Candelaria asiatica, New record from Pakistan

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ABSTRACT

An investigation of the morphology, chemistry, and molecular phylogenetics of a *Candelaria* specimen was undertaken when the lichens of Malakand were being surveyed. With a pulverulent surface, a tiny yellow lobate thallus, and a delicate lobe edge featuring lobules that resemble phyllidia, the taxon that is new to Pakistan is identified as *Candelaria asiatica*. The morpho-anatomical, ecological, and distribution details are provided.

KEYWORDS Biota, Candelariaceae, taxonomy, phylogeny

INTRODUCTION

Pakistan is widely recognized for its diverse landscape and climate, which is associated with an abundance of biological diversity (IUCN, 2006). Because there haven't been many studies conducted in many areas, it's likely that the lichen diversity in this region is quite great. (Aptroot & Iqbal, 2012).So far, up to 400 lichens have been reported from Pakistan (e.g. Aptroot & Iqbal, 2012; Fatima *et al.*, 2021; Habib *et al.*, 2021; Kousar *et al.*, 2021; Fayyaz *et al.*, 2022; Nadeem *et al.*, 2022).

Candelaria A. Massal. is a small lichen genus characterized by a micro-foliose to micro-fruticose yellowish thallus and 8- or polyspored asci (Westberg & Frödén, 2007). *Candelaria concolor* (Dicks.) Arnold, is reported from Pakistan (Aptroot & Iqbal, 2012). A *Candelaria* specimen was collected during a survey of the lichen flora in Malakand. The specimen, when subjected to morphological, chemical, and phylogenetic investigations, showed a strong resemblance to *Candelaria asiatica*.

MATERIALS AND METHODS

Morphological and Chemical Investigations

Lichen specimen was collected from Malakand, Khyber Pakhtunkhwa in 2018. Meiji Techno's EMZ-5TR stereomicroscope and the Swift M4000-D compound microscope, both equipped with a 9MP camera system, were used to study the specimen macro and micromorphologically. Hand-made sections of apothecia were investigated in 10% KOH and water for anatomical study. Each diagnostic feature from specimen was measured in the water at least twenty times. Spot tests using sodium hypochlorite solution (C) and KOH (10%; K) were used to assess the secondary chemistry. Standard procedures for Thin Layer Chromatography were followed while utilizing Solvent System C. (Orange *et al.*, 2010).

Extraction of DNA, amplification by PCR, and sequencing

Using a modified 2% CTAB technique, genomic DNA was directly isolated from thallus (Gardes & Bruns, 1993). The ITS-nrDNA region (Internal Transcribed Spacer of the nrDNA) was amplified using the primer pair ITS1F (forward primer) (Gardes & Bruns, 1993) and ITS4 (reverse primer) (White *et al.*, 1990. The PCR results were visualized using ethidium bromide on a 1% agarose gel. (Sambrook & Russel, 2001). PCR products were sent to Tsingke, China for sequencing.

BLAST analysis was used to obtain extremely comparable ITS region sequences (Altschul, 1990). Related taxa were indicated, along with the sequence maximum query coverage and percent identity. Sequences from published literature and GenBank were used in an initial alignment, and then web-PRANK was used to realign the data using its default parameters. (Dong Liu & Jae-Seoun Hur, 2018). Phylogenetic relationships were investigated using Maximum Likelihood bootstrapping, as implemented in RAxML-HPC2 v. 8.1.11 (Stamatakis, 2014), hosted on the CIPRES Science Gateway. The HYK+G+I substitution model and quick bootstrapping with 1000 iterations were employed in the analyses. Version 1.4.3 of FigTree (Rambaut *et al.*, 2014) was used for displaying trees from the ML analysis.

RESULTS

Phylogenetic analysis

In this study, the ITS on nrDNA of *Candelaria* species obtained from Buner was successfully amplified and generated. The phylogenetic analysis sample of taxa was based on collecting closely related sequences from GenBank. MAFFT v7.450 (Katoh & Standley, 2013) The ITS sequences were aligned using Mafft v7's 'automatic' option, with all other parameters set to default. Each one-locus alignment's ambiguous spots were deleted. Maximum Likelihood was used to do the phylogenetic analysis (ML). ML trees were generated using RAxML 8.2.11 on GENEIOUS (Stamatakis, 2014), with 1000 bootstrap pseudoreplicates and the GTR+G+I substitution model. FigTree 1.4.2 was used to visualise the tree files (Rambaut, 2014).

Analysis involved 26 nucleotide sequences. Pakistani collection *C. asiatica* is clustered with *C. asiatica* (MG694269, MG694270).

Taxon	Origin	Specimen Voucher	GenBank Number (ITS)	
Amandinea efflorescens	Sweden	SE-901 87	AY143409	
Candelaria asiatica	Pakistan	Bot-99	PP091273	
Candelaria asiatica	South Korea	D. Liu 171446 (KoLRI)	MG694269	
Candelaria asiatica	South Korea	D. Liu 171454 (KoLRI)	MG694270	
Candelaria concolor	Albany	APBP017A	MK966426	
Candelaria concolor	Canada	BIOUG24047-G11	KT695365	
Candelaria concolor	USA	PUL F27065	MW448573	
Candelaria concolor	Mexico	Westberg 454 (LD)	EF535205	
Candelaria concolor	Italy	Arup L07018 (LD)	GU929921	
Candelaria concolor	Sweden	SAR07	FJ959355	
Candelaria concolor	Italy	Arup L07001 (LD)	GU929922	
Candelaria fibrosa	USA	Worthington 21240 (COLO)	EF535206	
Candelaria fibrosa	Argentina	Froeden 1670 (LD)	GU929923	
Candelaria fibrosoides	Peru	Froden 1513 (LD)	EF535211	
Candelaria fibrosoides	Peru	Froden 1512 (LD)	EF535212	
Candelaria fruticans	Ecuador	ECU352	EF535207	
Candelaria pacifica	Turkey	28.IV.1992 John EF535210 [John, Lich. Anatol. Exs. no. 16] (ASU)		
Candelaria pacifica	USA	Westberg 967 (LD)	GU929918	
Candelariella aurella	USA	Westberg 150 (LD)	EF535163	

Table 1. Sequences used in the ITS phylogenetic analyses of *Candelaria* species.

