

# GENETIC IMPACT OF THE REPRODUCTION DYNAMICS IN THE EUROPEAN FLAT OYSTER *Ostrea edulis*

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# *Ostrea edulis*



**+ introduction in USA, Canada, Japan...**

# ***O. edulis* population genetics:**

*How is the genetic variability distributed...*

*... in space ?*

- Within vs among populations ?
- Atlantic vs Mediterranean populations ?
- Isolation by distance ?

*... among genomes ?*

- Nuclear vs Cytoplasmic ?

*... over time and/or generations ?*

- Adults vs juveniles
- Females vs larvae ?

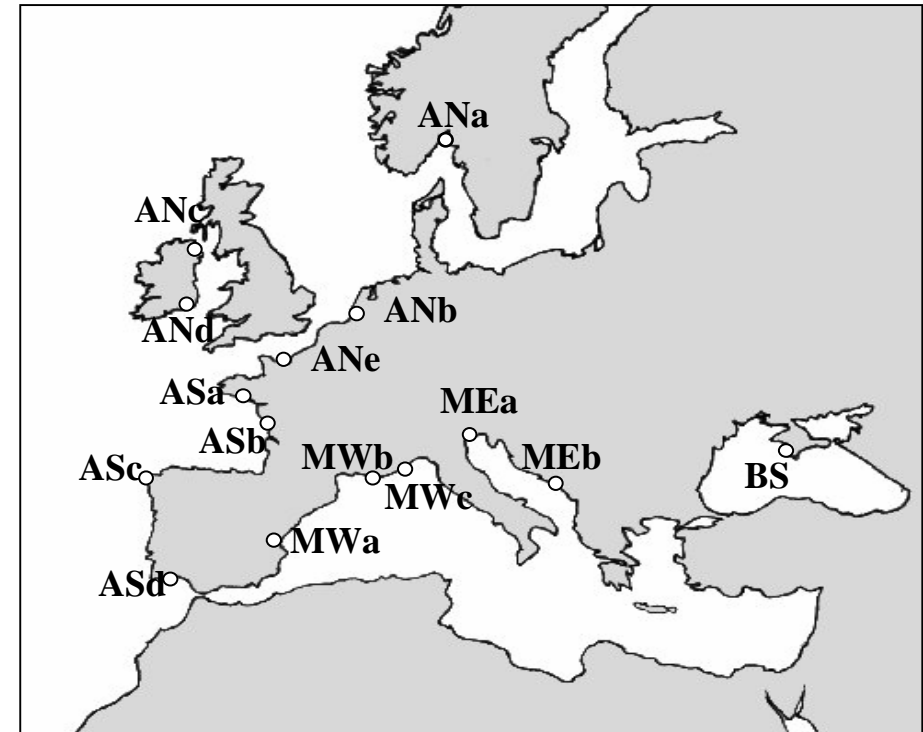
# Spatial distribution ?

## Sampling:

15 populations sampled  
14 to 50 individuals per location

## Markers:

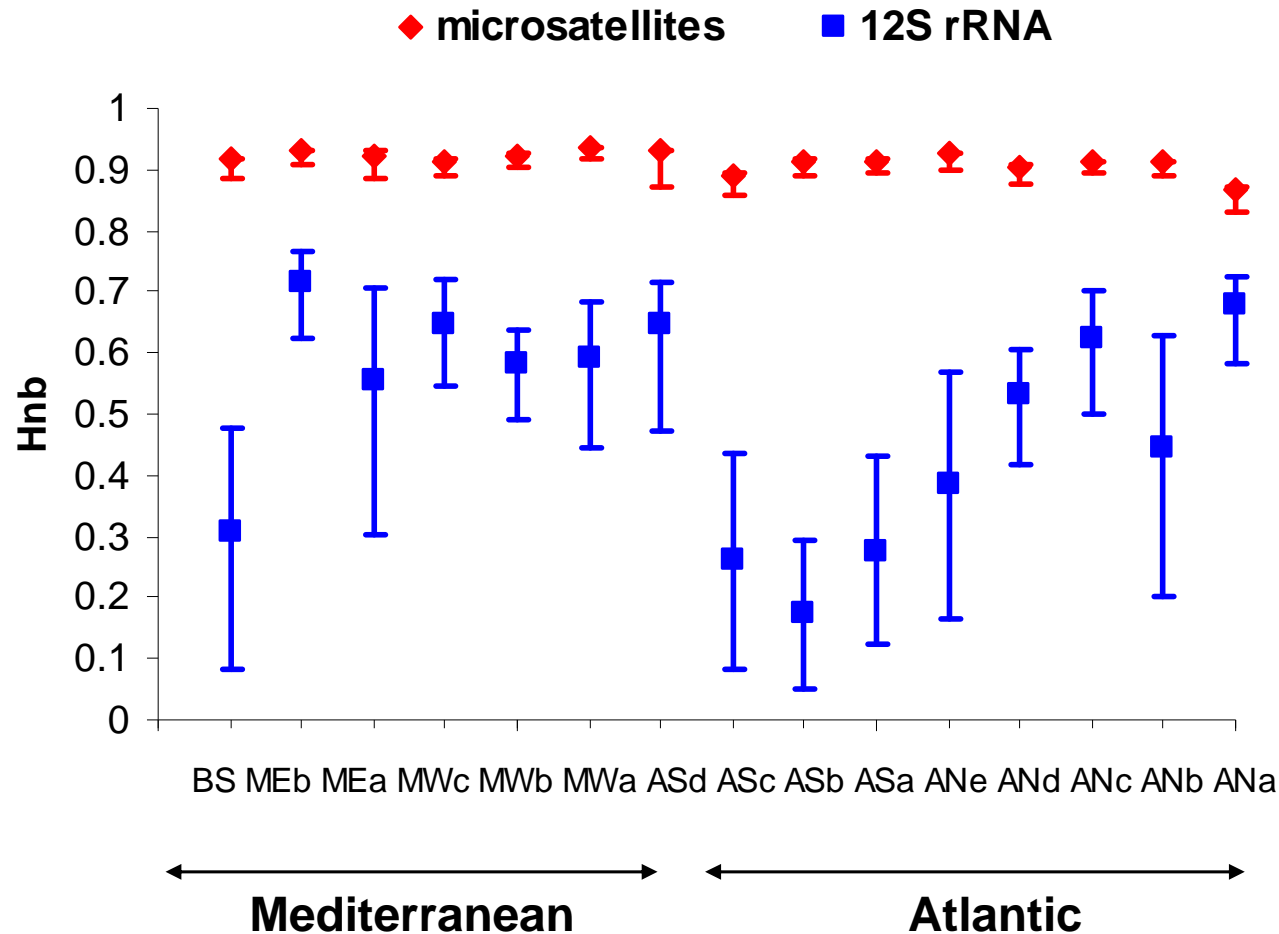
- Allozyme loci (Saavedra *et al.*, 1993, 1995)
- microsatellite loci (Launey *et al.*, 2002)
- 12S rRNA SSCP (Diaz Almela *et al.* in press)



