Isolation of microfungi from Arctic and Antarctic soils and their identification using ITS, LSU and SSU sequences.

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Abstract

This study presents the biodiversity results of microfungi of soils from Arctic and Antarctica. The Arctic soil samples were collected from Hornsund, whereas the Antarctic soils were sampled from King George Island. Warcup's soil plating method was used for fungal cultivation. Fungal species were identified based on ITS, LSU and SSU sequences using BLAST searches and phylogenetic analysis. Based on the ITS barcoding sequences, fungal species *Cosmospora* sp., *Tolypocladium inflatum* (*Elaphocordyceps subsessilis*), *Isaria farinosa* (*Paecilomyces* farinosus), *Oidiodendron truncatum*, *Aspergillus pseudodeflectus*, *Preussia borealis*, *Thelebolus* sp., and *Phialocephala* sp. were identified only from the Arctic soils. Meanwhile, *Penicillium* sp., *Geomyces* sp., *Guehomyces pullulans* were isolated from both the Arctic and Antarctic soils. *Leuconeurospora* sp. and *Pseudeurotium* sp. were isolated only from the Antarctic soils. Although there were limited barcoded species using LSU and SSU sequences in GenBank, some of the species identification corresponded with those of ITS sequences.

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Outline of talk

- Malaysia's involvement in polar research
- Diversity of soil microfungi from polar regions
- Barcoding of Fungi: ITS gene and other genetic markers
- This study aims to
 - Isolate microfungi from Arctic and Antarctic soil samples
 - Identify microfungi species using ITS, LSU and SSU sequences
- Materials & Method
- Results & Discussion
- Conclusion

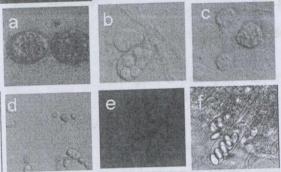


Malaysia's involvement in polar research

- Malaysian Antarctic Research Programme (MARP) was initiated in November 1997.
- First Malaysian scientific expedition to Antarctica was successfully held in October 1999. Malaysia joined the ATCM in 2011.
- The National Antarctic Research Centre to coordinate the research activities of the members of the programme.
- In the year 2006, MARP has extended the interest to Arctic. At present, three scientific expeditions has been carried out in Arctic.

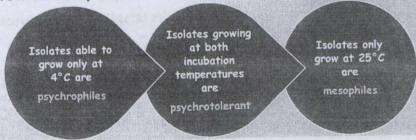


Diversity of soil microfungi from polar

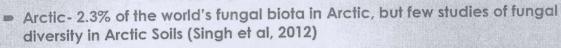


- a) Mortierella sp. b) Antarctomyces sp. 4 c) Mrakia sp. d) Yeast sp. 16
- e) Aureobasidium sp. f) Thelebolus sp.

- Antarctica studies mostly in the continental rather than the maritime Antarctic.
- Easily accessible, benign locations and environmentally challenging areas such as the Victoria Land Dry Valleys (Friedmann et al. 1985), Beaufort Island (Alias et al. 2013).
- Occurrence of fungi associated with human associated habitats and artifacts have been studied extensively.
- >80% belong to the Ascomycota, mostly perithecial with few discomycetes



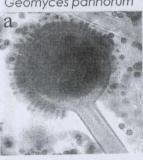
Diversity of soil microfungi from polar regions



- Diversity of fungi in soils of Bellsund, Svalbard, has been studied (Kurek et al., 2007), and new genera and species have been described from the region (Pang et al., 2008, 2009)
- Singh et al 2012 identified 19 species (14 genera) from Ny-Ålesund, Spitsbergen soils.
- Alias and Suhaila (2007) reported 89 microfungal taxa from the soil of Ny-Alesund
- Pang et al. (2009, 2011) reported six marine fungi isolated from wood debris collected at Longyearbyen.
- Hornsund, Spitsbergen ornithogenic influenced soils showed the highest species diversity (eg. Mortierella macrocystis, M. elongata, Mortierella sp., Cudoniella sp., Varicosporium elodeae, Beauveria bassiana, Geomyces pannorum, Penicillium sp. and Atradidymella muscivora) – (Ali et al 2014).



