

Oyster population genetics: Tracing introductions and understanding natural populations

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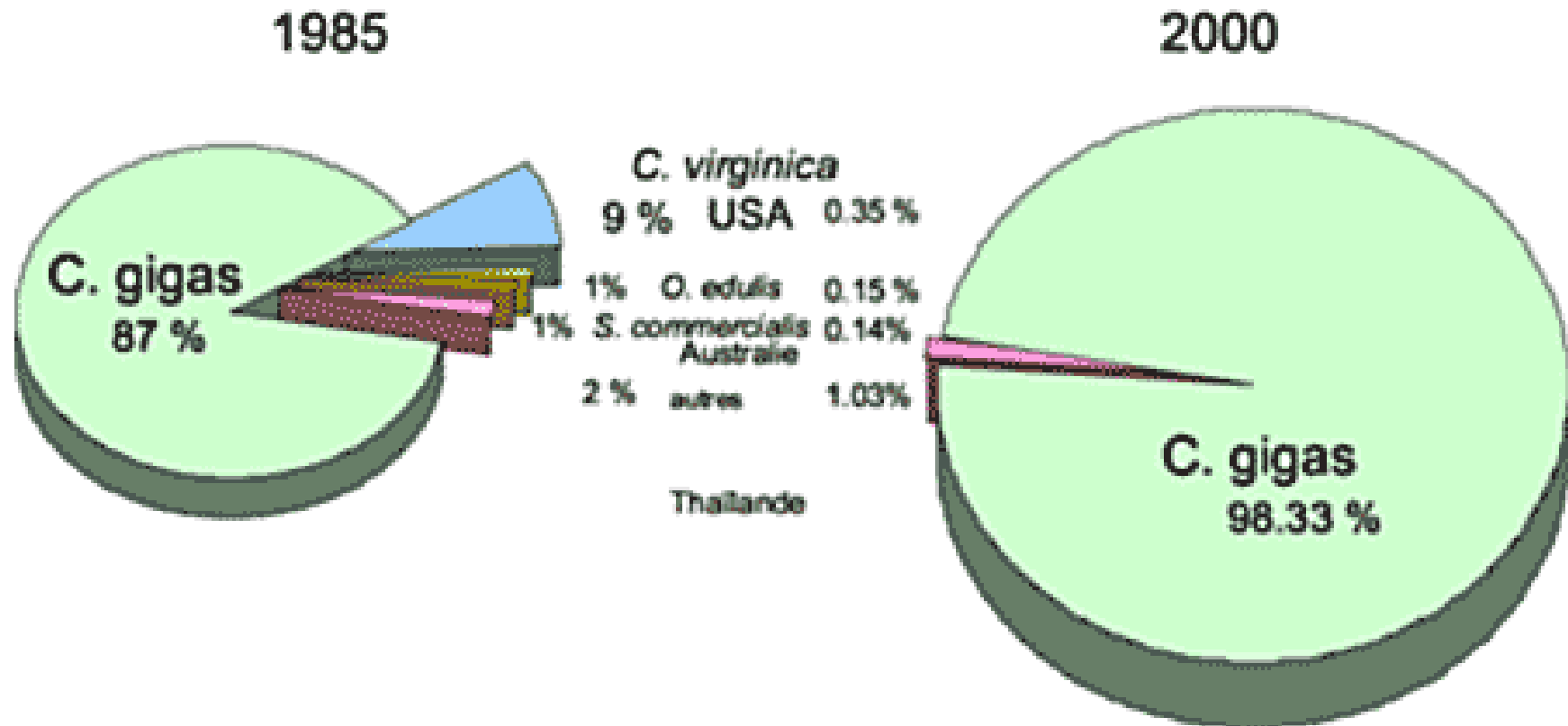
AQUA2006, Fish Genetics and Breeding, May 11th



Oysters in the world

About 100 leaving Ostreidae species
At least 18 species consumed by humans

Graphe 17



Introductions

Elton, 1958: "The greatest agency of all that spreads marine animals to new quarters of the world must be the business of oyster culture."

Ruesink et al. (2005) review:

182 records of introduction in 73 countries or smaller regions

66 were of *C. gigas*

France has been the recipient of the most introduced species

Population genetics tracing not "recorded" introductions for reliable oyster farming