# Within population diversity :

**Microsatellites:** allele/locus/pop =  $18.5 \pm 4.5$   
mean  $H_e = 0.914 \pm 0.018$

**12S rRNA:** 14 SSCP haplotypes

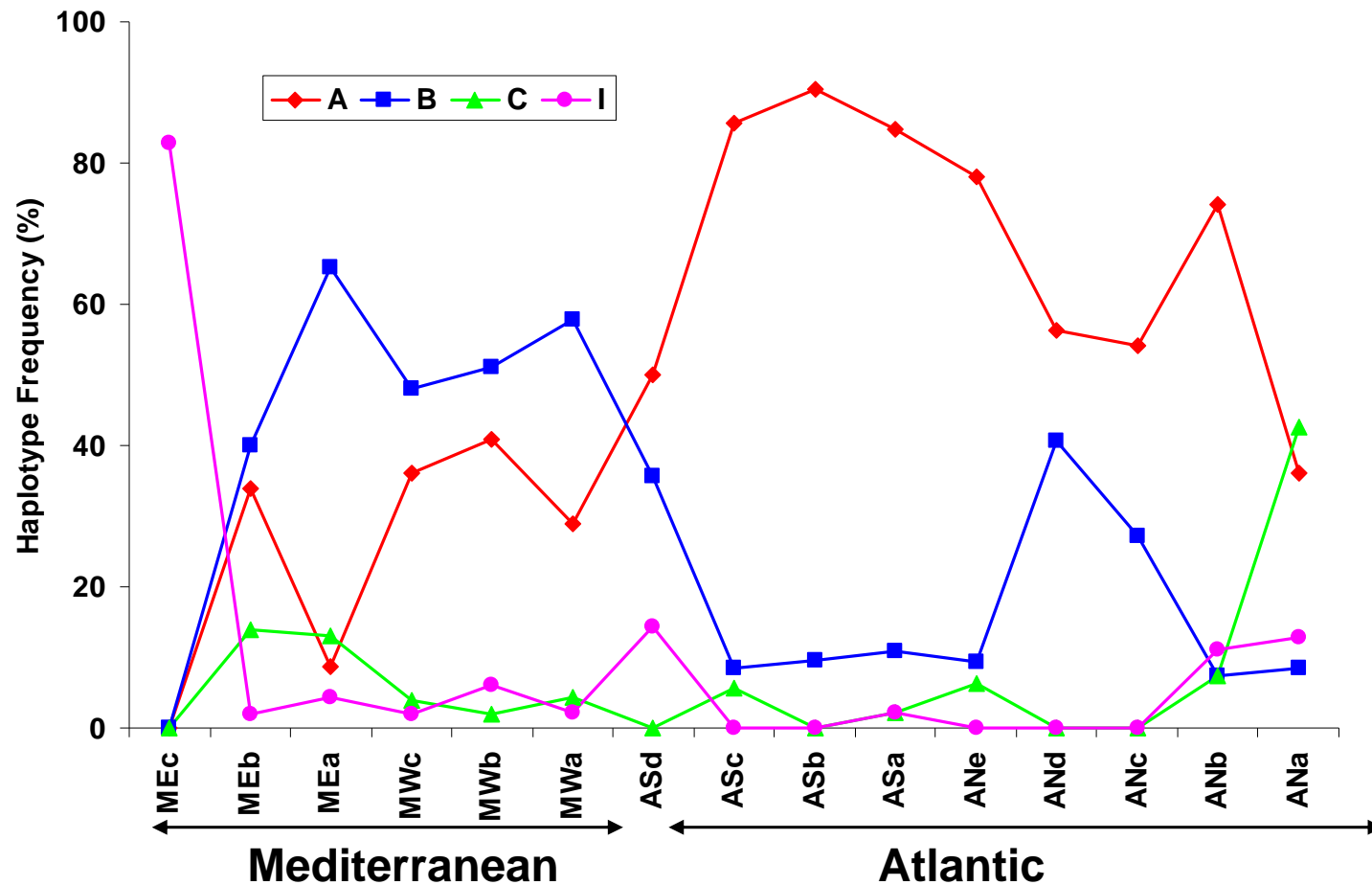


# Among population differentiation :

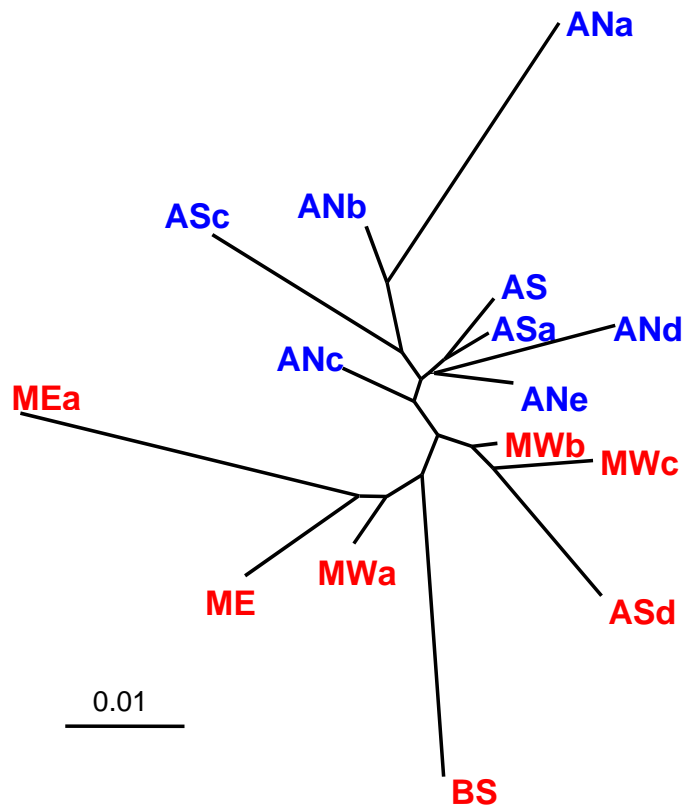
Microsatellites :  $F_{st} = 0.019^{***}$

