# Next-generation population genomics of native and aquacultured Mediterranean mussel, *Mytilus galloprovincialis*

Anamaria Štambuk





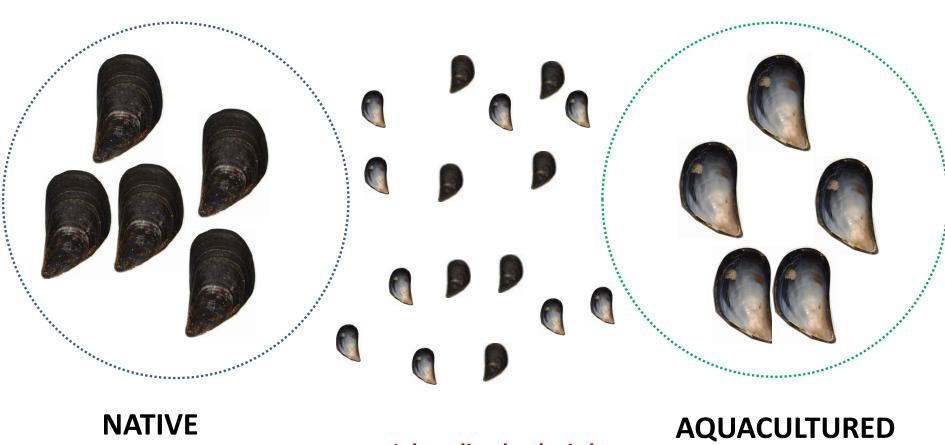
"Genotyping by sequencing: new tool for population genomics of native and aquacultured Mediterranean mussel, *Mytilus galloprovincialis*"

"The effects of pollution on rapid evolution and ecological change in the Mediterranean mussel (Mytilus galloprovincialis)"

