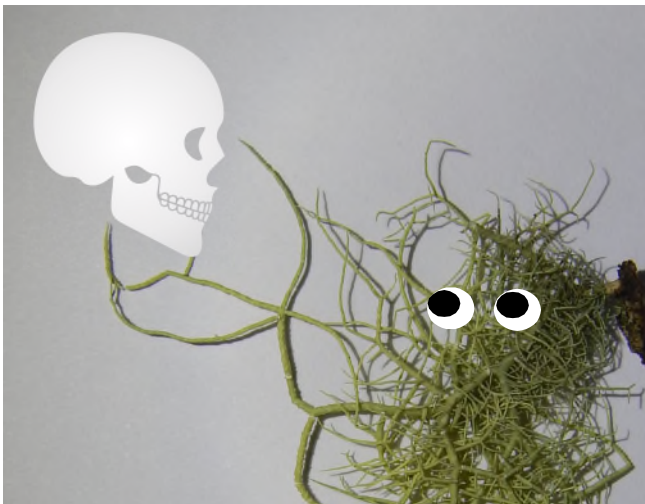


~the 7th Sweden-Japan Academic Network seminar~

To be or not to be a lichen?

Using genomics to interpret lifestyles in fungi

Mieko Kono



The Royal Swedish Academy of Sciences
01.02.2019

~ Aug. 2018



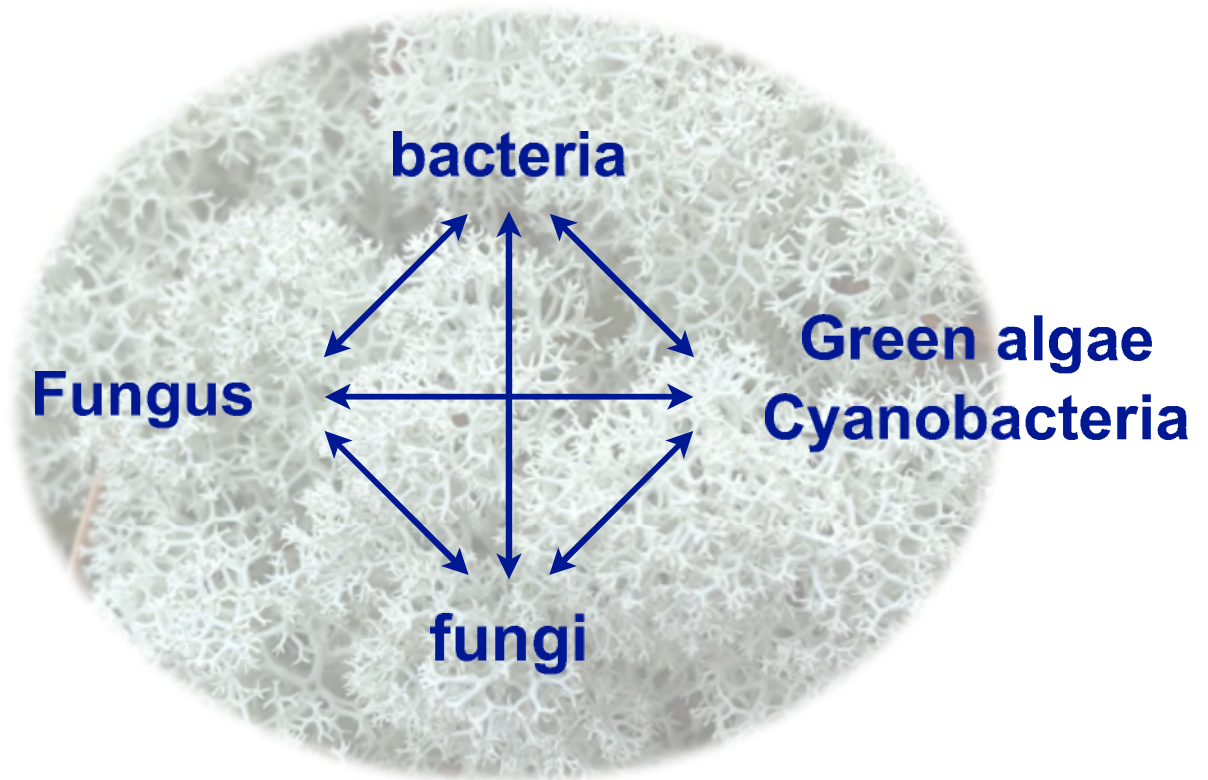
Sept. 2018~



SOKENDAI S O K E N D A I
The Graduate University for Advanced Studies



What are lichens?

