

# Genetic effect of domestication selective pressures on Pacific oyster at larval stage



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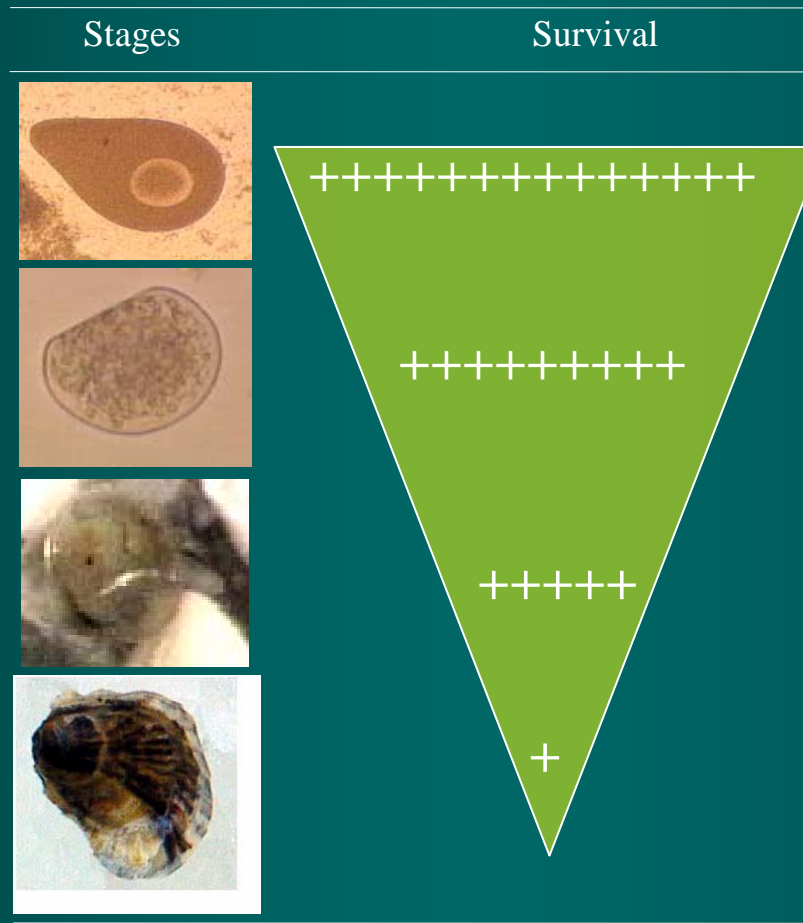
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# *Crassostrea gigas* life cycle

## The “elm-oyster model”

*Williams 1975*



High fecundity and high mortality at early stages

# Which consequences of such a life history strategy for hatchery production ?

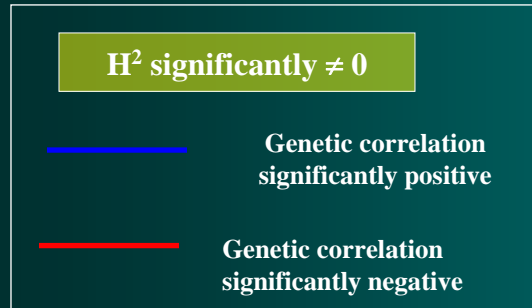


- ☺ Few genitors needed for mass production of juveniles
- ☺ Culling (size selection)
- ☹ Low effective population size (Hedgecock *et al.*, 1992)
- ☹ Risks of rapid loss of genetic variability and inbreeding in closed populations

**Can specific rearing practices (culling) and/or environmental conditions (high temperature) lead to a specific genetic adaptation in *C. gigas* larvae ?**