Examples

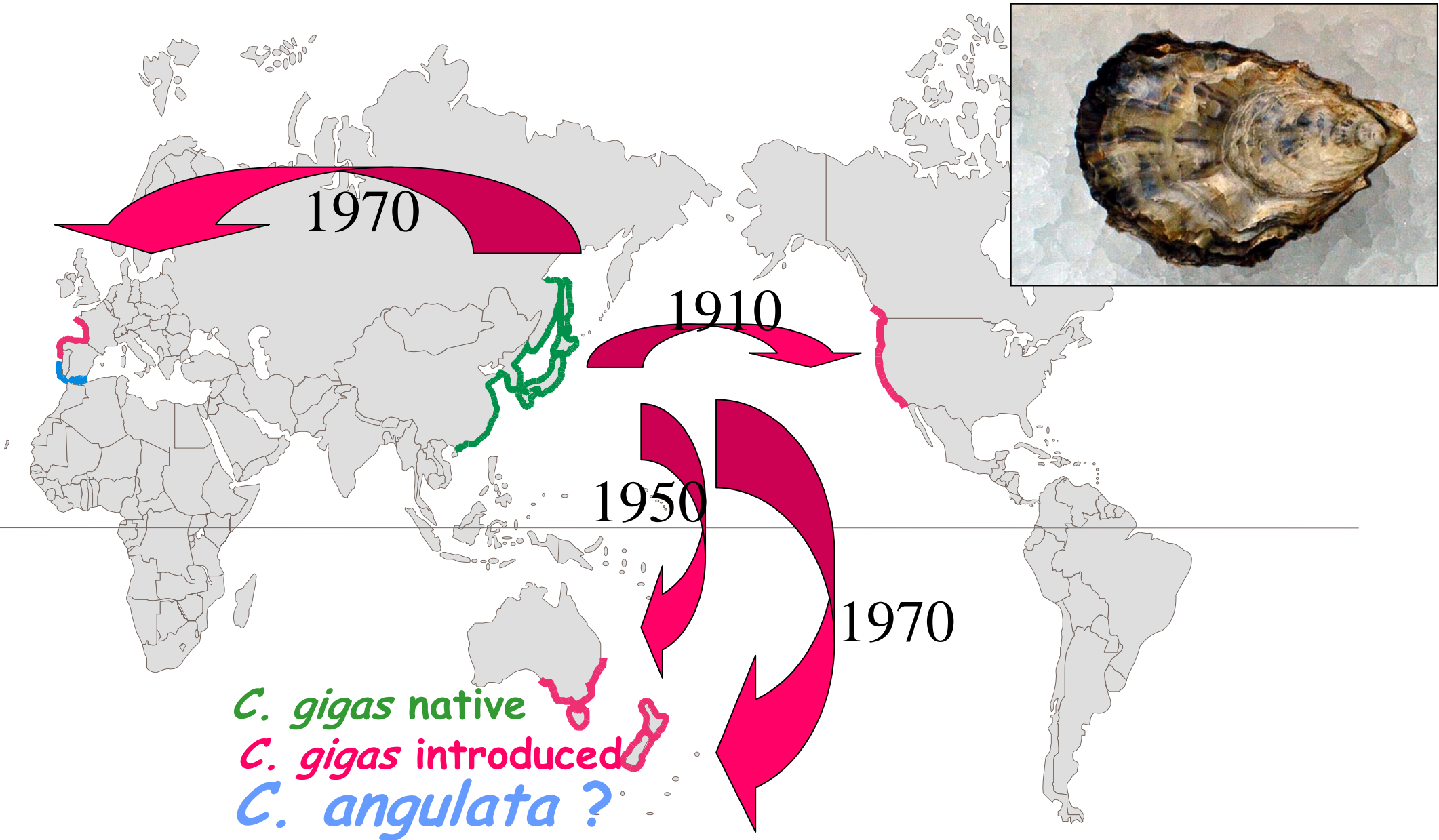
Crassostrea gigas and *C. angulata*

Crassostrea gasar

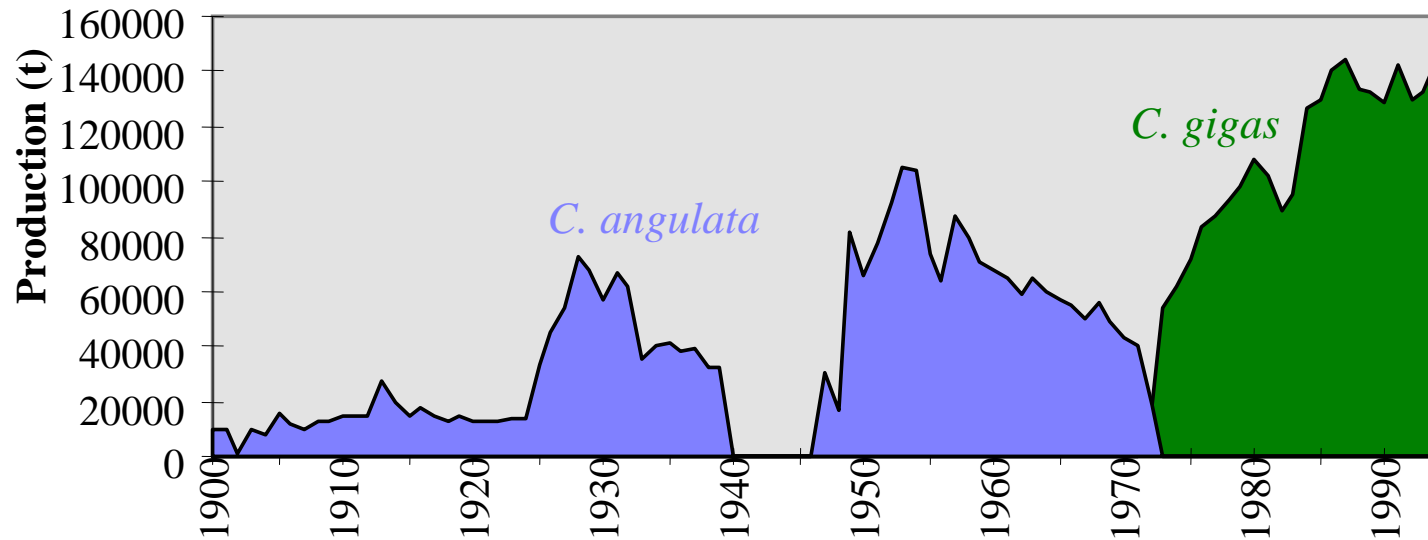
Ostrea stentina

Ostrea edulis

Crassostrea gigas and *C. angulata*



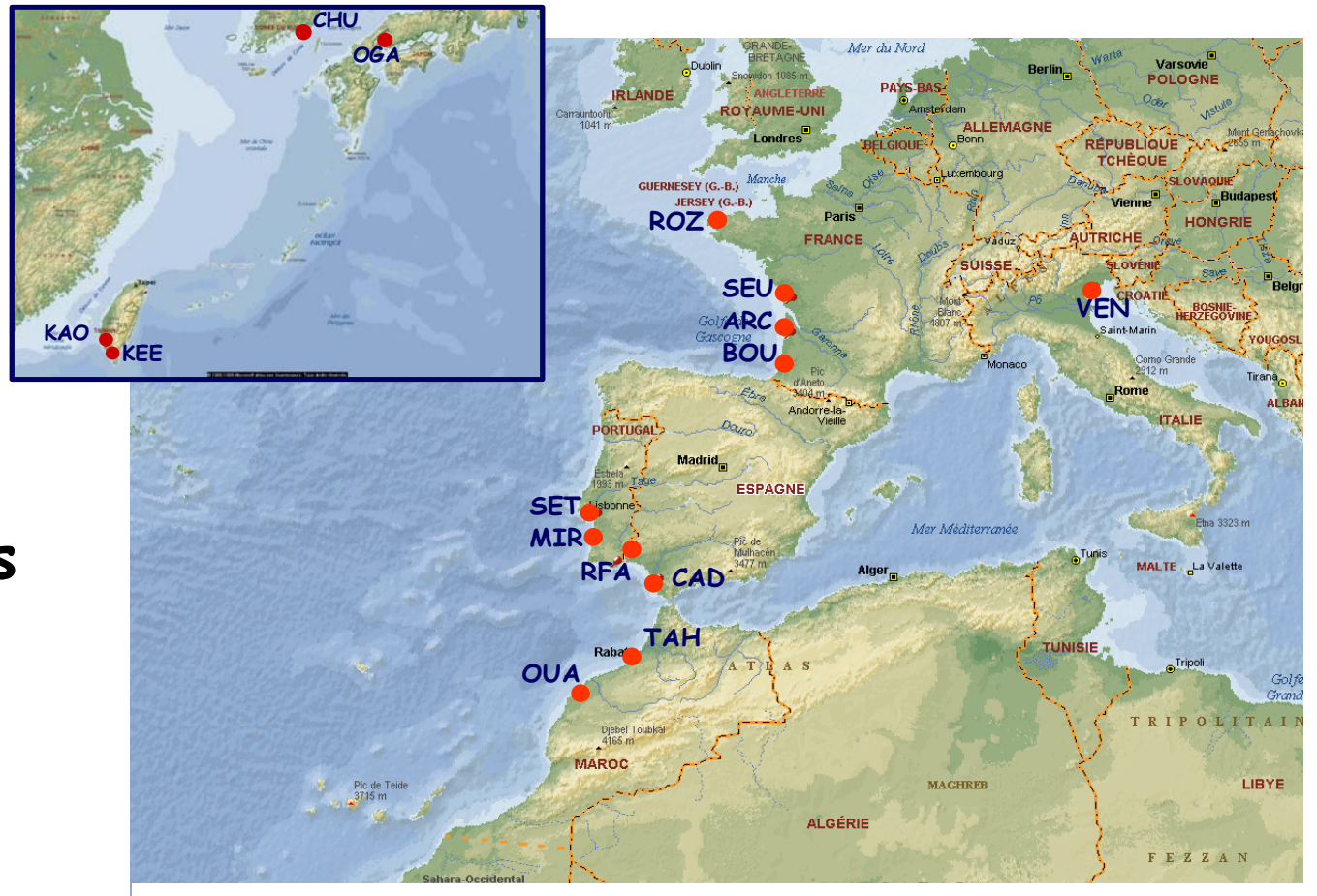
Crassostrea angulata



- 1819** : Lamarck named the Portuguese oyster « *C. angulata* »
- 1868** : Introduction of *C. angulata* from Portugal
- 1970** : Gill disease and subsequent introduction of *C. gigas*
- 1974** : "Portuguese and Japanese oysters are the same species" (Menzel) (Buroker et al., 1979) (Huvet et al., 2001)

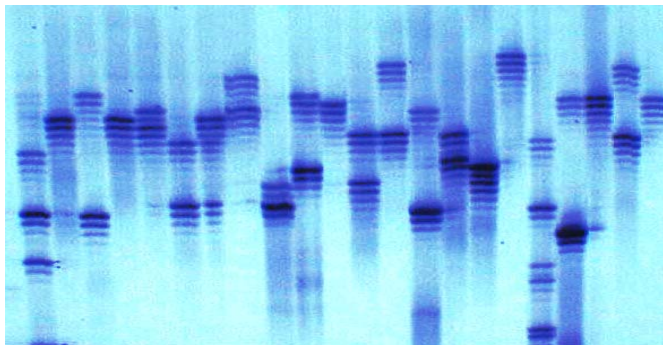
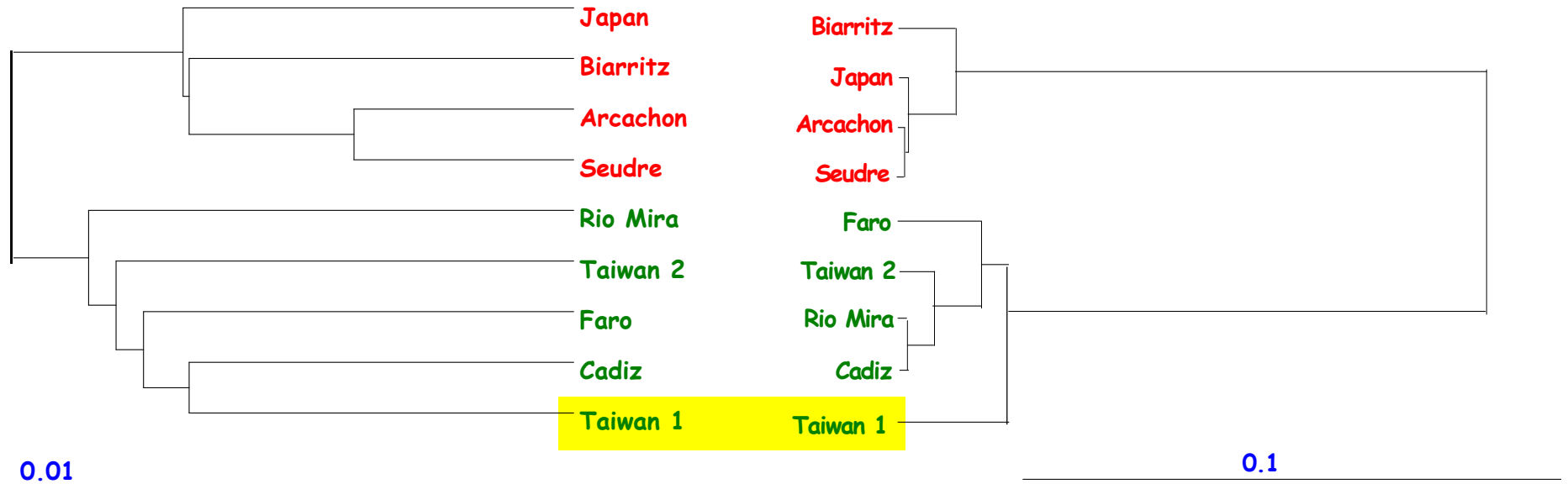
Pacific and Portuguese oysters: origin and relationships

- 15 populations
- 595 individuals
- 6 mtDNA haplotypes
- 3 microsatellite loci



C. angulata ≠ *C. gigas*

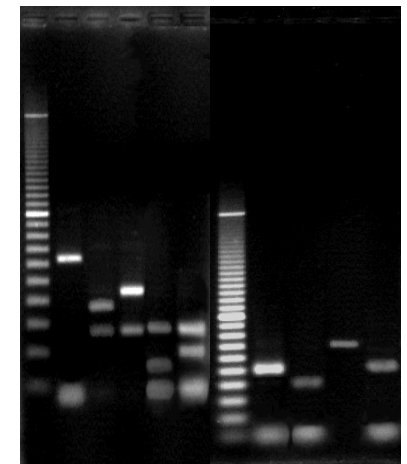
the Asian origin of *C. angulata*



15 populations
595 individuals

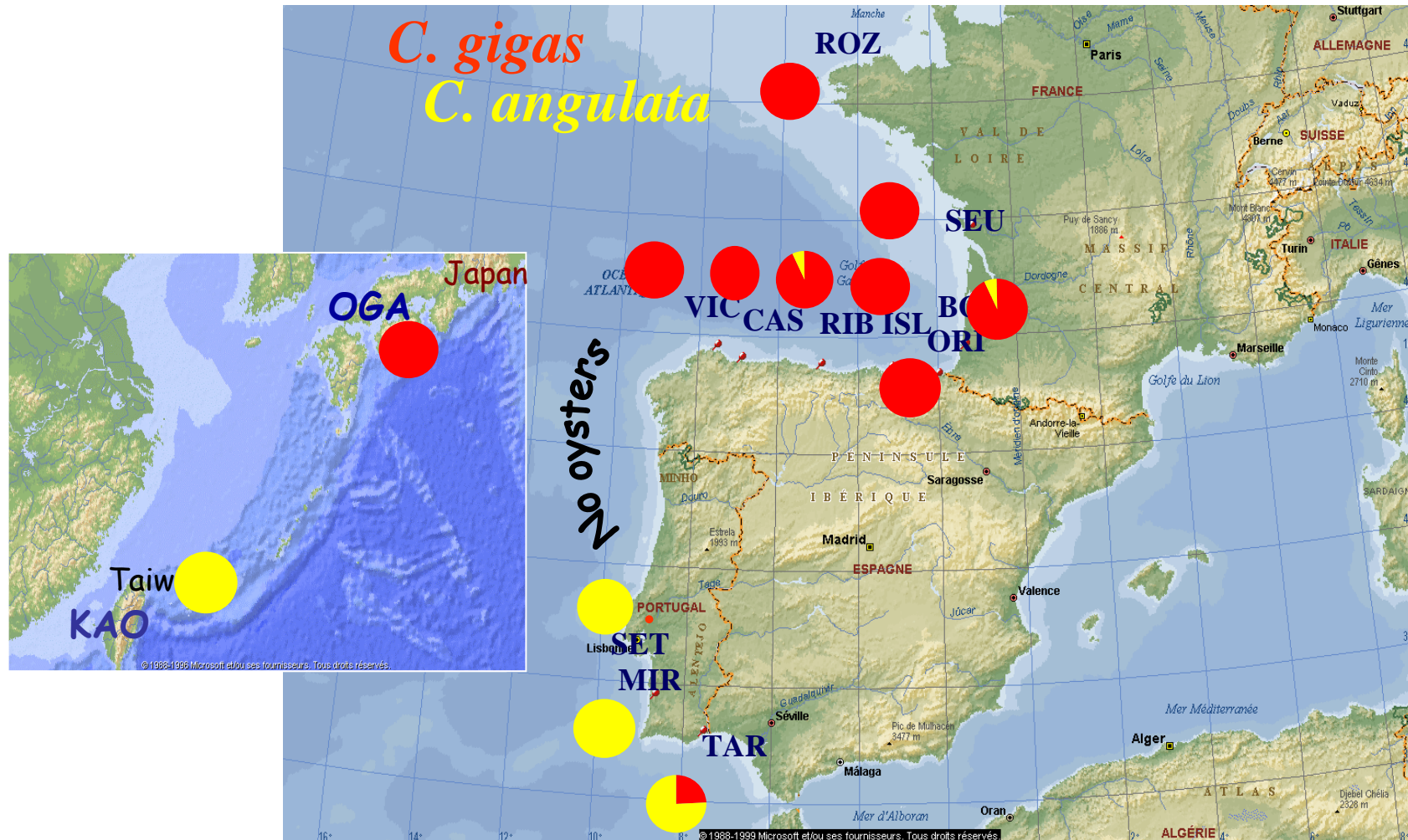
6 mtDNA haplotypes
3 microsatellite loci

