	Near Threatened
96	Egyptian Vulture Neophron percnopterus (Linnaeus, 1758)	Accipitridae	А	С	Resident	Endangered
97	Montagu's Harrier Circus pygargus (Linnaeus, 1758)	Accipitridae	w	С	Migrant	Least Concern
Order	Strigiformes					
98	Spotted Owlet Athene brama (Temminck, 1821)	Strigidae	А	С	Resident	Least Concern
Order	: Bucerotiformes					
99	Eurasian Hoopoe Upupa epops (Linnaeus, 1758)	Upupidae	А	I	Resident	Least Concern
100	Indian Grey Hornbill Ocyceros birostris (Scopoli, 1786)	Bucerotidae	А	F	Resident	Least Concern
Order	: Coraciiformes					
101	Common Kingfisher Alcedo atthis (Linnaeus, 1758)	Alcedinidae	А	AqA	Resident	Least Concern
102	White-throated Kingfisher Halcyon smyrnensis (Linnaeus, 1758)	Alcedinidae	А	С	Resident	Least Concern
103	Pied Kingfisher Ceryle rudis (Linnaeus, 1758)	Alcedinidae	А	AqA	Resident	Least Concern
104	Green Bee-eater Merops orientalis (Latham, 1801)	Meropidae	А	I	Resident	Least Concern
105	Blue-tailed Bee-eater Merops philippinus (Linnaeus, 1766)	Meropidae	S	I	Migrant	Least Concern
106	Indian Roller Coracias benghalensis (Linnaeus, 1758)	Coraciidae	А	С	Resident	Least Concern
Order	: Piciformes					
107	Brown-headed Barbet Psilopogon zeylanicus (Gmelin, 1788)	Megalaimidae	А	F	Resident	Least Concern
108	Coppersmith Barbet <i>Psilopogon haemacephalus</i> (Muller, 1776)	Megalaimidae	A	F	Resident	Least Concern
109	Black-rumped Flameback <i>Dinopium benghalense</i> (Linnaeus, 1758)	Picidae	А	I	Resident	Least Concern

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	Taxon	Family	Season	Feeding habit	Residential status	IUCN Status
Order	Psittaciformes					
110	Alexandrine Parakeet Psittacula eupatria (Linnaeus, 1766)	Psittaculidae	А	F	Resident	Near Threatened
111	Rose-ringed Parakeet Psittacula krameri (Scopoli, 1769)	Psittaculidae	А	F	Resident	Least Concern
112	Plum-headed Parakeet <i>Psittacula cyanocephala</i> (Linnaeus, 1766)	Psittaculidae	А	F	Resident	Least Concern
Order	Passeriformes				<u>.</u>	
113	Indian Golden Oriole Oriolus kundoo Sykes, 1832	Oriolidae	S	0	Migrant	Least Concern
114	Black Drongo Dicrurus macrocercus (Vieillot, 1817)	Dicruridae	А	Ι	Resident	Least Concern
115	Hair-crested Drongo Dicrurus hottentottus (Linnaeus, 1766)	Dicruridae	w	I	Migrant	Least Concern
116	Indian paradise Flycatcher Terpsiphone paradisi (Linnaeus, 1758)	Monarchidae	S	I	Migrant	Least Concern
117	Long-tailed Shrike Lanius schach (Linnaeus, 1758)	Laniidae	А	I	Resident	Least Concern
118	Bay-backed Shrike Lanius vittatus (Valenciennes, 1826)	Laniidae	А	Ι	Resident	Least Concern
119	Brown Shrike Lanius cristatus (Linnaeus, 1758)	Laniidae	W	Ι	Migrant	Least Concern
120	Common Raven Corvus corax (Linnaeus, 1758)	Corvidae	А	С	Resident	Least Concern
121	Rufous Treepie Dendrocitta vagabunda (Latham, 1790)	Corvidae	А	0	Resident	Least Concern
122	Grey-headed Canary Flycatcher Culicicapa ceylonensis (Swainson, 1820)	Stenostiridae	w	I	Migrant	Least Concern
123	Crested Lark Galerida cristata (Linnaeus, 1758)	Alaudidae	А	I	Resident	Least Concern
124	Yellow-bellied Prinia Prinia flaviventris (Delessert, 1840)	Cisticolidae	W	I	Migrant	Least Concern
125	Common Tailorbird Orthotomus sutorius (Pennant, 1769)	Cisticolidae	А	I	Resident	Least Concern
126	Rufous–fronted Prinia Prinia buchanani (Blyth, 1844)	Cisticolidae	А	I	Resident	Least Concern
127	Ashy Prinia Prinia socialis (Sykes, 1832)	Cisticolidae	А	Ι	Resident	Least Concern
128	Plain Prinia <i>Prinia inornata</i> (Sykes, 1832)	Cisticolidae	А	Ι	Resident	Least Concern
129	Zitting Cisticola Cisticola juncidis (Rafinesque, 1810)	Cisticolidae	А	Ι	Resident	Least Concern
130	Blyth's Reed Warbler Acrocephalus dumetorum (Blyth, 1849)	Acrocephalidae	W	Ι	Migrant	Least Concern
131	Barn Swallow Hirundo rustica (Linnaeus, 1758)	Hirundinidae	S	I	Migrant	Least Concern
132	Plain Martin Riparia paludicola (Vieillot, 1817)	Hirundinidae	S	I	Migrant	Least Concern
133	Red-Rumped Swallow Cecropis daurica (Laxmann, 1769)	Hirundinidae	S	I	Migrant	Least Concern
134	Wire-tailed Swallow Hirundo smithii (Leach, 1818)	Hirundinidae	S	Ι	Migrant	Least Concern
135	Streak-throated Swallow Petrochelidon fluvicola (Blyth, 1855)	Hirundinidae	А	Ι	Resident	Least Concern
136	Red-vented Bulbul Pycnonotus cafer (Linnaeus, 1766)	Pycnonotidae	А	F	Resident	Least Concern
137	Himalayan Bulbul Pycnonotus leucogenys (Gray, 1835)	Pycnonotidae	А	F	Resident	Least Concern
138	Common Chiffchaff Phylloscopus collybita (Vieillot, 1817)	Phylloscopidae	w	I	Migrant	Least Concern
139	Siberian Chiffchaff Phylloscopus tristis (Blyth, 1843)	Phylloscopidae	А	I	Resident	Least Concern
140	Indian White-eye Zosterops palpebrosus (Temminck, 1824)	Zosteropidae	w	0	Migrant	Least Concern
141	Common Babbler Argya caudate (Dumont, 1823)	Leiothrichidae	А	I	Resident	Least Concern
142	Jungle Babbler Argya striata (Dumont, 1823)	Leiothrichidae	A	I	Resident	Least Concern
143	Large Grey Babbler Argya malcolmi (Sykes, 1832)	Leiothrichidae	A	I	Resident	Least Concern
144	Asian Pied Starling Gracupica contra (Linnaeus, 1758)	Sturnidae	A	0	Resident	Least Concern
145	Bank Myna Acridotheres ginginianus (Latham, 1790)	Sturnidae	А	0	Resident	Least Concern
146	Brahminy Starling Sturnia pagodarum (Gmelin, 1789)	Sturnidae	A	Ι	Resident	Least Concern
147	Common Myna Acridotheres tristis (Linnaeus, 1766)	Sturnidae	А	0	Resident	Least Concern
148	Common Starling Sturnus vulgaris (Linnaeus, 1758)	Sturnidae	А	0	Resident	Least Concern
149	Black Redstart Phoenicurus ochruros (Gmelin, 1774)	Muscicapidae	w	I	Migrant	Least Concern
150	Bluethroat Luscinia svecica (Linnaeus, 1758)	Muscicapidae	w	I	Migrant	Least Concern
151	Brown Rock Chat Oenanthe fusca (Blyth, 1851)	Muscicapidae	А	I	Resident	Least Concern

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	Taxon	Family	Season	Feeding habit	Residential status	IUCN Status
152	Indian Robin Copsychus fulicatus (Linnaeus, 1766)	Muscicapidae	А	I	Resident	Least Concern
153	Oriental Magpie Robin Copsychus saularis (Linnaeus, 1758)	Muscicapidae	А	I	Resident	Least Concern
154	Pied Bushchat Saxicola caprata (Linnaeus, 1766)	Muscicapidae	А	I	Resident	Least Concern
155	Siberian Stonechat Saxicola maurus (Pallas, 1773)	Muscicapidae	w	I	Migrant	Least Concern
156	White-tailed Stonechat Saxicola leucurus (Blyth, 1847)	Muscicapidae	А	I	Resident	Least Concern
157	Purple Sunbird Cinnyris asiaticus (Latham, 1790)	Nectariniidae	А	I	Resident	Least Concern
158	Baya Weaver Ploceus philippinus (Linnaeus, 1766)	Ploceidae	А	0	Resident	Least Concern
159	Streaked Weaver Ploceus manyar (Horsfield, 1821)	Ploceidae	А	I	Resident	Least Concern
160	Indian Silverbill Euodice malabarica (Linnaeus, 1758)	Estrildidae	А	G	Resident	Least Concern
161	Scaly-breasted Munia Lonchura punctulata (Linnaeus, 1758)	Estrildidae	А	G	Resident	Least Concern
162	House Sparrow Passer domesticus (Linnaeus, 1758)	Passeridae	А	G	Resident	Least Concern
163	Citrine Wagtail Motacilla citreola (Pallas, 1776)	Motacillidae	w	I	Migrant	Least Concern
164	Grey Wagtail Motacilla cinerea (Tunstall, 1771)	Motacillidae	w	I	Migrant	Least Concern
165	Paddyfield Pipit Anthus rufulus (Vieillot, 1818)	Motacillidae	А	I	Resident	Least Concern
166	Western Yellow Wagtail Motacilla flava (Linnaeus, 1758)	Motacillidae	w	1	Migrant	Least Concern
167	White Wagtail Motacilla alba (Linnaeus, 1758)	Motacillidae	w	I	Migrant	Least Concern

A-Throughout year | S-Summer | W-Winter | O-Omnivorous | AqA- Aquatic animal Feeder | C-Carnivorous | F-Frugivorous | G-Granivorous | I-Insectivorous | H-Herbivorous.

poaching activities, as well as the regulation of land reclamation and fishing practices, as detailed in the work by Brraich & Saini (2019). The intricate relationship between wetlands and avian populations is influenced by various physical and biological attributes of wetland environments. Birds rely on wetlands for their daily and seasonal needs, including food and other lifesustaining activities. Therefore, effective conservation and management measures are essential to attract migratory birds to wetland areas. A comprehensive examination of wetland ecosystems and migratory bird populations is necessary to better understand their interaction. Public awareness regarding the vital role of wetlands in supporting migratory birds is crucial, and support for restoration and maintenance initiatives aimed at safeguarding these vital habitats is highly encouraged.

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Faunistic overview of the freshwater zooplankton from the urban riverine habitats of Pune, India

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Abstract: Urbanization modifies the physical, chemical, and biological nature of all ecosystems including rivers. Such changes negatively impact all aquatic biodiversity including the freshwater zooplankton. Given the fast pace of urbanization in all the major cities across India, the aim is to provide a faunistic overview of Rotifera, Cladocera, and Ostracoda from two polluted rivers flowing through Pune, one of the rapidly growing cities in the state of Maharashtra, India. A one-year survey of three localities on the rivers Mula & Mutha and data from published literature on another locality revealed the presence of 73 species which includes 47 rotifers, 15 cladocerans, and 11 ostracods. A higher species number of rotifers was seen at lesser polluted localities while cladocerans and ostracods occurred even in the most urbanized sampling locality. Many of the species found were commonly observed species from the region. Epizoic associations of cladocerans and rotifers and red coloration in the former group were observed during a low dissolved oxygen phase in both rivers. Such observations underscore the potential bioindicator value of these small animals to the impacts of urbanization.

Keywords: Biodiversity, Cladocera, Epizoic, Mula-Mutha river, Ostracoda, pollution, Rotifera, urbanization.

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Author contributions: All the authors contributed equally to the idea, methodology, field work and analysis. AlV and SMP contributed to writing of the manuscript and its subsequent revisions

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INTRODUCTION

Urbanization refers to the mass migration of human populations from rural to urban settings (Kuddus et al. 2020). More than half the world's population (~4.3 billion) lives in urban areas which may increase to six billion by the year 2041 (Ritchie & Roser 2018; UNDESA 2018). Thus, urban areas, especially in developing countries like India, are expanding at an exponential rate assisted by the ever-increasing population (Henderson 2002; Cohen 2006; Onda et al. 2019; Kuddus et al. 2020). Such rapid urbanization can have adverse effects on different ecosystems by way of native species loss and/or an increase in the number of non-native species (McMichael 2000; McKinney 2002, 2006).

Rivers are an important component of many urban centres providing water, power, and means of transport besides harbouring high biodiversity (McMichael 2000; Everard & Moggridge 2012; Tran Khac et al. 2018). Many studies have shown that anthropogenic activities like modification of the river channel/bank and untreated waste disposal impact riverine biodiversity in multiple ways which include cultural eutrophication and biotic homogenization to name a few (Blair 2001; Ouyang et al. 2002; Dudgeon et al. 2006; Schindler 2012; Braghin et al. 2018; Du et al. 2023).

Freshwater zooplankton is a well-represented group of invertebrates in rivers and forms an important component of aquatic food chains (Dumont & Negrea 2002; Liu et al. 2020). Zooplankton communities respond to physical and chemical changes in the riverine habitats by displaying variations in their growth, community composition, density, diversity, and distribution (Bērziņš & Pejler 1987, 1989; Duggan et al. 2001; Nogrady et al. 1993; Hulyal & Kaliwal 2008; Jeppesen et al. 2011; Adamczuk et al. 2015; Du et al. 2023).

Literature exists on the different limnological aspects of lotic and lentic habitats in India, though, several of them, especially in the case of zooplankton, have issues like species misidentifications (Sharma & Sharma 2021). Data from reliable studies point to species losses occurring in response to changes in environmental variables like nutrients (phosphorus and nitrogen), dissolved oxygen, turbidity and water flow (Padmavati & Goswami 1996; Arora & Mehra 2003; Rajaram & Das 2008; Padhye & Dahanukar 2015).

Pune is a rapidly growing city in India where its population has grown exponentially within the last 70 years from 3.75 lakhs (1941) to 5 million (2011) and is expected to be >9 million by 2035 (see Butsch et al. 2017; UNDESA 2018). Mula & Mutha, the two rivers that provide water to this urban centre are highly polluted within the city limits due to various anthropogenic activities (Wagh & Ghate 2008; Padhye 2020). Existing faunal literature on these rivers suggests decreasing species numbers across animal groups like odonates, molluscs, fish and birds due to this urbanization effects (Gole 1983; Kharat et al. 2001, 2003; Wagh & Ghate 2008; Kulkarni & Subramanian 2013; Kulkarni et al. 2021). Studies on zooplankton from a single locality on the Mula River have also shown a similar trend (Vanjare et al. 2010; Padhye 2020).

The present study aims to provide a faunistic overview of Rotifera, Cladocera, and Ostracoda of Mula & Mutha rivers passing through the urban part of Pune, Maharashtra. Additionally, peculiar observations and habits of some of the species found in the study are also commented upon.

MATERIALS AND METHODS

Site

Mula & Mutha are tributaries of the Bhima River (Pune, Maharashtra) and are heavily polluted within the city limits, receiving large amounts of untreated waste. Both rivers originate in the Western Ghats, meet in Pune and then join the Bhima River outside the city limits. Floating vegetation (*Pistia* sp., *Eichhornia* sp., and *Lemna* sp.) is observed here frequently and in high densities, after post-monsoon in the urban regions while submerged (*Hydrilla* sp.) and emergent vegetation (*Typha* sp.) is also seen at many places.

Two sampling sites were selected along the Mula River (Ram-Mula confluence & Aundh Bridge) and Mutha River (Vitthalwadi & Garware College) within Pune City for the study (Figure 1). Urbanization around Ram-Mula confluence and Vitthalwadi is comparatively lower than Aundh Bridge and Garware College sites (authors pers. obs. 19 November 2017).

Field and laboratory work

Qualitative sampling was carried out at Ram-Mula confluence, Vitthalwadi, and Garware College between post monsoon and winter season in 2017–18. Sample aliquots were taken (~3–4) from a stretch of ~100 m at each site and concentrated in a single container (100 ml). Effort was made to collect the sample once in each season. A plankton net of 53-micron and hand net of 100-micron mesh size was used for the collection. The sediment was gently disturbed, and water was filtered subsequently with the hand net for better

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Figure 1. The collection localities along the Mula & Mutha rivers, Pune.

representation of meiobenthic species. The samples were preserved in 4% formalin. Dissolved Oxygen was taken using a DO probe (Hanna) and salinity, pH, water temperature were taken at each sampling station using a multiparameter probe (Eutech). Identifications were done under light microscope (Olympus CH20i) and stereo microscope (Magnus MS 24). Identification was done using standard literature available for the respective groups (Supplementary list 1). Zooplankton data for the Aundh Bridge site was taken from Vanjare et al. (2010) and Padhye & Dahanukar (2015) since the site was inaccessible during the sampling period. Urbanization extent was assessed qualitatively by visual inspection.