# Genetic variability of early life traits in *C. gigas* ?

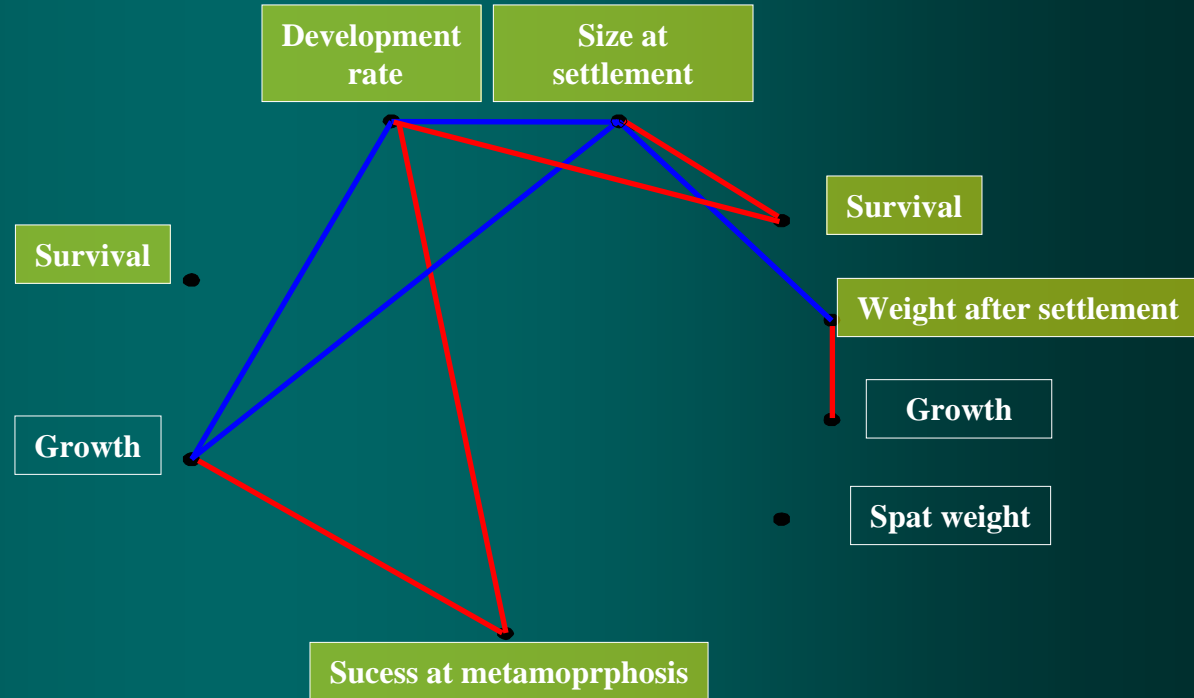
Ernande *et al.*, 2003



Larval traits

Metamorphic traits

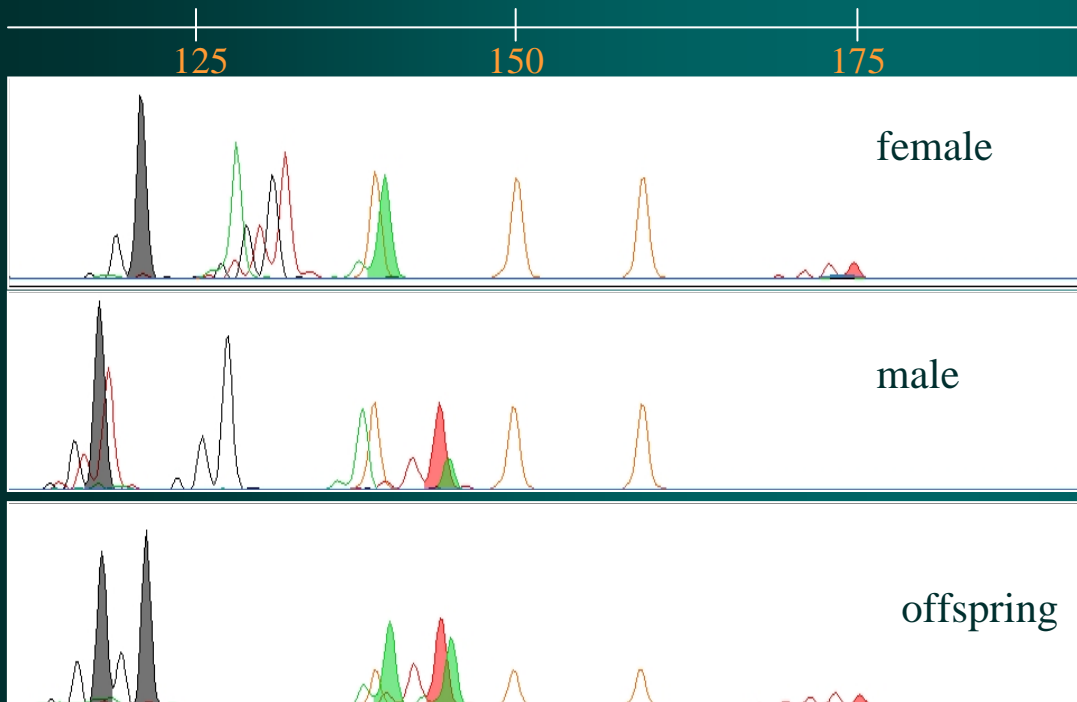
Post-metamorphic traits



- limited number of families
- no family replicates
- a single environment

# Mixed-family approach

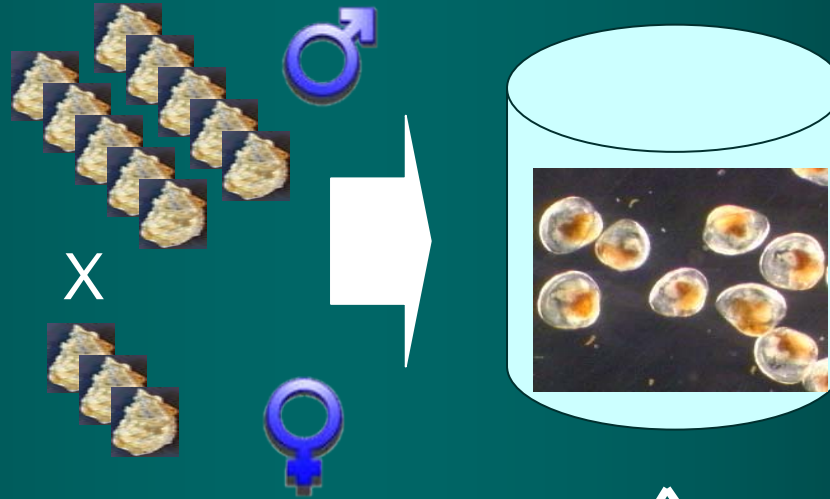
One set of 3 PCR-multiplexed markers allowing efficient parental assignment of larvae (Taris *et al.*, 2005)



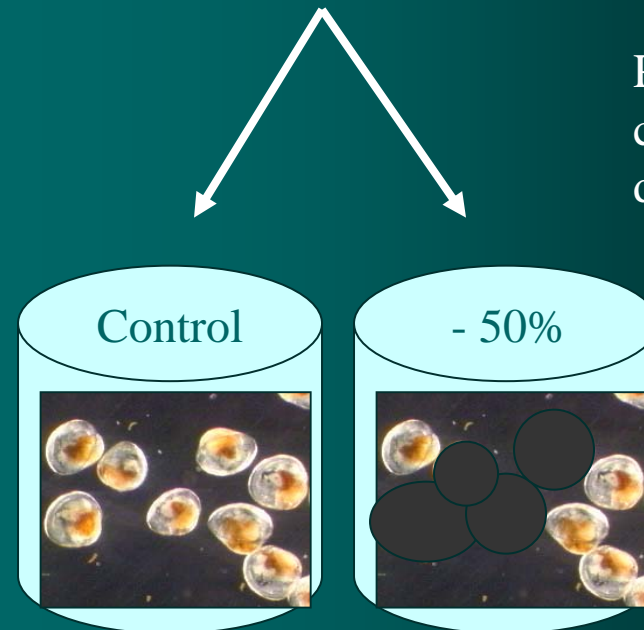
- More families
- Homogeneous rearing conditions
- G x E ?

# 1. Effect of culling

Crossing of 3 females x 10 males with equal gametic contribution within each sex



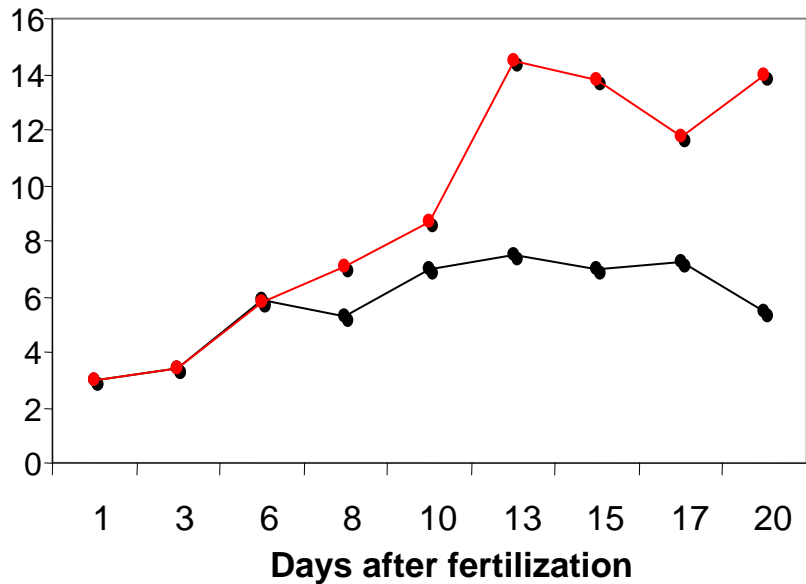
Progressive culling from day 4 to day 15



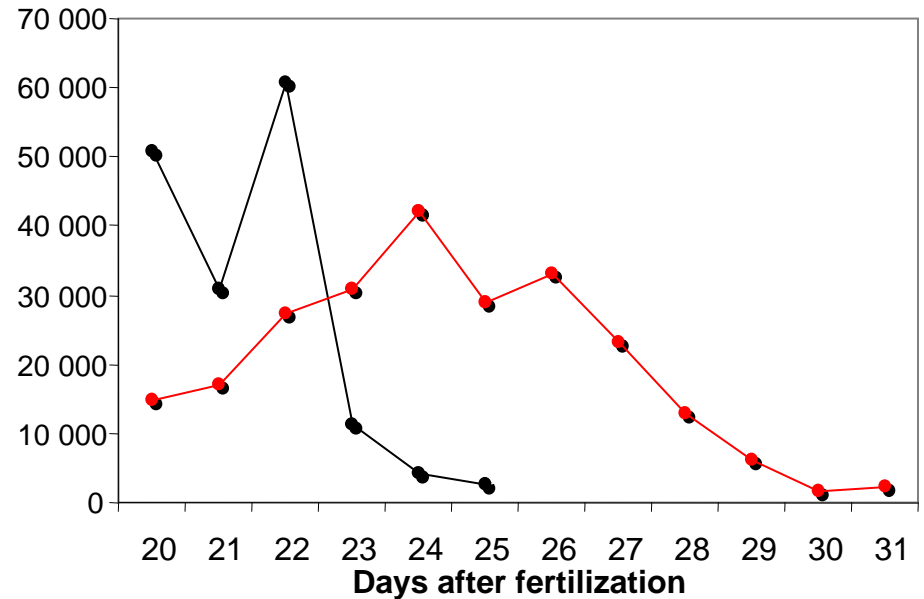
# 1.1 Phenotypic effect of culling 50% of the smallest larvae