(Huvet et al., 2000)



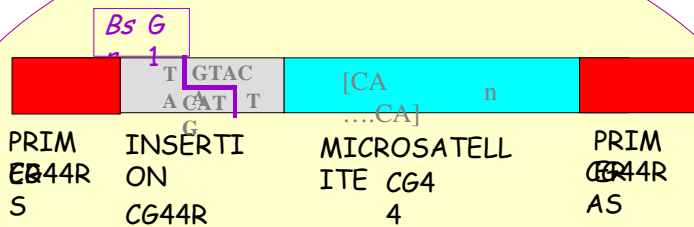
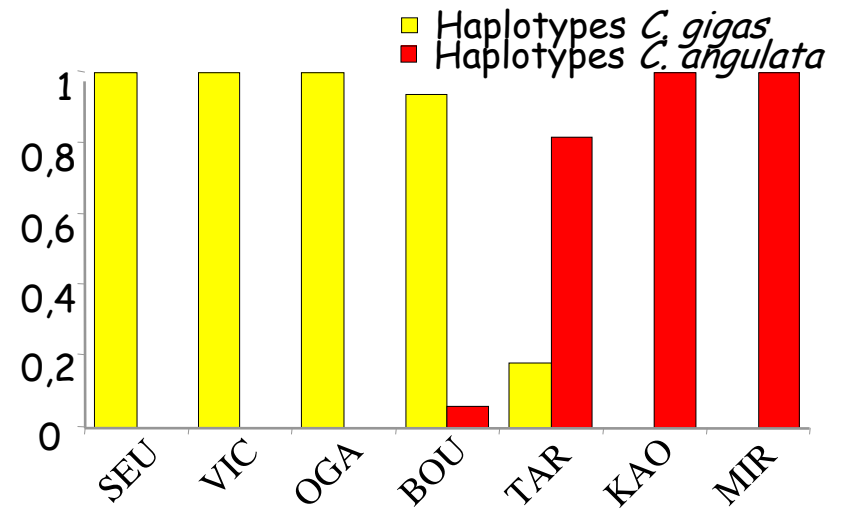
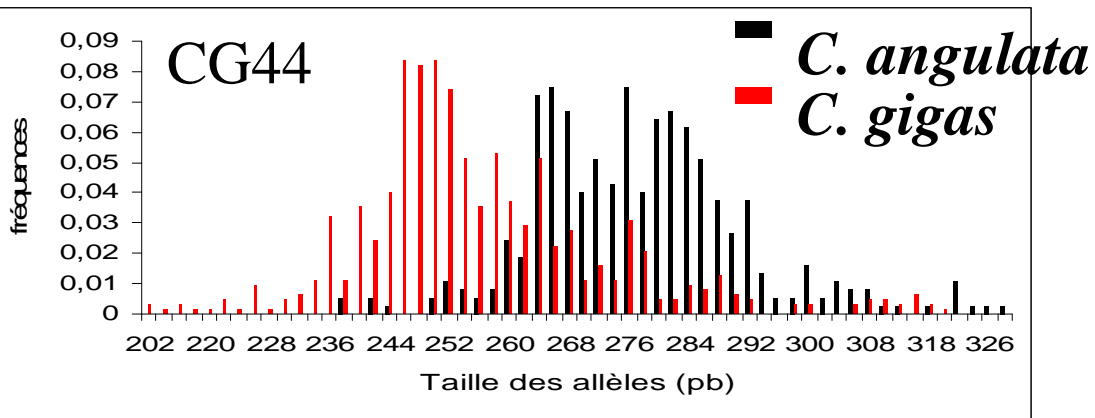
(Boudry et al., 1998)

Crassostrea angulata endangered in Europe ?

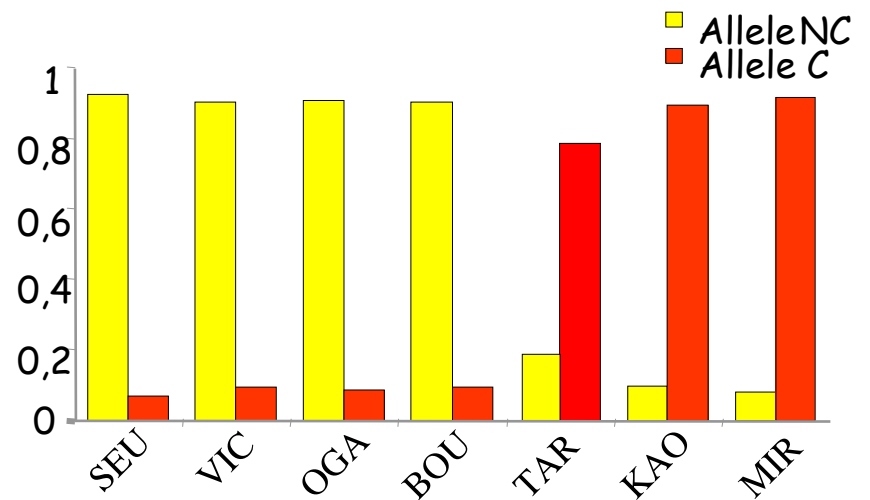
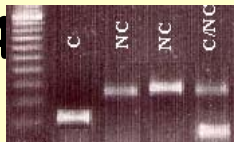


(Fabioux et al., 2002)

C. angulata and *C. gigas* hybridizing in natural populations in Europe ?

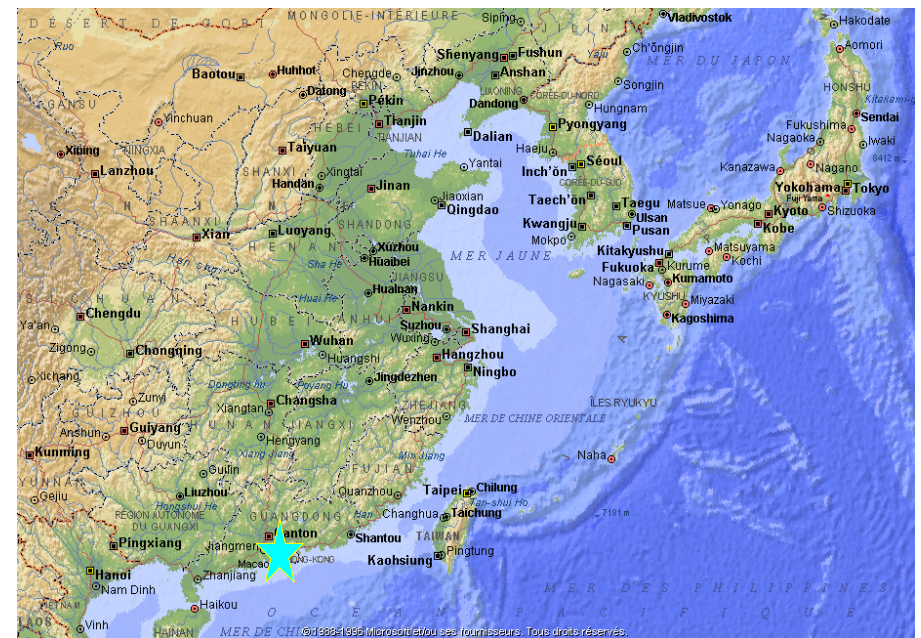
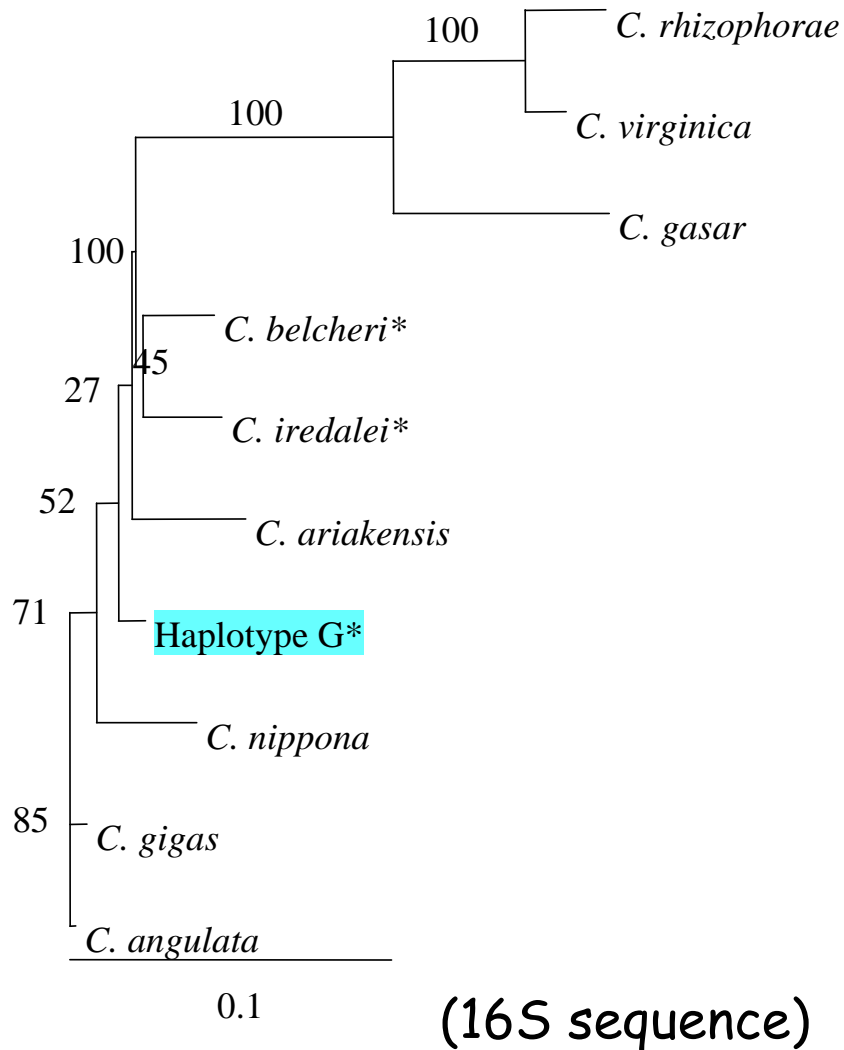


RFLP marker:
CG44



Crassostrea angulata in China ?

