## From genes to ecosystems

Integrating genetic diversity to study freshwater ecosystems under stress





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## Background



Time

## Background

#### We are using molecular genetics to

- ...assess species diversity with greater precision
- ...understand patterns of gene flow
- ...disentangle the roles of random genetic drift and selection in speciation / adaptation processes
  - ...assess the effects of stressors on the genetic diversity of species





## **Genetic diversity**



Genetic Diversity and Connectivity

**DNA Barcoding** 

ddRAD Sequencing

Genome Sequencing





Hannah Schweyen,



Jan Macher

Martina Weiss,

## **Genetic diversity**



## **Genetic diversity and connectivity**





Hohenlohe et al. 2010

## **Genetic diversity and connectivity**

"Next generation sequencing" (NGS) offers new tools for analyzing loci across the genome even in non-model species

RAD-seq (Restriction-site associated DNA sequencing)



## **RAD** sequencing

- Population genetic analyses of poorly studied freshwater insect species with:
  - a) one mitochondrial gene (CO1)
    - (658 bp Folmer region)
  - b) thousands of RAD-loci (few spec.)
    - ddRAD (1.9 million reads per specimen on average)
    - >13,000 loci (>16,000 SNPs)

Phylogenetic networks + Permutation tests Hydropsychidae orientalis & Cheumatopsyche brevilineata (Trichoptera)



# Yasu River Sampling Sites

# Kiso River Sampling Sites

12

Kiso Biological Station

12

## Genetic diversity and connectivity







#### Deleatidium (Ephemeroptera)

#### Thremma gallicum (Trichoptera)



# Genetic diversity and connectivity : Summary

Population genomic approaches (RAD) provide the necessary resolution to study genetic diversity and connectivity in "natural" and stressed environments

Few specimens needed

Costs per basepair very low







# From genes to ecosystems

# What is the benefit of integrating the level of genetic diversity to study freshwater ecosystems under stress?

- Faster, more complete and reliable identification of true species diversity in assessment / conservation programs
- . DNA Metabarcoding for assessments: Possible and affordable!
- . Evaluation of the impact of stressors on all levels (loss of genotypes)
- Linking ecosystem functioning to individual genotypes provides unique insights into ecosystem processes

## Thank you!