Geomyces pannorum



Aspergillus aculeatus

(photos taken from Singh et al, 2012)

Barcoding of Fungi: ITS region and other genetic markers

- Universal DNA barcode marker for Fungi internal transcribed spacer (ITS) region.
- High PCR amplification and sequencing success, broadest range of fungi, defined barcode gap between inter- and intraspecific variation.
- 28S nuclear ribosomal large subunit rRNA gene (LSU)
- 18S nuclear ribosomal small subunit rRNA gene (SSU)
- Subunits of RNA polymerase II (RPB1, RPB2)
- Mini chromosome maintenance protein (MCM7)

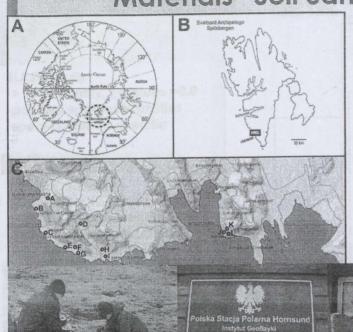
Used in fungi multigene phylogenies

This study aims to

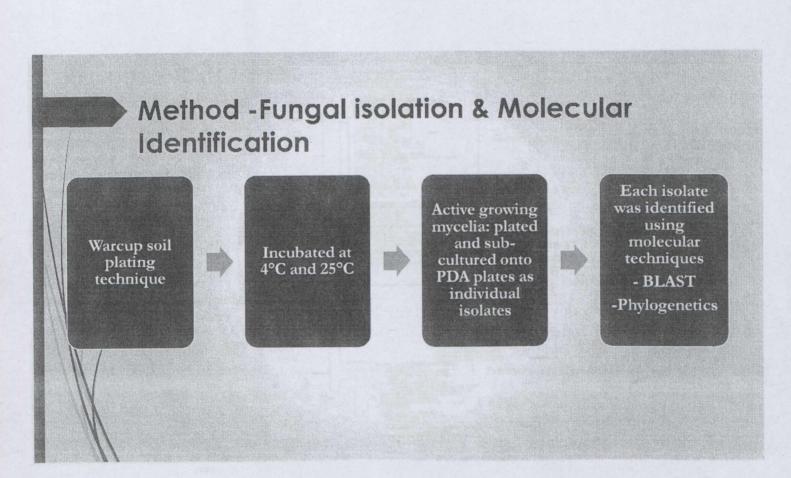
- Isolate microfungi from Arctic and Antarctic soil samples
- ►Identify microfungi species using ITS, LSU and SSU sequences

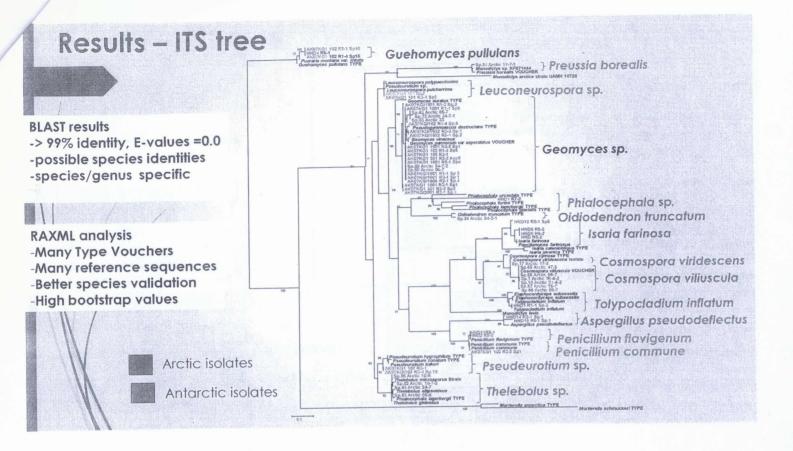
Materials – Soil Samples King George Island (Antarctica) Austral summer 2006/2007 Penguin rookery Pristine area Human impacted area Ornithogenically-influenced vegetated site