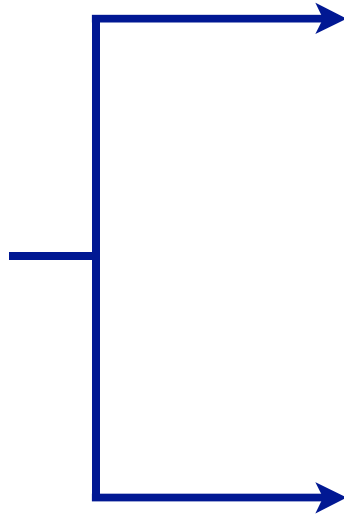


How are they genetically organized?

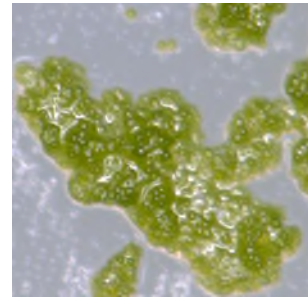
Symbiosis-specific genes?



Usnea hakonensis



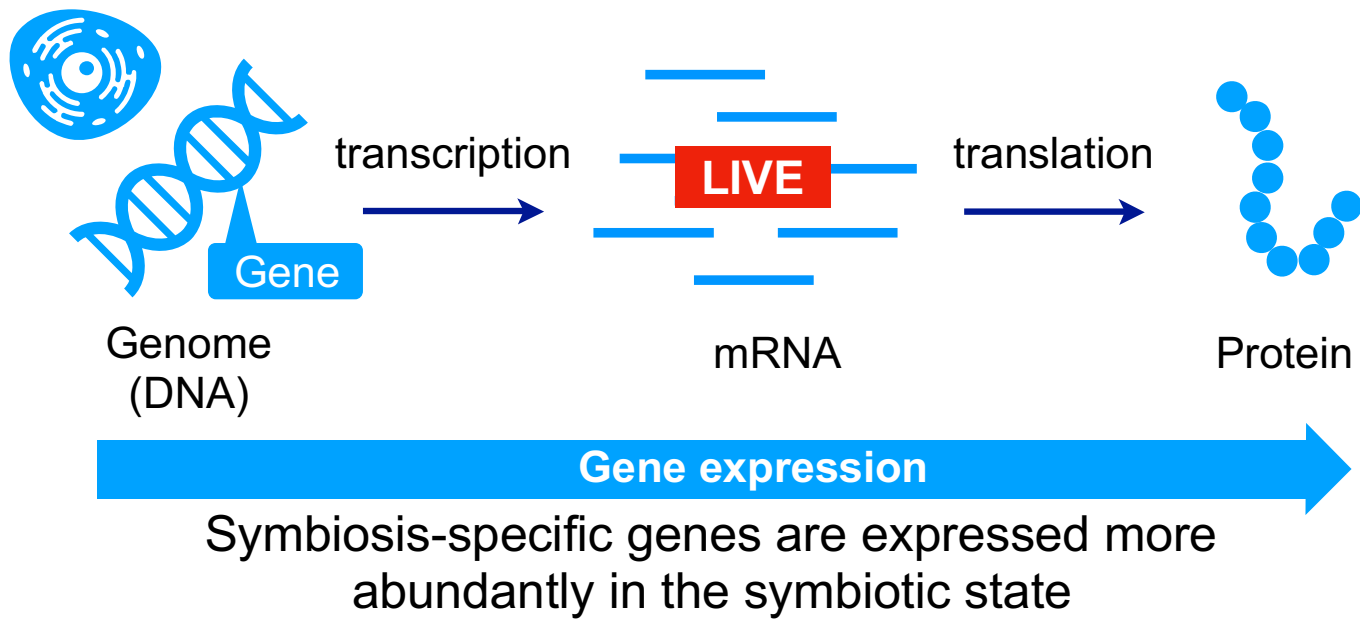
Fungus



Alga

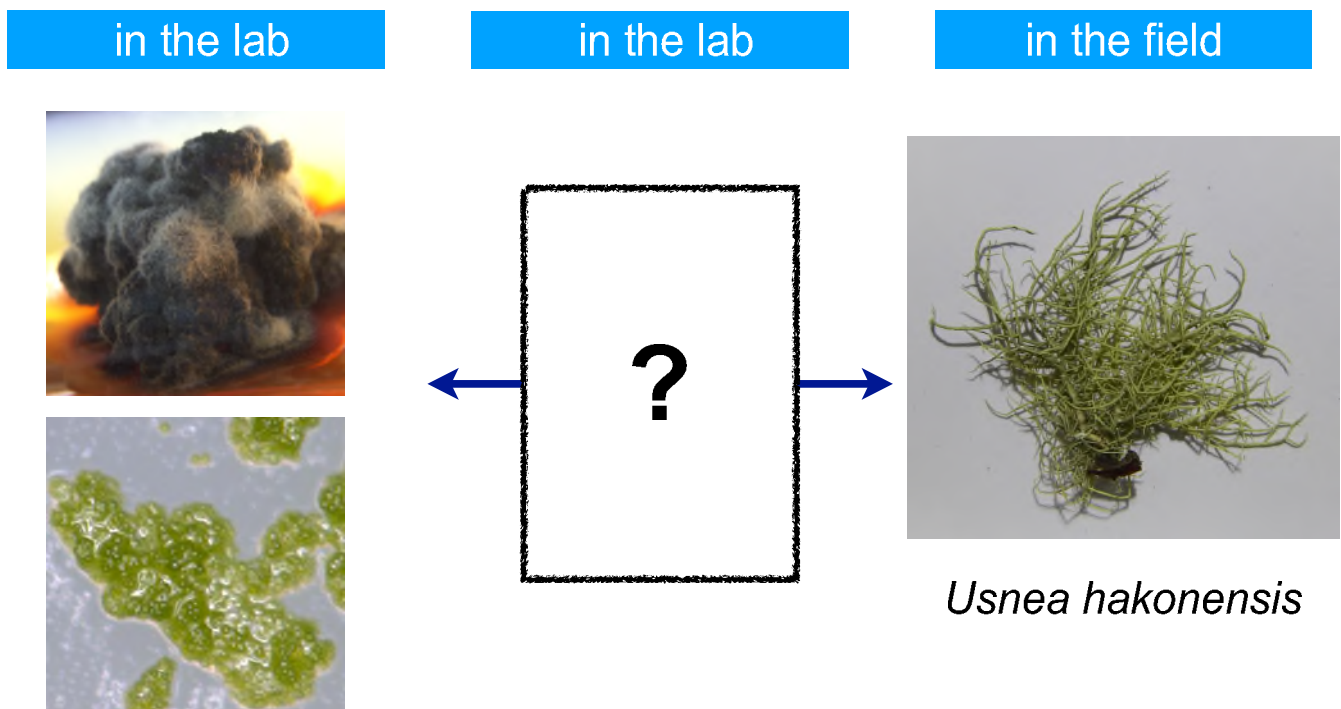
Symbiotic lifestyle requires morphological, physiological, biochemical changes

How can we identify symbiosis-specific genes?



Compare the gene expression between **before** and **after** the symbiosis

Compare the gene expression between...?



Thalli in the field are exposed to a complex environment distinct from the culture condition in the lab.

Can we reproduce lichens in the lab?