C. hongkongensis ? (Lam & Morton, 2003)

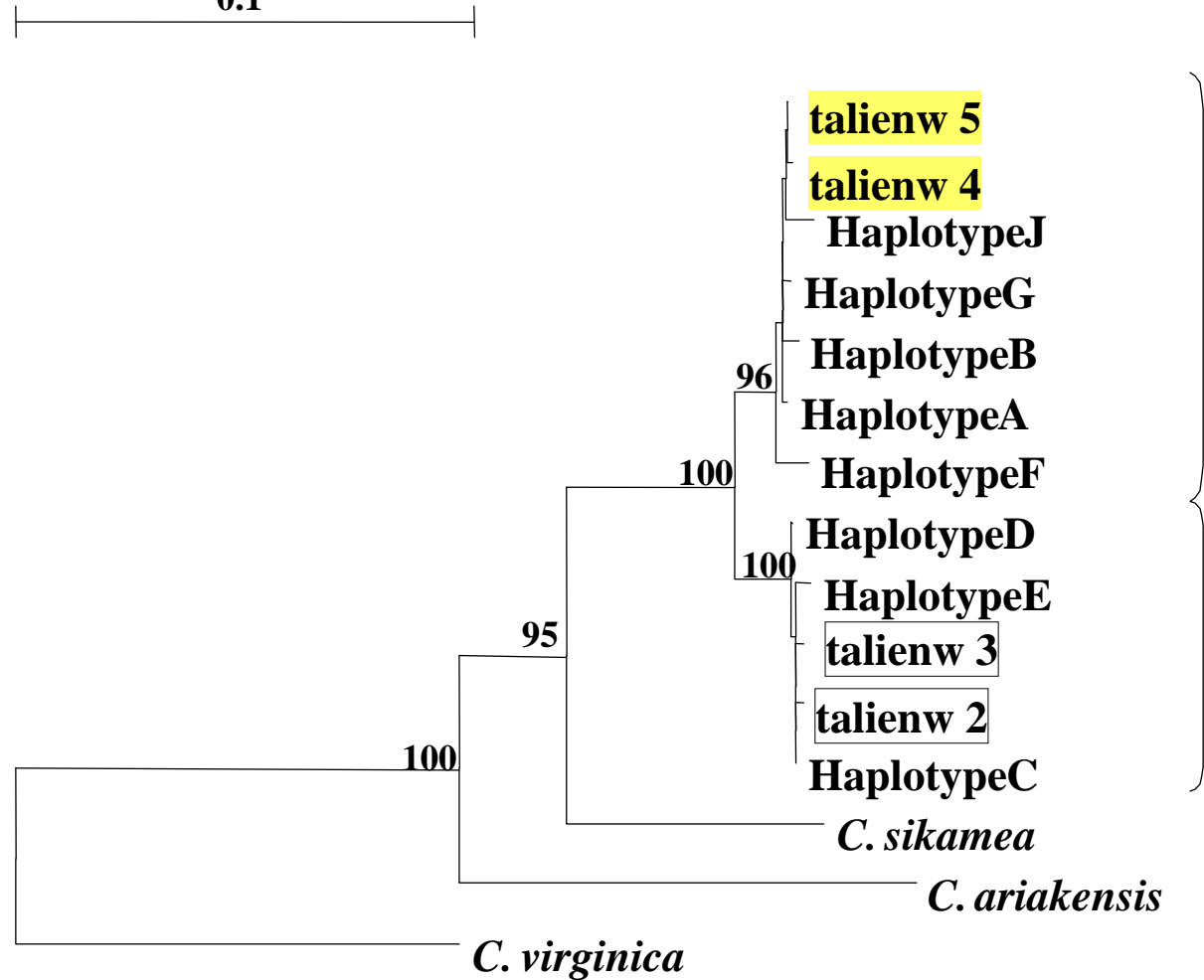


(Boudry et al., 2003)

Crassostrea angulata in China ?

« Daliawan oyster » *C. talienwhanensis* (Yu et al., 2003)

0.1



C. angulata

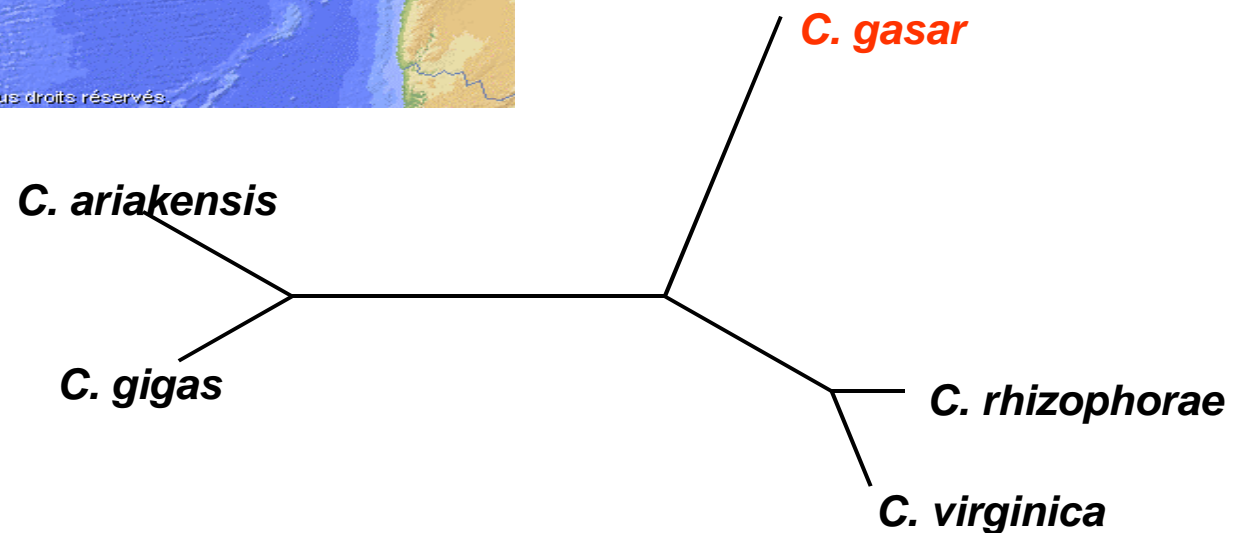
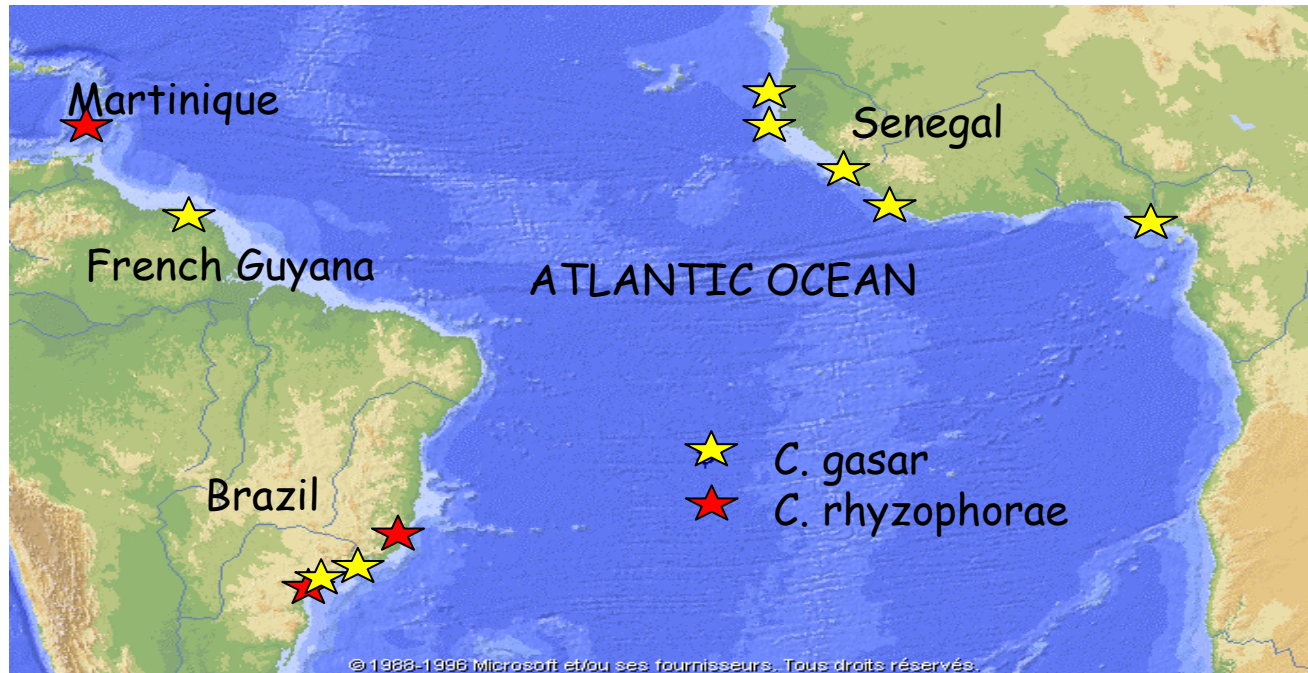
C. gigas



(COI sequence)

(Lapègue et al., 2004)

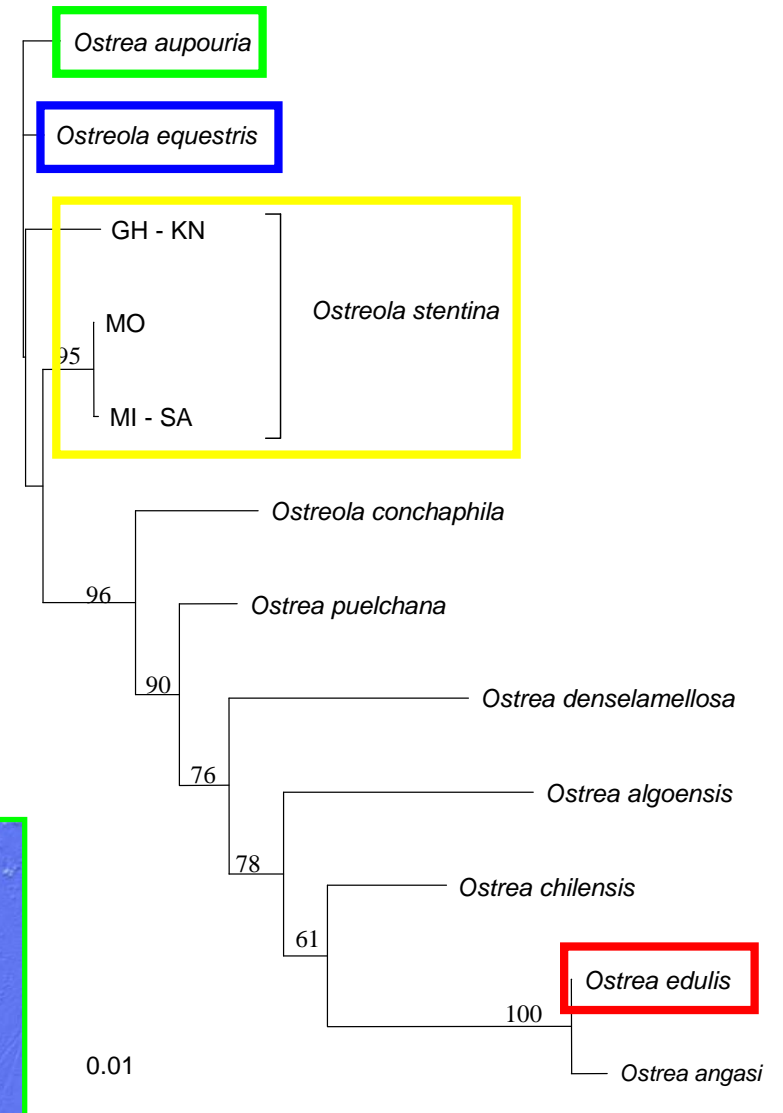
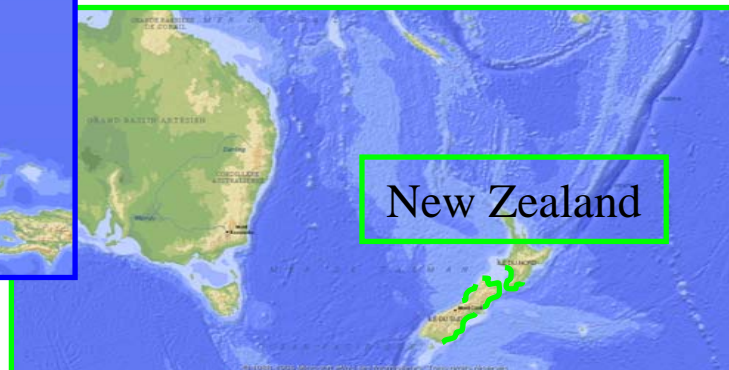
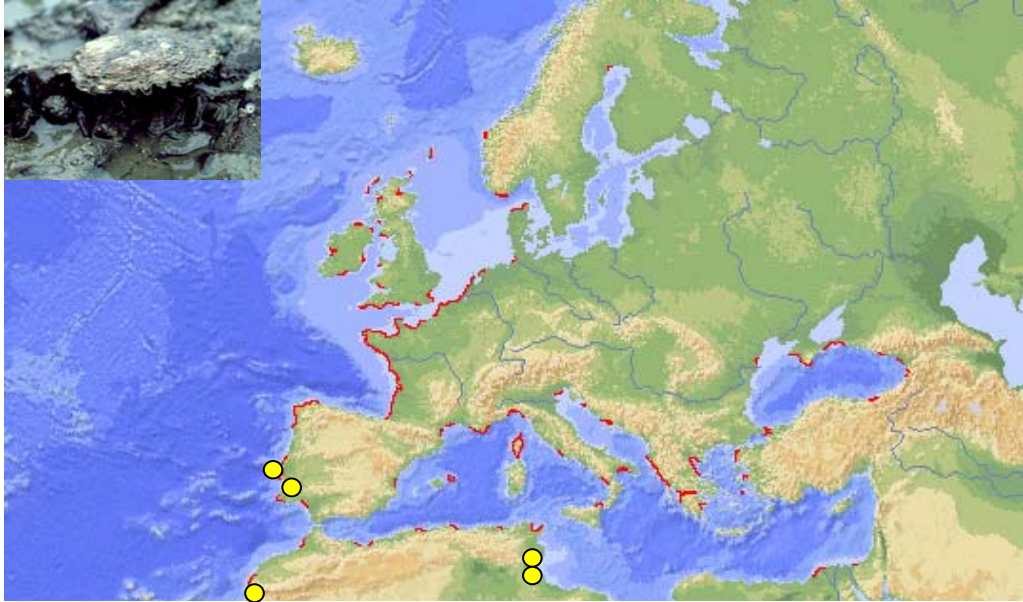
The « mangrove » oyster *Crassostrea gasar* is not only present in Africa



