

Cyanoprokaryotes of genus *Petalonema* - taxonomically and molecularly interesting taxa



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Introduction

The genus *Petalonema* contains six subaerial habitats associated endo/epilithic cyanobacteria requiring special substrate conditions for growth in the nature. The cyanoprokaryotes can be described as filamentous, heterocytous, falsely branching with yellowish-brown lamellated mucilage sheaths. Their cells are widely rounded, heterocytes are intercalary or basal. On the rocks and stones, they form soft gelatinous colonies. The areal of this genus is zonal, they can be found in Antarctica, Australia, India, North America and several places in Europe. At least one species, *P. crustaceum*, can be found also in the Czech Republic (Komárek & Hauer 2007).

Apart from the type species *Petalonema alatum*, other known *Petalonema* species are *P. crustaceum*, *P. densum*, *P. involvens*, *P. pulchrum* and *P. velutinum*.

This study focuses mainly on *Petalonema alatum*, which was first described from Scotland by Berkeley in 1833. Since that time, the taxonomical position of the species (and genus) changed several times. Borzi in 1879 considered *Petalonema* to be synonymous to *Scytonema*. Other opinion was published by Itzigsohn in 1855, who didn't accept the genus *Petalonema* at all and considered *P. alatum* to be just morphotypes inside *Scytonema myochrous*. During the 133 years since *Petalonema alatum* description, authors chose one of the three taxonomical possibilities. Although *Petalonema* exhibits high morphological similarity with *Scytonema*, Komárek & Anagnostidis (1989) didn't place it to the family *Scytonemataceae*, but to the family *Microchaetaceae*. The only molecular study done on the genus *Petalonema* (species *P. cf. involvens* in particular) was published by Taton et al. (2006) who sequenced 16S and 23S rRNA gene and placed *Petalonema* to the *Nostoc* clade.

The aim of the study was and is to confirm the validity and clarify the taxonomical position of genus *Petalonema*, based both in morphological variations and molecular taxonomical data.