Stage 1: Pre-contact

Stage 2: Contact

Stage 3: Growth together

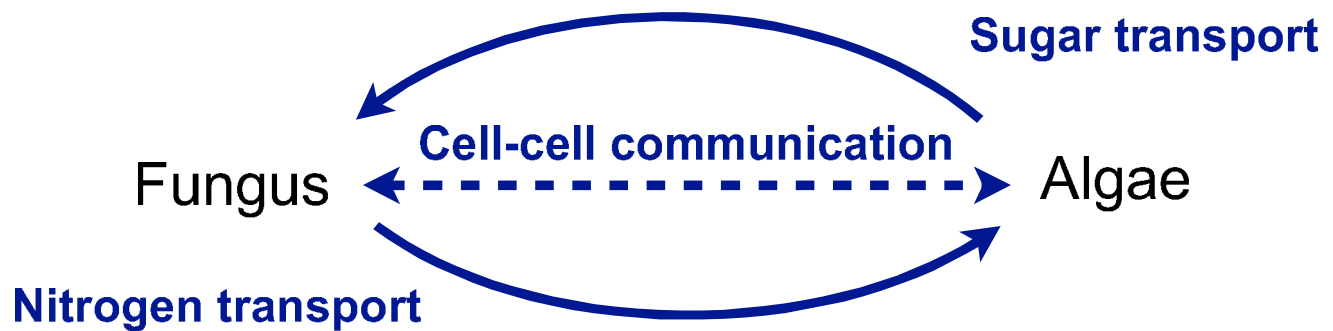
Stage 4: Incorporation of symbionts into a
common matrix

Stage 5: Thallus differentiation

Many attempts have been made but...

Predicted functions of symbiosis-specific genes

Stage 1 & 2	<i>Cladonia grayi</i>	Joneson <i>et al.</i> (2011) <i>Mycologia</i>
Stage 1	<i>Endocarpon pusillum</i>	Wang <i>et al.</i> (2014) <i>BMC Genomics</i>



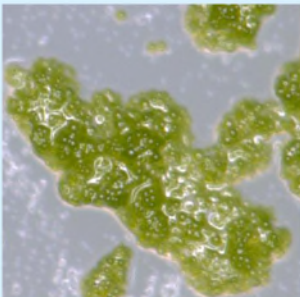
What kind of genes are expressed in the later stages and in the natural habitat?

Symbiotic model of *Usnea hakonensis*

in the lab



Usnea hakonensis



Trebouxia sp.



Stage 5

Thallus differentiation

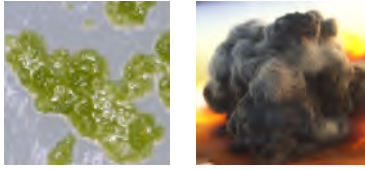


Usnea hakonensis

Established by Dr. Yoshiaki Kon

Searching for symbiosis-specific genes by using the *Usnea hakonensis* model

Independent culture



Reproduced



Natural



**Symbiosis-
specific genes**

Dr. Yoshihito Ohmura

Dr. Yoshiaki Kon

Dr. Yoko Satta

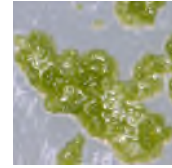
Dr. Yohey Terai

Whole genome sequencing

Fungus



Alga



	<i>U. hakovensis</i>	<i>Trebouxia sp.</i>
Genome size	41 Mb	69 Mb
Gene number	21,105	21,207