RESULTS AND DISCUSSION

Environmental data recorded during the study are shown in supplementary Table 1. The pH ranged from 7.12–8.6, dissolved oxygen from 0.25–11.8 mg/L, water temperature from 18.6–32°C and salinity from 105–386 ppm. No environmental data was collected for the Garware College site.

Seventy-three species of three different zooplankton groups were documented, of which rotifers being the most species rich with 47 species, followed by cladocerans—15 and ostracods—11, respectively (supplementary Table 2). Rotifers were reported from only three localities while cladocerans and ostracod species were observed at all the four sampling stations (Figure 2). Sampling stations having lesser urbanization had more species of rotifers (Ram Mula = 31 & Vitthalwadi = 34) with no species seen at the locality in the city centre (Garware College, Figure 1). Maximum species of cladocerans and ostracods were found at the Aundh Bridge (cladocerans = 11; ostracods = 7) though, representatives of these groups were also found at the Garware College site.

Among the 47 rotifer species, 41 were from the order Ploima, five from the order Flosculariaceae (subclass Monogononta) and one from the family Philodinidae within the subclass Bdelloidea. The Brachionidae family was the most species-rich with 13 species followed by Lecanidae with nine species while eight rotifer families

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were represented by a single species only. Rotifer genera *Brachionus* and *Lecane* (n = 9 each) were found in high numbers which is typical of these genera in tropical waters (Arora & Mehra 2002). Notable findings include rotifers with a restricted geographic distribution like *Brachionus durgae* (Ram Mula & Vitthalwadi), *Epiphanes brachionus spinosa* (all sites) and *Lecane stenroosi* (Vitthalwadi) were also seen in the study. Three predatory rotifers from the family Asplanchnidae, viz. *Asplanchnopus multiceps, Asplanchna brightwellii and Asplanchna priodonta* were also observed at three of the four sites (Image 2). Most of the recorded rotifer species are common and cosmopolitan in distribution.

Chydorids were the most species rich cladoceran group with five species followed by Daphniidae with three. Both the moinid species were observed in high densities at the site located on river Mutha in the most urbanized region of Pune city (Garware College). Some of the species such as *Simocephalus mixtus, Macrothrix spinosa*, and *Ilyocryptus spinifer* are known to occur seasonally at one of the sites on Mula River (Aundh Bridge). Most of these species are commonly known from the region with *Leydigia* (*Neoleydigia*) *ciliata* and *I. spinifer* being the most commonly occurring species in Pune (Padhye et al. 2023) (Image 2).

Only one species from the ostracod genus, *llyocypris* sp., was seen at all the four sampling points while the oriental endemics like *Stenocypris derupta* Vávra, *Plesiocypridopsis* cf. *dispar* (Hartmann, 1964) and

Chrissia formosa (Klie, 1938) were seen at the Ram-Mula confluence only. *Heterocypris incongruens* (Ramdohr, 1808) reported from Garware college is a cosmopolitan species known to tolerate high levels of pollution (Karakaş-Sarı & Külköylüoğlu 2008). *Plesiocypridopsis* cf *dispar* and *Stenocypris* sp. were seen near a natural spring pouring into the Mutha River at the Vitthalwadi site (Image 1).

Seventy-three species of rotifers, cladocerans, and ostarcods from just four locations in Pune City is a good number as compared to riverine fauna documented from some other urban zones of India. Arora & Mehra (2003) documented 89 rotifers from river Yamuna in Delhi, Hulyal & Kaliwal (2007) found 10 rotifers and six cladocerans in Almatti Reservoir of Bijapur, while Kamboj & Kamboj (2020) observed 10 rotifers and eight cladocerans in the Ganga River, Uttarakhand, and Rao (2001) reported 17 rotifers and six cladocerans from the river Ganga between Rishikesh and Kanpur, Uttar Pradesh. Reliable faunistic studies providing species numbers of ostracods from such habitats are not available. The variation observed in the species numbers between these studies could be explained by many possible differences in the geomorphological and geochemical features of the rivers, the local environmental conditions and biotic conditions such as predation pressure. Still, the trend in species numbers concerning the specific taxonomical groups was consistent with other studies, i.e., rotifers having the most number of species as



Figure 2. Number of species occurring at each of the study sites.



Image 1. Ostracods: A—*Cyprinotus cingalensis* Brady, 1886 | B—*Chrissia formosa* (Klie, 1938) | C—*Physocypria* sp. | D—*Plesiocypridopsis* cf. *dispar* (Hartmann, 1964). A–B images taken at 40x and C–D at 100x final magnification. © Yugandhar Shinde.

compared to cladocerans and ostracods (Sharma & Naik 1996; Arora & Mehra 2003; Sharma 2011). Similarly, the species number distribution between the order/families of each group was also in agreement with the studies available in India and other regions (Ploima being the most species-rich order in rotifers, Chydoridae being the most species-rich cladoceran family)

Occurrence of common species such as, Moina micrura, Brachionus spp., Polyarthra sp., and Heterocypris incongruens at such sites imply the ability of these organisms to tolerate a wide range of environmental conditions (Nogrady et al. 1993; Külköylüoğlu et al. 2018). Cultural eutrophication which can happen due to rapid urbanization (Dudgeon et al. 2006; Schindler 2006; Lodi et al. 2011) is known to affect zooplankton richness and increase the dominance of such common species in many cases (Nogrady et al. 1993; Dodson et al. 2000; Yuan & Pollard 2018; Kambhoj & Kambhoj 2020). Certain zooplankton species are known to evolve rapidly to cope with such environmental change to persist in unfavourable conditions like Daphnia magna adapting to an increased water temperature (Brans et al. 2017). The presence of such generalist species can lead to biotic homogenization, i.e., reduction in β diversity

across space (Wang et al. 2021), which is a worldwide phenomenon noticed in disturbed ecosystems (Blair 2001; McKinney 2002, 2006, 2008; Liu et al. 2020). Kulkarni et al. (2021) showed that the species richness of aquatic gastropods decreased along an urbanization gradient including the rivers studied here with only invasive species reported in the most urbanized locality. This locality was very close to the Garware College site for which we recorded the lowest species numbers (Supplementary Table 1; Figure2).

The presence or absence of certain species/ groups like zooplankton can be applied to indicate environmental disturbances (Duggan et al. 2001; Hulyal & Kaliwal 2008; Du et al. 2023). Certain species from the genus *Brachionus* are indicators of eutrophic conditions of water (Mäemets 1983) and we found seven species in our collections especially *B. angularis*, *B. rubens*, *B. calyciflorus*). An epizoic association of the rotifer *Brachionus rubens* and cladoceran *Moina macrocopa* was observed at the Aundh Bridge site with over 40 individuals of *B. rubens* attached to a single individual of *M. macrocopa*. This association was seen during the peak summer months (April/May) when the DO level was very low (Vanjare et al. 2010) and not observed at any



Image 2. Representative rotifers and cladocerans: A—Brachionus calyciflorus Pallas, 1766 | B—Mytilina ventralis (Ehrenberg, 1830) | C— Asplanchnopus multiceps (Schrank, 1793) | D—Lecane luna (Müller, 1776) | E—Lecane curvicornis (Murray, 1913) | F—Brachionus falcatus Zacharias, 1898 | G—Moina micrura Kurz, 1874 | H—Epizoic interaction of Brachionus rubens with Moina macrocopa (Straus, 1820) | I—Ilyocryptus spinifer Herrick, 1882 | J—Leydigia ciliata Gauthier, 1939 | F—Ovalona cambouei (Guerney et Richard, 1893) | L. Kurzia longirostris (Daday, 1898). H—image taken at 400x final magnification | A–G & I–L—images taken at 100x final magnification. © (A–F)—Avinash Vanjare, (G–L)—Sameer Padhye.

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other studied site. Dark red-coloured cladoceran species (*Moina micrura, M. macrocopa,* and *K. longirostris*) were spotted at the Aundh bridge site during the summer months when the DO was the lowest (Vanjare et al. 2010). We also observed a faint red coloration in the *Moina* species collected at the other localities in the winter samples. This colour change occurs due to haemoglobin production as a response to low dissolved oxygen in the water (Fox 1949).

Our study was based only on four sites on rivers Mula Mutha with no data available for upstream patches of both the rivers where the urbanization is relatively lower than the main city. Exhaustive sampling including more upstream localities would certainly increase the species number. Studying the environmental change indicator potential of these zooplankton groups along with longterm monitoring of their community dynamics will surely help us understand and devise ways of monitoring the impacts of urbanization. Conducting such studies is crucial in light of biodiversity loss happening as a consequence of increasing urbanization (Kharat et al. 2001, 2003).

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Supplementary Table 1. Environmental parameters at the site.

Site/ Parameter	Temperature (°C)	рН	Salinity (ppm)	DO (mg/L)	Remarks
Aundh Bridge	23.8-32.0	7.3–8.2	227–386	1.2–7.3	See Vanjare et al. 2010
Aundh Bridge	24-31.2	7.12-8.05	105-386	0.81-4.15	See Padhye & Dahanukar 2015
Ram-Mula	18.6-24.0	7.2–7.8	227–382	0.44-8.80	Current study
Vitthalwadi	22.0-30.1	7.2–8.6	129-356	0.25-11.8	Current study

Supplementary Table 2. List of species observed at the sites.

Phylum Rotifera				
Asplanchnidae	Asplanchna brightwellii Gosse, 1850			
, spianei indae	Asplanchna priodonta Gosse, 1850			
	Asplanchnopus multiceps Schrank, 1793			
	Brachionus angularis (Gosse, 1851)			
	Brachionus bidentata (Anderson, 1889)			
Brachionidae	Brachionus calyciflorus (Pallas, 1766)			
	Brachionus caudatus (Barrois & Daday, 1894)			
	Brachionus diversicornis Daday, 1883			
	Brachionus durgae (Dhanapathi, 1974)			
	Brachionus falcatus (Zacharias, 1898)			
	Brachionus quadridentatus (Hermann, 1783)			
	Brachionus rubens (Ehrenberg, 1838)			
	Keratella cochlearis (Gosse, 1851)			
	Keratella tropica (Apstein, 1907)			
	Plationus patulus (Müller, 1786)			
	Platyias quadricornis (Ehrenberg, 1832)			
Eninhanidae	Epiphanes brachionus spinosa (Rousselet, 1901)			
Epipilanidae	Beauchampiella eudactylota (Gosse, 1886)			
	Euchlanis dilatata (Ehrenberg, 1832)			
Euchlanidae	Tripleuchlanis plicata (Levander, 1894)			
	Lecane bulla (Gosse, 1851)			
Lecanidae	Lecane closterocerca (Schmarda, 1859)			
Lecanidae	Lecane hamata (Stokes, 1896)			
	Lecane leontina (Turner, 1892)			
	Lecane luna (Müller, 1776)			
	Lecane lunaris (Ehrenberg, 1832)			
	Lecane papuana (Murray, 1913)			
	Lecane curvicornis (Murray, 1913)			
	Lecane stenroosi (Meissner, 1908)			
	Colurella obtusa (Gosse, 1886)			
Lepadellidae	Lepadella (Heterolepadella) ehrenbergii (Perty, 1850)			
	Lepadella (Lepadella) ovalis (Müller, 1786)			
	Squatinella lamellaris (Müller, 1786)			
Mytilinidae	Mytilina bisulcata (Lucks, 1912)			
	Mytilina ventralis (Ehrenberg, 1830)			
Notommatidae	Cephalodella sp.			
Notominatione	Monommata sp.			
	Taphrocampa annulosa (Gosse, 1851)			

Phylum Rotifera					
Scaridiidae	Scaridium longicaudum (Müller, 1786)				
Synchaetidae	Polyarthra vulgaris (Carlin, 1943)				
Trichocercidae	Trichocerca cylindrica (Imhof, 1891)				
Hexarthridae	Hexarthra mira (Hudson, 1871)				
Filiniidae	Filinia longiseta (Ehrenberg, 1834)				
Testudinellidae	Testudinella patina (Hermann, 1783)				
Flosculariidae	Sinantherina semibullata (Thorpe, 1893)				
	Lacinularia elliptica Shephard, 1897				
Philodinidae	Rotaria neptunia (Ehrenberg, 1830)				
	CLADOCERA				
Sididae	Latonopsis australis Sars, 1888 s.lat.				
	Diaphanosoma sarsi (Richard, 1895)				
	Daphnia (Ctenodaphnia) lumholtzi (Sars, 1885)				
Daphniidae	Ceriodaphnia cornuta (Sars, 1885)				
	Simocephalus (Simocephalus) mixtus (Sars, 1903)				
Moinidae	Moina macrocopa (Straus, 1820)				
	Moina micrura (Kurz, 1874)				
Macrothricidae	Macrothrix spinosa (King, 1853)				
	Macrothrix triserialis (Brady, 1886)				
Ilyocryptidae	Ilyocryptus spinifer (Herrick, 1882)				
	Flavalona cheni (Sinev, 2001)				
Chydoridae	Ovalona cambouei (Guerney et Richard, 1893)				
	Kurzia longirostris (Daday, 1898)				
	Leydigia (Neoleydigia) ciliata (Gauthier, 1939)				
	Chydorus eurynotus (Sars, 1901)				
	OSTRACODA				
	Chrissia formosa (Klie, 1938)				
Cyprididae	Stenocypris hislopi (Ferguson, 1969)				
-)	Stenocypris derupta (Vávra, 1906)				
	Cypris granulata (Daday, 1898)				
	Plesiocypridopsis cf dispar (Hartmann, 1964)				
	Heterocypris incongruens (Ramdohr, 1808)				
	Hemicypris pyxidata (Moniez, 1892)				
	Hemicypris ovata Sars, 1903				
	Cyprinotus cingalensis Brady, 1886				
Candonidae	Physocypria sp.				
llyocyprididae	Ilyocypris sp.				

Supplementary List 1. List of references used for identification, classification and nomenclature used for identifying Rotifera, Claodocera, and Ostracoda.

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Utilization of a new restoration technique for the rehabilitation of a degraded mangrove ecosystem: a case study from Koggala Lagoon, Sri Lanka

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Abstract: Mangrove ecosystems, amongst the most productive and biologically complex on Earth, are being degraded worldwide, and their widespread decline during the past decades has affected vital ecosystem services. Mangroves at Koggala lagoon on the southern coast of Sri Lanka have been degraded at an alarming rate due to agricultural practices, coastal zone development, and tourism activities. Most of the banks are heavily eroded due to boat and sea plane activities, and the mangrove ecosystem has been significantly damaged. Implementation of a scientific rehabilitation project was needed to restore this degraded mangrove ecosystem, and research was carried out to enrich the mangrove community by re-establishing mangroves on the eroded banks using corrosion-resistant plastic barrels. The sustainability of replanted mangroves was monitored under phase one and the re-establishment of ecological functions in the mangrove community was monitored under phase two. The accumulated biomass carbon during the period of two and half years was calculated by an allometric equation suggested for calculating biomass carbon of mangroves using the girth and height of individuals. The highest rate of girth increment was observed by the 24th month from establishment, whilst the growth rate declined between the 18th and the 30th months. During the study period, the average above-ground and below-ground biomass per barrel showed a linear increment. Our case study showed that the new method used for restoration is successful in establishing mangroves in sites with high erosion. This restoration technique was successful in coping with the situation in Koggala lagoon where previous restoration attempts were failures. Thus, we recommend this restoration method for sites facing the threat of severe erosion.

Keywords: Allometric equations, carbon stocks, ecological function, eroded banks, growth rate, mangrove establishment, restoration technique.

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Our interest is in you

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Author contributions: M.G.G Dhanushka contributed to the study conception and design. Experimental arrangements and data collection were conducted by M.G.G Dhanushka. Data analysis was performed by K.M.G.G. Jayasuriya. The first draft of the manuscript was written by M.G.M. Prasanna, I.H. Vitanage and K.M.G.G. Jayasuriya and all authors commented on previous versions of the manuscript. All authors read and approved the final manuscript.