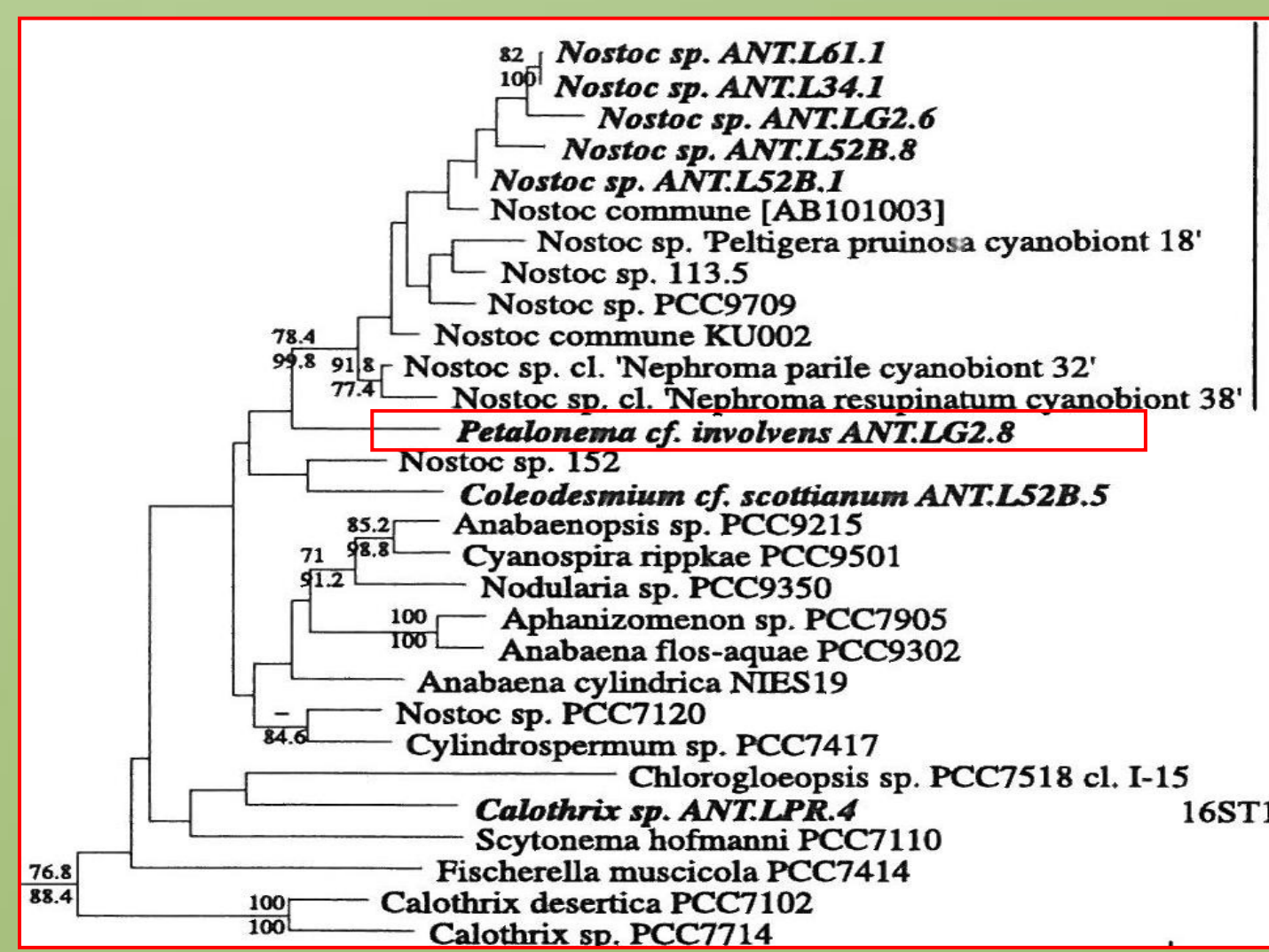


Fig. 1: Position of *Petalonema cf. involvens* in the phylogeny tree. Taken from Taton et al. (2006)

Fig. 2: Iconographs of *Petalonema alatum*: (a-i) natural samples; (j-s) culture; (a-c, e-g, j-m, o-q) heteropolarity of filaments and sheath stratification; (d, n) hormogonia formation; (r-s) trichome spiral-curved in culture. Orig. B. Uher.