3 Gb = 3000 Mb

human

138 Mb



fruit fly

12 Mb



baker's yeast

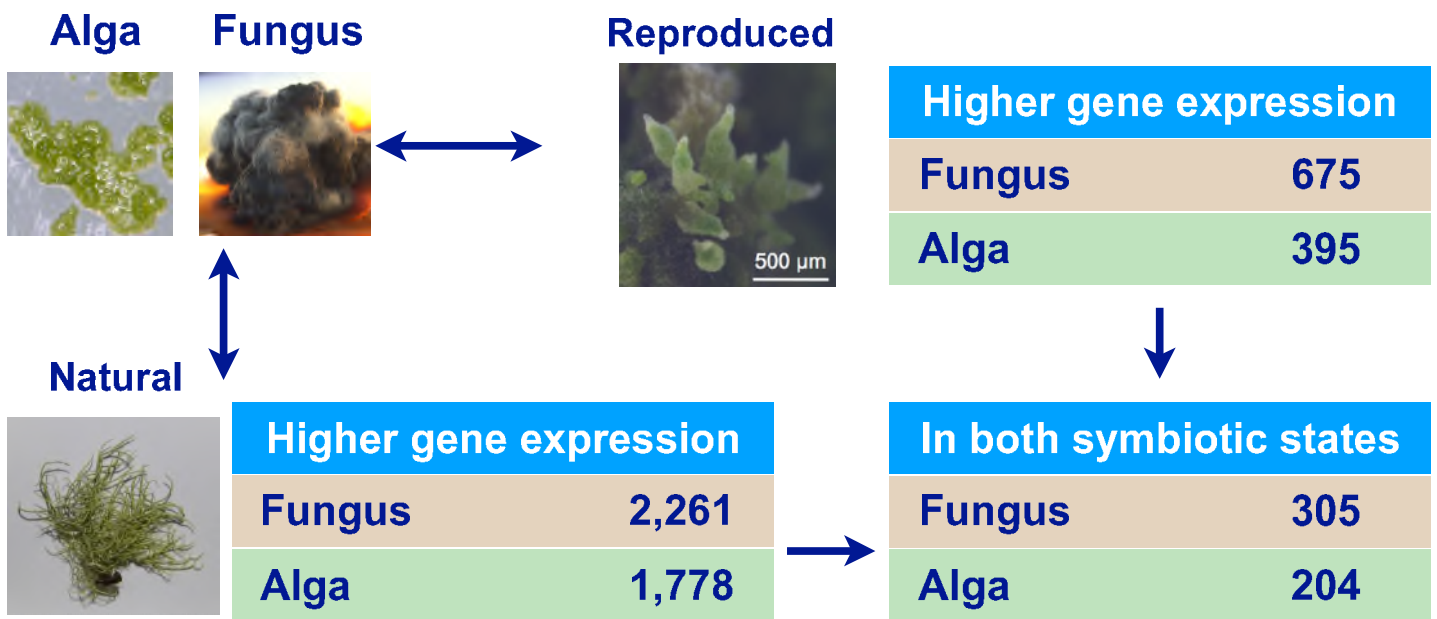


1 base pairs (bp)