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INTRODUCTION

Mangroves are among the world's most productive and biologically complex ecosystems, acting as bridges between land and sea. Thriving in intertidal areas along tropical and subtropical coastlines, mangroves consist of salt-tolerant woody plant species and are commonly found in lagoons, bays, and estuaries (Prasanna & Ranawana 2014), including several locations in Sri Lanka (Ministry of Environment 2012). Mangroves provide a multitude of essential goods and services crucial for human well-being and survival. They play a crucial role in maintaining the ecological integrity of coastal zones and provide many ecosystem services categorized as provisioning, regulatory, supportive as well as cultural services (Donato et al. 2011; Feller et al. 2017). Carbon sequestration is one of the most significant services provided by mangroves, as they possess a remarkable capacity to capture and retain high amounts of carbon in the soil owing to high productivity compared to other terrestrial ecosystems (Alongi 2014). Consequently, mangroves contribute significantly to the reduction of greenhouse gases and aid climate change mitigation efforts. Despite their importance, mangrove ecosystems have suffered global degradation (Duke et al. 2017; Feller et al. 2017), resulting in the decline of their ecosystem services over the past decades. According to Mukherjee et al. (2014), approximately 60% of major global ecosystem services have been degraded either totally or partially.

Mangrove Ecosystems of Sri Lanka

Sri Lanka, a tropical island with numerous estuaries and lagoons, possesses a diverse range of mangrove ecosystems along its coastline (Edirisinghe et al. 2012). These mangrove ecosystems consist of two types of plant communities: true mangroves and mangrove associates. True mangroves are woody plants that exclusively occur in mangrove forests, displaying specific adaptations to the environment and physiological mechanisms to tolerate high salinity levels (Tomlinson 2016). In contrast, mangrove associates are primarily herbaceous plants found in terrestrial or aquatic habitats, but they can also be present within the mangrove ecosystem (Tomlinson 1986). Most Sri Lankan mangrove areas (92.6%) are confined to the dry and intermediate zones. Similar to the global situation, Sri Lankan mangroves were also degraded especially during the last four decades due to various anthropogenic activities (Wickramasinghe et al. 2022).

Nevertheless, mangrove research and rehabilitation

efforts have progressed steadily over the last several decades as the importance of mangrove ecosystems been better understood and documented. has Furthermore, the mangrove ecosystems are considered a predominantly important ecosystem for coastal communities due to their provision of ecosystem services, such as supplying timber and fuel wood, supporting fisheries, sediment trapping, coastal defence and carbon storage (Donato et al. 2011; Feller et al. 2017). Amongst all the ecosystems across the tropics, mangrove ecosystems are considered one of the most threatened (Duke et al. 2017) mainly due to impacts from anthropogenic activities including conversion to agriculture and aquaculture as well as urbanisation and pollution (Feller et al. 2017). Under these circumstances, huge efforts are being put into mangrove rehabilitation and restoration in degraded areas. Though, such largescale efforts are generally unsuccessful due to various reasons such as poor species selection, inappropriate location selection and poor knowledge of mangrove ecology as well as physiology (Kodikara et al. 2017). However, when elements of species biology and hydrological requirements are incorporated into the design and implementation of rehabilitation projects with an appropriate knowledge base, some efforts are becoming more successful (Feller et al. 2017).

Status of Koggala Lagoon mangrove ecosystem

The Koggala lagoon is situated in the Southern province of Sri Lanka, specifically between 5°58'-6°20' N & 80°17'-80°22' E. It encompasses an area of 727 ha and consists of 14 islets (IUCN and Central Environmental Authority, 2006; Gunaratne et al. 2010). Several tributaries, including the Koggala Oya, provide freshwater input to the lagoon. The hydrology and water quality of the lagoon, including salinity and pH, are influenced by heavy rainfall and the characteristics of the lagoon mouth due to its location in the wet zone of the country. Previous studies have reported the presence of 10 true mangrove species in the lagoon. However, the classification of Acrosticum aurium and Dolchandrone spathacia as true mangroves by IUCN and Central Environmental Authority (2006) is disputed by the experts' team of the National Red List (2012), who considers them as mangrove associates. Therefore, the number of true mangrove species identified in the lagoon is recognized as eight. Mangroves are found in a narrow strip surrounding the lagoon's islands and along the stream banks. Unfortunately, due to activities such as boating and sea plane landing, and take-off, many of the banks have undergone degradation and significant

erosion, leading to substantial damage to the mangrove ecosystems in the area.

The structure of the lagoon mouth has changed since 1990 due to the removal of the natural sand barrier (Gunarathne 2011). Consequently, sand started to deposit on the river mouth and the bridge over Pol Oya in Galle-Matara main road, blocking the water flow. A rubble mound groyne system (old groyne) was built in 1997 to prevent the issue. Due to this artificial construction, erosion of the lagoon bank became threatened as the Galle-Matara main road and bridge became vulnerable to sea erosion. Another groyne (new groyne) (Image 1) was established in 2005 to control the said situation (Gunarathne 2011). The outlet (Image 2) has been diverted westward creating an approximately 30–40 m wide open passage to the sea consequently (Gunawickrama & Chandana 2006).

The construction of an artificially built groyne in the Koggala lagoon initially resulted in a reduction of sand deposition. However, it also led to seawater intrusion into the lagoon (Gunawickrama & Chandana 2006). Over time, sand deposition resumed at the river mouth, causing water blockage and a subsequent decrease in water salinity and a rise in water level. The increased water level further contributed to bank erosion within the lagoon. These degraded banks, characterized by high erosion and stream flow, present challenges for natural regeneration and make it impossible to rehabilitate the mangrove community. Additionally, the degradation and heavy erosion of the banks caused by boating and seaplane activities further exacerbate the problem.

Despite previous attempts at planting mangrove seedlings in the Koggala lagoon, the general approach has failed multiple times in recent years. Natural regeneration has not been observed in the degraded banks of the lagoon, necessitating a new restoration approach and the implementation of a continuous monitoring mechanism to ensure the success of mangrove restoration. Therefore, the primary objective of the study was to enhance the mangrove community in the Koggala lagoon using a technique suitable for the prevailing conditions in the lagoon.

METHODS

Establishment of the restoration trail

A controlled plot using general restoration processes could not be established due to the unsuitable ground conditions and heavy erosion of the lagoon banks. A new restoration approach was designed to support restored plants to withstand the bank erosion. In this approach mangrove saplings were planted in plastic barrels.

Empty and well-cleaned chemical plastic barrels (~38 cm diameter and ~79 cm height) were gathered from factories located in the area. The top and bottom of all the barrels were removed. Thirty seven of these barrels were placed in holes excavated in eroded banks of three islands: Thalathuduwa, Kuruluduwa (Image 3) and Ganduwa. Barrels were placed with 60-90 cm spacing between each other, covering ~600 m stretch of the banks. The barrels were filled with soil excavated from the same restoration site. Two true mangrove species occurring in the area, Rhizophora mucronata and R. apiculta were selected as restoring species for this pilot study. These two species were selected as they contain large numbers of prop and stilt roots which assist in the proper establishment of the plant in the planted site. The availability of diaspores at the time of nursery establishment was also considered. Four R. mucronata saplings (~20–35 cm height) and one R. apiculta sapling (~20–35 cm height) were planted in each barrel. Saplings were raised in a nearby nursery using the diaspores collected from trees in the existing vegetation of the Koggala lagoon.

Maintenance and monitoring

Planted seedlings were observed weekly during the first six months, and later monthly. Dead saplings were not replaced as it would affect the final analysis. There was no need to replenish the soil, as the soil in the barrels was not eroded during the period (Image 4). The diameter at breast height (dbh) and height of each sapling in each barrel were recorded on the first day of planting and then after every six months for two and half years. Monthly measurements were not taken as the changes in girth and height were not significant within a month.

Data analysis

Height and dbh increments were separately plotted against time. A logistic four-parameter sigmoidal curve was fitted to determine the pattern of growth (Tsoularis 2001). The growth rate based on height and dbh was calculated separately for six months period from the initial planting date to August 2020. Accumulated biomass carbon during the period of two and half years was calculated using the dbh and height of the individuals with an allometric equation suggested for calculating biomass carbon of mangroves.

Above ground biomass (AGB) for *Rhizophora* mucronata,



Image 1. Satellite image of lagoon outlet with the existing groyne structures in 2010 (Source: Google Earth Pro 2022).



Image 2. The present situation of Koggala Lagoon mouth (Source: Google Earth Pro 2022).



Image 3. First mangrove planting in Kurulu Duwa island in July 2018.



Image 4. The soil in the barrels was not eroded even after a year from planting, another mangrove hedge was planted in barrels parallel to the previous mangrove hedge.

 $\log_{e}(AGB) = 6.247+2.64 \log_{e}(dbh)$ (Amarasinghe & Balasubramaniam 1992b)

and for Rhizophora apiculata,

AGB = 0.251 ρ dbh ^{2.46} (Komiyama et al. 2005)

Bellow ground biomass (BGB) for both species,

BGB = $0.199 \rho^{0.899}$ dbh ^{2.46} (Komiyama et al. 2005)

RESULTS

Survival of plants during the two-and-half years of the monitoring period

After the first six months of establishment, all the *R. mucronata* and *R. apiculata* saplings survived in the study site. Within the next six months period, more *R. apiculata* saplings died compared to *R. mucronata* saplings. After 18 months of establishment, 85 % of the R. *mucronata* and 67 % of R. *apiculata* saplings survived (Figure 1). Thereafter, none of the remaining saplings died during the observation period of 30 months.

Growth of established saplings

The height of both *R. mucronata* and *R. apiculta* increased gradually with time, following a sigmoidal curve as expected (Figure 2). However, the height increment of *R. apiculata* was slightly higher than that of *R. mucronata*. The dbh of the saplings of both species increased with time in a similar pattern (Figure 3). dbh increment of *R. apiculata* was also higher than that of *R. mucronata*.

The height increment rate of *R. apiculata* was higher than that of *R. mucronata* throughout the observational period (Figure 4A). However, during the first 12 months period, the dbh increment rate of *R. mucronata* was higher than that of *R. apiculta*, whereas, during the rest of the period, the dbh increment rate of *R. apiculata* was slightly higher than *R. mucronata* (Figure 4B). The rate of height increment of the two species increased with time until the 18th month from the establishment and started to decline thereafter. Thus, the highest rate of height increment was observed by the 18th month of the establishment. The highest rate of dbh increment was observed by the 24th month from establishment whilst the increment rate declined between the 18th–30th month from establishment.

Biomass Carbon accumulation by the established stand

The average above-ground and below-ground biomass per barrel showed a linear increment during the study period (Figure 5). At the end of the study period, the average above-ground biomass per barrel was 70.7 \pm 11.7 kg. This biomass included 29.7 \pm 4.9 kg of carbon and it is equivalent to 108.2 \pm 17.9 kg of CO₂. Bellow ground biomass content at the time of final observation was 35.0 \pm 5.8 kg per barrel. This included 14.7 \pm 2.4 kg of carbon and equivalent to 53.5 \pm 8.3 kg of CO₂. By the end of the study period, plants have accumulated 105.8 \pm 17.5 kg of biomass per barrel which is equivalent to 161.7 \pm 26.8 kg of CO₂. Thus, these plants have sequestrated 217.15 tonnes of carbon per hectare, which is equivalent to 788.1 tonnes of CO₂ per hectare.

According to the calculations up to the final sampling date, the study site has accumulated 2,619.5 kg, 1294.5 kg and 3,914.9 kg of above-ground, below-ground and total biomass respectively. Furthermore, the total biomass accumulated up to the final monitoring date included 1,643.9 kg of carbon which is equivalent to 5,983.9 kg of CO₂.

However, up to the end of the monitoring period, no natural recolonization was observed in the restored area. Rehabilitation of degraded mangrove ecosystem: a case study



Figure 1. Survival percentage of *Rhizophora mucronata* and *R. apiculata* saplings during the 30th months of the observation period. Fourparameter logistic curves were fitted to observe the pattern of survival.



Figure 2. Height increment of *Rhizophora mucronata* and *R. apiculata* saplings against time, during the monitoring period. Four-parameter logistic sigmoidal curves were fitted to determine the pattern of height increment.



Figure 3. Diameter at breast height increment of *Rhizophora mucronata* and *R. apiculata* saplings during the monitoring period. Four parameter logistic sigmoidal curves were fitted to determine the pattern of height increment.



Figure 4. A—Height increment rate | B—girth increment rate of *Rhizophora mucronata* and *R. apiculata* were established in barrels in the Koggala lagoon.

DISCUSSION

The results of this study indicate the success of the restoration technique employed, as evidenced by the high survival rates of the restored species after a substantial period since the establishment ($2\frac{1}{2}$ years). The observed survival rates of 85% for *R. mucronata* saplings and 65% for *R. apiculta* suggest the effectiveness of the restoration approach.

Comparison with previous trials conducted without a controlled plot revealed a significant improvement in sapling survival. In contrast to previous attempts, where none of the saplings survived for more than a year, the current restoration technique demonstrated higher success rates. These findings align with research conducted by Kodikara et al. (2017) on mangrove restoration projects in Sri Lanka, where most restored sites exhibited less than 50% survival, and only a small number surpassed this threshold. Thus, the higher sapling survival rates observed at the Koggala mangrove restoration site indicate a comparative success compared to other restoration efforts.

Sapling growth analysis showed that saplings of both species used have normal sigmoidal growth patterns and they were reaching the maturity level. Especially, the reduction in growth rate during the 24th-30th month of



Figure 5. Average biomass and carbon accumulation of the established plant with time. Error bars are \pm SE.



Image 5. Mangrove plants after forty-five months from planting.

establishment shows that these saplings were gradually reaching the matured stage. Thus, it seems that the plants have well established within the restored sites.

The restoration of the mangrove site demonstrated a significant potential for carbon sequestration, with an observed carbon sequestration rate of 217.15 tonnes per ha (equivalent to 788.1 tonnes of CO_2 per ha), highlighting its contribution to reducing atmospheric CO_2 levels.

However, it cannot be compared with the total carbon content reported in other mangroves. However, the above-ground biomass carbon content of the restored site (128.8 t per ha) was higher than the average aboveground carbon content for global mangroves (78 t of carbon per ha; Estrada & Soares 2017), Mahanadi Mangrove, India (Sahu et al. 2016), and Negambo estuary (80.5 t of carbon per ha; Perera et al. 2018). This value
is slightly lower than that was reported for Batticaloa lagoon (131 t of carbon per ha; Perera et al. 2018) in Sri Lanka. These unusually high values may have been caused due to lower planting spacing of the restoration site than the usual spacing of a natural mangrove community. Further, the used spacing in the current study is less than the recommended spacing between mangrove seedlings planting for restoration (80–120 cm recommended [Intenational Coral Reef Initiative and Pole-Relais, Zones Humides Tropicales, 2020] vs. 60–90 in the current study). Thus, thinning of the mangrove vegetation of the restored site may be required to allow the saplings to

Our analysis showed a higher growth rate in *R. apiculata* compared to *R. mucronata* when considering the dbh and height. This could be due to the genetic potential of the two species as the same type of observation has been reported by Nit et al. (2011). However, further studies are needed to conclude the growth rates of the two species.

grow in their usual manner.

Our case study showed that the new method of mangrove restoration is successful in establishing mangroves in sites facing high erosion (Image 5). Especially, it seems that the new method is successful in coping with the situation in the Koggala lagoon as previous normal restoration trials conducted on this site failed. Thus, we recommend this restoration method for sites facing the threat of severe erosion.