12S rRNA :  $F_{st} = 0.224^{***}$

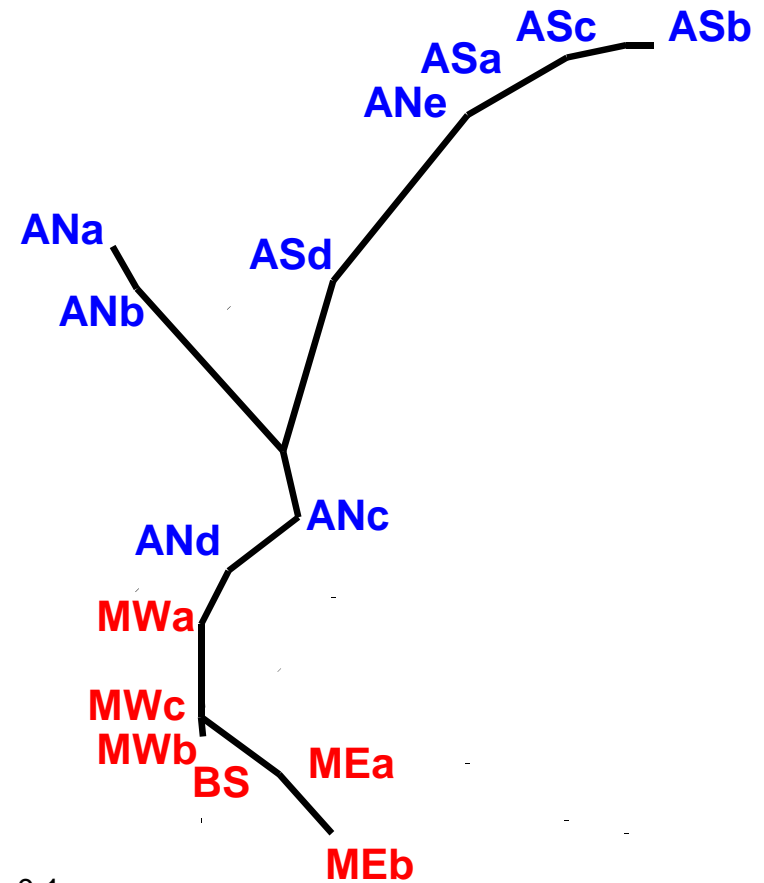
↻ 10 X



# Among population differentiation :

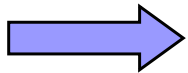
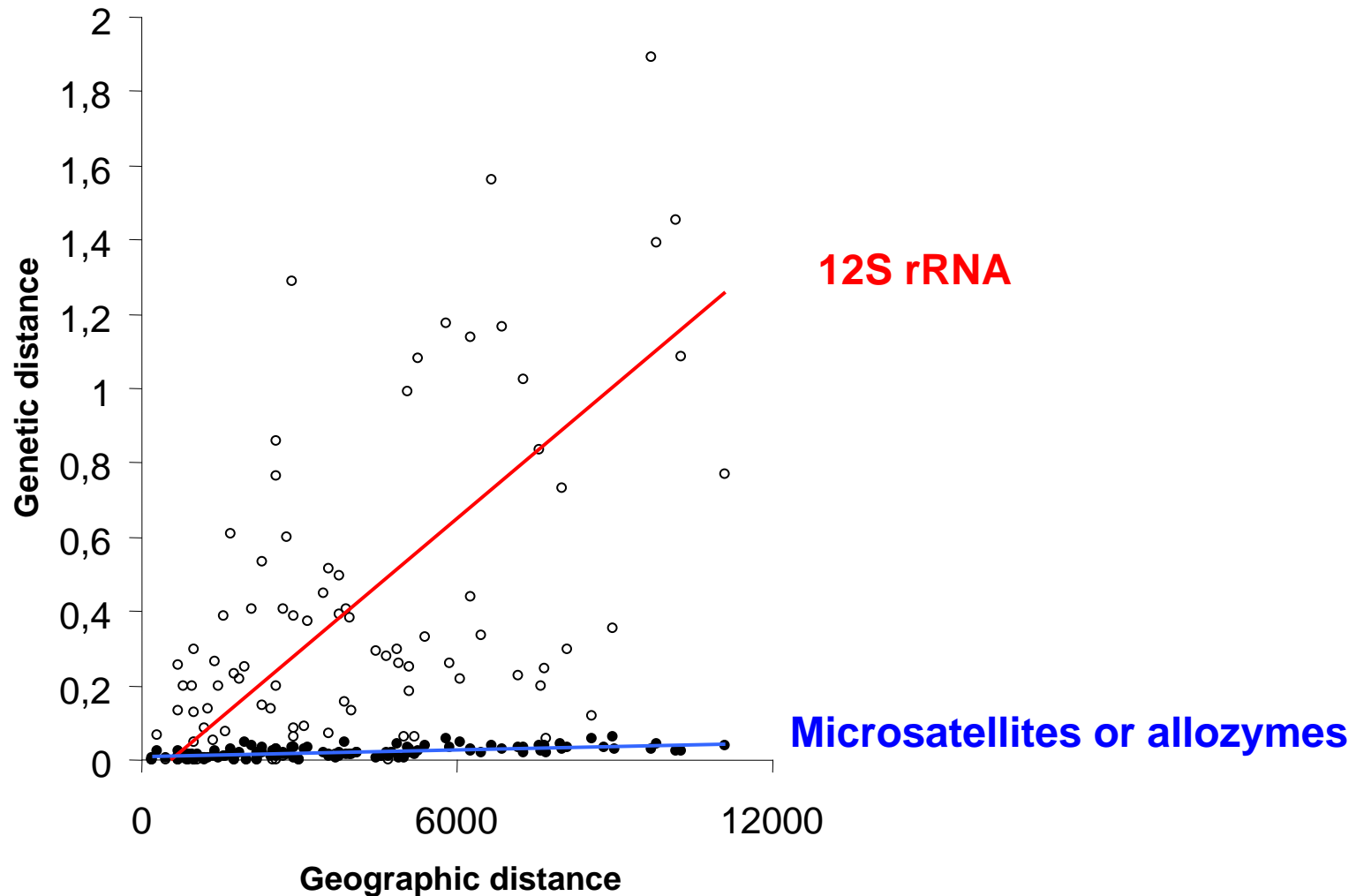


