

Population differentiation of the protist Synura (Chrysophyceae, Stramenopiles)



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Introduction

- driving speciation • Factors are still enigmatic, especially among protists.
- •In macrobes, geographical barrier was considered as the most important factor.
- •On the contrary, microbes are often thought to have unlimited dispersal, therefore speciation would takes place in sympatry.

Goals

- •To infer population structure of Synura petersenii using RADseq
- evaluate •To importance Of geographical and ecological factors shaping the population structure using environmental data and gradient experiments with cultured strains

Sample map



Phylogeny IQTREE, 85626 positions

Bioclimatic variables

Max growth rate - temperature



- Strains from the same regions can prefer different values of conductivity.
- Conductivity was measured at the time





- Ancestral state reconstruction of maximal growth rate of cultures in a gradient of temperature.
- •Two groups first adapted to all temperatures and second adapted to moderate and high.

of sampling.

Structure – population structure



- Similar pattern revealed for both assumed numbers of populations (K).
- Strains were mostly separated into two populations.
- Few strains had mixed origin, which suggests a possible gene flow among them.

Conclusions

• There are two major groups within *S. petersenii*

Variation partitioning - sampling site data

• How much of a variability is explained by each factor or together?



Methods

• Group 1 – Czechia and Sweden and group 2 – France, Canada and Norway

•Both groups contain samples with high and low conductivity. Variation partitioning showed that climate and geography are the most important. Gradient experiments with cultures revealed that group 1 is adapted to all temperatures, while group 2 is adapted and moderate to high temperatures.

•Organism – Synura petersenii (Chrysophyceae, Stramenopiles) •Sampling, environmental condition measurement – conductivity, pH, temperature • Culturing in gradient of conductivity and temperature – growth rate. RADseq Phylogeny, Structure and ancestral state reconstruction