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Sinhala abstract:

ලෝකයේ පරිසර පද්ධතීන් අතර ජීවවිදාහත්මකව සංකීර්ණතම සහ ජෛව නිෂ්පාදීතාවය අධිකතම පරිසර පද්ධකියක් වන කඩොලාන පරිසර පද්ධකිය, ලොවපුරා දිනෙන් දින විනශය කරා ඇදීයයි. පසුගිය දශක ගණනාවක් පුරා සිදුවූ කඩොලාන විනශය නිසා, කඩොලාන මගින් ලබාදුන් බොහොමයක් පාරිසරික සේවාවන් මිනිසට අහිමිවෙමින් පවතී. ශුී ලංකාවේ දකුණු මුහුදු තීරයේ කොග්ගල කළපුව ආශුිත කඩොලාන පරිසරයද, සංචාරක ව්යාපාරයට සම්බන්ධ කටයුතු, කෘෂිකාර්මික කටයුතු සහ නාගරික සංවර්ධන කටයුතු නිසා එසේ සීගුලෙස විනාශවන කඩොලාන පරිසර පද්ධතියකි. මෙම කළපුවේ ඉවුර, කළපු ජලතලය මත මුහුදු ගුවන්යානා ගොඩබෑම නිසා අධික ලෙස බාදනයට ලක්වී ඇති අතර, කඩොලාන පරිසරයද අධිකව විනාශයට ලක්වී ඇත. මෙම කඩොලාන පරිසර පද්ධතිය නැවත පුනරුත්තාපනය කිරීම සඳහා, විදාානුකූල වාාපෘතියක අවශාතාවය පැනනැගී ඇතිහෙයින්, ඒ සඳහා පෙරහුරුවක් ලෙසද, මෙම පද්ධතියෙකි කඩොලාන පුතිස්තාපනය කිරීම අරමුණු කොටගෙනද, පර්යේෂණයක් ආරම්භ කරනලදි. බාදනය වූ ඉවුරෙහි සමනාঃ ආකරයට කඩොලාන පුනිස්තාපනය කල නොහැකි හෙයින්, කඩොලාන පැල ඉවුරෙහි ස්ථාපිත කිරීම සඳහා විබාදනයට ලක් නොවන ප්ලස්ටික් බැරල් යොදා ගන්නාලදි. පුනිස්තාපනයෙන් අනතුරුව, පළමු පියවර යටතේ, පරිසර පද්ධතියේ කඩොලාන ශාක ස්ථීරව ස්ථාං ාවීම අධිකෘණය කරනුලැබූ අතර, දෙවන පියවර යටතේ ස්තාපිත පරිසර පද්ධතිය මගින් ලබාදෙන පාරිසරික සේවාවන්ගේ පුතිස්තාපනයවීම අධිකෂණය කරනු ඇත. කඩොලාන පුතිස්තාපනය වීමෙන් අවුරුද 2.5 කට පසුව කඩොලාන ජෛව ස්කන්ධයතුල තිරකර ඇති කාබන් පුමණය, කඩොලාන ශාක විශේෂවල ජෛව ස්කන්ධ කාබන් පුමාණය ගණනය කිරීම සඳහා ඉදිරිපත් කර ඇති සමීකරණ යොදාගෙන ගණනය කරන ලදි. කඩොලාන ශාක පුතිස්තාපනයෙන් මාස 24කට පසුව කඩොලාන ශාක කඳේ වට පුමාණය වැඩිවීමේ වැඩිම වේගය වාර්තාවිය. එමෙන්ම, ශාකවල වර්ධන වේගය, මාස 18 සිට 30 දක්වා කුම්කව අඩුවිය. පර්යේෂණ කාලය තුළදී, මධාාන අපි-භෞමික සහ අධෝ-භෞමික ජෛව ස්කන්ධ කාබන් පුමාණය රේඛීයව වැඩිවිය. සිදුකරන ලද මෙම පර්යේෂණයේ පුරීඵල වලට අනුව බාදනය වූ කළපු ඉවුරුවල කඩොලාන පුතිස්තාපනය ක්රීම සඳහා භාවිතා කරන ලද සුමය ඉතා සුදුසු බවට නිගමනය කලහැක. විශේෂයෙන්, බදනය වූ කොග්ගල කළපුවෙ ඉවුරුවල කඩොලාන පුතිස්තාපනය කිරීම සඳහා, ව්බාදනය නොවන ප්ලස්ටික් බැරල්වල කඩොලාන සිටුවීම ඉතාම යෝගාා වූ හෙයින්, මෙවැනි බාදනය වූ ඉවුරු සහිත වෙනත් කළපුවල කඩොලාන පුතිස්තාපනය කිරීම සඳහා මෙම කුමය නිර්දේෂ කල හැකිය

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Diversity of powdery mildew fungi from protected areas of Jizzak region, Uzbekistan - a checklist

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Abstract: The first-ever compilation of a checklist for powdery mildew fungi within the protected areas of Uzbekistan's Jizzakh region, specifically Nuratau Nature Reserve, Zaamin National Nature Park, and Zaamin Nature Reserve is presented through this study. Field research spanning from 2009 to 2022 facilitated this comprehensive endeavor. A comprehensive identification process revealed a total of nine groups, consisting of 57 distinct types, 69 variations, and four different strains of powdery mildew. Two species, namely Erysiphe platani and E. syringae, were newly identified within the mycobiota of the country. Additionally, the plant index of Uzbekistan documented 11 powdery mildew species and ten variations on 18 previously unrecorded host plant species. Most species of powdery mildew fungi belong to the genera Erysiphe and Leveillula. The annotated checklist includes data on the host plant, location, date and collection number of every species.

Keywords: Disease, Erysiphaceae, host plants, Nuratau Nature Reserve, Zaamin Nature Reserve, Zaamin National Nature Park.

Uzbek: Ushbu maqolada O'zbekistonning Jizzax viloyatida joylashgan Nurota qo'riqxonasi, Zomin milliy bog'i va Zomin davlat qoʻriqxonasi hududlarida tarqalgan un-shudring zamburugʻlarining dastlabki roʻyxati keltirilgan. Natijalar 2009 yildan 2022 yilgacha bo'lgan tadqiqot natijalari asosida shakllantirilgan. Dala tadqiqotlari davomida olib kelingan mikologik gerbariy namunalarini identifikatsiya qilish jarayonida un-shudring zamburugʻlarining 57 tur, 69 forma va 4 variatsiyasi qayd etildi. Shundan, Erysiphe platani va E. syringae Oʻzbekiston mikobiotasi uchun yangi ekanligi aniqlangan. Bundan tashqari, 11 tur un-shudring zamburugʻlari xoʻjayin oʻsimliklarning 18 turida ilk bor qayd etilgan. Tadqiqot hududlarida un-shudring zamburugʻlari orasida Erysiphe va Leveillula turkumi vakillari yetakchilik qiladi. Ushbu maqolada xar bir zamburugʻ turning xoʻjayin oʻsimligi, qayd etilgan yilllari keltirilgan.

Editor: Anonymity requested.

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Author contributions: I.M.M. - reviewing and editing; I.Z.O. - writing, reviewing and editing; K.K.N. - data collection; D.S.K. - reviewing and data collection. All authors have read and agreed to the published version of the manuscript.

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INTRODUCTION

Powdery mildews belong to the Erysiphaceae. The main geographical regions of the distribution are in the temperate zone of the northern hemisphere. They are obligate biotrophs responsible for diseases in many plants. They are cosmopolitan, widespread on various hosts including vegetables, trees, herbs, shrubs, grasses, ornamental plants and weeds (Braun 1987). In many cases, powdery mildew fungi are more common in cultivated plant hosts (Pawar & Patil 2011). The fungi infect almost every group of plants, i.e., from grasses to higher angiosperms (Braun & Cook 2012). Powdery mildews are easily visible on infected plant leaves, stems, and fruits. The first symptoms appear as white powdery spots on leaves, branches and sometimes on fruits. Over time, disease severity increases under favorable conditions. Disease symptoms usually appear with the onset of summer and begin to disappear during the scorching heat and rainy season (Pap et al. 2013).

A considerable number of powdery mildew samples were collected and examined during exploring of the fungal diversity of Uzbekistan; contributions have been published in 1926–2023 (Zaprometov 1926; 1928; Golovin 1949; Rotkevich 1960; Gaponenko 1965; Gaponenko et al. 1983; Solieva 1989; Kamilov 1991; Nuraliev 1998; Gafforov 2004; Mustafaev 2018; Mustafaev et al. 2019; Abdurazakov et al. 2021).

Recently, invasive species of powdery mildew, originating from other regions of the world, have spread extensively in Uzbekistan (e.g., *Erysiphe australiana* on *Lagerstroema indica*); they are mostly found on introduced trees and shrubs that are widely used for landscaping (Nabieva et al. 2021).

The current study aims to compile a checklist of powdery mildew of vascular plants of the Nuratau Nature Reserve, Zaamin Nature Reserve, and Zaamin National Nature Park in the Jizzakh region. The present checklist serves as one of the sources to make the complete modern list of powdery mildew of the Rebuplic of Uzbekistan.

Study Area

The present compilation brings together as much information, related to powdery mildew reported from protected areas of Jizzakh region (Nuratau Nature Reserve 40°30'34.8" 66°44'18.4", Zaamin Nature, Reserve 39°34'41.4" 68°24'21.5", Zaamin National Nature Park 39°39'56.4" 68°24'18.3") (Fig. 1), as possible.

METHODS

About 1,000 powdery mildew fungi were collected in 2009–2022 years in different habitats of protected natural areas. Samples were collected from the affected parts of the plants and either used fresh or stored dry as herbarium material until analysis, as described by Heluta (1989). These are designated in the checklist by the following abbreviations: NNR -Nuratau Nature Reserve, ZNR - Zaamin Nature Reserve, ZNNP - Zaamin National Nature Park. Specimens: Chasmothecia, conidiophores, and conidia were observed and measured in oil immersion, using a Moticam 5N-300M microscope at a magnification up to ×400 and identified using the relevant literature (Zaprometov 1926; 1928; Golovin 1941; Gaponenko 1965; Panfilova & Gaponenko 1983; Gaponenko et al. 1983; Heluta et al. 2004; Braun 2012; Raximova et al. 2014). (1968-1993) and 'Flora of Uzbekistan' (1941-1962). All collected specimens are stored in the Fungarium of the Institute of Botany in Tashkent. Voucher specimens for each species of powdery mildew fungi are cited in the checklist below. The name of the collector is designated by abbreviation: IM - Ilyor Mustafaev, IOZ - Islom Ortiqov, XD -Xo`jaqulova Durdona. The taxonomy and nomenclature of powdery mildew fungi in the checklist follow international databases Index Fungorum (2023) and Mycobank (2023). The nomenclature of host plants is given according to powo.science.kew.org/results (2023). All collected specimens are stored in the Tashkent Mycological Herbarium (TASM) of the Institute of Botany.

RESULTS

In the protected areas of the Jizzakh region, including NNR, ZNR, and ZNNP, an extensive survey uncovered a diverse range of powdery mildew fungi. These findings encompass nine genera and 57 species, with 69 forms and four varieties, all identified through morphological characteristics. These fungi were documented on 137 host plant species, further grouped into 107 genera and 34 families (Table 1). Erysiphe emerged as the most species-rich genus, boasting 19 species, 26 forms, and four variations, closely followed by Leveillula with 16 species and 27 forms. Podosphaera accounted for nine species and three forms, while Phyllactinia exhibited four species and four forms. Additionally, Golovinomyces and Sphaerotheca each comprised four and three species, respectively, while Uncinula, Neoerysiphe, and Blumeria were each confined to one host plant species.



Figure 1. The map illustrates the location of the protected areas in the Jizzakh region.

Erysiphe species demonstrated the broadest host range, infecting 57 host plant species (41.6 % of the total), followed by *Leveillula* with 35 host plant species (25.4 %), and *Podosphaera* with 19 host plant species (13.8 %). *Sphaerotheca* was found on 11 host plant species (8.0 %), while the remaining genera—*Blumeria*, *Phyllactinia*, *Golovinomyces*, *Neoerysiphe*, and *Uncinula*—showed isolation to fewer host species. This comprehensive investigation highlights the wide distribution of powdery mildew fungi in the protected areas of the Jizzakh region, underscoring the need for further research to uncover their ecological roles and potential impacts.

Two species *Erysiphe platani* – on *Platanus orientalis* and *E. syringae* – on *Ligustrum* sp. were found for the first time for the mycobiota of Uzbekistan. Eleven species, 10 variaties of powdery mildew were recorded on 18 new host plant species index of Uzbekistan: *Blumeria* graminis – *Poa trivialis, Erysiphe cichoracearum* f. *lactucae* – *Lactuca orientalis, E. cichoracearum* f. *tanaceti- Lepidolopha komarowii, E. cruciferarum* – *Descurainia sophia, E. heraclei* – *Ferula penninervis, Ferula angreni* and *Physocaulis nodosus, Leveillula* boraginacearum f. lappulae – Pseudoheterocaryum szovitsianum, L. compositarum f. artemisiae – Artemisia oliveriana, L. compositarum f. helichrysi – Helichrysum nuratavicum, L. labiatarum f. dracocephali – Dracocephalum nuratavicum, L. labiatarum f. scutellariae – Scutellaria ramosissima, L. labiatarum f. phlomidis – Phlomis nubilans, L. umbelliferarum f. heraclei – Semenovia pimpinelloides, Podosphaera fugax – Geranium rotundifolium, Sphaerotheca fuliginea f. lophanthi – Lophanthus schtschurowskianus, Sph. fuliginea f. sedi – Pseudosedum lievenii, Sphaerotheca sp. – Geum kokanicum.

Among them, *Helichrysum nuratavicum*, *Phlomis nubilans* are endemic species for Nuratau ridge and *Phlomis nubilans*, *Helichrysum nuratavicum* and *Platanus orientalis* are listed in the Red Data Book of Uzbekistan (2019).

Thirty-two hosts of family Asteraceae were infected with powdery mildew followed by Rosaceae (19), Lamiaceae (14), Poaceae (13), Brassicaceae (12), Apiaceae (10), Fabaceae Boraginaceae (8 each), Dipsacaceae, Polygonaceae (5 each), Scrophulariaceae, Geraniaceae Table 1. In this section, the scientific name of host plants along with the powdery mildew fungi occurring on them are provided.

Host plant (scientific name)	Powdery mildew fungi (scientific name)
Achillea sp.	Leveillula taurica f. achilleae
Aegilops crassa	Blumeria graminis
Aegilops cylindrica	Blumeria graminis
Aegilops triuncialis	Blumeria graminis
Alcea nudiflora	Erysiphe cichoracearum f. althaeae
Alhagi maurorum	Leveillula alhagi
Alhagi sp.	Leveillula alhagi
Alyssum alyssoides	Erysiphe cruciferarum
Alyssum minutum	Erysiphe cruciferarum
Alyssum dasycarpum	Erysiphe cruciferarum
Alyssum desertorum	Erysiphe cruciferarum
Arctium umbrosum	Erysiphe cichoracearum f. cousiniae
Arctium lappa	Golovinomyces depressus
Artemisia oliveriana	Leveillula compositarum f. artemisiae
Artemisia vulgaris	Golovinomyces cichoracearum
Asperugo procumbens	Erysiphe horridula
Astragalus sp.	Leveillula leguminosarum f. astragali
Atraphaxis pyrifolia	Erysiphe atraphaxis
Avena fatua	Blumeria graminis
Bidens tripartita	Sphaerotheca fuliginea f. bidentis
Capparis spinosa	Leveillula capparidacearum f. capparidis
Carthamus lanatus ssp. turkestanicus	Leveillula compositarum f. carthami
Centaurea besseriana	Leveillula compositarum f. centaureae
Cichorium intybus	Golovinomyces cichoracearum Sphaerotheca fuliginea
Chaerophyllum nodosum	Erysiphe heraclei
Convolvulus arvensis	Erysiphe convolvuli var. convolvuli
Cousinia microcarpa	Erysiphe cichoracearum f. cousiniae
Cousinia sp.	Erysiphe cichoracearum f. cousiniae
Cousinia coronata	Leveillula compositarum f. cousiniae
Crambe cordifolia ssp. kotschyana	Erysiphe communis f. crambes
Crataegus songarica	Phyllactinia guttata
Crataegus turkestanica	Podosphaera oxyacanthae f. crataegi Phyllactinia guttata
Crepis pulchra	Sphaerotheca fuliginea f. crepidis
Crepis sp.	Erysiphe cichoracearum f. crepidis
Cydonia oblonga	Podosphaera oxyacanthae f. cydoniae
Datisca cannabina	Leveillula datiscacearum f. datiscae
Daucus carota	Leveillula umbelliferarum f. dauci
Delphinium semibarbatum	Leveillula ranunculacearum f. delphinii
Descurainia sophia	Erysiphe cruciferarum