**Microsatellites**



**12S rRNA**

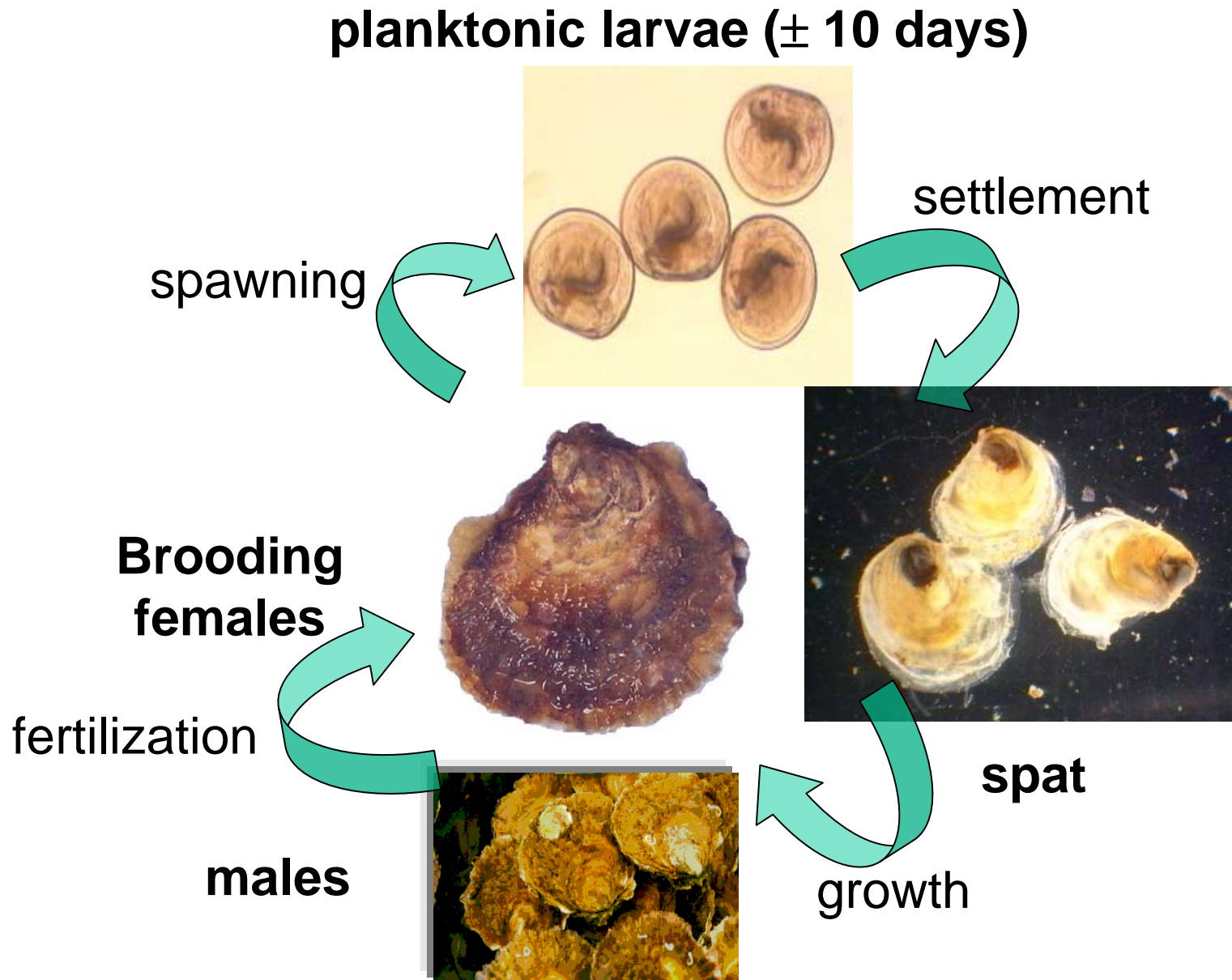
# Isolation by distance :



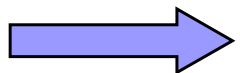