(Lapègue et al., 2002)

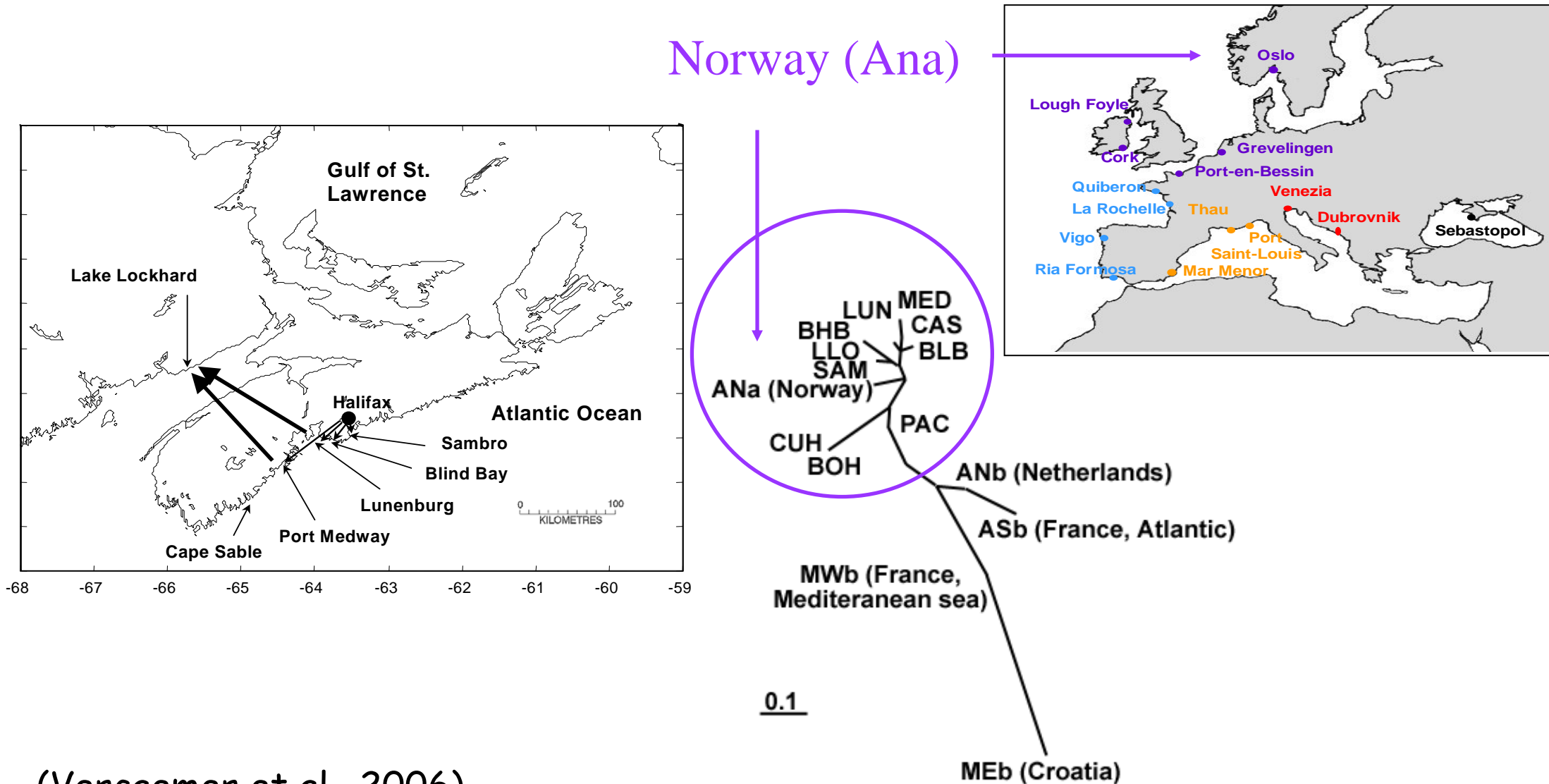
0.01

O. edulis et *O. stentina*, two very different « European » flat oyster



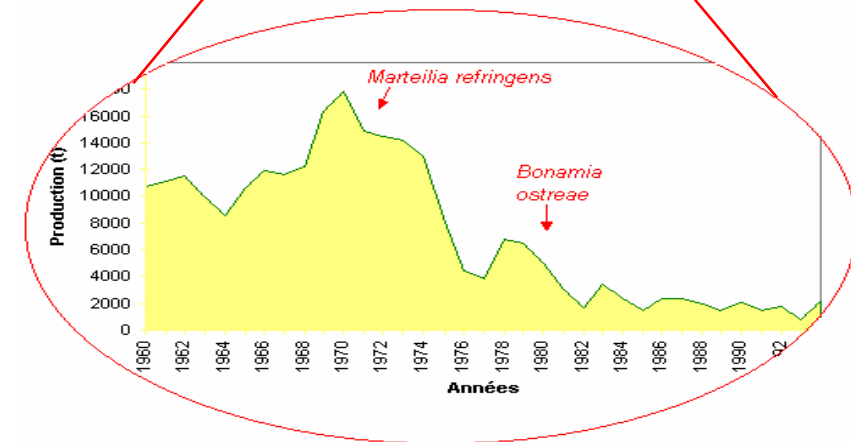
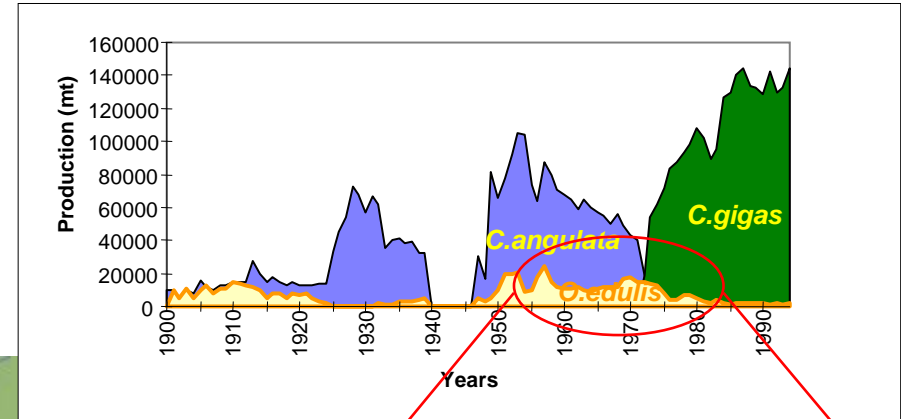
(Lapègue et al., 2006)

The European flat oyster introduced in Nova Scotia, Canada



(Vercaemer et al., 2006)

Understanding natural populations as potential resources for breeding



Ostrea edulis in 2004 :