Host plant (scientific name)	Powdery mildew fungi (scientific name)
Dipsacus azureus	Sphaerotheca fuliginea f. dipsaci
Dodartia orientalis	Leveillula scrophulariacearum f. dodartiae
Dracocephalum nuratavicum	Leveillula labiatarum f. dracocephali
Erysimum sp.	Erysiphe cruciferarum f.erysimi
Euphorbia sp.	Podosphaera euphorbiae
Eremogone griffithii	Leveillula caryophyllacearum f. arenariae
Ferula angreni	Erysiphe heraclei
Ferula penninervis	Erysiphe heraclei
Fraxinus sogdiana	Phyllactinia fraxini
Galium aparine	Neoerysiphe galii
Galium pamiroalaicum	Neoerysiphe galii
Geranium linearilobum	Podosphaera fugax
Geranium pusillum	Podosphaera fugax
Geum kokanicum	Sphaerotheca sp.
Hedysarum sp.	Erysiphe communis f. hedysari
Helichrysum nuratavicum	Leveillula compositarum f. helichrysi
Heracleum lehmannianum	Leveillula umbelliferarum f. heraclei
Hieracium sp.	Erysiphe cichoracearum f. euhieracium
Hippophae rhamnoides	Phyllactinia suffulta f. hippophaes
Hordeum spontaneum	Blumeria graminis
Hordeum bulbosum	Blumeria graminis
Hypericum perforatum	Erysiphe hyperici
Hypericum scabrum	Leveillula guttiferarum
Juglans regia	Erysiphe juglandis
Lactuca orientalis	Erysiphe cichoracearum f. lactucae
Lepidium draba	Leveillula cruciferarum f. lepidii
Lepidolopha komarowii	Erysiphe cichoracearum f. tanaceti
Ligustrum sp.	Erysiphe syringae
Lindelofia anchusoides ssp. macrostyla	Erysiphe horridula f. lindelofiae
Lomelosia micrantha	Leveillula datiscacearum f. scabiosae
Lomelosia songarica	Sphaerotheca fuliginea f. scabiosae
Malus domestica	Podosphaera leucotricha
Marrubium anisodon	Erysiphe labiatarum f. marrubii
Medicago sativa	Erysiphe communis f.sp. medicaginis–sativae Leveillula leguminosarum f. medicaginis
Melilotus officinalis	Erysiphe trifoliorum
Morus alba	Phyllactinia suffulta f. moricola
Nepeta schtschurowskiana	Sphaerotheca fuliginea f. lophanthi
Ornithogalum gulnariense	Podosphaera fugax
Onobrychis chorassanica	Leveillula leguminosarum f. onobrychidis
Origanum vulgare	Erysiphe labiatarum f. origami

Host plant (scientific name)	Powdery mildew fungi (scientific name)			
Peganum harmala	Leveillula taurica f. pegani			
Pentanema britannica	Erysiphe cichoracearum f. inulae			
Phlomis nubilans	Erysiphe labiatarum f. phlomidi Leveillula labiatarum f. phlomidis			
Phlomis thapsoides	Leveillula labiatarum f. phlomidis			
Phlomoides labiosa	Erysiphe labiatarum f. eremostachydis			
Pistacia vera	Phyllactinia suffulta f. pistaciae			
Plantago lanceolate	Leveillula plantaginis, Podosphaera plantaginis			
Plantago major	Erysiphe cichoracearum var. plantaginis			
Platanus orientalis	Erysiphe platani			
Poa trivialis	Poa trivialis			
Poa versicolor	Blumeria graminis			
Poa bulbosa	Blumeria graminis			
Polygonum aviculare	Erysiphe polygoni f. polygoni			
Populus afghanica	Phyllactinia populi			
Potentilla pedata	Sphaerotheca macularis			
Potentilla reptans	Erysiphe thuemenii			
Prangos pabularia	Leveillula umbelliferarum f. prangotis			
Pseudosedum lievenii	Sphaerotheca fuliginea f. sedi			
Psychrogeton cabulicum	Podosphaera fusca			
Prunus persica	Podosphaera pannosa			
Prunus cerasus	Podosphaera tridactyla f. cerasi			
Prunus erythrocarpa	Phyllactinia suffulta f. pruni Podosphaera tridactyla f. cerasi			
Prunus bucharica	Podosphaera pannosa, Phyllactinia suffulta f. pruni			
Pseudopodospermum inconspicuum	Leveillula compositarum f. scorzonerae			
Pseudoheterocaryum szovitsianum	Leveillula boraginacearum f. Iappulae			
Ranunculus sericeus	Erysiphe aquilegiae var. ranunculi			
Rochelia cardiosepala	Erysiphe horridula f. rocheliae			

Host plant (scientific name)	Powdery mildew fungi (scientific name)
Rochelia sp	Erysiphe horridula f. rocheliae
Rosa canina	Podosphaera pannosa
Rosa persica	Podosphaera pannosa
Rosa beggeriana var. beggeriana	Podosphaera pannosa
Rosa ecae	Podosphaera pannosa
Rumex acetosa	Erysiphe polygoni var. rumicis
Rumex chalepensis	Erysiphe polygoni var. rumicis
Salvia austriaca	Golovinomyces salvia
Salvia scrophulariifolia	Leveillula compositarum f. perovskia
Scandix pecten–veneris	Erysiphe heraclei
Scutellaria ramosissima	Leveillula labiatarum f. scutellariae
Semenovia pimpinellioides	Leveillula umbelliferarum f. heraclei
Solenanthus circinnatus	Erysiphe horridula f. solenanthi Leveillula umbelliferarum f. seseli
Solenanthus turkestanicus	Erysiphe horridula f. solenanthi
Sonchus asper	Erysiphe cichoracearum f. sonchi
Tanacetopsis mucronata	Erysiphe cichoracearum f. tanaceti
Taraxacum officinale	Erysiphe cichoracearum f. taraxaci
Taraxacum maracandicum	Podosphaera erigerontis– canadensis
Trifolium pretense	Erysiphe trifolii
Thinopyrum intermedium ssp. intermedium	Blumeria graminis
Ulmus laevis	Uncinula ulmi
Urtica dioica	Erysiphe urticae
Verbascum songaricum	Leveillula verbasci
Verbena officinalis	Erysiphe cichoracearum f. verbenae
Veronika argute–serrata	Erysiphe cichoracearum f. veronicae
Vitis vinifera	Erysiphe necator
Vickifunkia thomsonii	Sphaerotheca fuliginea f. senecionis
Ziziphora tenuior	Erysiphe labiatarum f. ziziphorae

(4 each), Euphorbiaceae, Oleaceae, Plantaginaceae, Ranunculaceae, Rubiaceae, Hypericaceae, Malvaceae (2 each), Capparaceae, Caryophyllaceae, Crassulaceae, Datiscaceae, Elaeagnaceae, Juglandaceae, Anacardiaceae, Moraceae, Peganaceae, Platanaceae, Salicaceae, Ulmaceae, Urticaceae, Verbenaceae, Vitaceae (1 each).

The highest number of powdery mildew species was reported in the following host genera: *Plantago*, *Prunus* (3 species each, 5.26% of the total species number), *Cichorium*, *Arctium*, *Artemisia*, *Cousinia*, *Crataegus*, *Crepis*, *Lomelosia*, *Medicago*, *Phlomis*, *Potentilla*, *Salvia*, *Solenanthus* and *Taraxacum* (2 species each, 3.50%), while other plant genera host one powdery mildew species per genus.

The recorded powdery mildew species were found on 31 medicinal species such as *Alcea nudiflora, Alhagi maurorum, Capparis spinosa, Cichorium intybus, Crataegus songarica, C. turkestanica, Crambe cordifolia* ssp. kotschyana, Hippophae rhamnoides, Hypericum perforatum, Juglans regia, Malus domestica, Medicago sativa, Morus alba, Nepeta schtschurowskiana, Pistacia vera, Plantago lanceolate, P. major, Prunus erythrocarpa, P. bucharica, Helichrysum nuratavicum, Rosa canina, R. persica, R. beggeriana var. beggeriana, R. ecae, Rumex acetosa, R. chalepensis, Taraxacum

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officinale, T. maracandicum, Urtica dioica, Vitis vinifera, Ziziphora tenuior etc.

This work represents itself as a checklist of powdery mildew fungi in the Jizzakh region of protected areas. The fungal species listed below have been arranged in alphabetical order.

Blumeria graminis (DC.) Speer

(= Erysiphe communis f. graminis (DC.) Fr., Alphitomorpha communis var. graminearum Wallr., Erysiphe graminis DC., Tigria graminis (DC.) Trevis.) Host: Poaceae Poa trivialis (2015) NNR (Xayatsay V.) - First record on Poa trivialis in Uzbekistan. Poa versicolor (2015) NNR (Tikchasay V.) Poa bulbosa (2010), NNR (Beshbarmoq V.) (2019-2022) ZNNP (Usmanlisay, Qorongisay V.) Thinopyrum intermedium subsp. intermedium trichophorum) (2022) (=Agropyron 7NNP (Ettikechuvsay V.) Aegilops crassa (2010) NNR (Xayatsay V.) Aegilops cylindrica (2012), NNR (Qarisay V.) Aegilops triuncialis (2022) ZNNP (Usmanlisay V.) Hordeum bulbosum (1983) ZNR (Kulsay V.), (2009-20120) NNR (Xayatsay V.), (2019) ZNNP (Usmanlisay V.), (2021) ZNNP (Boytepa V.) Hordeum spontaneum (2021) ZNNP (Qo'riq V.) Avena fatua (2022) ZNNP (Usmanlisay V.)

Erysiphe aquilegiae var. **ranunculi** (Grev.) R.Y. Zheng & G.Q. Chen.

Host: Ranunculaceae Ranunculus sericeus (2016) NNR (Xayatsay V.).

E. atraphaxis (Golovin) U. Braun & S. Takam.
 Host: Polygonaceae
 Atraphaxis pyrifolia (2019) ZNNP (O'riklisay V.).

E. communis f. crambes Jacz.

Host:Brassicaceae Crambe cordifolia ssp. kotschyana (Boiss.) Jafri. (2014) NNR (Gurdara V.), (2021) ZNNP (Qorong'isay V.).

E. communis f. *dianthi* Jacz. Host: Caryophyllaceae *Dianthus helenae* (2017) NNR (Beshbarmaq V.).

E. communis f. hedysari Jacz.

Host: Fabaceae Hedysarum sp.- (2010) NNR (Beshbarmaq V.). *E. communis* f. sp. *medicaginis-sativae* Hammarl. Host: Fabaceae *Medicago sativa* (2019) ZNNP (O'riklisay V.).

E. cichoracearum f. althaeae Jacz.

Host: Malvaceae Alcea nudiflora (2010) NNR (Xayatsay V.), (2019) ZNNP (O'riklisay V.).

E. cichoracearum f. lactucae Jacz.

Host: Asteraceae

Lactuca orientalis (2010) NNR (Xayatsay V.), (2012) NNR (Majrumsay V.) - Note: First record on *Lactuca* orientalis in Uzbekistan.

E. cichoracearum f. tanaceti Jacz.

Host: Asteraceae

Lepidolopha komarowii (2015) NNR (Xayatboshi V.) - Note: First record on *Lepidolopha komarowii* in Uzbekistan.

E. cichoracearum f. veronicae Jacz.

Host: Scrophulariaceae

Veronica argute-serrata (2010) NNR (Beshbarmoq V.).

E. cichoracearum f. verbenae Jacz.

Host: Verbenaceae Verbena officinalis (2010) NNR (Xayatboshi V.), (2017) NNR (Xayatsay V.).

E. cichoracearum f. *crepidis* Jacz.

Host: Asteraceae Crepis sp. (2019) ZNNP (Ettikechuvsay V.).

E. cichoracearum f. euhieracium S. Blumer.

Host: Asteraceae Hieracium sp. (2019) ZNNP (Sherbulaq V.).

E. cichoracearum f. tanaceti Jacz.

Host: Asteraceae Tanacetopsis mucronata (2019) ZNR (Qizilmazar V.).

E. cichoracearum f. taraxaci Jacz .

Host: Asteraceae Taraxacum officinale – (2021) ZNNP (Boytepa V.).

E. cichoracearum f. sonchi Jacz .

Host: Asteraceae Sonchus asper (2021) ZNNP (Sherbulaq, Irgʻaylisay V.).

E. cichoracearum f. *cousiniae* (Jacz). Golov. Host: Asteraceae

Cousinia sp. (2021) ZNNP (Boytepa V.). Cousinia microcarpa (2022) ZNNP (O'riklisay).

E. cichoracearum var. *plantaginis* (Link). Host: Plantaginaceae *Plantago major* (2021) ZNNP (Qorong'isay V.).

E.cichoracearum f. inulae Jacz .

Host: Asteraceae

Pentanema britannica (2022) ZNNP (Irg'aylisay V.)

E. cruciferarum Opiz ex L. Junell.

(= Erysiphe communis (Wallr.) Link., Erysiphe communis (Wallr.) Schltdl., Erysiphe cruciferarum var. longispora G.J.M. Gorter., Erysiphe pisi var. cruciferarum (Opiz ex L. Junell) Ialongo., Erysiphe radulescui Docea.) Host: Brassicaceae

Alyssum dasycarpum (2010–2012) NNR (Tikchasay, Qarisay V.).

Alyssum alyssoides (2010) NNR (Xayatsay V.).

Alyssum desertorum (2020) ZNNP (Sufa V.), (2022) ZNNP (Qorong'isay V.).

Alyssum minutum (2020) ZNNP (Usmanlisay V.) Descurainia sophia (2022) ZNNP (Usmanlisay V.) - Note: First record on Descurainia sophia in Uzbekistan.

E. cruciferarum f. erysimi (Jacz).

Host: Brassicaceae Erysimum sp. (2022) ZNNP (Usmanlisay V.).

E. convolvuli var. *convolvuli*

Host: Convolvulaceae Convolvulus arvensis (2010) NNR (Xayatsay V.), (2021) ZNNP (Usmanlisay V.).

E. horridula f. *asperuginis* S. Blumer.

Host: Boraginaceae Asperugo procumbens (2010) NNR (Xayatsay V.), (2019) ZNNP (O'riklisay V.).

E. horridula f. lindelofiae Golovin.

Host: Boraginaceae Lindelofia anchusoides ssp. macrostyla (2010) NNR (Beshbarmoq Mt.), (2018) ZNNP (Oʻriklisay).

E. horridula f. *rocheliae* Golovin. Host: Boraginaceae Rochelia sp. (2009) NNR (Tikchasay V.) Rochelia cardiosepala (2012) NNR (Qarisay, Gurdara V.).

E. horridula f. *solenanthi* Jacz.

Host: Boraginaceae

Solenanthus circinnatus (2010) NNR (Parandoz V.). Solenanthus turkestanicus (2011) NNR (Majrumsay V.).

E. heraclei DC.

(= *Erysiphe heraclei* var. *himalayensis* Y.S. Paul & V.K. Thakur.)

Host: Apiaceae

Ferula penninervis (2017) NNR (Xayatsay, Beshbarmaq V.) - Note: First record on *Ferula penninervis* in Uzbekistan.

Ferula angreni (2017) NNR (Xayatsay V.)- First record on *Ferula angreni* in Uzbekistan.

Chaerophyllum nodosum (2014) NNR (Gurdara V.).

Scandix pecten veneris (2012) NNR (Xayatsay V.), (2019) ZNNP (Boytepa V.), (2022) ZNNP (Qorong'isay V.).

E. thuemenii U. Braun (= E. communis f. potentillae Jacz). Host: Rosaceae Potentilla reptans (2019) ZNNP (Usmanlisay V.).
E. trifolii Grev. (= Erysiphe martii Lév., Microsphaera martii (Lév.) YS Paul & VK Thakur., Microsphaera trifolii (Grev.) U. Braun). Host: Fabaceae

Trifolium pretense (2021) ZNNP (Usmanlisay V.).

- *E. trifoliorum* (Wallr.) U. Braun. Host: Fabaceae *Melilotus officinalis* (2019) ZNNP (O'riklisay V.).
- *E. hyperici* (Wallr.) S. Blumer. Host: Hypericaceae *Hypericum perforatum* (2012) NNR (Qarisay V.).
- *E.juglandis* (Golovin) U. Braun & S. Takam. (= *Microsphaera juglandis* Golovin.). Host: Juglandaceae *Juglans regia* (2019) ZNNP (Usmanlisay V.).
- *E. labiatarum* f. *origani* (H.A. Dietr.) Jacz. Host: Lamiaceae *Origanum vulgare* (2014) NNR (Parandoz V.).
- *E. labiatarum* f. *phlomidis* Jacz. Host: Lamiaceae

Phlomis nubilans (2014), NNR (Majrumsay V.).

E. labiatarum f. leonuri Jacz.

Host: Lamiaceae Leonurus turkestanicus (2017), NNR (Xayatboshi V.).

E. labiatarum f. *ziziphorae* Pospelov.

Host: Lamiaceae Ziziphora tenuior (2019) ZNNP (Yettikechuvsay V.).

E. labiatarum f. eremostachydis Golovin.
 Host: Lamiaceae
 Phlomoides labiosa (1983) ZNR (Guralashsay V.).

E. labiatarum f. marrubii Jacz.

Host: Lamiaceae Marrubium anisodon (2021) ZNNP (Qorong'isay V.).

E. necator Schwein.

(= Erysiphe necator var. ampelopsidis (Peck) U. Braun & S. Takam., Oidium tuckeri Berk., Uncinula americana Howe., Uncinula ampelopsidis Peck., Uncinula necator (Schwein.) Burrill., Uncinula necator var. ampelopsidis (Peck) U. Braun.). Host: Vitaceae Vitis vinifera (2016) NNR (Qarisay V.), (2022) ZNNP (Qorong'isay V.).

E. platani (Howe) U. Braun (= *Microsphaera platani* Howe.)

Host: Platanaceae Platanus orientalis (2022) ZNNP (Usmanliysay V.) -Note: First report in Uzbekistan.