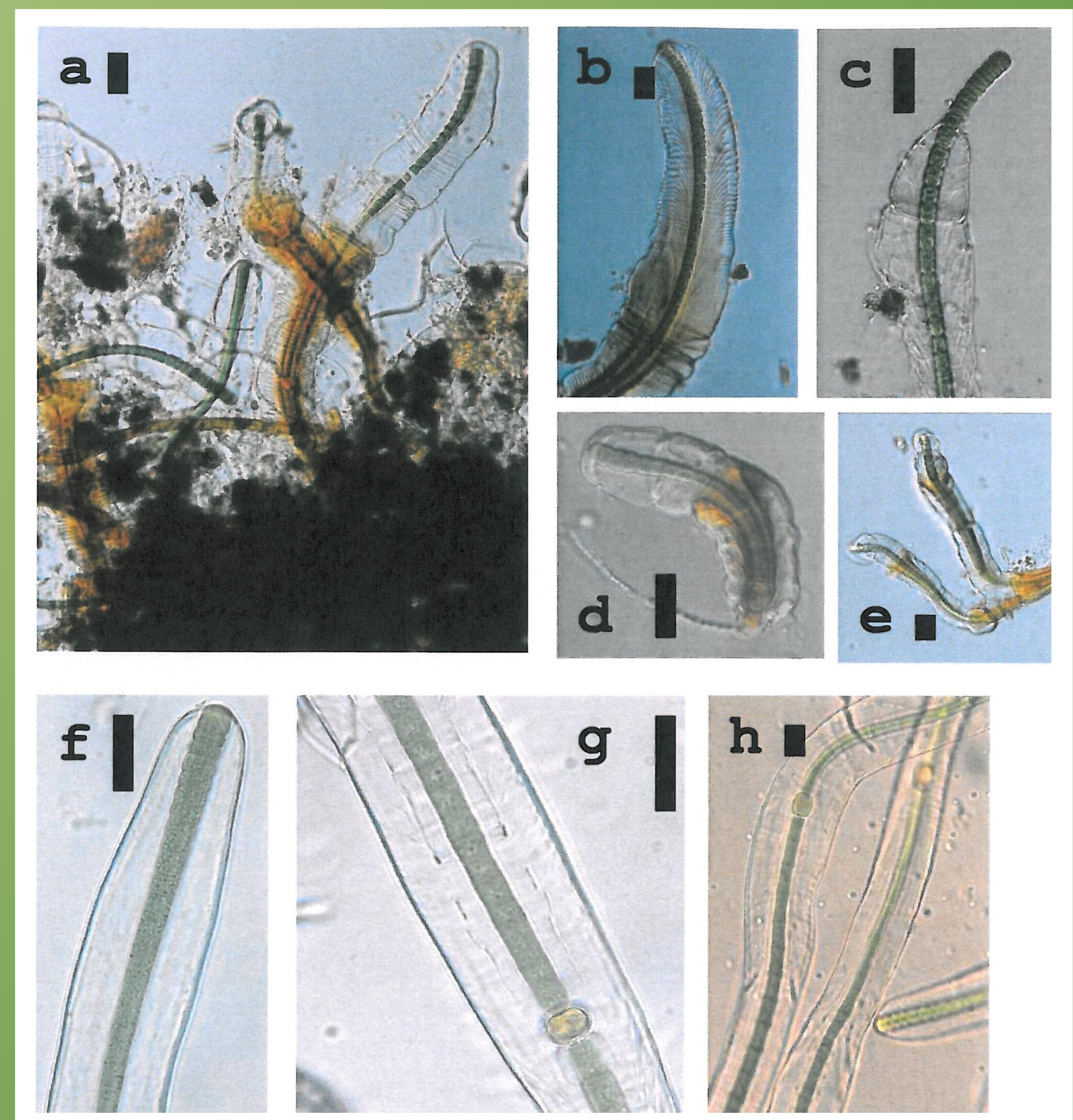
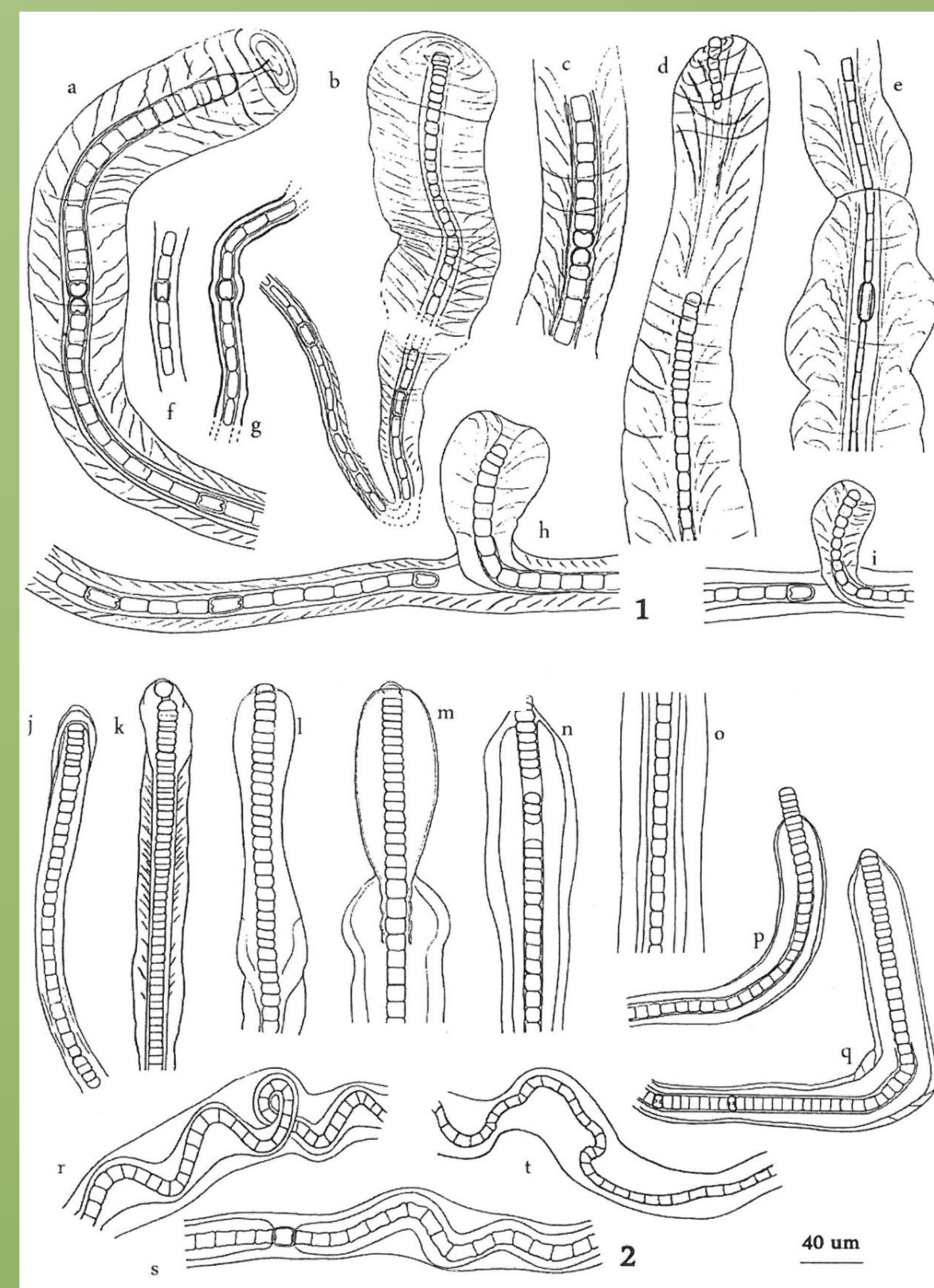