— culled population — Control

### Coefficient of variation of larval length



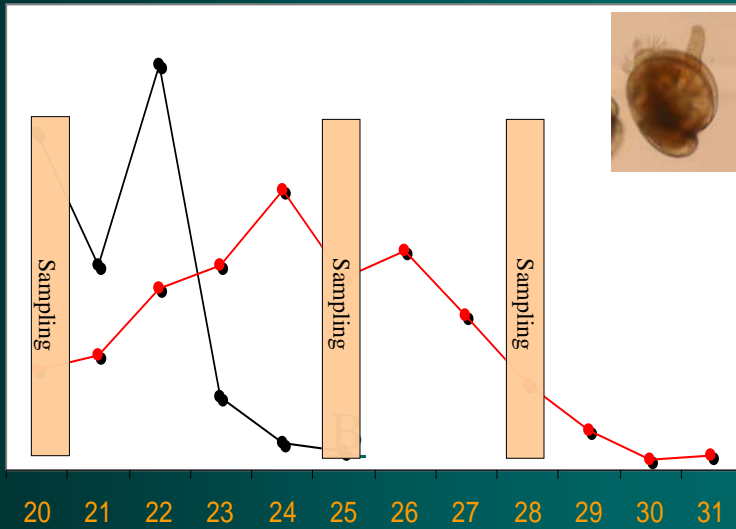
### Number of pediveliger larvae



## Limited effect on yield:

- 30 % of ready-to-settle larvae (higher survival of fast growing larvae)
- 15 % of spat (higher settlement success of fast growing larvae)

# 1.2 Genetic effect of culling

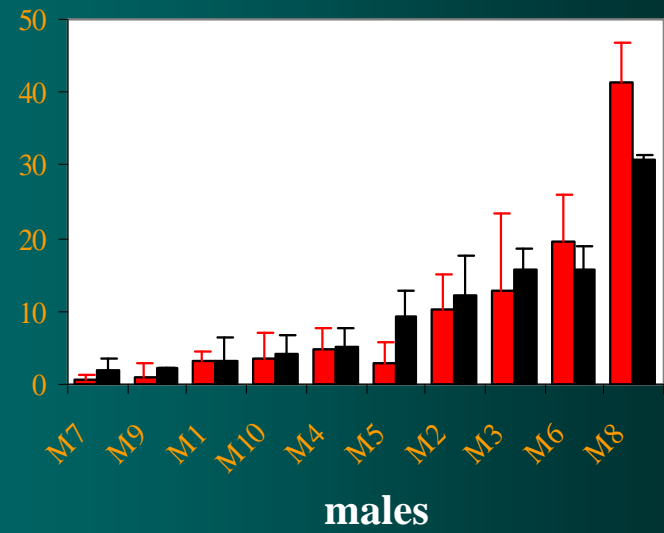


$N_e = 8.2$   
6.3

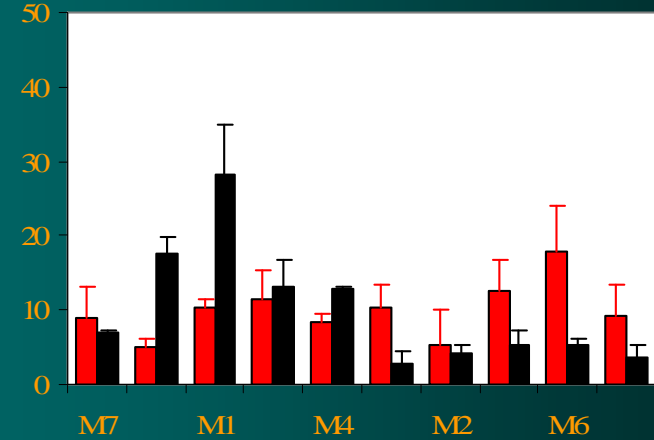
12.3  
15.2 15.9

The effect of culling on genetic diversity is mediated through its effects on the timing of settlement

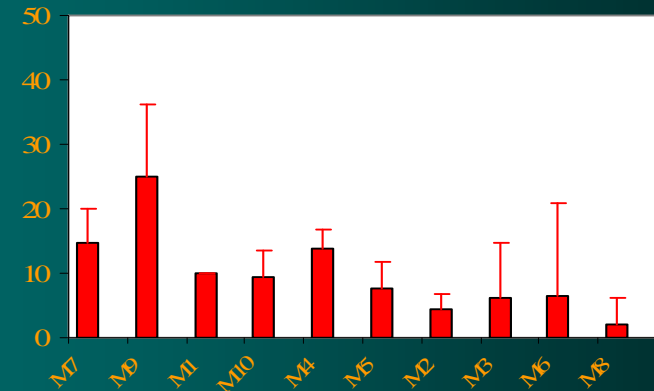
D20



D25



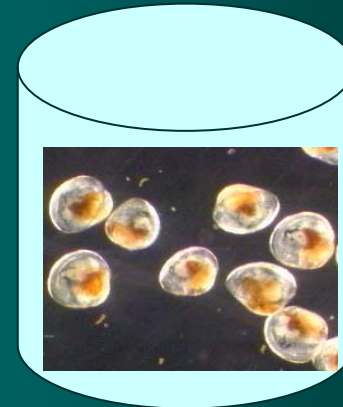
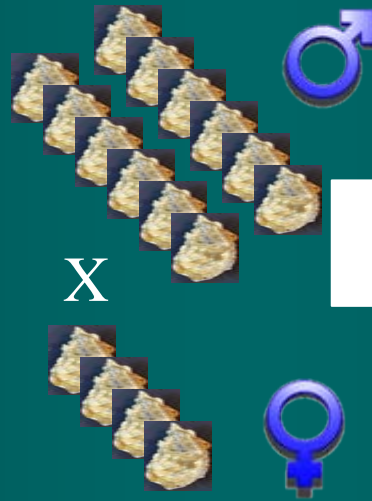
D28



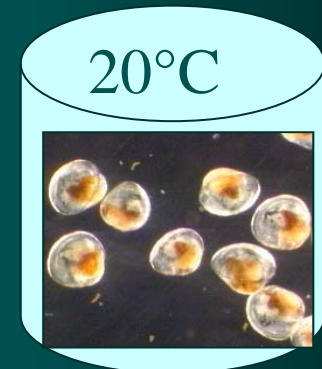
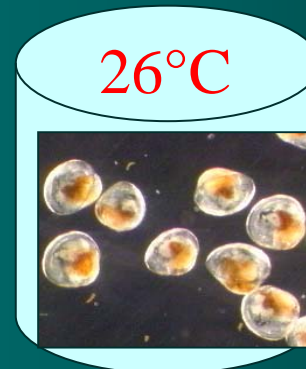