E. polygoni f. polygoni DC.

Host: Polygonaceae Polygonum aviculare (2010) NNR (Qarisay V.), (2019) ZNNP (O'riklisay V.), (2022) ZNNP (Usmanliysay V.).

E. polygoni var. rumicis Y.S. Paul & V.K. Thakur.

Host: Polygonaceae Rumex chalepensis (2017) NNR (Xayatsay V.). Rumex acetosa (2021) ZNNP (Usmanliysay V.).

E. syringae Schwein. (= *Microsphaera syringae* (Schwein.) H. Magn.).

Host: Oleaceae

Ligustrum sp. (2022) ZNNP (Qorong'isay V.) - Note: First report in Uzbekistan. *E. urticae* (Wallr.) S. Blumer Host: Urticaceae *Urtica dioica* (2014) NNR (Xayatsay V.).

Golovinomyces cichoracearum (DC.) VP Heluta.

(= Erysiphe communis var. cichoracearum (DC.) Link., Erysiphe cichoracearum var. luvungae M.S. Patil & Maham., Erysiphe cichoracearum var. saussureae Y.S. Paul & V.K. Thakur., Erysiphe cichoracearum var. transvaalensis G.J.M. Gorter & Eicker., Golovinomyces cichoracearum var. latisporus (U. Braun) U. Braun., Golovinomyces cichoracearum var. transvaalensis (G.J.M. Gorter & Eicker) U. Braun., Oidium asterispunicei Peck., Oidium tabaci Thüm.). Host: Asteraceae

Artemisia vulgaris (2019) ZNR (Qashqasuv V.).

G. salviae (Jacz.) M. Scholler, U. Braun & Anke Schmidt.
(= Erysiphe biocellata var. salviae (Jacz.) VP Heluta., Erysiphe labiatarum f. salviae Jacz., Erysiphe salviae (Jacz.) S. Blumer., Leveillula labiatarum f. salviae (Jacz.) Golovin.).
Host: Lamiaceae

Salvia austriaca (2017) NNR (Xayatsay V.).

G. depressus (Wallr.) VP Heluta.

(=Erysiphe cichoracearum f. bardanae (Wallr.) Jacz., Erysiphe communis f. depressa (Wallr.) Fr., Erysiphe depressa (Wallr.) Link., Erysiphe depressa var. artemiciae Link., Erysiphe depressa var. bardanae Wallr.).

Host: Asteraceae Arctium lappa (2009) NNR (Xayatsay V.).

Golovinomyces cichoracearum (DC.) VP Heluta.

(= Erysiphe communis var. cichoracearum (DC.) Link., Erysiphe cichoracearum DC., Erysiphe cichoracearum f. cichorii S. Blumer., Erysiphe cichoracearum var. latispora U. Braun., Erysiphe cichoracearum var. luvungae M.S. Patil va Maham., Erysiphe cichoracearum var. saussureae Y.S. Pol va VK Thakur., Erysiphe cichoracearum var. transvaalensis G.JM Gorter & Eicker., Golovinomyces cichoracearum var. latisporus (U. Braun) U. Braun., Golovinomyces cichoracearum var. transvaalensis (GJM Gorter & Eicker) U. Braun., Oidium asteris-punicei Peck., Oidium tabaci Thüm.).

Host: Asteraceae

Cichorium intybus (2017) NNR (Xayatsay V.).

Leveillula alhagi (Sorokīn) U. Braun.

(= Erysiphe alhagi Sorokīn., Leveillula leguminosarum f. alhagi (Sorokīn) Golovin.). Host: Fabaceae Alhagi sp. (2009) NNR (Xayatsay V.), (2017) NNR (Majrumsay V.). Alhagi maurorum (2021) ZNNP (Boytepa V.).

L. boraginacearum f. lappulae (Jacz). Golov.

Host: Boraginaceae

Pseudoheterocaryumszovitsianum(2022)ZNNP(O'riklisayV.)-Note:FirstrecordonPseudoheterocaryum szovitsianum in Uzbekistan.

L.capparidacearum f. capparidis (Jacz.) Golovin.

Host: Capparaceae Capparis spinosa (2019–2021) ZNNP (O'riklisay, Usmanlisay V).

- *L. caryophyllacearum* f. *arenariae* Golovin. Host: Caryophyllaceae *Arenaria griffithii* (2015–2016) NNR (Xaytbashi V.).
- L. compositarum f. scorzonerae (Kuprev.) Golovin.

Host: Asteraceae *Pseudopodospermum inconspicuum* (2022) ZNNP (Ettikechuvsay V.).

L. compositarum f. artemisiae (Jacz.) Golovin.

Host: Asteraceae Artemisia oliveriana (2017) NNR (Majrumsay V.) - Note: First record on Artemisia oliveriana in Uzbekistan.

L. compositarum f. helichrysi (Jacz.) Golovin.

Host: Asteraceae

Helichrysum nuratavicum (2015) NNR (Fargun V.) - Note: First record on Helichrysum nuratavicum in Uzbekistan.

L. compositarum f. centaureae (Jacz.) Golovin.

Host: Asteraceae

Centaurea besseriana (2010-2014) NNR (Xayatbashi - Majrumsay V.).

L. compositarum f. *carthami* (Jacz.) Golovin.

Host: Asteraceae

Carthamus lanatus ssp. turkestanicus (2015) NNR (Xayatsay V.).

L. compositarum f. cousiniae (Jacz.) Golovin.
 Host: Asteraceae
 Arctium umbrosum (2016) NNR (Tikchasay V.).
 Cousinia coronata (2019) ZNNP (Yettikechuvsay V.).

- L. compositarum f. perovskiae Kurbana.
 Host: Lamiaceae
 Salvia scrophulariifolia (2017) NNR (Beshbarmaq V.).
- *L. cruciferarum* f. *lepidii* (Jacz.) Golovin. Host: Brassicaceae *Lepidium draba* (2014) NNR (Xayatsay V.).
- L. datiscacearum f. datiscae (Jacz.) Golovin.
 Host: Datiscaceae
 Datisca cannabina (2014) NNR (Majrumsay V.), (2016) NNR (Tikchasay V.).
- *L. dipsacacearum* f. *scabiosae* (Jacz.) Golovin. Host: Dipsacaceae *Lomelosia micrantha* (2022) ZNNP (Usmanlisay V.).

L. guttiferarum Golovin.

Host: Hypericaceae Hypericum scabrum (2016) NNR (Tikchasay V.).

L. labiatarum f. *dracocephali* Golovin.

Host: Lamiaceae Dracocephalum nuratavicum (2015) NNR (Andibaraut V.) - Note: First record on Dracocephalum nuratavicum in Uzbekistan.

L. labiatarum f. scutellariae (Jacz.) Golovin.
 Host: Lamiaceae
 Scutellaria ramosissima (2015) NNR (Xayatbashi-Xayatsay V.) - Note: First record on Scutellaria ramosissima in Uzbekistan.
 Scutellaria sp.- (2016) NNR (Tikchasay V.).

L. labiatarum f. *phlomidis* (Jacz.) Golovin.

Host: Lamiaceae Phlomis thapsoides (2010) NNR (Xayatsay V.). Phlomis nubilans (2015) NNR (Fargun V.) - Note: First record on Phlomis nubilans in Uzbekistan.

L. leguminosarum f. *astragali* (Jacz.) Golovin. Host: Fabaceae *Astragalus* sp. (20221) ZNNP (Sherbuloq V.).

L. leguminosarum f. *onobrychidis* Golovin. Host: Fabaceae

Onobrychis chorassanica (2015) NNR (Fargun V.).

L. leguminosarum f. *medicaginis* (Jacz.) Golovin. Host: Fabaceae *Medicago sativa* (2017) NNR (Xayatsay V.).

L. plantaginis Golovin. Host: Plantaginaceae *Plantago lanceolata* (2010) NNR (Xayatsay V.).

L. scrophulariacearum f. *dodartiae* (Jacz.) Golovin. Host: Scrophulariaceae *Dodartia orientalis* (2010) NNR (Xayatbashi V.).

L. taurica f. pegani Jacz.

Host: Peganaceae Peganum harmala (2010) NNR (Tikchasay V.).

L. taurica f. achilleae Jacz.

Host: Asteraceae Achillea sp. (2019) ZNNP (O'riklisay V.).

L. umbelliferarum f. *dauci* Golovin.

Host: Apiaceae Daucus carota (2010) NNR Xayatsay V.).

L. umbelliferarum f. seseli Golovin.

Host: Boraginaceae Solenanthus circinnatus (2015) NNR (Xayatbashi V.).

L. umbelliferarum f. *prangotis* Golovin. Host: Apiaceae

Prangos pabularia (2017) NNR (Xayatsay V.).

L. umbelliferarum f. heraclei Golovin.

Host: Apiaceae

Heracleum lehmannianum (2018) ZNNP (O'riklisay V.).

Semenovia pimpinelloides (2022) ZNNP (Qorong'isay V.) - Note: First record on Semenovia pimpinelloides in Uzbekistan.

L. ranunculacearum f. delphinii Golovin.

Host: Ranunculaceae

Delphinium semibarbatum (2019) ZNNP (Yettikechuvsay V.).

L. verbasci (Jacz.) Golovin. (= *Leveillula taurica* f. *verbasci* Jacz.).

Host: Scrophulariaceae Verbascum songaricum (2015) NNR (Tikchasay V.), (2016) NNR (Xayatbashi V.).

Neoerysiphe galii (S. Blumer) U. Braun. (= Erysiphe galii S. Blumer). Host: Rubiaceae Galium aparine - (1983) ZNR (Kulsay V.), (2022) ZNNP (O'riklisay V.). Galium pamiroalaicum (2016) NNR (Parandoz V.).

Phyllactinia suffulta f. pistaciae Jacz. Host: Anacardiaceae

Pistacia vera (2015) NNR (Xayatsay V.), (2021) ZNNP (Boytepa V.).

P. populi (Jacz.) Y.N. Yu.

(= Phyllactinia suffulta f. populi Jacz.). Host: Salicaceae Populus afghanica (2014) NNR (Majrumsay V.), (2019–2021) ZNNP (Qorong'isay-O'riklisay V.).

P. suffulta f. pruni Golov.

Host: Rosaceae Prunus bucharica (2019–2021) ZNNP (O'riklisay – Usmanliysay V.). Prunus erythrocarpa (2018) ZNNP (O'riklisay V.).

P. suffulta f. hippophaes Jacz.

Host: Elaeagnaceae Hippophae rhamnoides (2022) ZNNP (Umanlisay V.).

P. suffulta f. moricola Jacz.

Host: Moraceae *Morus alba* (2009–2010–2016) NNR (Xayatsay, Tikchasay V.), (2021) ZNNP (Qorong'isay V.).

P. guttata (Wallr.) Lév.

(= Phyllactinia berberidis Palla., Phyllactinia betulae (DC.) Fuss., Phyllactinia corylea (Pers.) P. Karst., Phyllactinia suffulta (Rebent.) Sacc.). Host: Rosaceae Crataegus turkestanica (2017) NNR (Xayatsay V.). Crataegus songarica (2022) ZNNP (Yettikechuvsay V.).

P. fraxini (DC.) Fuss.

(= Erysiphe communis f. lamprocarpa (Wallr.) Fr., Erysiphe fraxini DC., Erysiphe lamprocarpa (Wallr.) Link.). Host: Oleaceae Fraxinus sogdiana (2017) NNR (Qarisay V.).

Podosphaera euphorbiae (Castagne) U. Braun & S. Takam. (= Sphaerotheca euphorbiae (Castagne) E.S. Salmon., Sphaerotheca tomentosa G.H. Otth.). Host: Euphorbiaceae Euphorbia sp. (2016) NNR (Tikchasay V.).

P. fugax (Penz. & Sacc.) U. Braun & S. Takam.

(=Erysiphe communis var. geranii Klotzsch., Sphaerotheca fugax Penz. & Sacc.).
Host: Geraniaceae
Geranium linearilobum (2013) NNR (Gurdara V.).
Geranium pusillum (2010) NNR (Tikchasay V.).
Geranium rotundifolium (2012) NNR (Gurdara V.)
Note: First record on Geranium rotundifolium in Uzbekistan.

P. fusca (Fr.) U.Braun va Shishkoff.

(= Podosphaera phaseoli (Z.Y. Zhao) U. Braun & S. Takam., Podosphaera xanthii (Castagne) U.Braun & Shishkoff., Sphaerotheca astragali var. phaseoli Z.Y.Zhao., S. calendulae (Malbr. & Roum.) Malbr., S. cucurbitae (Jacz.) Z.Y.Zhao., S. fuliginea f. calendulae (Malbr. & Roum.) Jacz., S. fuliginea f. cucurbitae Jacz., S. fusca (Fr.) S. Blumer., S. fuscata (Berk. & M.A.Curtis) Serbinow., S. indica Patw., S. melampyri L. Junell., S. microcarpa Hazsl., S. phaseoli (Z.Y.Zhao) U.Braun., S. verbenae Săvul. & Negru., S. xanthii (Castagne) L. Junell.).

Host: Asteraceae Psychrogeton cabulicum (2016) NNR (Tikchasay V.).

P. tridactyla f. cerasi Jacz.

Host: Rosaceae

Prunus cerasus (2017) NNR (Xayatsay V.). Prunus erythrocarpa (2018) ZNNP (Oʻriklisay V.).

P. leucotricha (Ellis & Everh.) E.S.Salmon.

(= Sphaerotheca leucotricha Ellis & Everh.). Host: Rosaceae Malus domestica (2009) NNR (Majrumsay V.), (2021) ZNNP (Ettikechuvsay V.).

P. erigerontis-canadensis (Lév.) U.Braun & T.Z.Liu. (= *Erysiphe erigerontis-canadensis* Lév., *Sphaerotheca erigerontis-canadensis* (Lév.) L. Junell.).

Host: Asteraceae

Taraxacum maracandicum (1983) ZNR (Guralashsay V.).

P. euphorbiae (Castagne) U. Braun & S. Takam.
 (= Sphaerotheca euphorbiae (Castagne) E.S. Salmon.,

S. tomentosa G.H. Otth.). Host: Euphorbiaceae Euphorbia sp. (2016) NNR (Xayatsay V.).

P. oxyacanthae f. cydoniae Jacz.

Host: Rosaceae Cydonia oblonga (1956) ZNR (Guralashsay V.).

P. oxyacanthae f. crataegi Maurizio.

Host: Rosaceae *Crataegus turkestanica* (2021) ZNNP (Yetikechuvsay V.).

P. plantaginis (Castagne) U.Braun & S.Takam.

(= Erysiphe plantaginis Castagne., Sphaerotheca plantaginis (Castagne) L.Junell.) Host: Plantaginaceae Plantago lanceolata (2016) NNR (Tikchasay V.).

P. pannosa (Wallr.) de Bary.

(= Erysiphe pannosa (Wallr.) Link., Leucothallia pannosa (Wallr.) Trevis., Oidium forsythiae Bunkina., O. leucoconium Desm., O. leuconium Desm., Sphaerotheca macularis f. rosae Jacz., S. pannosa (Wallr.) Lév., S. pannosa var. persicae Woron., S. pannosa var. rosae Woron., S. persicae (Woron.) Erikss., S. rosae (Jacz.) Z.Y.Zhao.).

Host: Rosaceae

Rosa canina (2010-2012) NNR (Xayatsay - Gurdarasay V.).

Rosa beggeriana var. tilanchi (2011) NNR (Majrumsay V.), (2021) ZNNP (Sufa-Usmanlisay V.).

Rosa persica (2011–2017) NNR (Xayatsay V.). Rosa ecae (2021–2022) ZNNP (Yetikechuvsay -

Oʻriklisay V.). Prunus bucharica (2012) NNR (Tikchasay V.). Prunus persica (2009) NNR (Xayatsay V.).

Sphaerotheca fuliginea f. scabiosae Jacz.

Host: Dipsacaceae Lomelosia songarica (1983) ZNR (Guralashsay V.), (2010) NNR (Beshbarmaq V.).

S. fuliginea f. crepidis Jacz.

Host: Asteraceae Crepis pulchra (2019) ZNNP (O'riklisay V.).

S. fuliginea f. senecionis Jacz.

Host: Asteraceae Vickifunkia thomsonii (2022) ZNNP (Irgaylisay V.).

S. fuliginea f. bidentis Jacz.

Host: Asteraceae Bidens tripartite (2018) ZNNP (O'riklisay V.).

S. fuliginea f. dipsaci Jacz.

Host: Dipsacaceae Dipsacus azureus (2010) NNR (Beshbarmaq V.), (2021) ZNNP (Usmanlisay V.).