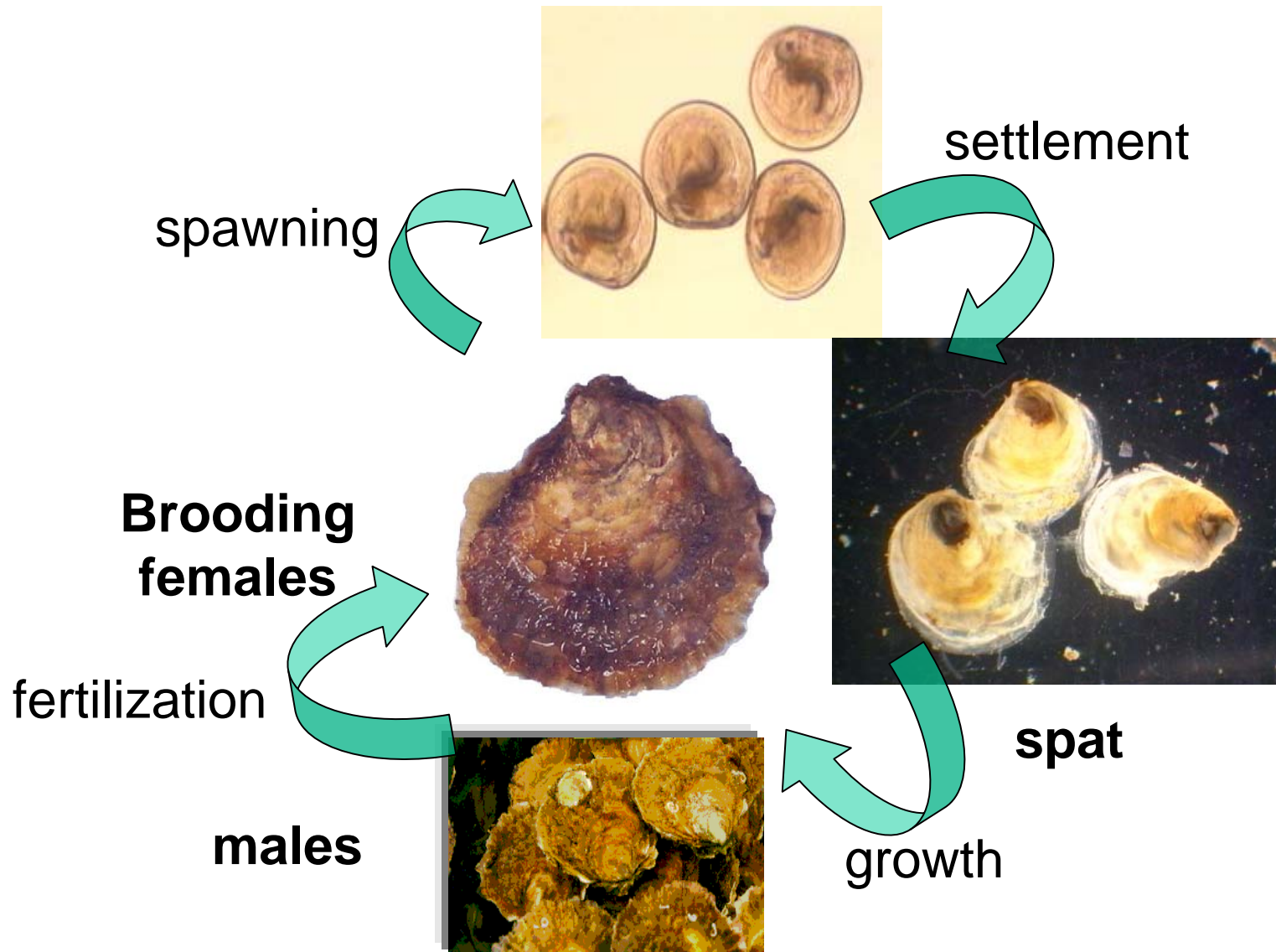
World production = 5.100 t/y
French production = 1.500 t/y

(Jaziri, 1985)



Reproductive cycle

planktonic larvae (± 10 days)



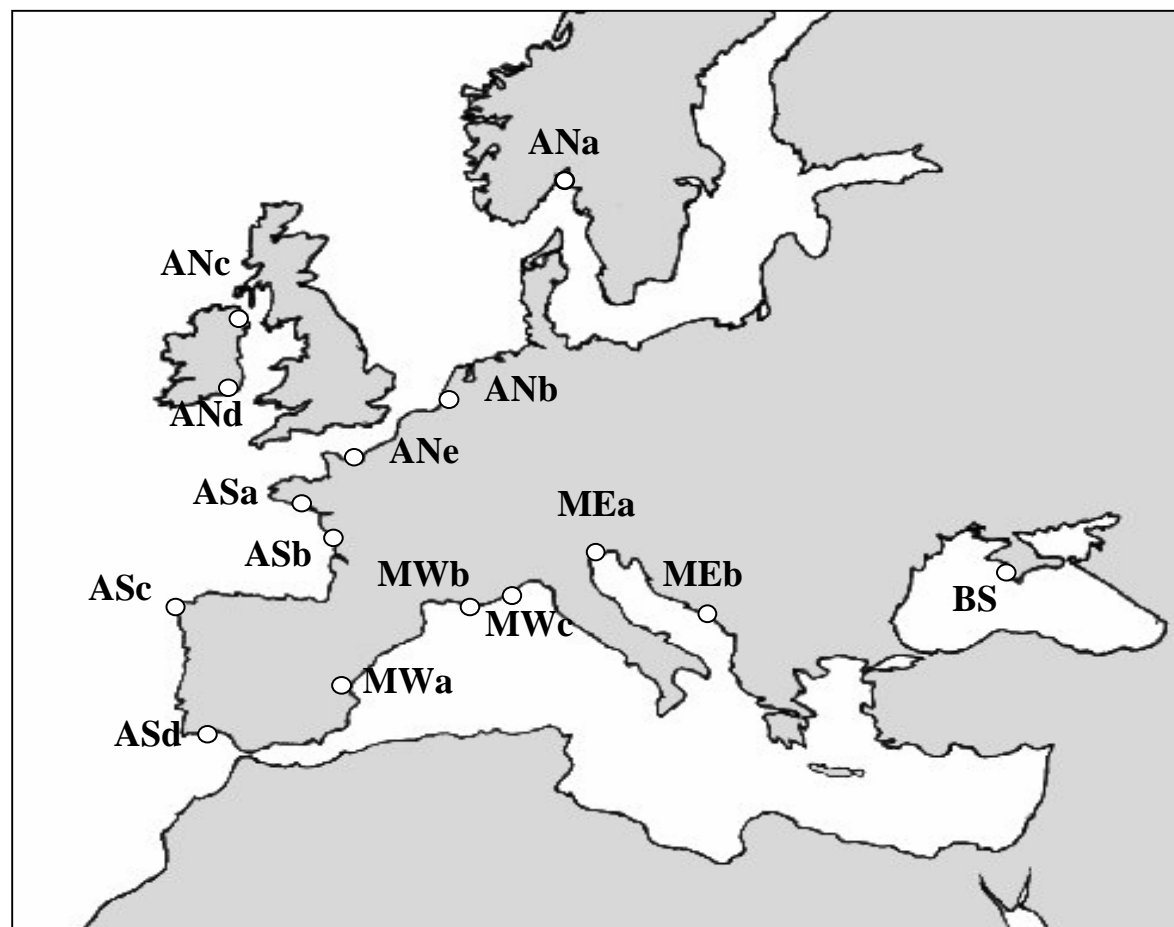
European sampling

Markers

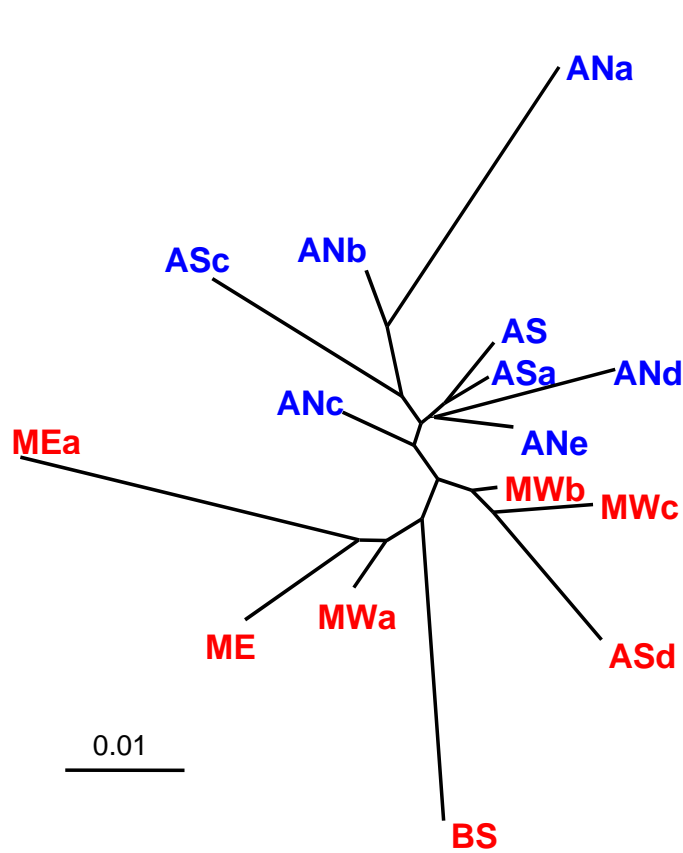
- Allozyme loci (Saavedra *et al.*, 1993, 1995)
- microsatellite loci (Launey *et al.*, 2002)
- 12S rRNA SSCP (Diaz Almela *et al.*, 2004)

Sampling

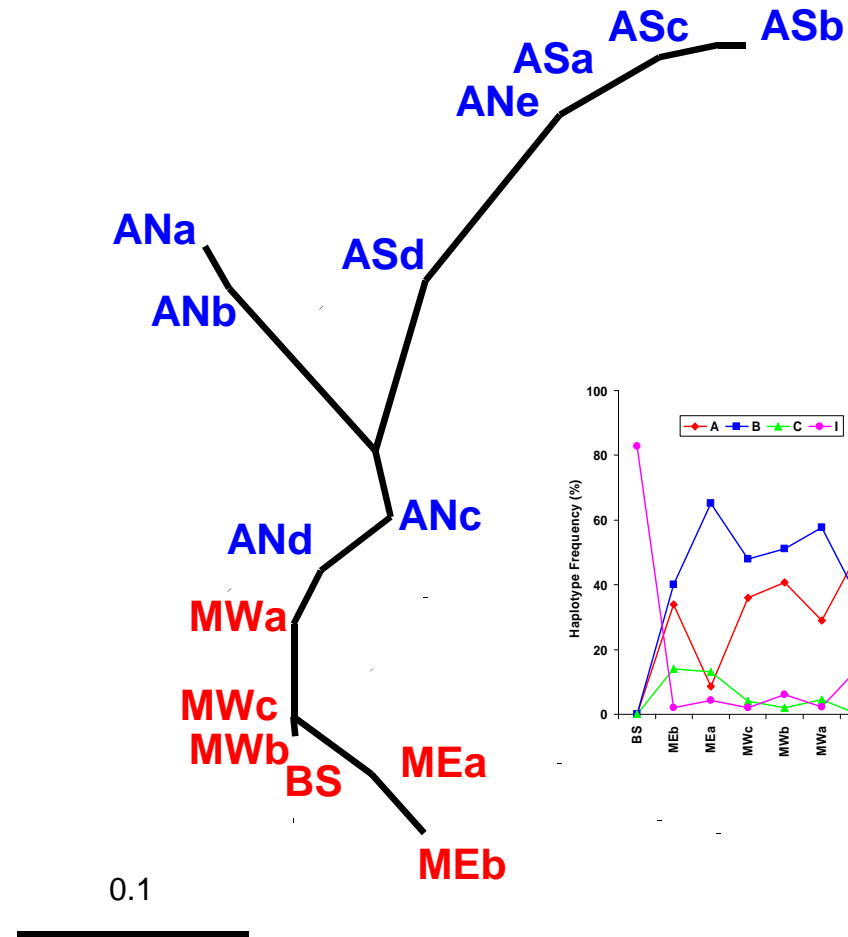
- 15 populations sampled
- 14 to 50 individuals per location



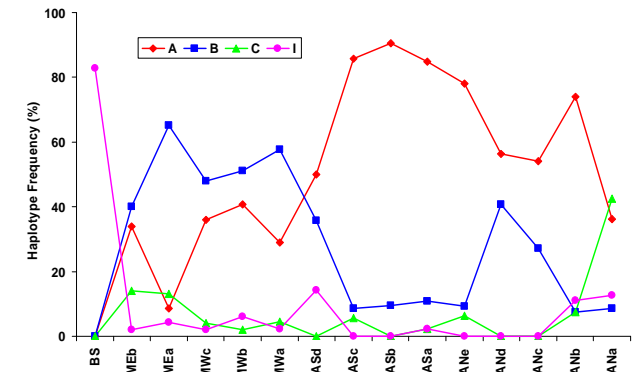
Mediterranean sea versus Atlantic ocean differentiation