## 2. Effect of temperature

Crossing of 4 females x 12 males with equal gametic contribution within each sex

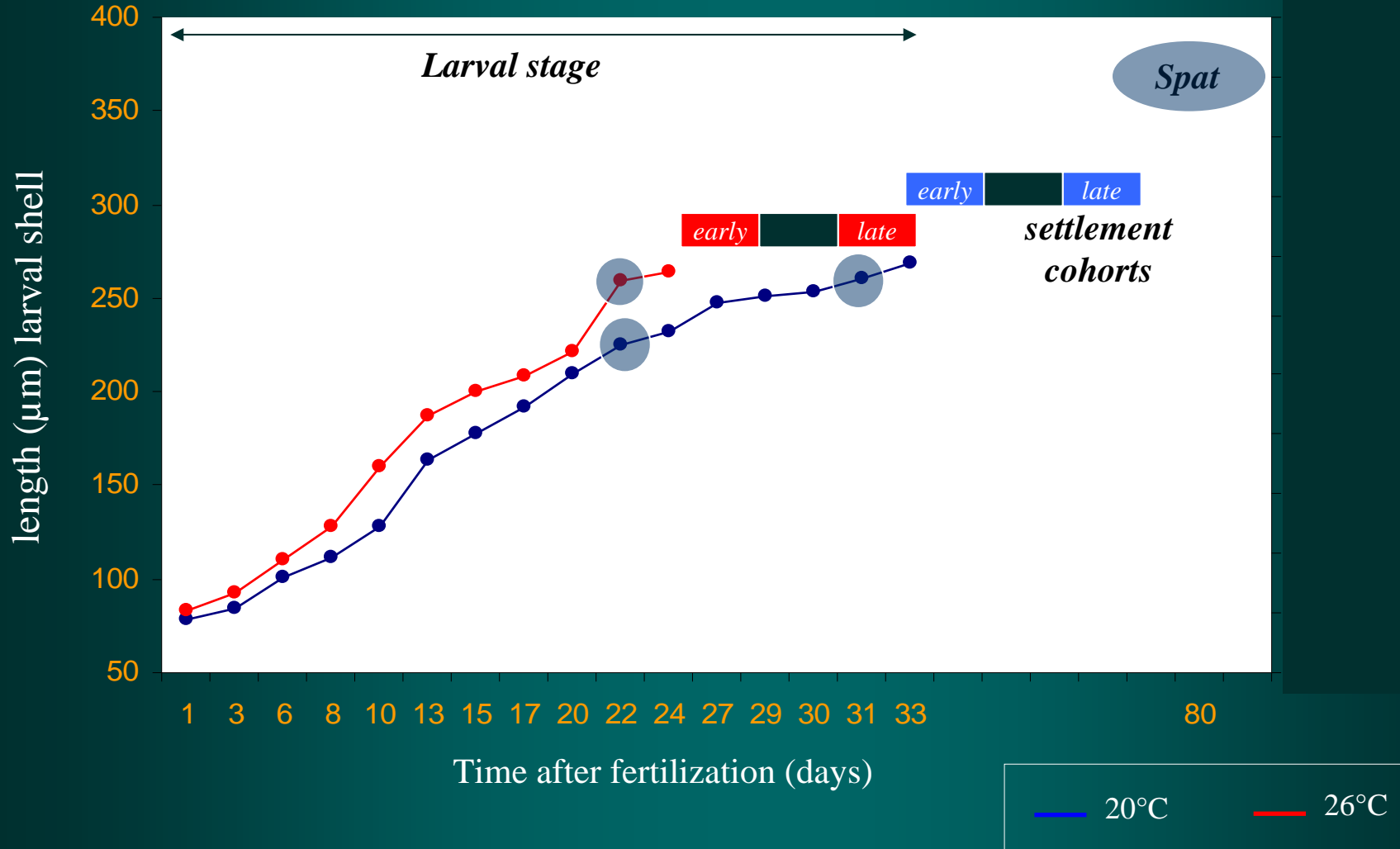


Estimation of hatching rate at day 1



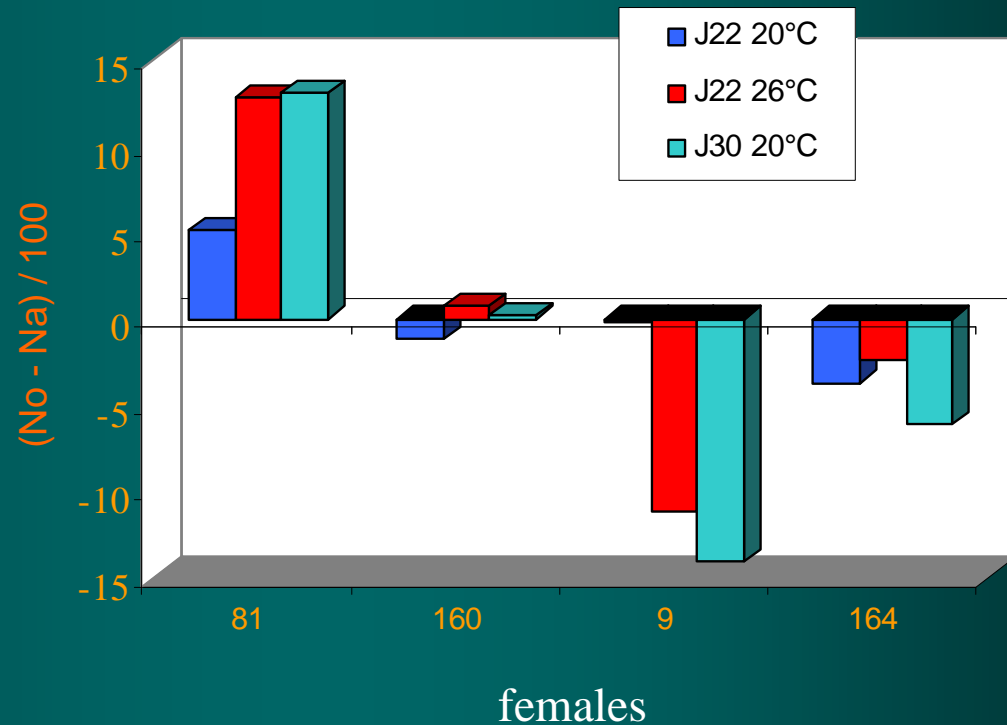
Individual measurements of larvae prior to genotyping

# 2.1. Phenotypic effect & sampling



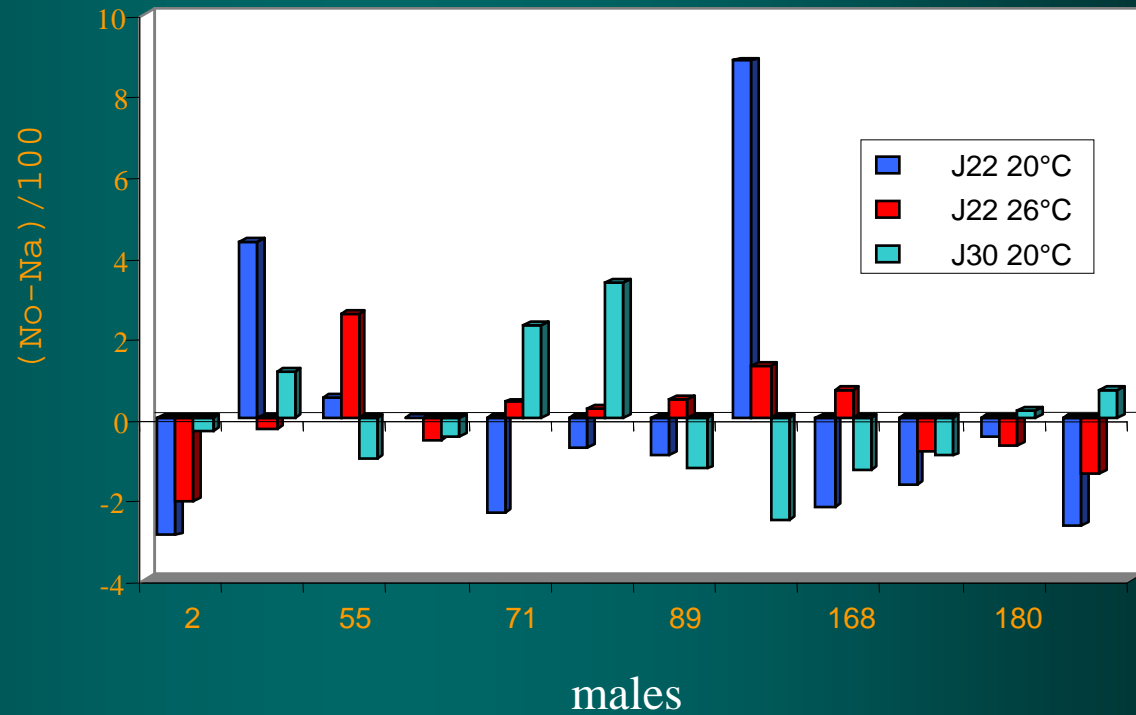
## 2.2. Variance of female reproductive success