1 kb = 1000 bp

1 Mb = 1000000 bp

Symbiosis-specific genes

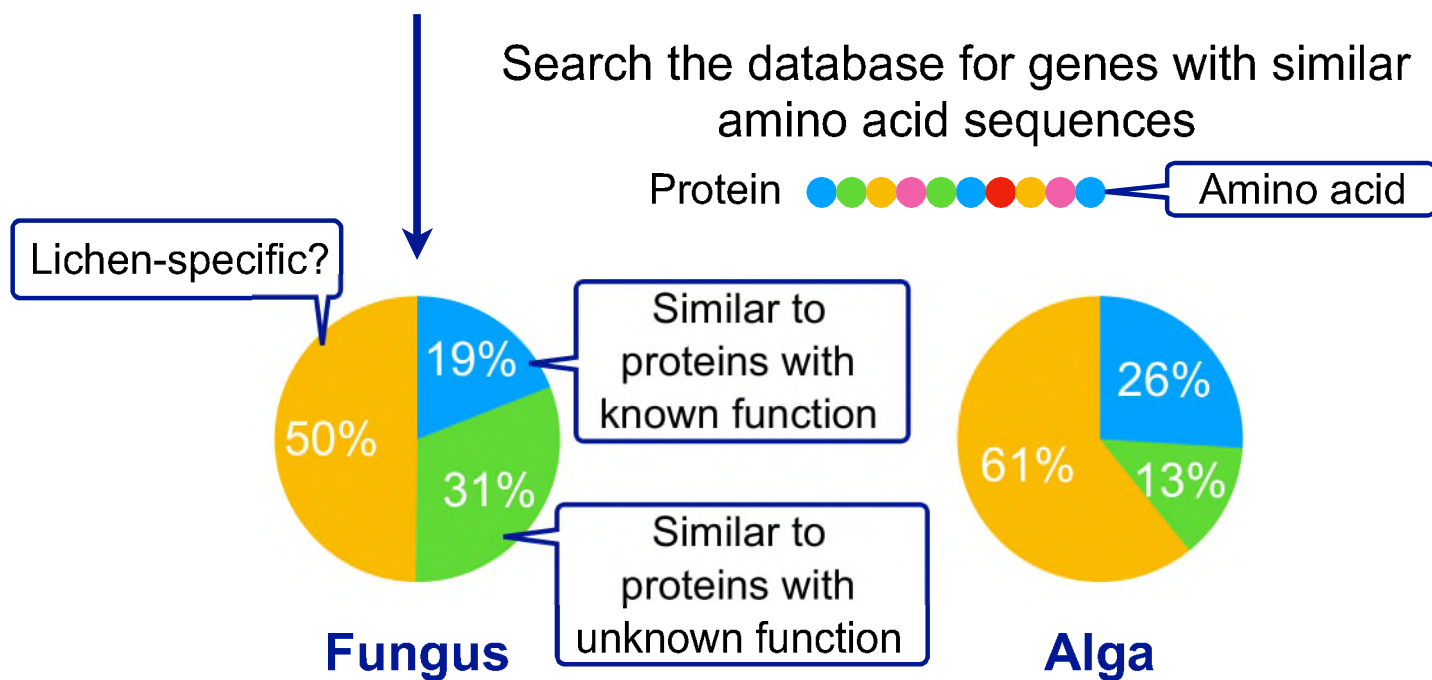


305 fungal and 204 algal genes were identified as “symbiosis-specific” genes

Functions of symbiosis-specific genes

Symbiosis-specific genes	
Fungus	305
Alga	204

Are there similar genes with known function?



Perspectives

- Identify the role of the bacterium in the symbiosis.
- Identify other symbiotic partners required to reproduce a lichen thallus in the lab.

How did lichens evolve?

Is there an universal set of symbiosis-specific genes?

Lichens or Saprotrophs or Both?

Stictidaceae

Closely related species with various lifestyles

What is the genetic background of each lifestyle?

Compare the genetic features among the three fungal species in Stictidaceae

Optional lichenization

Schizoxylon albescens

Choose the lifestyle depending on the environment (substrate)