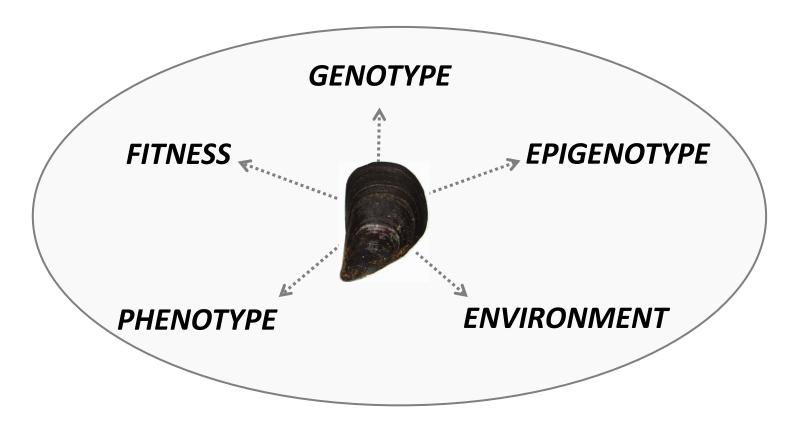




# HIGH CONNECTIVITY BETWEEN NATIVE AND AQUACULTURED POPULATIONS



gametes → long lived pelagic larvae



NGS – GENOMICS AND EPIGENOMICS FITNESS AND MORPHOMETRY GWAS

#### **ENVIRONMENTAL VARIABLES**

- Native and aquacultured populations
- Transplant and mesocosms experiments



### **AQUACULTURE PRACTICES**

- 1. Collection of natural spats mooring ropes
- 2. Nylon nets
- a) cotton nets



b) anti-predator protection nets

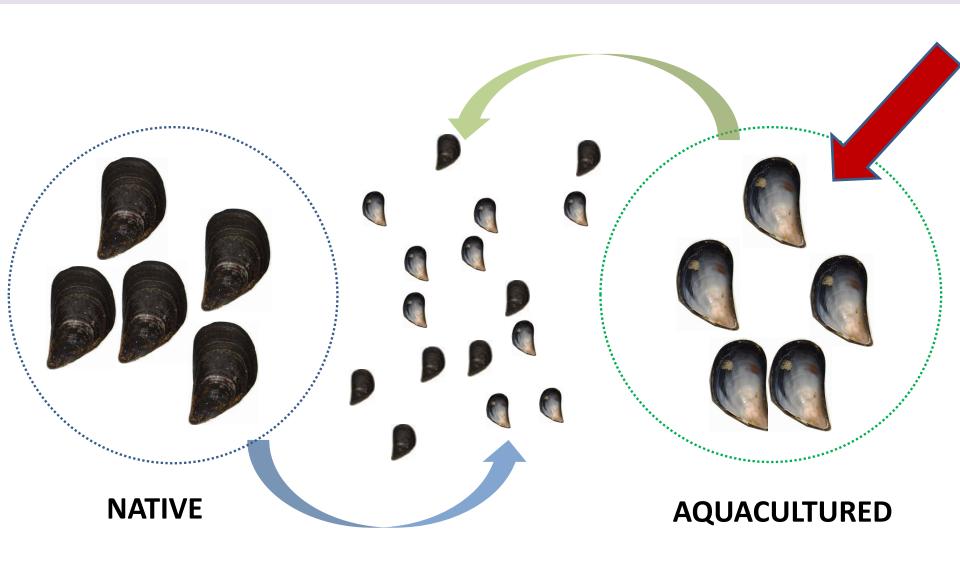


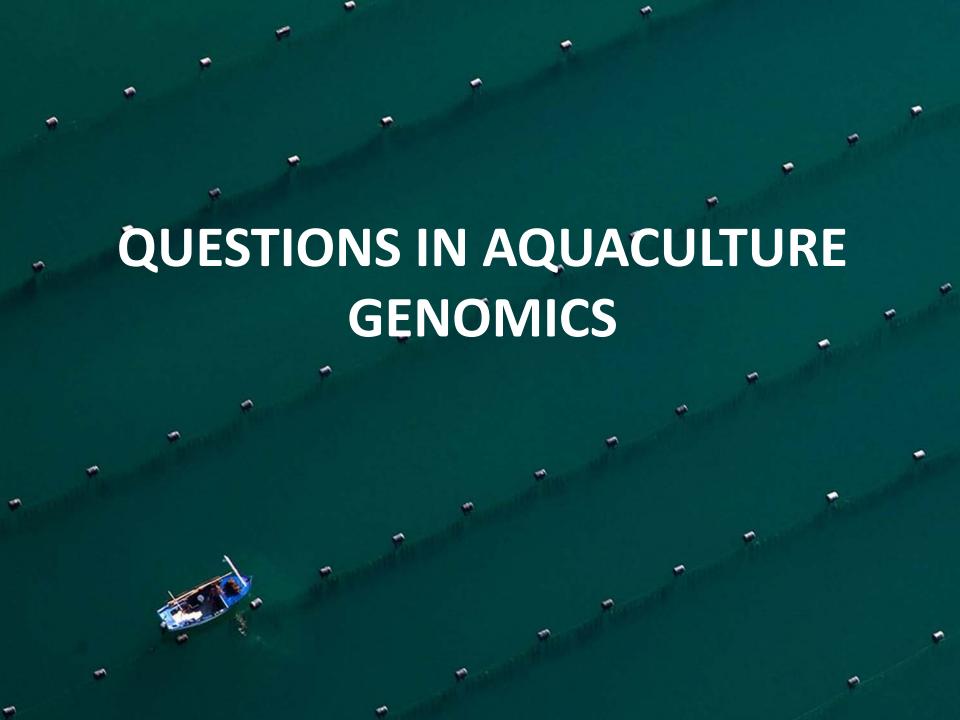
c) without anti-predator nets



#### **DIFFERENT SELECTIVE REGIMES!**

# HIGH CONNECTIVITY BETWEEN NATIVE AND AQUACULTURED POPULATIONS





### **QUESTIONS IN AQUACULTURE GENOMICS**

- Effect of aquaculture practices on Mytilus genomics?
- Adaptation and acclimatization to local environmental conditions?