Candelariella	USA	Westberg 1181 (LD)	EF535164
biatorina			
Candelariella	Russia	ALTB:E.A. Davydov	KX853128
blastidiata		7716	



Figure. 1. Molecular phylogenetic analysis of ITS sequences of *Candelaria* inferred by using the maximum likelihood method.

TAXONOMY

Candelaria asiatica D. Liu & J.S. Hur, Mycobiology 46(4): 308 (2018).

Thallus: corticolous, scattered, and squamulose to foliose or subfruticose, 0.3–1.5 cm, irregular, rosettes, aggregated into extensive colonies covering the substrate. **Lobes**: linear and irregular branched, 0.1–0.59 mm, adnate to erect, lobe tip slightly inverted, crenate, lobe surface color yellow to greenish yellow in the center, then becoming bright yellow toward the lobe tip, phyllidia-like lobules. **Margins:** 50–130 mm wide loosely adnate lobe tips. **Upper cortex:** 11–25 µm thick. paraplectenchymatous, texture globularis. **Medula:** white but not well-developed; the lower cortex white, covered with white rhizines; **Algae cells:** chlorococcoid, cells 8-12 µm in diam. **Medulla**: white but undeveloped, the lower cortex is present at the center of the thallus. **Lower surface:** white, covered with white rhizines. **Rhizines:** white, densely present. **Apothecia:** not observed.

Chemistry: K-, KC-, C-, PD-; Secondary metabolites: calycin and pulvinic acid detected.

Habitat: On bark in coniferous forest, an altitude is 994 m a.s.l., dominant plants are *Picea smithiana* (Wall.) Boiss. *Cedrus libani* subsp. Stenocoma. *Juniperus communis* subsp. Hemisphaerica, Taxus wallichiana Zucc. *Quercus dilatata* Royle. *Acer cappadocicum* Gled. And *Betula utilis* var. jacquemontii (Himalayan Birch). Himalayan region, maximum and minimum temperature of 35°C and -10°C respectively, annual rainfall varying between 600-800 mm.

Material Examined: PAKISTAN. Khyber Pakhtunkhwa: District Buner, Daggar; at 994 m a. s. 1., May, 04, 2018, H. Wahab, A. N. Khalid. HB–5. ITS GenBank accession number PP091273.

REMARKS:

A small lichen genus *Candelaria* A. Massal. is distinguished by its 8- or polyspored asci and microfoliose to micro-fruticose yellowish thallus. There are eight species in this genus: *Candelaria Antarctica* (Js. Murray) Poelt, *C. crawfordii* (M€ull. Arg.) P.M.Jørg. & D.J. Galloway, *C. concolor* (Dicks.) Arnold, C. fibrosa (Fr.) M€ull. Arg., *C. fibrosoides* M. Westb.& Fr€oden, *C. fruticans* Poelt & Oberw., *C. murrayi* Poelt, and *C. pacifica* M. Westb. & Arup, around the world (Dong Liu & Jae-Seoun Hur, 2018). Only *C. concolor* is reported from Pakistan by Aptroot and Iqbal in 2012. The Pakistani collection of *Candelaria asiatica* is grouped with South Korean specimens (MG694269, MG694270) in a branch with a strong bootstrap value. Their designation as *Candelaria asiatica* is confirmed by morpho-anatomical and chemical characters.



FIG. 2. *Candelaria asiatica* A, habitat; B, lobes margins after blastidia or phyllidia like lobules are dropped; C, young thalli; D, lower surface E, transverse section of the thallus.

Taxon	Lobe	Upper surface	Asexual	Distribution
			propagules	
C. asiatica	linear and	rough	phyllidia-like	Pakistan
	irregular		lobules present	
	branched			
C. asiatica	Shallowly to	rough	Blastidia or	South Korea
	deeply branched,		phyllidia-like	
	narrower (up to		lobules present	
	0.47 mm)			
C. concolor	Deeply	smooth	Soredia granular	Asia, America,
	branched,			and Europe
	narrower (up to			
	0.5 mm)			
C. crawfordii	Shallowly	smooth	Soralia	Asia, Australia
	branched, wider		labriform, with	
	(up to 0.7 mm)		soredia on the	
			upturned lower	
			surface.	
C. fibrosa	Wider (up to 2	Smooth or	Soredia absent	Pantropical
	mm)	wrinkled		region

Table 2. Mor	phological d	ifference a	mong the	species of	f <i>Candelaria</i>	species
			()			

CONCLUSIONS

The ITS rDNA phylogenetic trees had similar topologies with *Candelaria* species forming a separate clade with *C. asiatica* (Fig. 1). The morphological characters are similar to *Candelaria asiatica* (Dong Liu & Jae-Seoun Hur, 2018).

However, information regarding this species is still limited and the abilities of this species should be investigated further.

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