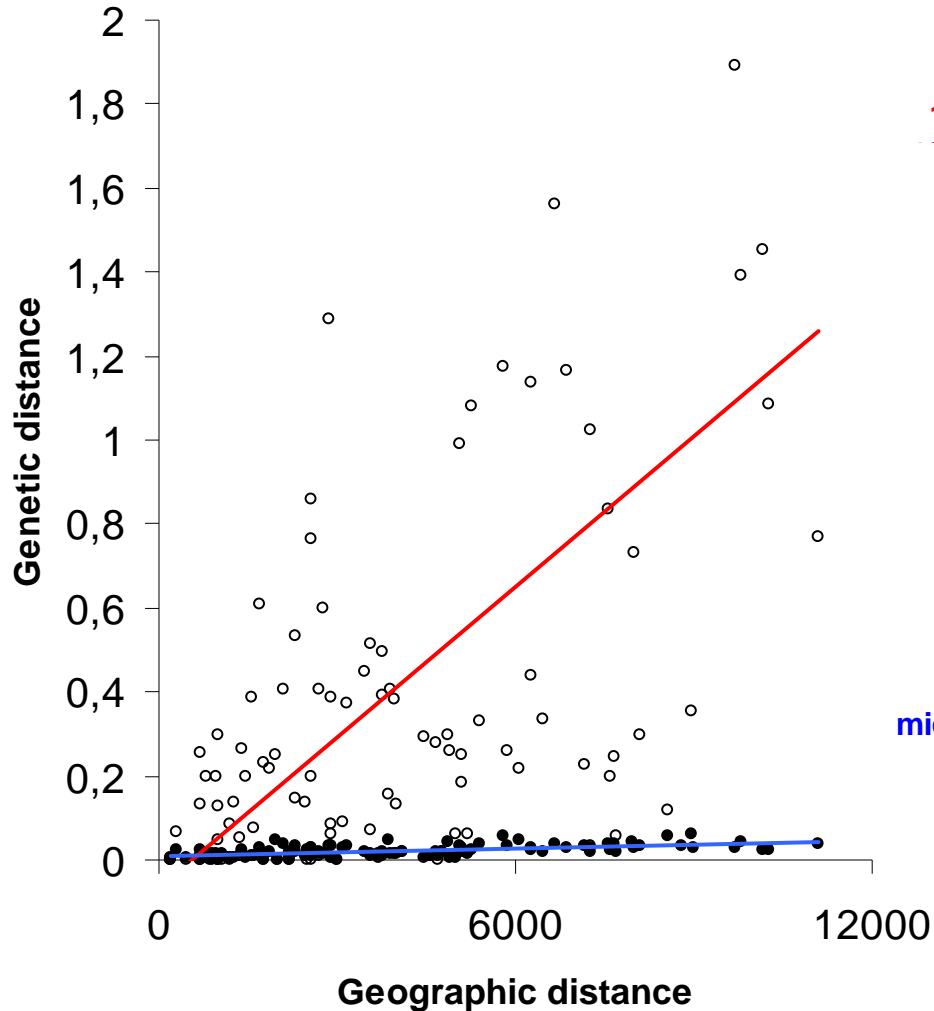
Microsatellites



12S rRNA

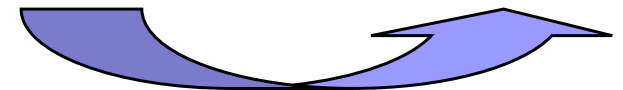


Isolation by distance

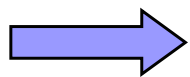


Microsatellites
 $F_{st} = 0.019^{***}$

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



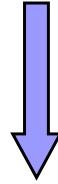
10 X



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

Sweepstake hypothesis

Great fecundity, very high larval mortality



variance in the number of offspring contributed by adults to subsequent generations.

Hedgecock et al., 2006 :

Cohort sampled during 10 days near Sète

	adult	cohort
number of alleles	23	14
		family structure

Parentage analyses

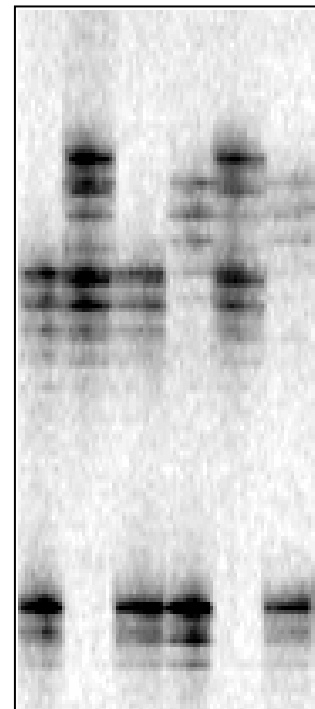


Natural population

Locus OeduU2

Allele 188
→

Allele 156
→

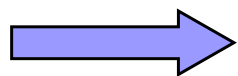
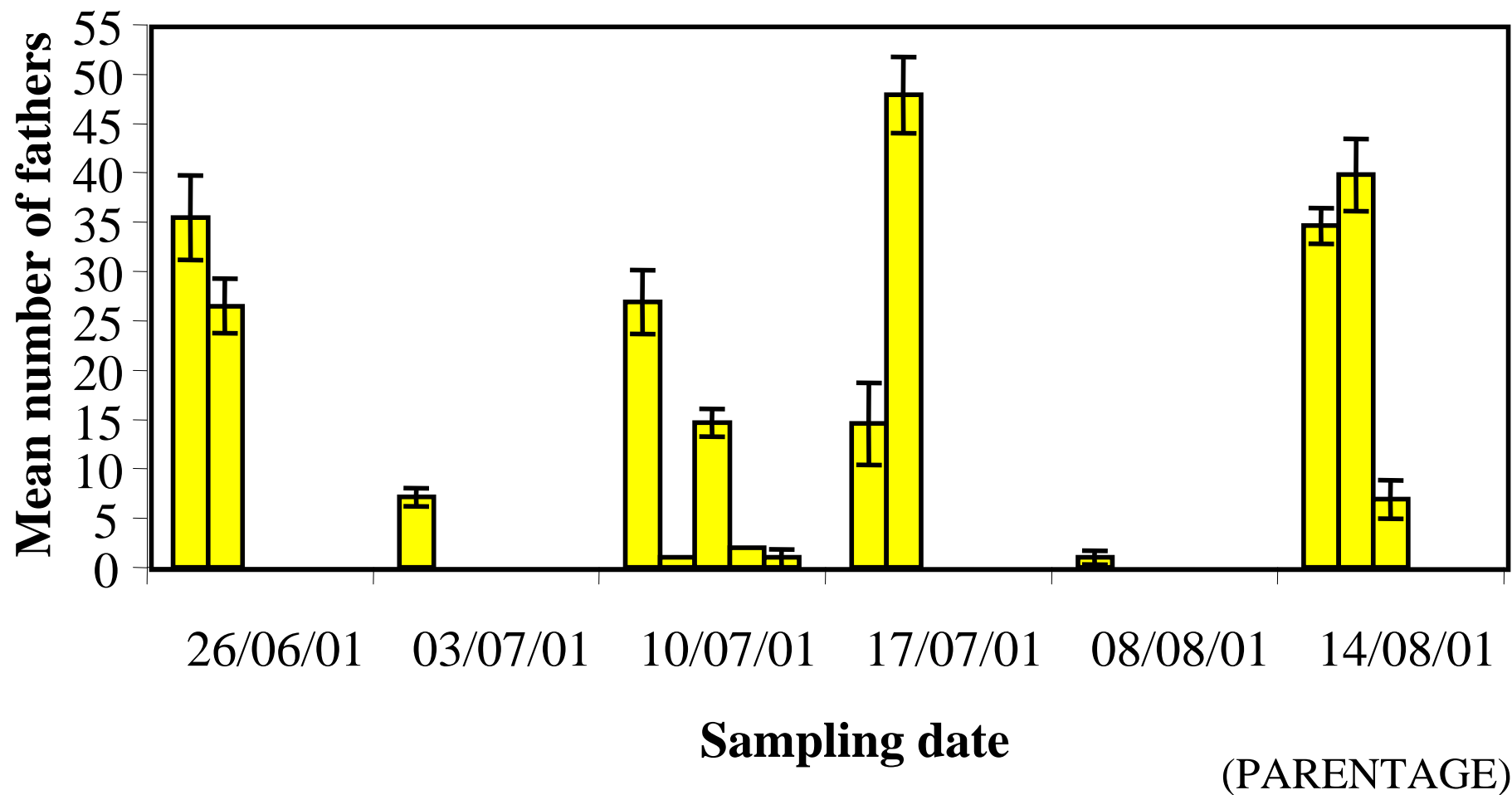


Experimental population



How many males / female ?

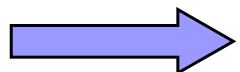
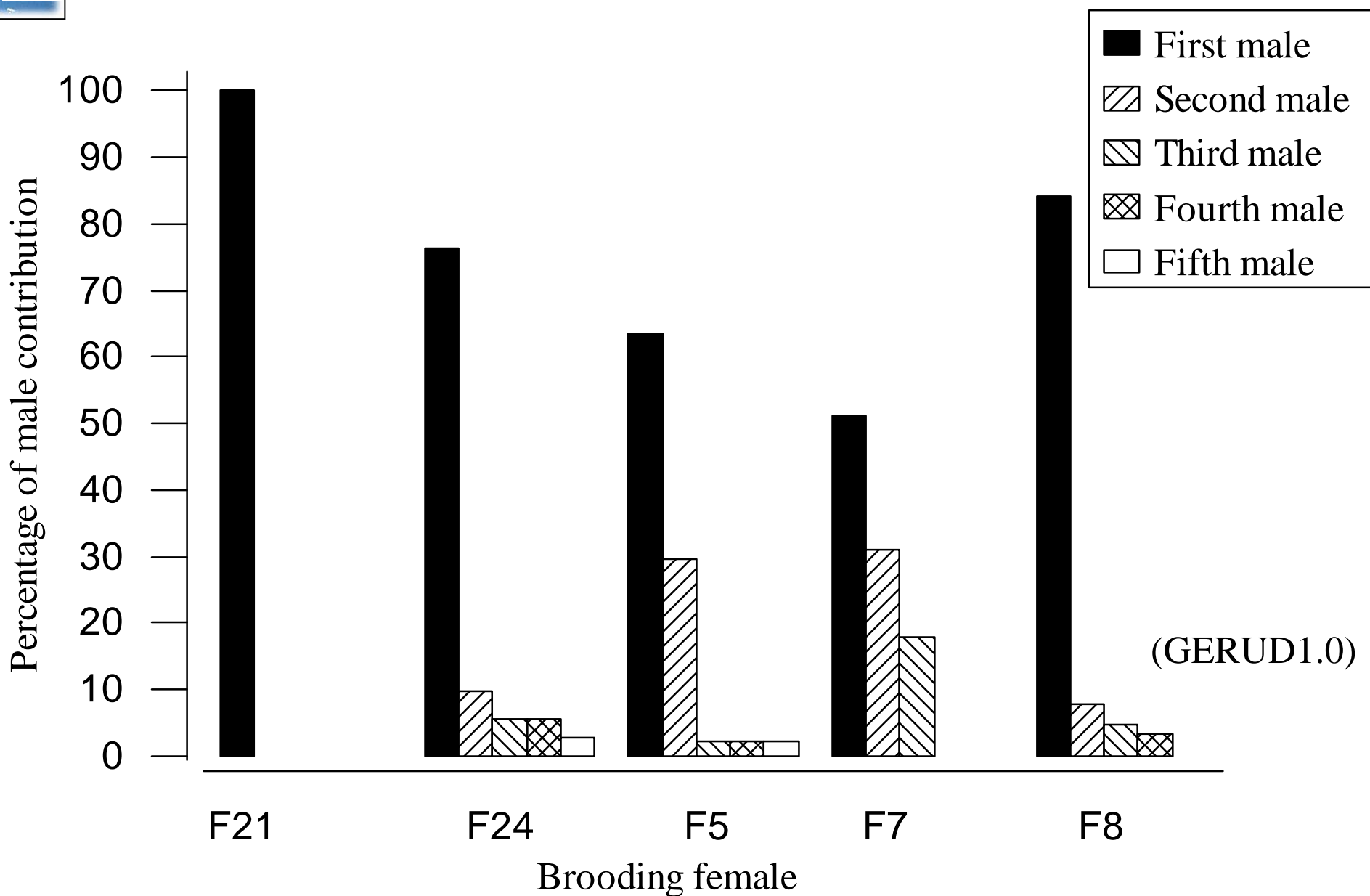
Number of males fertilizing
each of the 14 brooding females