• Genomic basis of fitness

Interaction between ecological factors and fitness

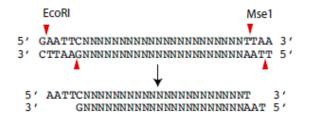
# Mytilus galloprovincialis

Non-model species

Genotyping by sequencing approach (RAD tags)

### **GENOTYPING BY SEQUENCING**

 Digest doublestranded DNA with EcoRI and Mse1.



2. Ligate adaptors to fragments. Adaptors include adaptor sequence, barcode, cutsite, and protector base (adaptors in color; EcoRl on left, Mse1 on right).

Illumina PCR primer I (Illpcr1)

- 5' AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCT 3
  - 5' CTCTTTCCCTACACGACGCTCTTCCGATCTATCAGACACGCAATTCNNNNNNNNTTACAGATCGGAAGAGCTCGTATGCCGTCTTCTGCTTG 3'
    3' TGTGAGAAAGGGATGTGCTGCGAGAAGGCTAGATAGTCTGTGCGTTAAGNNNNNNNNAATGTCTAGCCTTCTCGAGCATACGGCAGAAGACG 5'
    - 3' GAATGTCTAGCCTTCTCGAGCATACGGCAGAAGACGAAC 5'

ACACTCTTTCCCTACACGACGCTCTTCCGATCT
Illumina sequencing primer

Illumina PCR primer II (Illpcr2)

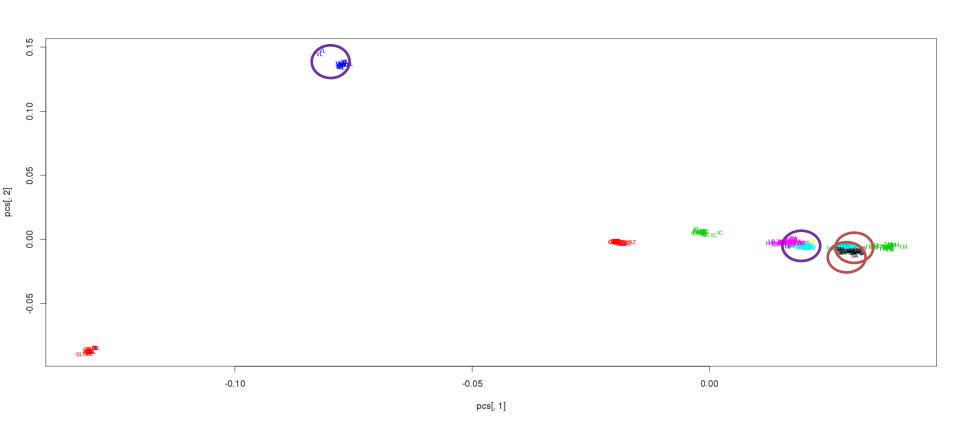
- 3. Amplify fragments with Illumina PCR primers.
- 4. Gel purify PCR product in the desired size range (300-400bp).
- Illumina sequencing.

#### **ANALYSIS OF GBS DATA**

- 1. Demultiplexing and quality control of reads
- 2. De novo assembly of pseudoreference Mytilus genome
- 3. Mapping the reads and SNP calling
- 4. Genotype likelihoods and probabilities
- > over 800,000 SNPs across the genome!

