Phylogeny of Aspergillus and Penicillium species: — engaging undergraduate students in science

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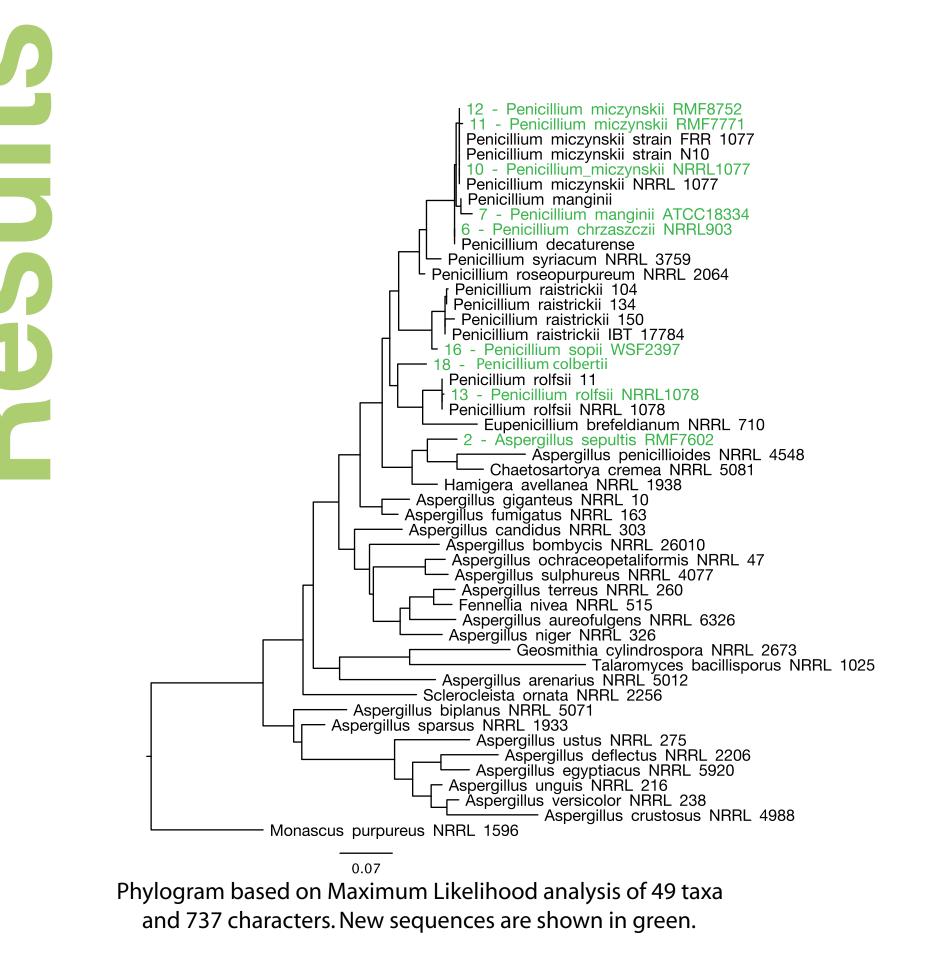