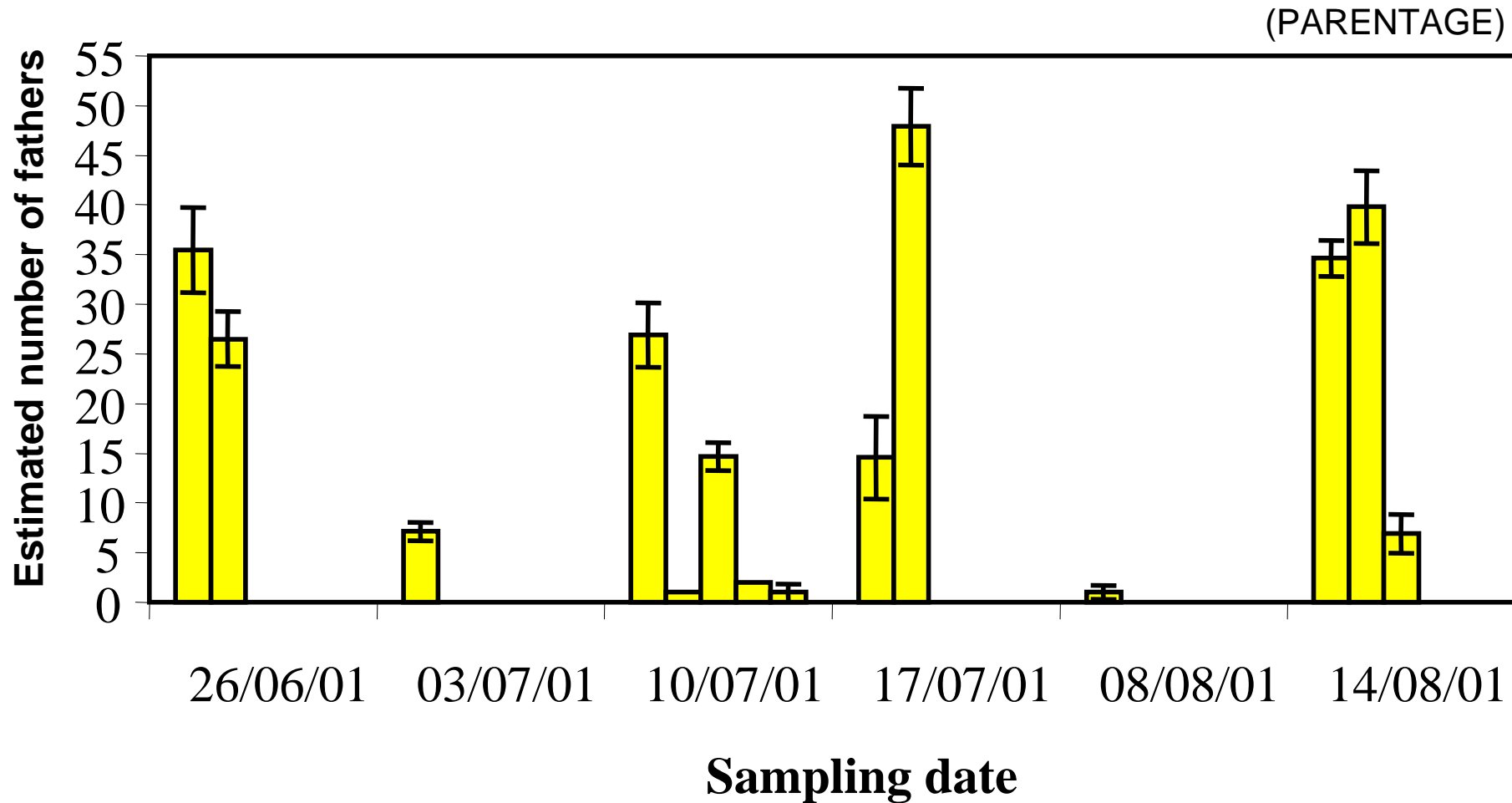
**Higher variance in reproductive success in the female than in the male ?**