Observed maternal contributions relative to mean hatching rate at Day 1

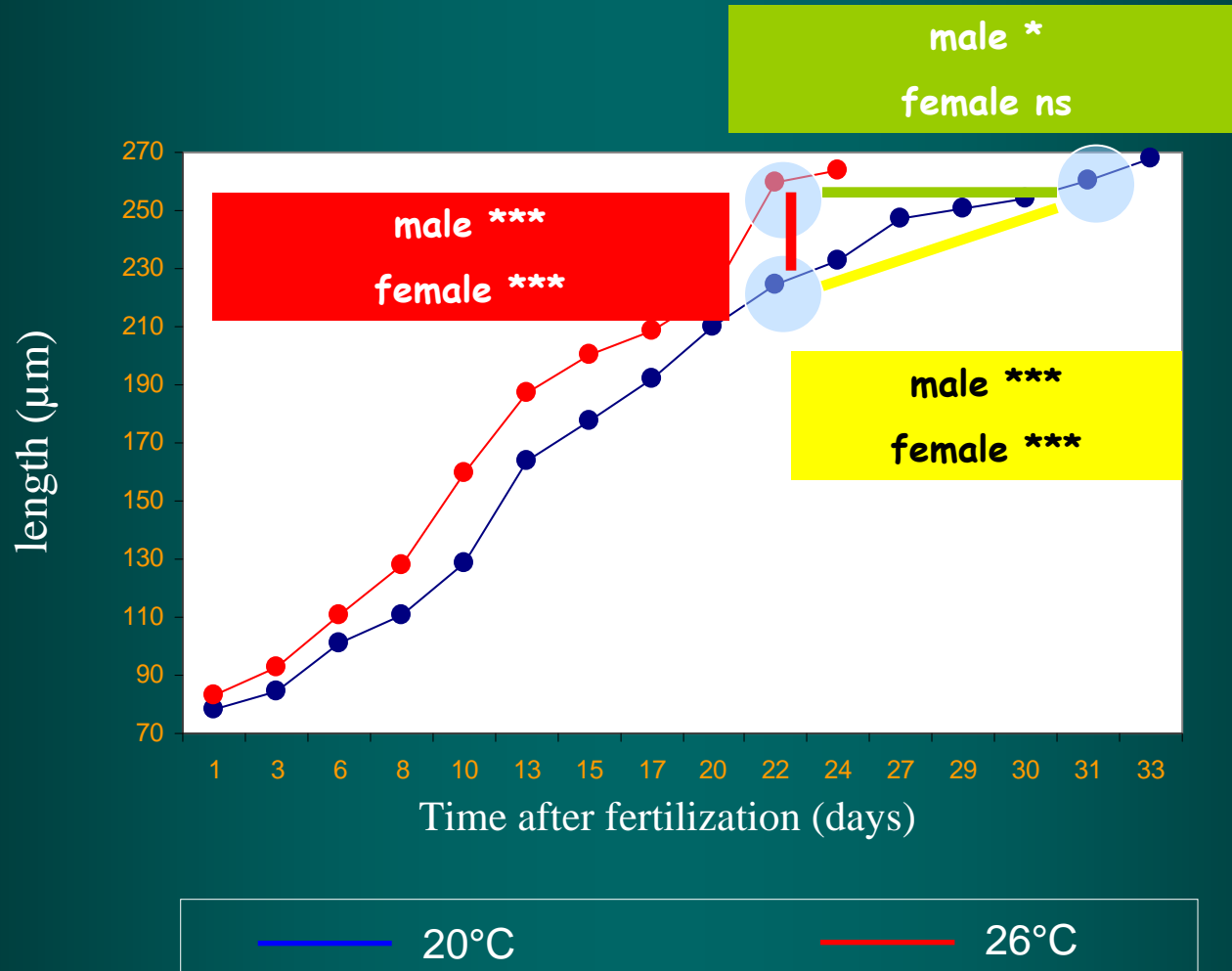


## 2.2. Variance of parental reproductive success

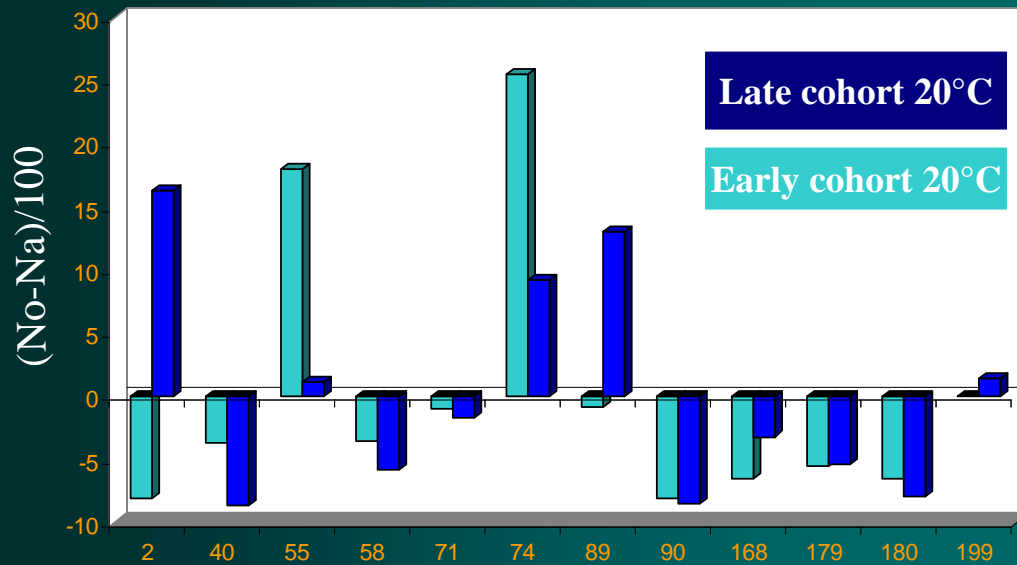
Observed paternal contributions relative to mean hatching rate at Day 1



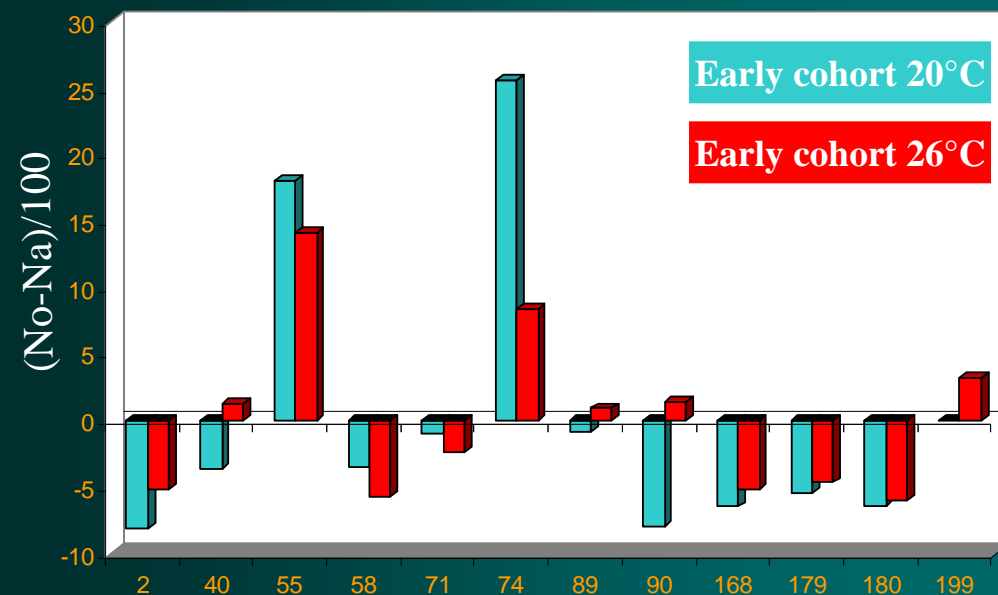
## 2.2. Variance of parental reproductive success