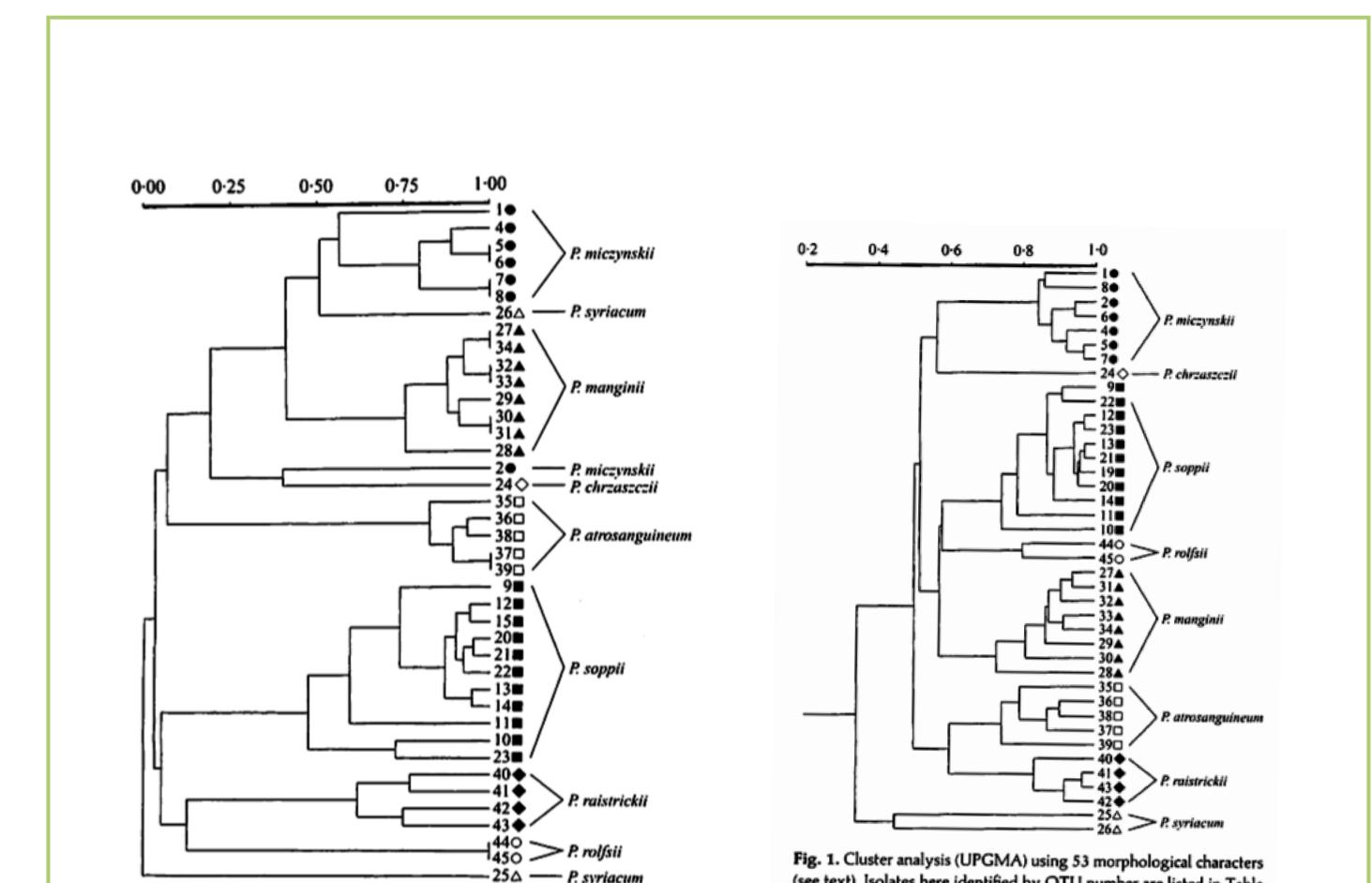
Teaching

While students in many fields experience the life of the profession for which they are headed, biology students are expected to take a leap of faith into their careers without any real-life exposure. **Comgen** is an NSF-funded project that provides biology students a mini-graduate school experience, including the frustrations and delights of scientific discovery.

Research

Over her career, Martha Christensen has collected new and diverse soil fungi from around the world with a special focus on their ecology and ecosystem roles. Her fungal collection <moldsforyou.org> is an international treasure including over 2,200 specimens.





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CZ.

This project is aimed at classifying fungi from this collection with the latest molecular techniques and introducing the next generation of students to microbial diversity. Our preliminary study focuses on the phylogeny of Aspergillus and Penicillium species, whose impact ranges from life-saving to lethal. We initially seek to compare phylogenies suggested by ITS rDNA sequences to those based on morphological or biochemical characters.

Teaching

Self-directed learning Prerequisite only a basic biology course Socratic method with almost no lectures Students collaborate on learning concepts and techniques Connect new protocols to underlying theories Real-world lab product assessment

Original Research

Authentic problems Unknown answers

Lab notebook as documentation

Our initial comparison of a maximum likelihood analysis of ITS rDNA with Christensen et al.'s we found that our DNA data confirms the general groupings of taxa in these trees. We will next expand our dataset to include sequences from other genes (Beta-tubulin introns and the LSU D1 and D2 loop regions), and extend our statistical analysis.

Fig. 1. Cluster analysis (UPGMA) using 53 morphological characters (see text). Isolates here identified by OTU number are listed in Table 1. Scale is Sørensen – Dice similarity. Cophenetic correlation value = 0.88.