# Reproductive cycle

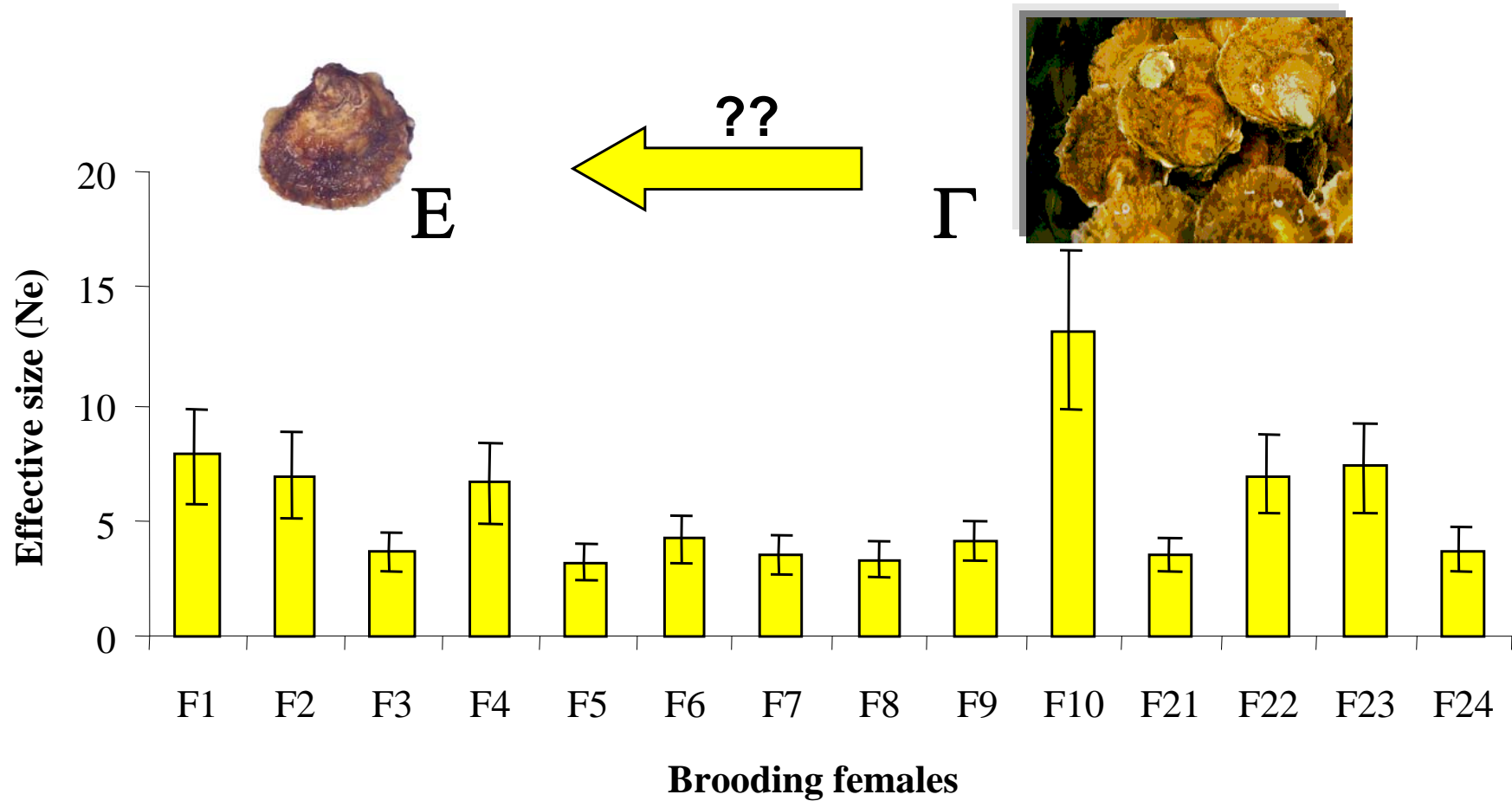


# how many males / female ?

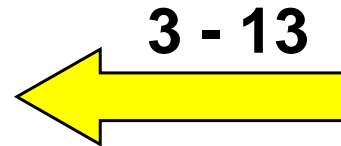
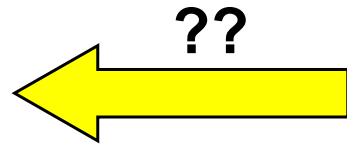