Fig. 4: *Petalonema alatum*: (a-e) natural sample; (f-h) culture; (a) filament arising from the substrate, (b, c, f) apices with apical rounded cell, (d) tolypotrichoid fragment of filament; (e) false branching; (g, h) well established sheath stratification. Scale bars 30 μ m. Orig. B. Uher

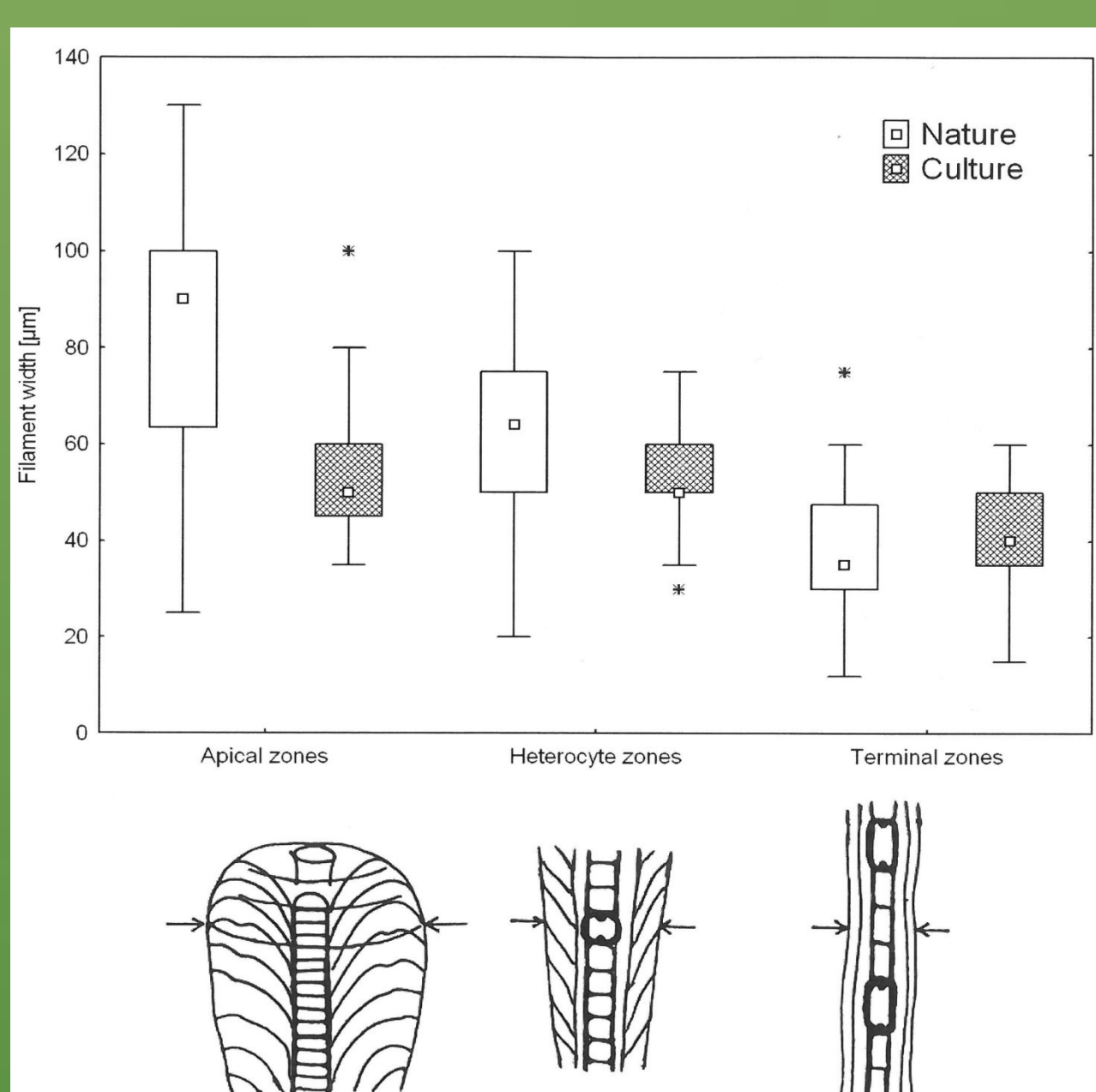
Materials and methods

Samples used for the study were taken from wet calcareous wall located in Piecky Gorge in the National Park Slovenský raj (Slovakia) by random scraping of the rock surface into sterile tubes, irregularly during summer seasons 1998 - 2007. At the same time, relative humidity and temperature were measured using a thermo-hygrometer and irradiance values using a radiometer with quantum sensor.

Prokaryotes taken from the nature were cultured in Petri dishes and tubes with BBM medium under controlled conditions (21 °C, humidity 60%, irradiance 10.3 μ m.s⁻¹.m⁻², light 660 lx, Phar 2.14 W.m⁻² and in a photoperiod 12:12 light:dark).

Both natural isolates and isolates from the culture were observed and documented using a stereomicroscope Olympus SZH and Olympus BX 50 light microscope equipped with Lucia Image Analysis.

After the initial cultivation, the clones were isolated. At both isolates from nature and culture (60 each) the morphometric data were measured: width of filament apex, width of the first heterocyte in the heterocyte zone and width of the terminal zone. These data were then statistically visualised using STATISTICA (Statsoft®) program. The dimensions of cells of both types of isolates were compared using Wilcoxon Signed Ranks Test.



Wilcoxon Signed Ranks Test
 significant value 0.05;
 Apical zones:
 Z = -5,62,
 P_{apex} < 0.0001;
 Heterocyte zones:
 Z = -3,16,
 P_{heterocyte} = 0.0016;
 Terminal zones:
 Z = -2,09,
 P_{terminal} = 0.0370



Fig. 5: *Petalonema alatum* – the queen of cyanobacteria. Orig. B. Uher.

Acknowledgement

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Discussion

The previous and recent taxonomical position of *Petalonema* is based only in morphological features. The oldest descriptions focused basically only on the structure types of trichome sheaths and filaments. Other features, such as presence of heterocytes or meristematic zones, were found crucial later.