Fig. 2. Cluster analysis (UPGMA) using 42 secondary metabolite characters. Cophenetic correlation value = 0.96.

Martha Christensen, Jens Frisvad, and Dorothy Tuthill. 1999. Mycological Research 103: 527-541.

Taxonomy of the *Penicillium miczynskii* group based on morphology and secondary metabolites.

"Science education should focus less on what instructors 'cover,' and more on what students learn and how well they can use their knowledge; less on vocabulary and facts that students memorize, and more on students' understanding of scientific concepts and how these concepts fit together in a **framework of knowledge** about a subject; less on what students can repeat back immediately in class, and more on their long-term retention and ability to **transfer knowledge** to contexts outside the classroom" (DeHaan, 2005)



Fungal Cultures

| \mathbf{O} | Isolate | Species |
|--------------|-----------|---------------------|
| | RMF7602 | Aspergillus sepul |
| | NRRL903 | Penicillium chras |
| | ATCC18334 | Penicillium mang |
| | NRR1077 | Penicillium miczy |
| | RMF7771 | Penicillium miczy |
| U | RMF8752 | Penicillium miczy |
| | NRR1078 | Penicillium rolfsii |
| | WSF2397 | Penicillium sopii |
| | | |

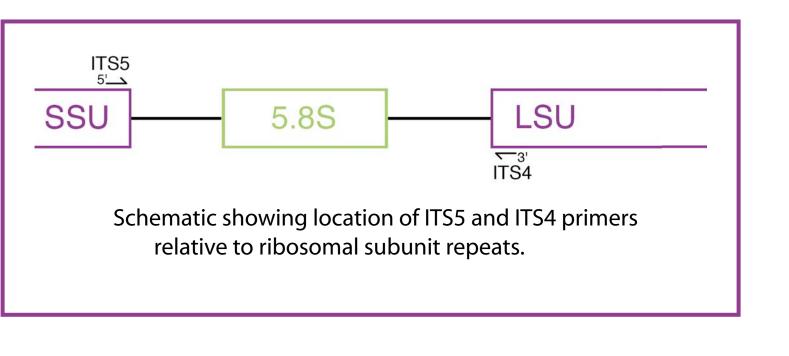
Geographical Location spergillus sepultis Ida County, Iowa USA enicillium chraszcii Picea forest soil, Poland enicillium manginii Fagus mycorrhizae, Italy enicillium miczynskii Conifer forest soil, Poland Pinus contorta forest soil, Wyoming USA 197 enicillium miczynskii Conifer forest soil, Oregon USA enicillium miczynskii enicillium rolfsii Pineapple fruit, Florida USA

Temperate forest, Wisconsin USA

| Isolation Date | C |
|----------------|---|
| 1984 | |
| 1927 | |
| 1963 | |
| 1927 | |
| 1977 | |
| 1988 | |
| | |

PCR

Bio-Rad Mycycler Program: 95°C 2 minutes 36 cycles: 95°C 30 seconds 50°C 30 seconds 72°C 1 minute Hold at 4°C



Final extension, 72°C 10 minutes

Through this project, students with minimal experience in molecular biology, and no experience in mycology or phylogenetics, gained confidence in their abilities as independent researchers, and are emerging as self-directed learners comfortable with ambiguity and able to handle the ever-increasing amount of knowledge.

PCR clean-up

Qiagen[®] QIAquick PCR Purification KitTM

PCR (sequencing)

Applied Biosystems XTerminator[®] Kit Program:

96°C 4 minutes 25 cycles: 96°C 10 seconds 50°C 5 seconds 60°C 4 minutes Hold at 4°C

DNA Sequencing

Applied Biosystems 3130 Genetic Analyzer

Sequence Analysis

BLASTn Sequences from GenBank, Alignment in Se-Al, Maximum Likelihood analysis Garli 0.96b8

"We have gone from simply following the protocols to adding to them or changing them in ways that helped our experiment."

1905

1960

"I never thought frustration could be fun."

Our teaching methods have wide applicability for fostering student interest in science. We are currently developing our teaching methods for dissemination to other educators.

Our goal is transformation of not just the classroom, but also the teaching experience.

"Overall we are more comfortable with doing original research — we can read papers, we can do sequence searches, we can do phylogenetic analyses."

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