**Very variable number of males / female**

# Variability over generation: how many males / female ?



# Variability over generations : $N_e = ?$



$\Gamma$



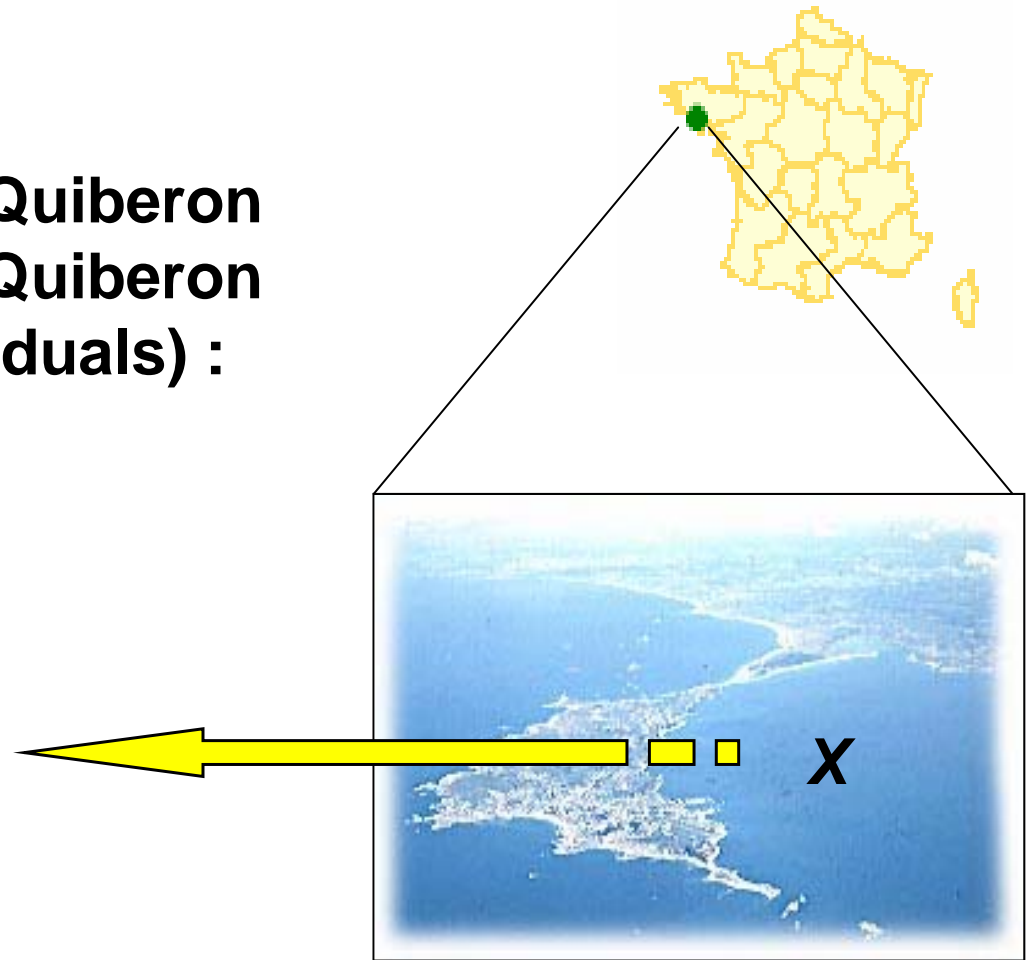
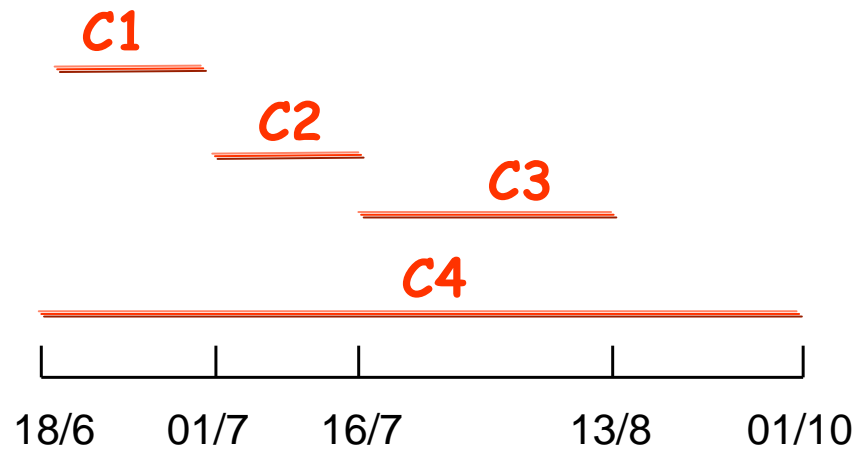
➤ **1994** : one cohort recruited over 15 days in Sète

alleles / locus : spat (16.4)  $\neq$  adults (21.8)

$N_e = 16$  [11,23] (Launey *et al.*, in prep)