## 2.3. Paternal contributions at day 80 (spat)

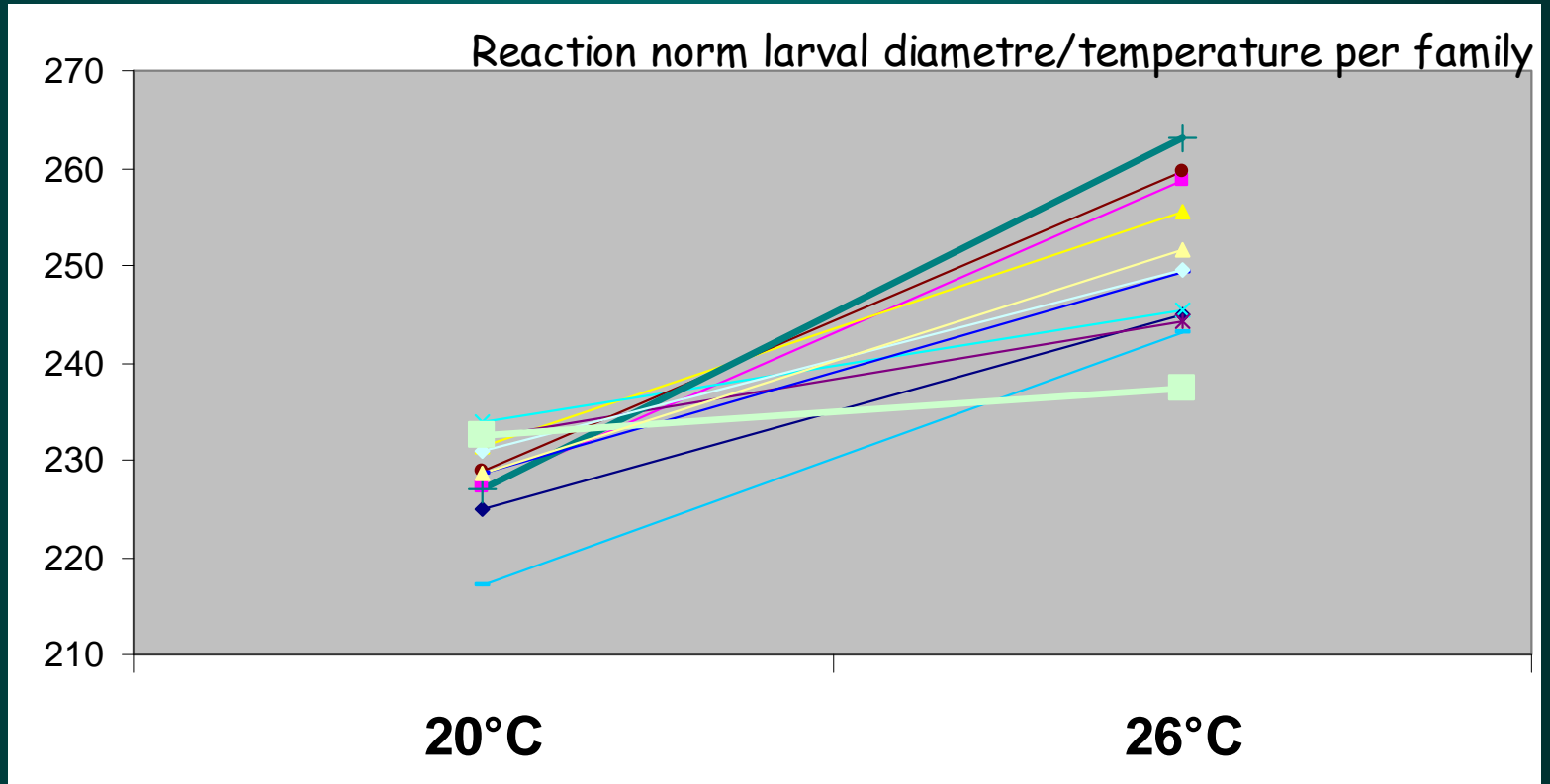


Significantly different contributions between early and late cohorts at 20°C (same result at 26°C)



Significantly different contributions between early cohorts at 20°C and 26°C (same result for the late cohort)

## 2.2. « G x E » interaction on larval size



Male

ns

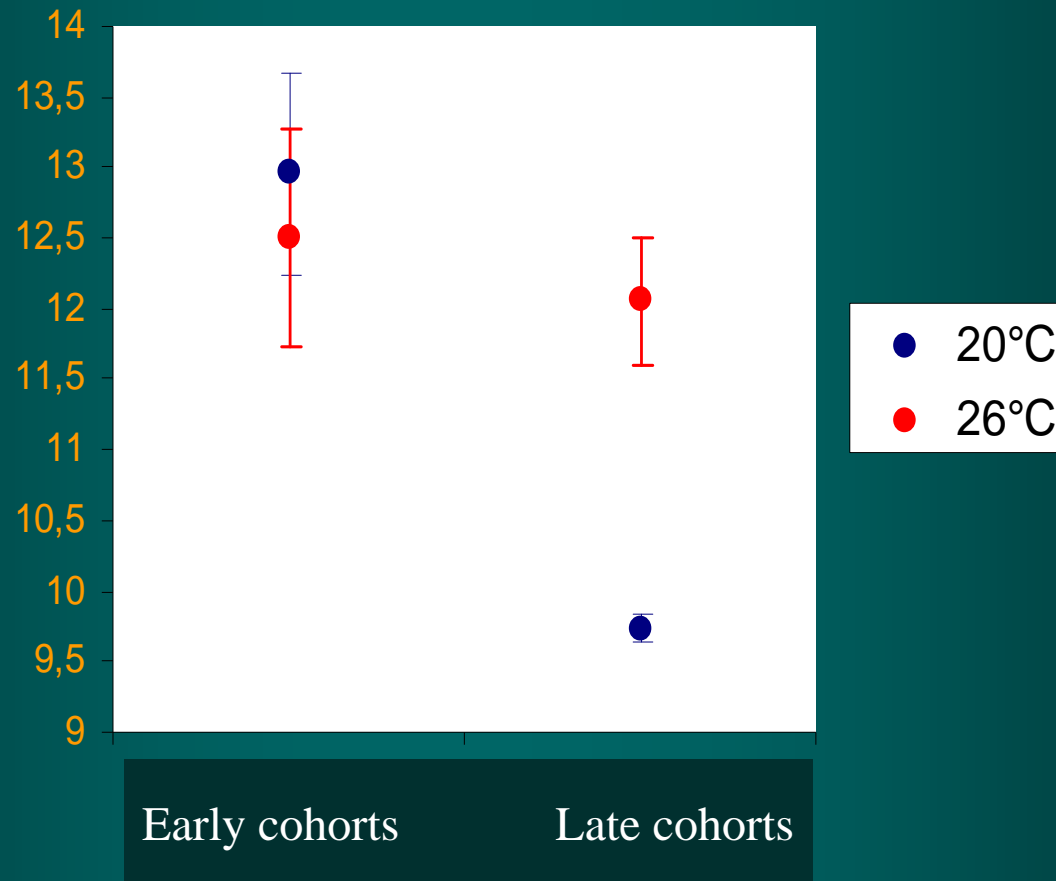
$p < 0.05$

Female

ns

$p < 0.05$

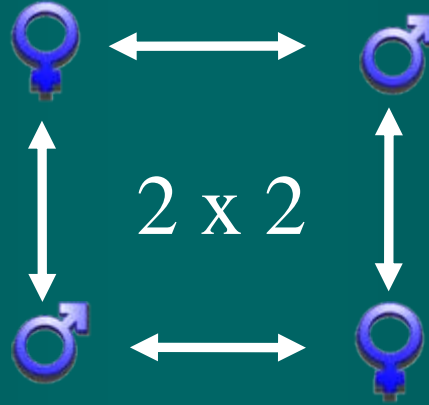
## 2.4. Effect of temperature during larval rearing on spat growth