Option 1: Lichenized style

In and around cracks in the bark

Associated with algae that
provide nutrients

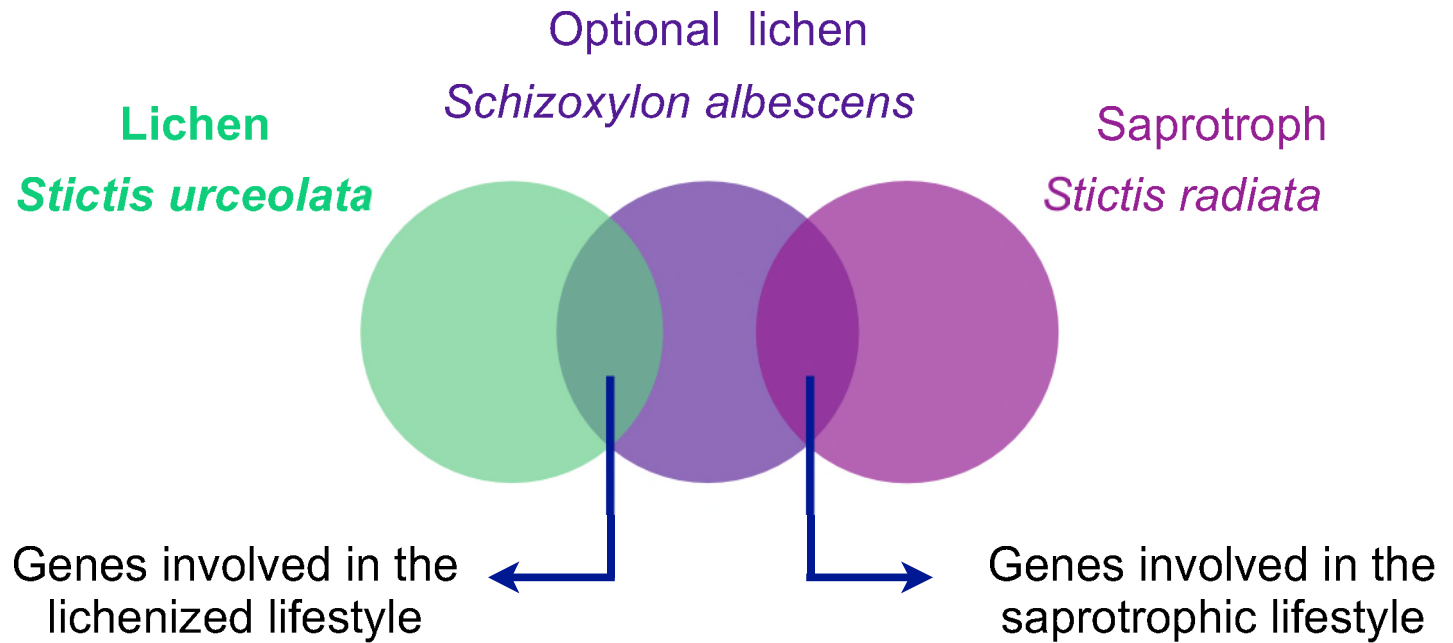
Option 2: Saprotrophic style

On dead, decorticated branches

No algae!
Exploit the substrates for
nutrients

Identify genes related to each lifestyle

Step 1 Compare the genomic structures



Identify genes related to each lifestyle

Step 2 Investigate the gene expression of *Schizoxylon albescens* in different lifestyles

Lichenized lifestyle ↔ **Saprotrophic lifestyle**

Genes involved in the fungal-algal recognition and physiological interaction.

Genes involved in the exploitation of substrate.

Our goal

Identify the genes that involve in the choices of lifestyles in fungi

Understand lichenization in the context of fungal evolution

How far have we gotten?

Culturing the fungi in the laboratory

Cut out one fruiting body



Culture 1 month on an agar plate



Transfer a fungal colony to a new plate



1~2 months

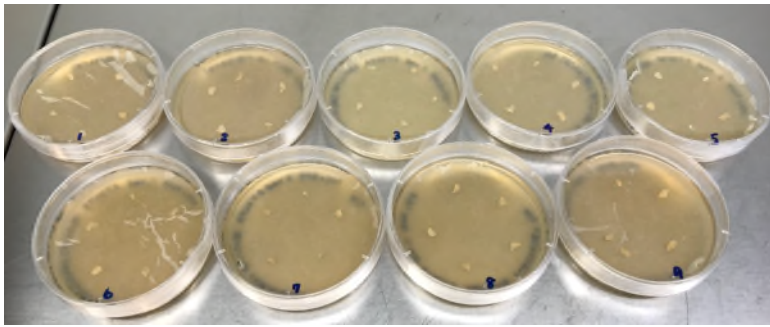
READY!

Stictis radiata

Schizoxylon albescens



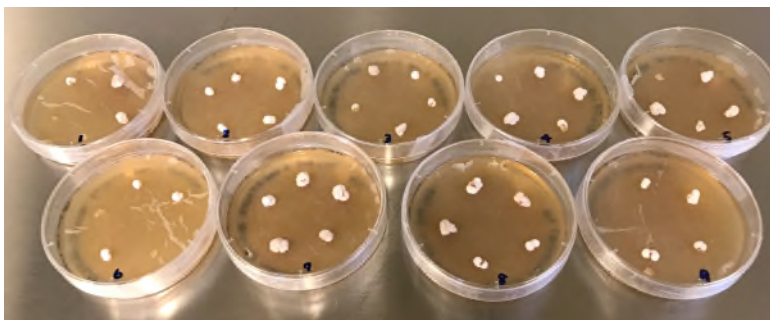
Preparing DNA for the whole genome sequencing



Stictis radiata



1 month later



DNA extraction

10 ~ 15 μg
required

ca. 100 ng /colony
at least 100 colonies

Growing fastest ever!

Acknowledgements

Work in Japan

Dr. Yoshiaki Kon Tokyo Metropolitan Hitotsubashi High School

Dr. Yoshihito Ohmura National Museum of Nature and Science

Dr. Yohey Terai

Dr. Yoko Satta

SOKENDAI (The Graduate University for Advanced Studies)

This research was granted by:

Sasagawa

Thank you for your attention

Work in Sweden

Professor Mats Wedin

Bodil Cronholm

Swedish Museum of Natural History

Professor Göran Thor Swedish University of Agricultural Sciences

Dr. Lucia Muggia University of Trieste

This research was granted by:

Swedish Research Council (Vetenskapsrådet) through the project “Fungal phylogeny and evolution”, grant VR 2016-03589