# Variability over generations: $N_e = ?$

**1995** : 49 adults collected in Quiberon  
**2001** : 68 adults collected in Quiberon  
+ 4 cohorts (672 individuals) :



**4 microsatellites**  
**12S sequence**

# Differentiation among cohorts (*Fst*)

	Cohort 1	Cohort 2	Cohort 3	Cohort 4	Adults
Cohort 1		-0.019	-0.014	<b>0.052*</b>	<b>0.058*</b>
Cohort 2	0.000		-0.024	0.041	0.049
Cohort 3	0.001	-0.001		0.021	0.029
Cohort 4	0.000	0.002	0.002		-0.019
Adults	0.001	0.002	0.002	0.000	

\*  $p < 0.05$

**No differentiation using microsatellites**

**Low mitochondrial differentiation (cohort 1)**

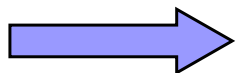
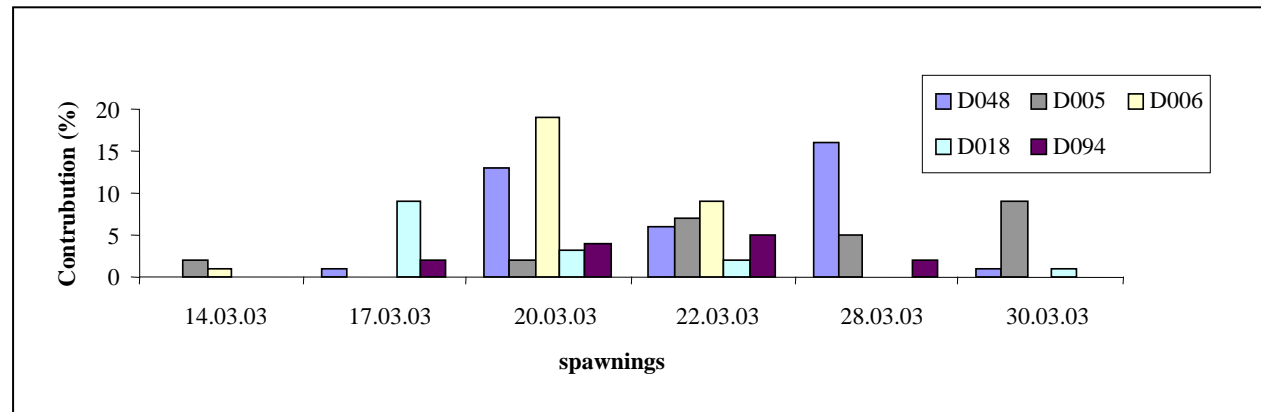
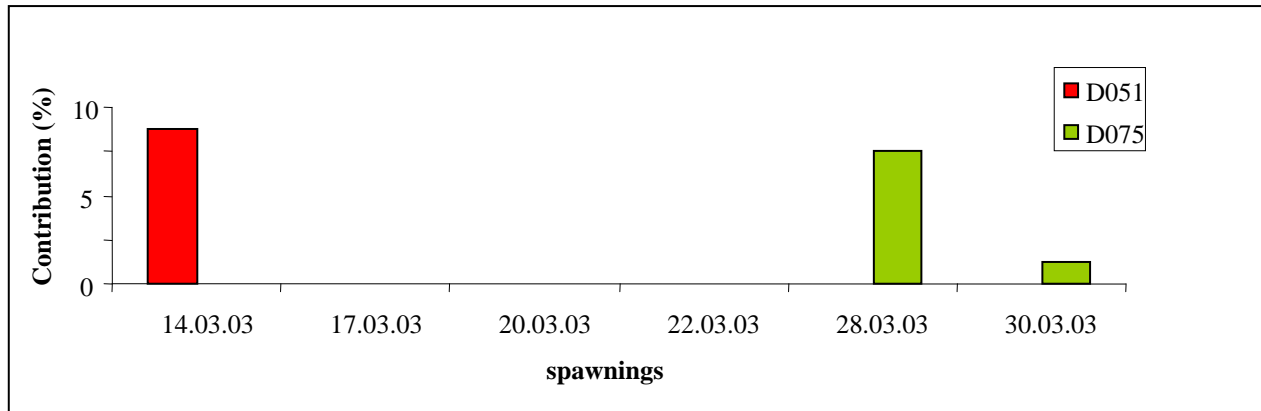
# Variability over generations: $N_e = ?$

- alleles / locus : spat = adults
- adults /spat :  $N_e = 135 [44, +\infty]$
- adults 1994 / adults 2001 :  $N_e = 137 [60, +\infty]$



# Spawning behaviour under controlled conditions :

60 individuals in a spawning tank  
Ne / spawning event = 7 to 17



Variation of reproductive success



# Risks and benefits of hatchery propagated spat to enhance production ?

- **Breeding programs have been initiated to improve disease resistance** (Naciri-Graven et al., 1998; Culloty et al., 2001)
- **Hatchery populations usually have low  $N_e$**  (Hedgecock et al., 1992) in *O. edulis* :
  - ✓ **Saavedra & Guerra (1996):  $N_e \approx 4$**
  - ✓ **Launey et al. (2001):  $N_e = 3$  to 20**
- **Today hatchery-based production is still very limited**
- **Its development could have a positive impact in terms of aquaculture but might have a strong negative impact on genetic diversity of wild populations.**