S. fuliginea f. lophanthi Jacz.

Host: Lamiaceae

Nepeta schtschurowskiana (2015–2016–2017) NNR (Xayatbashi, Tikchasay, Beshbarmaq V.) - Note: First record on *Lophanthus schtschurowskianus* in Uzbekistan.

S. fuliginea f. sedi Kalymb.

Host: Crassulaceae - *Pseudosedum lievenii* (2012) NNR (Gurdara V.) - Note: First record on *Pseudosedum lievenii* in Uzbekistan.

S. macularis f. potentillae Jacz.

Host: Rosaceae Potentilla pedata (2016) NNR (Tikchasay V.).

Sphaerotheca sp.

Host: Rosaceae

Geum kokanicum (2015) NNR (Xayatboshi V.) - Note: First record on *Geum kokanicum* in Uzbekistan.

Uncinula ulmi M.N.Kusnezowa.

Host: Ulmaceae Ulmus laevis (2017) NNR (Qarisay V.).

DISCUSSION

It is known that the diversity of the powdery mildew fungi is closely related to the diversity of the plant flora. Uzbekistan's plant flora consists of more than 4,500 species and 88 species (334 forms) of powdery mildew fungi were registered on 778 host plant species (Gaponenko et al. 1983). The plant flora from protected areas of the Jizzakh region includes 1986 species of 645 genera and 115 families (Tojibaev et al. 2015) and found 57 species, 69 forms and 4 variaties of powdery mildew on 137 host plant species from 34 families.

The distribution of powdery mildew species in host plants is given based on the regions of Jizzakh region in which they are distributed. The largest number of powdery mildew and host plants were found in the Nuratau Nature Reserve (8 genera, 40 species, 41 forms, 3 forms, of the total species number 70.17%), followed by the Zaamin National Nature Park (7 genera, 35 species, 32 forms, 3 varieties, of the total species number 61.40%), Zaamin Nature Reserve (8 genera, 20 species, 36 forms, 2 varieties, of the total species number 35.08 %).

It is noted that studies on powdery mildew fungi of many regions of Uzbekistan including the Anger river basin, and Fergana valley carried out by some mycologists (Panfilova & Gaponenko 1963; Gaponenko et al. 1983; Abdurazakov et al. 2021). The diversity of powdery mildew fungi in protected areas of the Jizzakh region has been compared with their research results (Table 2).

This diversity of powdery mildew fungi of protected areas of the Jizzakh region is represented by 9 genera and 57 species, 69 forms and 4 variaties species and has approximately 30.80% of the currently known mildew fungi biota of Uzbekistan. On the territory of protected areas of the Jizzakh region, the powdery mildew diseases occur frequently and severely damage plants belonging to *Aegilops, Alhagi, Artemisia, Alyssum, Capparis, Cousinia, Convolvulus, Ferula, Morus, Populus, Rosa, Rumex,* and *Trifolium*.

The present checklist is the third work about powdery mildew microfungi of Uzbekistan and serves as one of the sources for a complete list of powdery mildew microfungi of the Rebuplic of Uzbekistan.

Table 2. Comparison of the powdery mildew diversity of protected areas of the Jizzakh region with other regions of Uzbekistan.

Study areas	Plant flora species	Powdery mildew	Powdery mildew host plant species (Percentage towards overall flora)
Uzbekistan (Gaponenko et al. 1983)	4500	88 species, 334 forms	778 (17.28 %)
Angren river basin (Panfilova & Gaponenko 1963)	1500	25 species, 100 forms and 4 varieties	164 (10.93%)
Fergana Valley (Abdurazakov et al. 2021)	2625	67 species	153 (5.82%)
Jizzakh region (this research)	1986	57 species, 69 forms and 4 varieties	137 (6,89 %)

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A case report on chronic renal disease in a captive wild Leopard Panthera pardus (Mammalia: Carnivora)

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Leopards are found all throughout Africa and Asia, but due to isolated and declining populations, they have disappeared from a significant portion of their original range (Stein et al. 2020). The IUCN Red List of Threatened Species (2020) has categorised leopards as 'Vulnerable'. An ideal age distribution for a species population would include a large proportion of young animals and a progressively declining proportion of adults with increasing age. For a number of our captive felid species, the age distribution pattern is now biased toward older individuals. In older captive felids, chronic renal illness is a major cause of death and morbidity (Wack 2008). Age is one of the major contributing factors for glomerular and interstitial alterations in kidneys (Junginger et al. 2015). The kidneys in geriatric canines are often found to exhibit contracted, pale, and indented appearance (Kumar et al. 2020). Captive Leopards are now outliving their free ranging counterparts due to advancing husbandry and veterinary care (Longley 2011).

In the present investigation, on the same day of detection of collapse a Leopard carcass was sent for necropsy examination to the Department of Veterinary Pathology, DGCN COVAS, CSKHPKV, Palampur. A detailed postmortem examination was conducted (Image 1) and representative tissue samples of approximately 0.5 cm thickness were collected in 10% neutral buffered formalin (NBF) for histopathological examination. The fixed tissue sections were dehydrated in ascending grades of alcohol, cleared in benzene, and impregnated in molten paraffin. The tissue sections containing paraffin blocks were sectioned with microtometo 2-3 micron thickness and were stained with Haematoxylin and Eosin (H&E) stain and Masson's trichrome stain as per the standard protocol (Luna 1968) and were microphotographed (Olympus BX40).

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The necropsy examination of the animal showed enlargement of both kidneys, which showed irregular or rough surface along with completely adhered and tense capsule which was hard to peel (Image 2). The lungs were voluminous with oedematous fluid accumulation (Image 3). The small intestine showed the presence of blood mixed with catarrhal exudate (Image 4). The histological examination of the renal tissue exhibited severely congested vasculature with multiple areas of tubular necrosis along with hyaline and cellular degenerations. The glomerular tufts were occupied

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NOTE



Image 1. Leopard presented for necropsy examination. $\ensuremath{\mathbb{O}}$ Rakesh Kumar.



Image 2. Rough surface of kidney with tightly adhered renal capsule. © Rakesh Kumar.



Image 3. Tracheal lumen with mucus mixed oedematous contents. © Rakesh Kumar.



Image 4. Intestinal lumen showing mucus mixed hemorrhagic contents. © Rakesh Kumar.



Image 5. Renal tissue showing congested blood vessels and eosinophilic material in tubular lumen along with peri-vascular infiltration of lymphocytes. H&E x 100. © R.K Asrani.



Image 6. Peri-glomerular fibrosis, glomerulosclerosis and glomerular atrophy in kidney section. MSTx200. © R.K Asrani.

Chronic renal disease in captive Leopard

by collagenous fibrous tissue deposition with resulted atrophy and infiltrating inflammatory cells especially lymphocytes (Image 5). The fibrous tissue in kidneys observed on histopathological examination was further confirmed by Masson's trichrome staining which showed widespread peri-glomerular, inter-tubular fibrosis along with glomerulosclerosis (Image 6) and similar results are supported by a book compiled by Maxie & Newman (2007). Among geriatric dogs and felines, age-related systemic hypertension can contribute to the progression of CKD (Bidani et al. 2012). It has been speculated that environmental stressors like dehydration, psychological stress etc. coupled with aging produce pronounced detrimental impacts on renal perfusion. Furthermore, evidence suggests that inflammatory bowel disease or gastroenteritis in felines and human beings would be expected to cause mild to moderate renal injury due to inflammatory changes or drug therapy (Mitchell et al. 2018)

Based on gross and microscopic changes in kidneys, the leopard in the present investigation is speculated to have died of chronic lymphocytic tubulo-interstitial nephritis with associated lesions in intestine and lungs. The life expectancy of captive felids is longer compared to their free ranging counterparts owing to advances in management and treatment aspects. Animals in captivity are thus more prone to develop age-related degenerative diseases.

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The first photographic evidence of Ruddy Mongoose Herpestes smithii Gray, 1837 (Mammalia: Carnivora: Herpestidae) in Katerniaghat Wildlife Sanctuary, Uttar Pradesh, India

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The herpestid family of mongooses is highly adaptable and can be found in diverse environments, ranging from tropical forests to deserts. These mongooses have a diverse diet that includes insects, crabs, small mammals, birds, and reptiles, with a particular fondness for snakes, and they also consume bird eggs. Some mongoose also consume vegetable matter in the form of tubers, fruits, and berries (Feldliamer et al. 1999). The enormous variety of food they consume is produced in a wide range of environments, making it difficult to measure, even though they must respond to site productivity in some way. Hence, the apparent food abundance is not a good indicator of the distribution or abundance of the majority of species. Environmental and landscape parameters play a significant role in shaping the distribution of mongoose in India, as highlighted in the study by Kalle et al. (2012). The Indian Grey Mongoose Herpestes edwardsii is commonly sighted in disturbed areas, dry secondary forests, and thorn woods, as reported by Gupta (2011). However, it's worth noting that despite their natural habitat preferences, these mongooses are still under high demand in the wildlife trade. Trappers readily capture them to sell them as

pets, as documented by Hanfee & Ahmed (1999) and Kalle (2011).

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The black-tipped tail of the Ruddy Mongoose, which is slightly larger than the Indian grey mongoose, is a defining characteristic (Mudappa 2013). The IUCN's Red List of Threatened Species, categorised it as 'Least Concern' (Mudappa & Choudhury 2016). It is found in peninsular India and Sri Lanka, in the Western and Eastern Ghats, up to the open thorn forest of Rajasthan, and north to Bihar (Phillips 1984; Dookia 2013; Mudappa 2013). In more recent times, the species was documented in Nepal as reported by Subba et al. (2014).

Located on the Indo-Nepal border in the Bahraich District of Uttar Pradesh, the Katerniaghat Wildlife Sanctuary covers an area of 400.69 km² and is positioned between 28°56'72"N, 81°20'97"E. It lies in the Tarai-Bhabhar biogeographic subdivision of the upper Gangetic Plain and supports a variety of habitats (Kalam 2005). In the Katerniaghat Wildlife Sanctuary, the major rivers Karnali and Girwa from Nepal converge and give rise to the Ghaghara River, as documented by Bajpai et al. (2012).

Three distinct Ruddy Mongoose sightings were

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dae) in Katerniaghat Wildlife Sanctuary, Uttar Pradesh, India. Journal of Threatened Taxa 15(9): 23914–23916. https://doi.org/10.11609/jott.8358.15.9.23914-23916

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Photographic evidence of Herpestes smithii in Katerniaghat WS, India

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Figure 1. Distribution map of Ruddy Mongoose with recorded locations in the study area (Katerniaghat Wildlife Sanctuary) with sightings date highlighted.

	Date	Time	Latitude	Longitude	Altitude	Record	Individuals
1	14.v.2021	1444	28.01482	81.32987	134 m	Direct sighting	2
2	30.iv.2018	1255	27.99394	81.33528	138 m	Direct sighting	2
3	16.x.2018	1410	28.02814	81.32841	134 m	Direct sighting	1

Table 1. Distribution records of Ruddy Mongoose.

made in the Katerniaghat Wildlife Sanctuary. The photo that was taken during the direct observation was used to assist identify the species. All three observations were made directly. The species was sighted and photographed while crossing the forest road in the Motipur range of Katerniaghat WS. The Indian Grey Mongoose and Small Indian Mongoose, the other two species living in the study area, can be distinguished from the Ruddy Mongoose by their black-tipped tail. With its diverse vegetation, the Katerniaghat Wildlife Sanctuary is with tropical moist deciduous forest (Bajpai et al. 2012). *Ficus benghalensis, Ficus racemosa, Shorea* robusta, Tectona grandis, Syzygium cumini, and the shrub species Lantana camara, Glycosmis pentapjhylla, and Clerodendrum viscosu make up the majority of the vegetation at all three observation sites.

Previously known distribution range of the Ruddy Mongoose is in peninsular India, and the states of Rajasthan and Bihar, and Sri Lanka (Muddappa 2013). The Ruddy Mongoose is less tolerant toward humans and is considered to dwell in habitats with less human disturbance (Hussain 1999). It was previously recorded in Asola Wildlife Sanctuary in Delhi, but there were no previous records of Ruddy Mongoose from Katerniaghat Photographic evidence of *Herpestes smithii* in Katerniaghat WS, India



Image 1. Adult Ruddy Mongoose *Herpestes smithii* with a visible black-tail tip.



Image 3. Ruddy Mongoose in its recorded habitat in the Motipur range of Katerniaghat Wildlife Sanctuary.



Image 2. Ruddy Mongoose Herpestes smithii.

Wildlife Sanctuary; however, it was recently recorded from Banke National Park in Nepal (Subba et al. 2014). Thus, our record of Ruddy Mongoose in Katerniaghat Wildlife Sanctuary marks the distribution in gap regions of its known distribution range.

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New locality record of the Asiatic Long-tailed Climbing Mouse Vandeleuria oleracea (Bennett, 1832) (Mammalia: Rodentia: Muridae) from Kohora River Basin, Assam, India

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Vandeleuria oleracea (Bennett, 1832) is a 'Least Concern' species belonging to the genus Vandeleuria found in southern and southeastern Asia (Aplin & Molur 2017). Although Agrawal & Chakraborty (1980), Agrawal (2000), and Srinivasulu & Pradhan (2003) mention that Vandeleuria oleracea has two subspecies, namely, V. o. oleracea from southern India and V. o. dumeticola from northern India, the present taxonomic changes indicate that V. oleracea may be a species complex with subspecies recognized by earlier workers synonymized under V. oleracea, pending further studies (Musser & Carleton 1993, 2005; Srinivasulu & Srinivasulu 2012; Wilson et al. 2017).

On 5 June 2023, during our regular field survey, we encountered a small dead individual of a mouse at Rongtara Village of Kohora River Basin, (26.53–26.60 °N & 93.33–93.43 °E; covering an area of 31.50 km²) Karbi Anglong District, Assam (Figure 1). Morphological measurements were taken (Head and body length: 60.66 mm; tail length: 100.31 mm; hindfoot: 14.12 mm, and ear length: 11.35 mm). The mouse was identified as *Vandeleuria oleracea* by its rusty brown dorsum and white ventral coloration with the head and body length less than 100 mm, unicoloured tail, which was much longer, about one and a half times the head and body length; hallux and fifth toe clawless; fifth toe appeared to be opposable (Agrawal 2000) (Image 1).

The mouse was seen dead in the Jhum plantation near a bamboo clump. Later that day, we conducted informal interviews among the local community in Rongtara Village and showed community members photographs of the species (Image 1). According to the local community, the species is rarely seen in the forest (Image 2). The species was known to them and they call

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Editor: Anonymity requested.



Figure 1. Distribution map of Vandeleuria oleracea in Assam with the past record at Kamrup in 1961 and the present record at Rongtara in 2023.



Image 1. Dead Vandeleuria oleracea at Rongtara.

it 'Jukikso' in Karbi dialect.

The only known record of *Vandeleuria oleracea* from Assam was from Angarkhata, North Kamrup (Ellerman & Morrison-Scott 1951; Ellerman 1961; Molur et al. 2005; Chatterjee et al. 2020; Talukdar et al. 2021). The new site record from Rongtara Village is approximately 215 km from the earlier report by Ellerman (1961) (Figure 1). The species inhabits agricultural regions (Jhum) in close

New locality record of *Vandeleuría oleracea* from Kohora River Basin, Assam



Image 2. Habitat of Vandeleuria oleracea at Rongtara.

proximity to secondary bamboo forests. This species is arboreal and exhibits nocturnal behavior, primarily consuming fruits, buds, and flowers.

The present study reported the new locality record from Karbi Anglong that confirms its distribution in Assam and Northeast India with an addition of a second confirmed locality in Assam.

Further studies on the ecology and the habitat of the species are needed to understand the species' habitat requirements. Studies involving an integrative taxonomic approach including phylogenetic studies will help resolve this species complex.