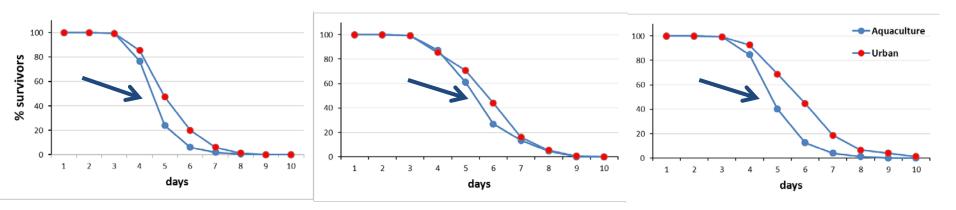
#### **LOW IBD - ROLE OF IBA?**

### **Preliminary results on native populations**

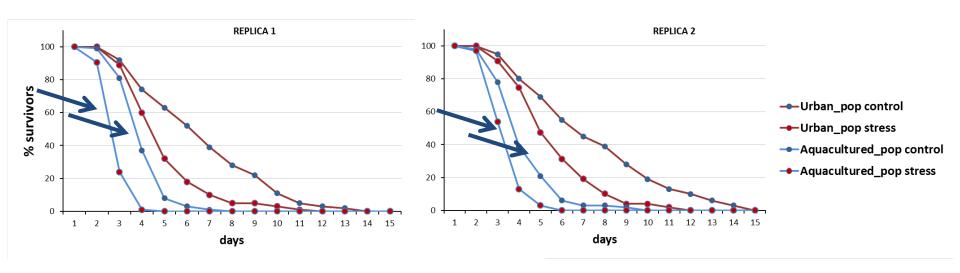


# ENVIRONMENT AND GENOTYPE DEPENDENT LOWER STRESS IN AQUACULTURE POPULATIONS

#### TRANSPLANT EXPERIMENT – ONE AQUACULTURE POP TO THREE PAIR SITES

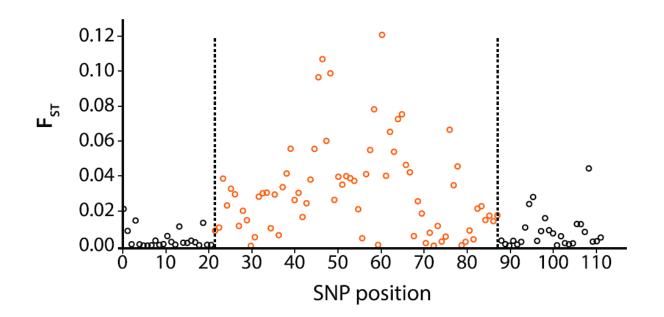


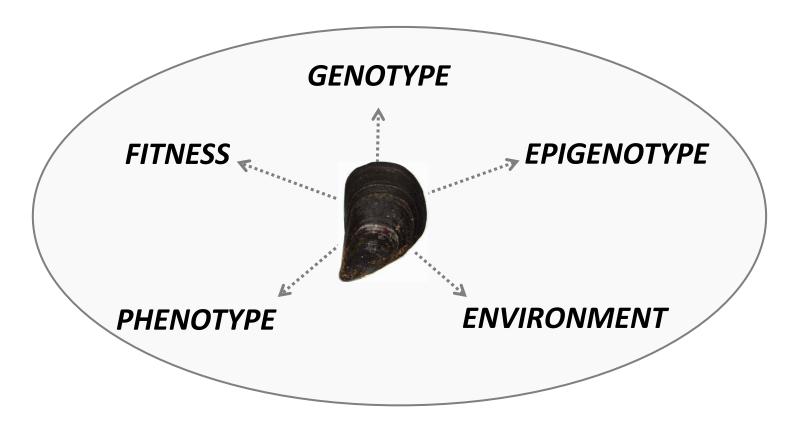
#### **MESOCOSM EXPERIMENT – AQUACULTURE AND URBAN POP EXPOSED TO STRESS**



#### **LOCI UNDER SELECTION**

# Bayesian approach to detect F<sub>ST</sub> outliers





NGS – GENOMICS AND EPIGENOMICS FITNESS AND MORPHOMETRY GWAS

#### **ENVIRONMENTAL VARIABLES**

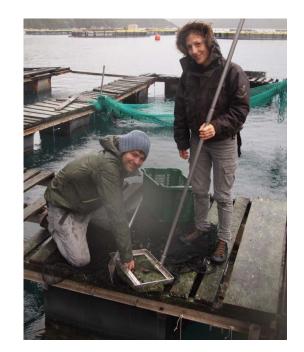
- Native and aquacultured populations
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