# 3. Selection for fast growing larvae in hatcheries ?

Hatchery broodstock



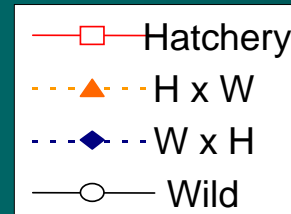
Wild broodstock



7 generations of breeding

Loss in allele diversity  $\approx 70\%$

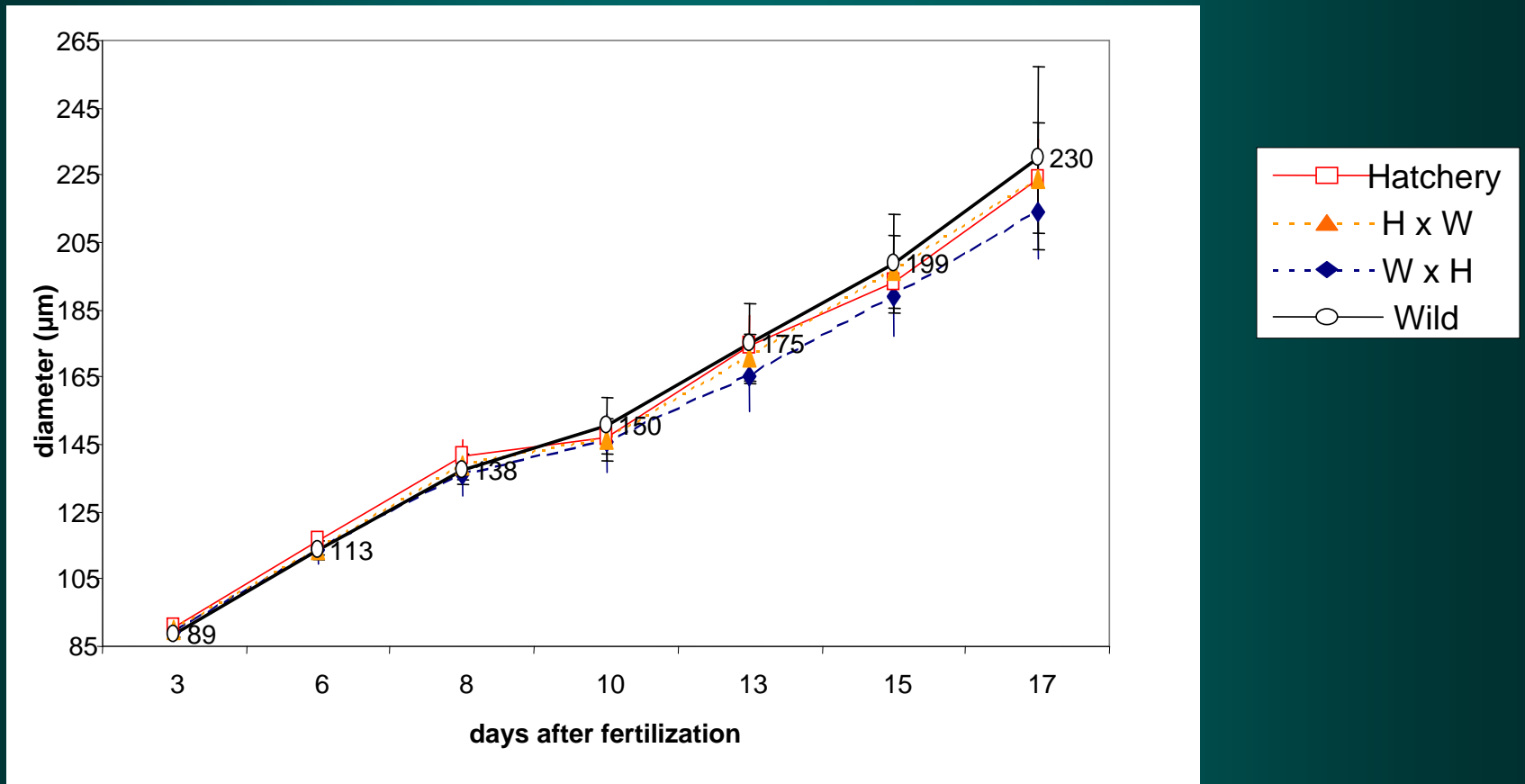
Loss in heterozygosity  $\approx 20\%$



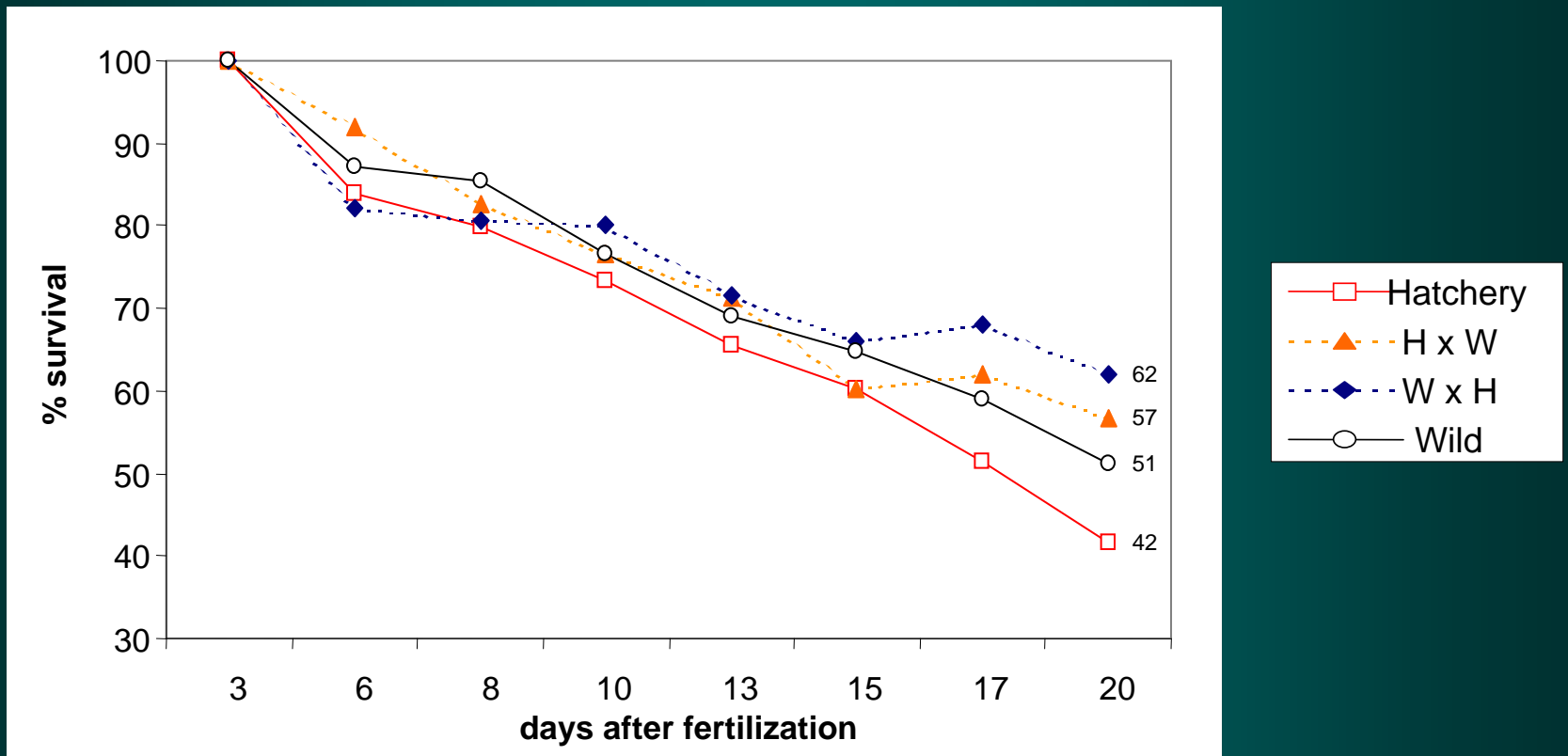
Rearing conditions:

- 24°C
- no culling

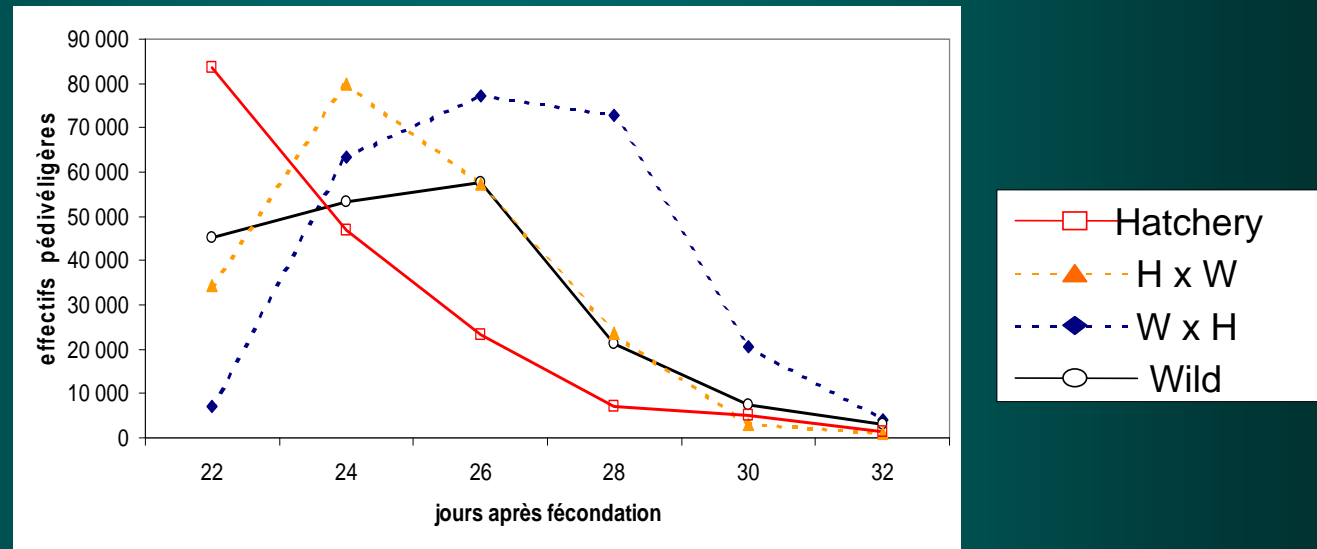
# 3.1. Larval growth



## 3.2. Larval survival



### 3.3. Timing to reach the pediveligere stage and settlement success

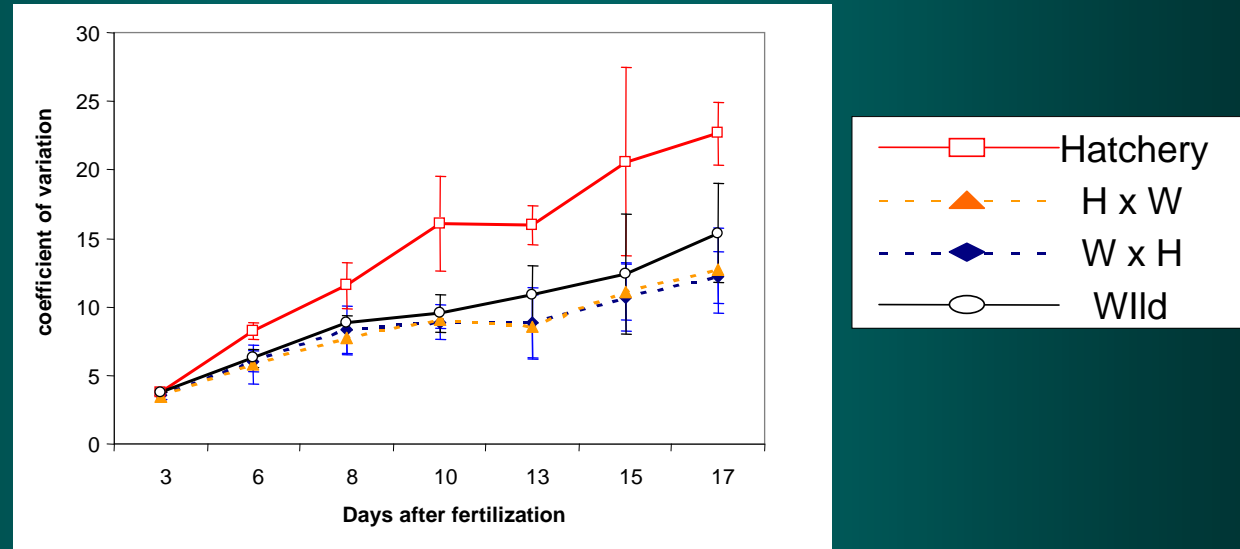


Settlement success (%):

Hatchery 90.7 > HxW 78.1 > Wild 72.3 > WxH 68.7

# 3.4. Within progeny variation for larval size

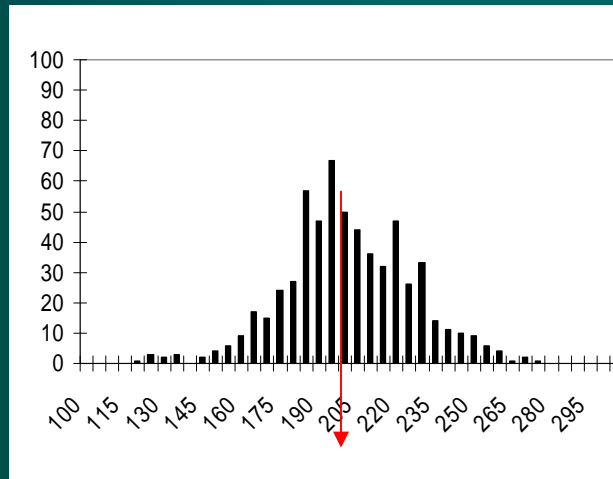
C.V. of larval length



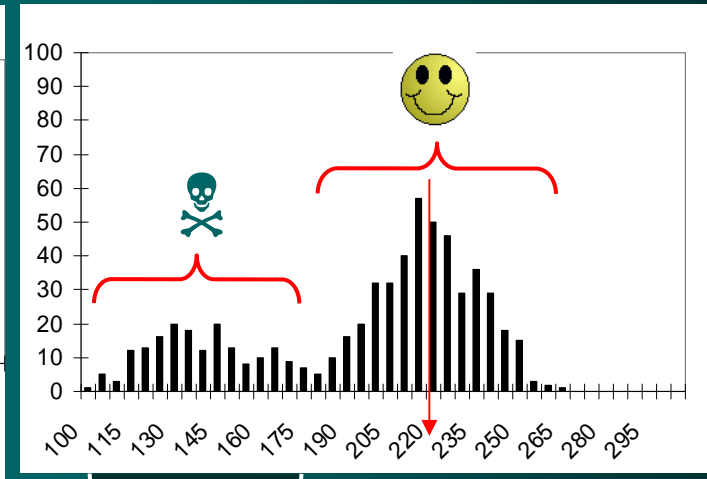
Wild progeny

Hatchery progeny

Distribution of larval length at Day 15



205 μm



225 μm \*\*  
Inbred larvae ?

# Conclusions

## Methodology

- As individual tagging is impossible at early life stages, marker-based parentage analysis of mixed families represents an efficient tool to study genetic variability of larval traits.

## Selection at larval stage

- Significant differences are observed between progenies, confirming the existence of genetic variation for several traits.
- Temperature influences the expression of genetic variability for growth and survival and therefore is likely to increase the genetic effect of culling.
- Intensive rearing practices can lead to the selection of faster growing / higher settlement larvae, despite inbreed depression.



Thanks