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New distribution record of fish *Clupisoma garua* (Hamilton, 1822) (Siluriformes: Ailiidae) from the Sarpang District in southern central part of Bhutan

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Clupisoma garua (Hamilton, 1822) falls under catfish group, which is widely known by different names as Neria (Assam in India), Ghaura (Bangladesh), Baikha/ Jalkapoor (Nepal) (Wang et al. 2016). It belongs to the Ailiidae family under Siluriformes order and it is widely distributed across the Indian rivers and reservoirs. Globally, Ailiidae familiey is native to Africa and Asia which comprises of 66 species that belongs to 14 genera. However, nearly half of the species (32 species) are known from Asian countries that comprises of five genera including Clupisoma, Ailia, Horabagrus, Laides, and Pseudeutropius (Wang et al. 2016). Among them, the Clupisoma genus has five species of which four are reported from the Indian region: garua, bastari, naziri, and montana (Hora 1937). Globally, C. garua is distributed around the Ganga River system in India and Nepal, Ganga-Brahmaputra River system in Bangladesh and Indus River system in Pakistan (Bhokta & Solanki 2020). In case of India, C. garua is widely distributed in Bihar, West Bengal, Odisha, Madhya Pradesh, and Assam (Brahmaputra and Barak drainage) (Bhokta &

Solanki 2020). However, this species is threatened in some localities such as southwestern Bengal due to overfishing (Verma et al. 2014) and decline from natural water bodies (Patra et al. 2005; Mishra et al. 2009). Meanwhile, the recent record of *C. garua* from Ayechu River had set new distribution record from the Bhutan. As per the IUCN Red List, the species is categorized under 'Least Concern' (LC) IUCN Red List (2020). However, both CAMP (Molur & Walker 1998) and CAFF (2006) had declared it as Vulnerable (VU), due to the reduction of populations in their natural habitats; while, in Bangladesh, the species is recently kept under Critically Endangered (Hanif et al. 2015) due to restricted geographical distribution fueled by the increasing anthropogenic and natural hazards (Siddik et al. 2017).

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Clupisoma garua is commercially important freshwater fish that has a potential species for aquaculture system (Saraswat et al. 2014). The studies of Bhuiyan (1964) and Memon et al. (2010) also reported that *C. garua* is mostly consumed by various group of people including the marginalized people due

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Figure 1. Bhutan map showing the location of Ayechu where Clupisoma garua was recorded in Sarpang District.

to high level of protein (18.40%) and fat (5.2%) content, followed by most abundant in some of the river system (Galib et al. 2009). Besides these, the species also has ornamental values that promotes livelihood of the coastal communities (Gupta et al. 2016).

Mouchhu basin in Gelephu is located between 26.923–26.847°N & 90.506–90.504°E that falls under Sarpang District in the southern central part of Bhutan (Figure 1). Mouchu basin drains from the Black Mountain which is located at the central part of Bhutan that flows through Gelephu and exit towards India, which finally confluences with Brahmaputra River. In the case of Ayechu (26.875°N, 90.501°E), which is the diverted river of Mouchhu basin whereby *C. garua* was opportunistically recorded for the first time from this river in September 2020 using cast net. Since *C. garua* is a migratory species, it could have migrated from India during monsoon season through Mouchu basin as it hasn't been recorded during the past survey. With this new addition, Bhutan now has 126 fish species (DoFPS

2020), out of which 28 fish species are found in Sarpang District that belongs to 11 families (Tenzin 2022). Among 28 species, three species were categorized under Vulnerable, three Endangered, two Near Threatened and rest are Least Concern as per Tenzin (2022). The district falls within the convergences of three ecologicallydiverse protected areas of Bhutan which are connected with each other by Biological Corridor No. 03 (Tenzin et al. 2021). Sarpang shares the southern border with the northeastern state of Assam in India which is further connected with the Royal Manas National Park (RMNP) and Indian Manas National Park (MNP) towards the east and Phibsoo Wildlife Sanctuary (PWS) in the west (Tenzin et al. 2021). Floristically, this area comprises of subtropical broad-leaved forests located at an elevation of 200 m and this area receives the average precipitation of 3,500-5,500 mm (Tenzin et al. 2022). The specimen was confirmed with professor D.B. Gurung from College of Natural Resources, Punakha District, Bhutan through morphometric measurements using digital caliper. The

New distribution record of *Clupisoma garua* from Sarpang District, Bhutan

studies of Bhokta & Solanki (2020) was also referred for confirmation. Meanwhile, the specimen was collected and euthanized using 0.001 percent clove oil and treated in 10% formalin for fixation as per Gurung et al. (2012) and Tenzin & Dhendup (2017) and it is currently deposited in the Laboratory of the Southern Wildlife Rescue and Rehabilitation Centre (SWRRC) in Sarpang District. Fin formula is the key feature been used for identification and comparison as tabulated (Table 1).

The studies of Jayaram (1977) found that *Clupisoma garua* is a herring-shaped fish that gradually tapers toward both ends and abdominal edge between pelvic fins and vent (Image 1, 2 & 3). Talwar & Jhingran (1991) also substantiated that the adipose dorsal fin is absent in adults, while, caudal fin is deeply forked with lower lobe longer than upper with black edged dorsal, pectoral and caudal. On other hand, the eyes are large with circular adipose eyelids and their mouth is wide and terminal.

Table 1	. Mo	rphometric	meas	surement	of F	in	formula	0	f <i>C. ga</i>	rua
species	and	compared	with	morphor	netri	ic s	studies	of	Bhokta	&
Solanki	(2020)).								

	Variables of fin formula (cm)	Specimen of current study	Specimen Fin formula of Bhokta & Solanki (2020)
1	D	1/7	1/6
2	А	27	3/28
3	Р	1/11	1/11
4	v	1/5	1/5
5	С	18	17–20
6	Maximum length (TL)	16.6 cm	60.9–100 cm
7	Colour of fin	Yellowish-orange	Yellowish-orange
8	Presence of adipose fin	No.	No.

D–Dorsal fin | A–Anal fin | P–Pectoral fin | V–Ventral/Pelvic fin | C–Caudal fin.



Image 1. Ventral view of Clupisoma garua species. © Sangay Dorji 2020.



Image 2. Lateral view, mouth and head view of Clupisoma garua species. © Sangay Dorji 2020.

New distribution record of Clupisoma garua from Sarpang District, Bhutan



Image 3. Head and mouth view of *Clupisoma garua* species. © Sangay Dorji 2020.

Nonetheless, body is coloured with silvery grey on the back which is lighter on the sides and abdomen with tinted grey color fins and black edged dorsal, pectoral and caudal fin respectively (Talwar & Jhingran 1991). The present *C. garua* specimen has a total length (TL) of 16.60 cm; however, it can grow up to maximum TL of 60.90 cm (Bhokta & Solanki 2020).

The species is mainly found in lacustrine habitat in larger rivers and reservoirs with stagnant impoundments (Bhokta & Solanki 2020). The studies of Froese & Pauly (2013) and Saraswat et al. (2014) reported that *C. garua* is potamodromous that migrates within streams & rivers and travels a long distance (>100 km) for feeding as well as for seeking suitable breeding habitat in new water bodies. Further, *C. garua* is a carnivorous fish that exploits food resources in the surface guild and also feeds along the margins of the river. Feeding intensity is higher during the September–October months (Bhokta & Solanki 2020).

As per IUCN Red List (2020), the species is categorized as 'Least Concern' (LC). However, it's categorized under Critically Endangered in Bangladesh IUCN Bangladesh (2000) and Vulnerable in India (Molur & Walker 1998; Lakra et al. 2010). In several parts of the range country, the populations are reported to be declining from the natural habitat (Bhokta & Solanki 2020). Biswa et al. (2018) also substantiated that over exploitation, habitat loss, human interference, climate change, pollution, and siltations are the main causes of the population decline, besides overfishing in range countries. However, indepth ecology and pertinent conservation threats from Bhutan is still unknown, due to recent occurrences which may require separate ecological studies along the Mouchu River in future.

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Kukumseri: a home to *Colchicum luteum* Baker (Colchicaceae), a rare and endangered medicinal herb

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Colchicum luteum Baker (Colchicaceae) commonly known as Hirantutiya (Hindi), Hiranyatutha (Sanskrit), Suranjantalkh (Urdu), and Kukum (locally) is an important medicinal herb, used in various traditional herbal medicine systems (Ayurveda, Siddha, Unani) (Rather et al. 2022). The species contains alkaloids including, colchicine & democolcine which have antimitotic properties and used for the treatment of solid tumors and leukemia (Ondra et al. 1995; Yue et al. 2010). C. luteum also reported to have analgesic, antiinflammatory, antimicrobial, aphrodisiac, carminative, laxative & wound healing properties, and especially useful in gout, rheumatism and Alzheimer disease (Aisen et al. 2001; Javed et al. 2005; Rather et. al. 2022). The said effects have been well proven in various studies through animal models (Ahmad et al. 2006; Akbar 2020)

Colchicum luteum is found in the Hindu Kush-Himalayan region and has very restricted distribution, endemic to certain places only, and considered as a rare species (CAMP report 2010; Rather et al. 2022). The habitat specificity; requirement of low temperature, less relative humidity and specific soil types for growth are being the reasons for their restricted distribution among others. Besides its restricted distribution and rarity, the indiscriminate over-exploitation for medicinal use has further endangered the survival and has been categorized as an endangered plant species (CAMP report 2010). In Indian Himalayan region (IHR) *C. luteum* has only been reported from certain places of Kashmir and Himachal Pradesh (Ved et al. 2003). The literature revealed, lack of systematic studies on diversity and distribution of *C. luteum*, the available data being either subjective or ethnobotanical. Further, the absence of species in detailed reports/ works on flora of the area including, flora of Lahaul-Spiti (Aswal & Mehrotra 1994) confirm the same. Rather et al. (2022) also observed the lack of information regarding distribution and population size of *C. luteum* from Kashmir Himalaya.

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NOTE

Therefore, to explore the distribution and population size of C. luteum in Lahaul, Himachal Pradesh, India, a survey was conducted during March-April, 2021 and 2022. The species was found on open moist slopes between 2,650-3,000 m elevation (Images 1 & 2). The species appeared soon after the melting of the snow in March and were the earliest plants to flower during spring, completing the growth cycle (vegetative, flowering, and seed set) within two to three months (March-May). During this period of the year the area remains mostly snow bound, less accessible, and has poor vegetation due to unfavorable or cold weather conditions. Thus, due to above listed facts (habitat specificity, rare and endangered population status and comparatively very short life cycle during unusual period of the year) people rarely visit the places and/

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Images 1 & 2. Colchicum luteum in its natural habitat in Lahaul, Himachal Pradesh. © Rajender Kumar Sharma.



Figure 1. Locations of *Colchicum luteum* population in Lahaul, Himachal Pradesh, India.

or notice the species during the said period. Therefore, despite the importance of this elite genotype (Wagh et al. 2015), the species has been poorly investigated.

In the present study area, the species were recorded at two locations: Mailing $(32.605^{\circ}N, 76.934^{\circ}E, 2,925)$ m elevation) and Kukumseri $(32.698^{\circ}N, 76.687^{\circ}E, 2675)$ m elevation) (Figure 1 & Image 3). The observed species density in Mailing and Kukumseri was 6.4 ± 5.7 and $3.08 \pm 4.15/m^2$, respectively, determined by counting the total number of individuals of a species in all quadrants $(1 \times 1 m, each)$, and divided by total number of quadrants studied. The extent of occurrence (EOO) of species in Kukumseri region was found higher than the Mailing. In Mailing, the species were only distributed in an area of about three Km². The distribution of species only to a certain specific place seems due to the requirement of specific soil condition, and the latter need to be explored. Very interestingly, it was observed that one location (Kukumseri) has been named after the species (*Colchicum luteum*) where, 'Kukum' means *Colchicum luteum* and 'Seri' means field. Thus, 'Kukumseri' means, the field of *Colchicum luteum*.



Images 3. A view of habitat of Colchicum luteum population at Kukumseri in Lahaul, Himachal Pradesh, India.

Furthermore, the location justifies its name due to the presence of a reasonably large number of species in the area. Recently, huge destruction of natural habitat (4.2 ha) in Kukumseri, has been done for the opening of Krishi Vigyan Kendra (2.6 ha) and College campus (1.6 ha). The latter would have been avoided considering the restricted distribution, habitat specificity, rarity, endangered status and medicinal/economic importance of *C. luteum*.

The information reported here will be of immense help to locate and visit the places at appropriate period of the year for studies, aimed at propagation, cultivation, and conservation of the species. Further, such data plays crucial role while assessing the population status of the species.

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First record of the Western Himalayan Yew *Taxus contorta* (Gymnosperms: Cupressales: Taxaceae) from Lumbini Province, Nepal

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Yews (Taxus spp.) are among the most threatened plant species within the Hindu Kush-Himalaya (HKH) region including Nepal (Mulliken & Crofton 2008). Due to over exploitation of the species for the production of anticancer drugs and in some areas intense local use for medicine, timber, and fodder the natural yew populations along the HKH region have been cleared up to 90% over the last few decades (Schippmann 2001; Mohapatra et al. 2009). Delayed germination (1.5-2 years) of its seeds and poor survival rate of seedlings have further accelerated the decline of yew population in the Himalaya (Rikhari et al. 1998). Despite its threatened (EN) status, little information is available regarding the size and status of its populations (Iqbal et al. 2019). Even the taxonomic identification of species remained controversial in Nepal till the year 2012 (Poudel et al. 2012). The Forest Regulation, 1995 named the taxus species found in Nepal as Taxus baccata in its annex till 12 October 2015, which was corrected and named as Taxus contorta Griff., Taxus wallichiana Zucc., and Taxus mairei (Lemée & H. Léveillé) S.Y. Hu ex T.S. Liu in its fifth amendment in the year 2015.

Currently, a total of 13 (four in North and South

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America, one in Europe and eight in Asia) species of *Taxus* have been identified in the world (Gao et al. 2007; Farjon 2010; Liu et al. 2011). The threatened Western Himalayan Yew *T. contorta* is one among three of the *Taxus* species (*T. contorta, T. mairei, T. wallichiana*) found in Nepal (Thomas 2011; Poudel et al. 2012, 2014; Bhatt et al. 2017). It is assumed to be distributed sparsely over Darchula District in the far western region to the northern belt of Gorkha District in the central region of Nepal (Poudel et al. 2014). It has been recorded from several districts of three of the seven provinces of Nepal (Bhatt et al. 2017). However, this is the first record from Lumbini Province, Nepal.

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T. contorta is a medium-sized dioecious evergreen tree species which grows naturally within the elevation range of 2,000–3,500 m in Nepal and the recorded height of *T. contorta* in Nepal is 25–30 m (Bhatt et al. 2017). The species has diagnostic characters such as bud scales few, ovoid, persistent at the base of branchlets; leaves arranged irregularly, pectinate, usually linear, equally wide throughout length, base cuneate, mostly symmetric, apex acute, midrib papillate, midrib & leaf margin underneath not shiny, loosely arranged (6–9)

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Fírst record of Taxus contorta from Lumbini Province, Nepal



Image 1. Map showing the Taxus contorta recorded locations in Rolpa, Lumbini Province, Nepal.

stomatal bands, margin revolute-incurving when dried (Poudel et al. 2012).

During the transect walk survey conducted on March-April 2021 along the four distinct elevation gradient of 2,000–2,400 m, 2,400–2,800 m, 2,800–3,200 m, and 3,200–3,600 m in Thawang Rural Municipality (28.50°N, 82.71°E) of Rolpa District in Lumbini Province, we recorded *T. contorta* in at least 55 locations (Image 1). Taxus contorta is recorded mostly (90%) in the northern aspect along the wet sites, a habitat also mentioned in earlier literature (Poudel et al. 2012; Bhatt et al. 2017), with slopes more than 35 degrees, crown cover more than 70% and elevation range of 2,424–3,002 m. The species was recorded at sites away from human dominated landscapes where anthropogenic activities are either minimum or totally absent. The species was found in association with Picea smithiana Wall., Tsuga dumosa (D.Don) Eichler, Quercus semecarpifolia Sm., Abies pindrow D.Don, Rhododendron arboreum Sm. (Freitag 1971; Rau 1974; Sapru 1975), Daphne bholua D.Don, Rhododendron barbatum Wall., and Himalayacalamus asper Stapleton in the temperate

forest.

In western Nepal, T. contorta populations are on a satisfactory level inside the protected areas (Api-Nampa Conservation Area, Khaptad National Park, Rara National Park, and Dhorpatan Hunting Reserve), however, it is sporadically distributed outside the protected areas (Bhatt et al. 2017). Furthermore, the T. contorta is an 'Endangered' species (Thomas 2011), and these newly located populations provide an opportunity for indepth study of their niche and associated site specific threats that will further aid in developing community engagement conservation programmes outside the protected area. Also, the species is facing a threat due to illegal felling for timber and leaves, improper harvesting methods, loss of its natural habitat, unmanaged grazing, delayed germination and lower survival rate. Therefore, it is comprehended to conserve them in-situ and promote its revival through nursery cultivation and plantation initiatives.

Fírst record of *Taxus contorta* from Lumbíní Provínce, Nepal

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Image 2–4. 2—Branching habit of healthy *Taxus contorta* sapling (X: 669769, Y: 3148319, Elevation: 2,810 m.) | 3—Tree stem of *Taxus contorta* (X: 665087, Y: 3149405, Elevation: 2,947 m.) | 4—*Taxus contorta* forest patch (X: 669637, Y: 3148356, Elevation: 2,720 m.). © Image 2—Deepak Raj Prakash Jung Shahi, 3–4—Santa Bahadur Thing.

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