Phycologist of 20th century represented different approaches to the taxonomy and also considered different features to be diagnostic for *Petalonema*; three main streams can be recognized. The most important is the first group, based on Berkeley's description of *Petalonema alatum* as separate taxon, but also among these people there were differences. For example Berkeley stressed the branching, filament lined by lamella, and envelope structure, on the other side, Komárek & Anagnostidis (1989) stressed the type of heterocytes and branching, apex of filament and structured sheath. This moved *Petalonema* to the family *Microchaetaceae*.

This taxonomical position seems to be correct and better than other approaches. Also presented results of this study confirms that the morphological variability inside the species is very high and there is no reason why to put *Petalonema* inside other species.

In the only molecular study, Taton et al. (2006) didn't include many *Microchaetaceae* species in the tree, and so *Petalonema* appeared in the *Nostoc* clade, which it probably doesn't belong to.

Conclusion

The morphological part of the study confirmed the validity of genus *Petalonema* and the differentiated it from other genera it used to belong to, such as *Scytonema* or *Tolypothrix*. Now the molecular improvement of these results is needed. New samples will be collected in Slovakia and other places in Europe, rRNA and *rbcl* genes will be sequenced, so that we will be able to place *P. alatum* more accurately in the phylogeny tree of cyanobacteria.

References

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 Taton, A., Grubisic, S., Ertz, D., Hodgson, D., Piccardi, R., Biondi, N., Tredici, M., Mainini, M., Losi, D., Marinelli, F., Wilmette, A. (2006): Polyphasic study of Antarctic cyanobacterial strains. J. Phycol. 42: 1257–1270.

Fig. 3: Comparison of filament measurements of *Petalonema alatum* – in the nature vs. culture, * outliers, median, \top non outlier, box border quantil 25% and 75%