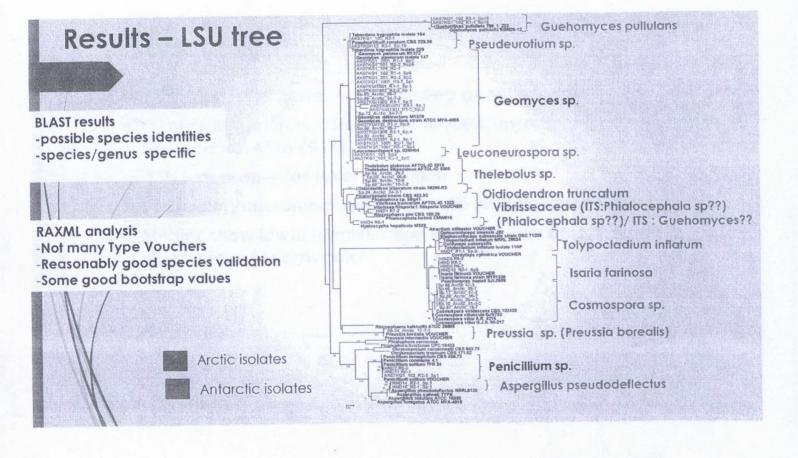
Materials -Soil Samples

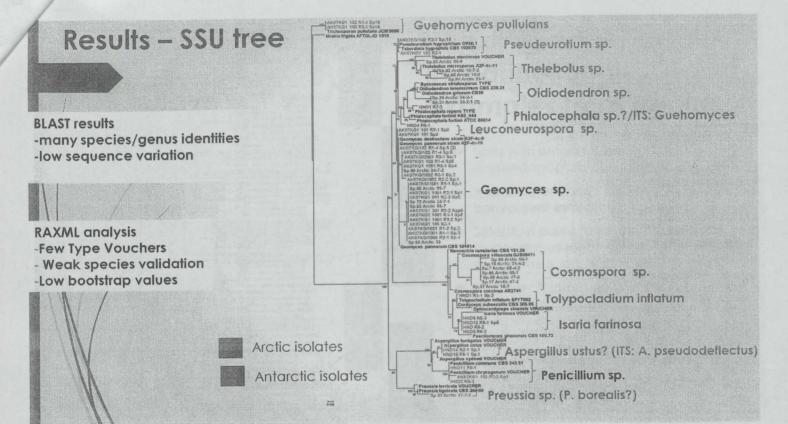


- Hornsund, Spitsbergen (Arctic)
- **⇒** Boreal summer (August 2010)
- Dry and bare fellfields
- Moist moss tundra,
- · Ornithogenic sites,
- Vertebrate-influenced pond shore
- · Glacier foreland



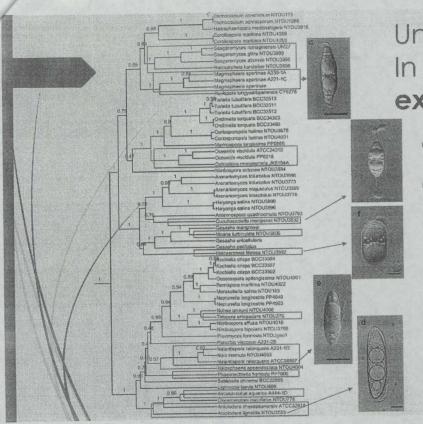






Discussion

- ITS region the preferred barcoding gene in fungi
- Primary fungal barcode marker to the Consortium for the Barcode of Life (Schoch et al 2012).
- SSU poor species level resolution
- Soil (polar) microfungi sequences still lacking in GenBank
- Studies show lower identification success can be expected in filamentous Ascomycota



Unfurling appendages In marine fungiexamples from tropics

- Taxa with unfurling ascospore appendages are polyphyletic
- A character possibly gained through convergent evolution
- Selection pressure for the evolution of appendages for attachment to substrata due to the scarcity of substrata in the sea
- May be an intermediate form between unappendaged ascospores to those with complex appendages

Multigene phylogeny of marine fungi -Taxonomy/Taxonomists is needed to avoid misidentification!!

Conclusion

- Vast majority of fungal species remains unknown.
- > 90% of Fungi awaits discovery from different environments
- Changes in Fungal Nomenclature
- Coordination in Field Sampling, Traditional Taxonomy & Barcoding

THANK YOU