Very variable number of males / female (1 to >40)



Male contribution within a female



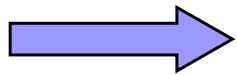
Variance in individual male contribution : male with the highest contribution fathered from 51 to 100% of larvae



From the individual to the population

These results are of limited value to answer our initial question because :

- The population size is poorly known
- The effective population size is difficult to estimate
- The spatial distribution of males and females is unknown
- The effective sex ratio unknown
- ...

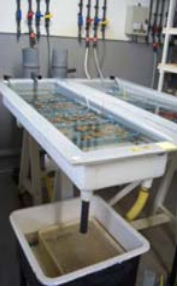


What about experimental (hatchery) populations ?



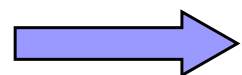
Differentiation of the 6 mass spawnings

$F_{st} \times 100$

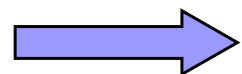


	14/03/03	17/03/03	20/03/03	22/03/03	28/03/03	30/03/03
<u>Adults</u>	3 ^{***}	5.5 ^{***}	3.1 ^{***}	1.5 ^{***}	1.3 ^{***}	1.2 ^{***}
14/03/03	-	11.9 ^{***}	7.1 ^{***}	5.9 ^{***}	5 ^{***}	5.1 ^{***}
17/03/03	-	-	9.8 ^{***}	6.8 ^{***}	7.9 ^{***}	7.8 ^{***}
20/03/03	-	-	-	0.7 ^{**}	4.2 ^{***}	4.5 ^{***}
22/03/03	-	-	-	-	3 ^{***}	3 ^{***}
28/03/03	-	-	-	-	-	1.8 ^{***}

	14	14+17	14+17+20	14+17+20+22	14+17+20+22+28	14+17+20+22+28+30
<u>Adult</u>	3 ^{***}	1.1 ^{***}	0.6 ^{**}	0.6 ^{**}	0.3 [*]	0.2 ^{NS}

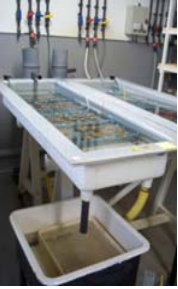


Highly significant genetic differentiations (1-12%)

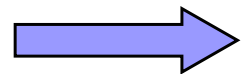


While pooling, genetic differentiation becomes blurred

Effective population size

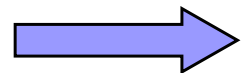


	Ne (temporal method)
14/03/03	21.0 [12.4 36.2]
17/03/03	12.5 [7.8 19.6]
20/03/03	21.0 [12.3 36.5]
22/03/03	22.3 [12.9 39.6]
28/03/03	33.2 [17.8 70.1]
30/03/03	29.6 [16.3 58.9]



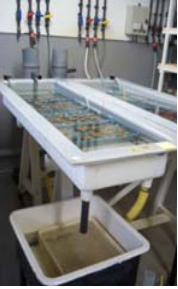
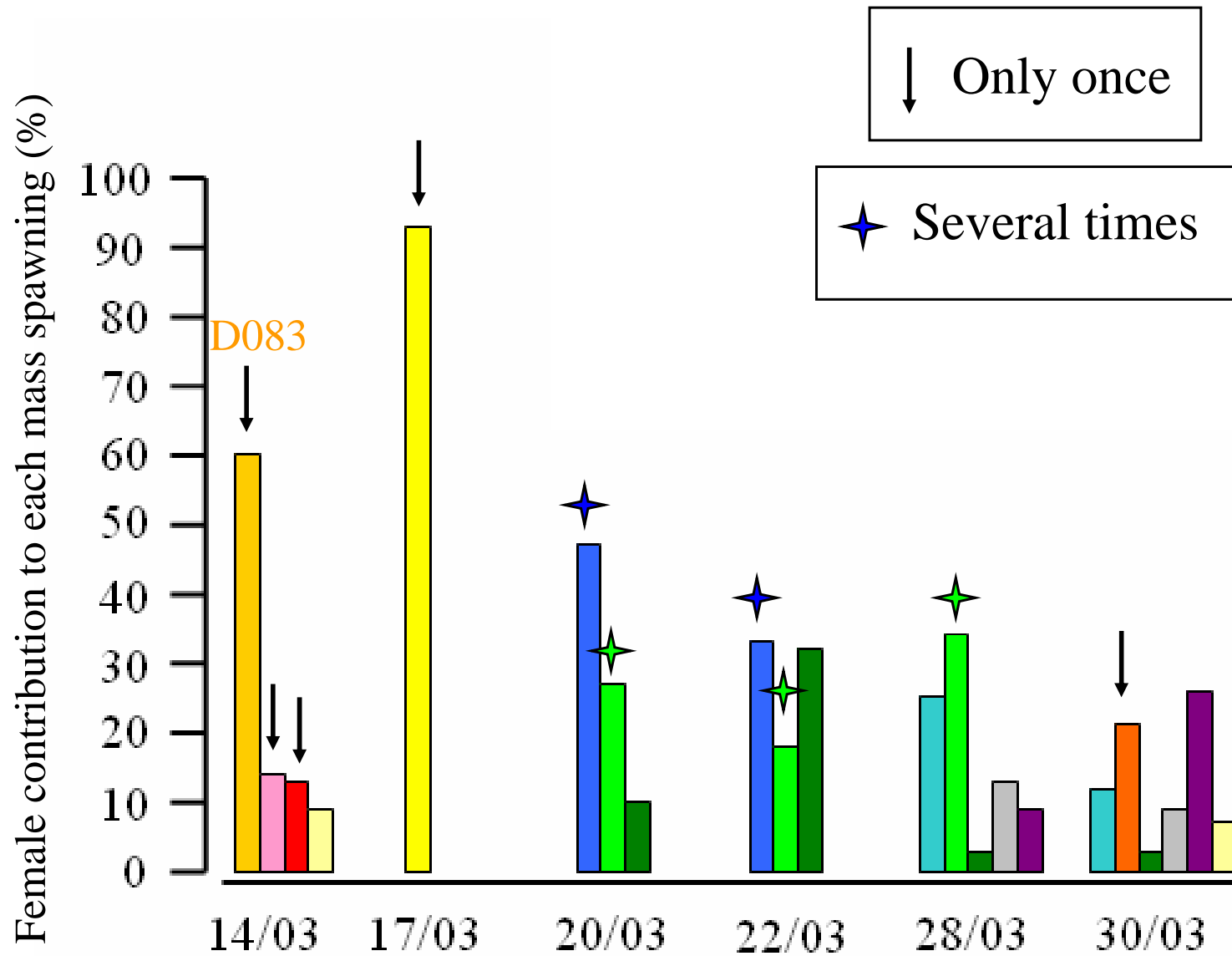
Relatively small Ne (12-33)

Genetic variability of a cohort : $N_e < 20$ (Hedgewood *et al.*, 2006)

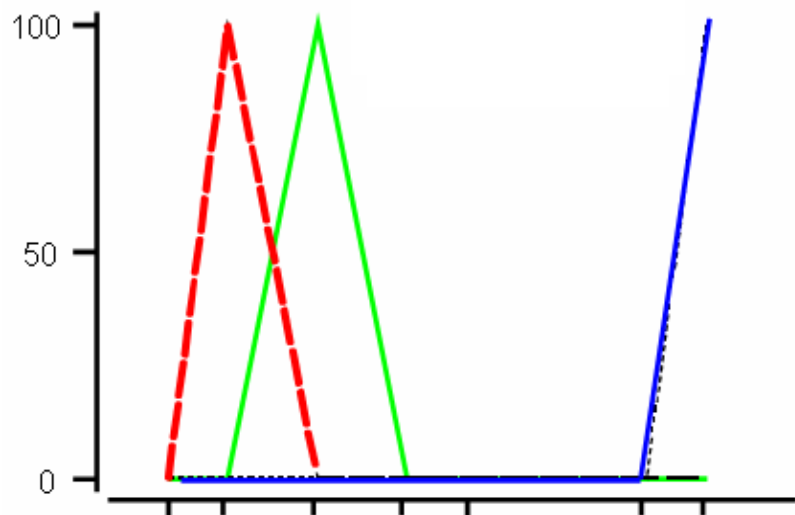
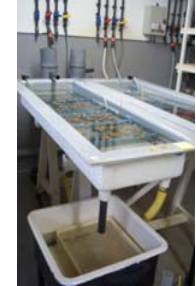


Large 95% CI : small number of markers

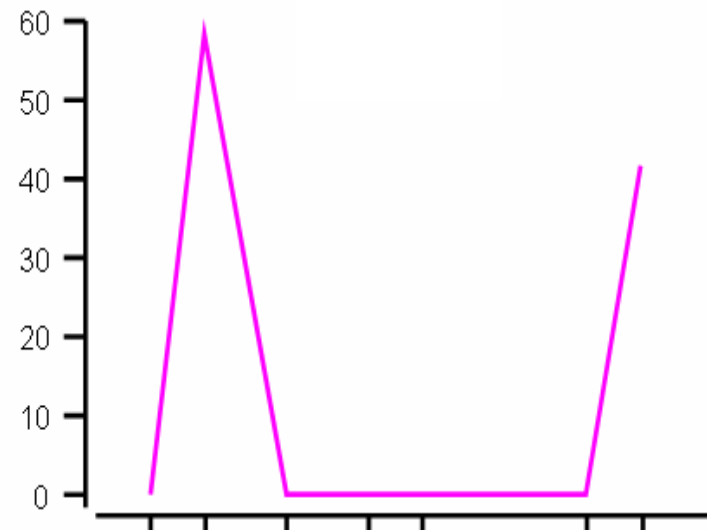
Female contribution



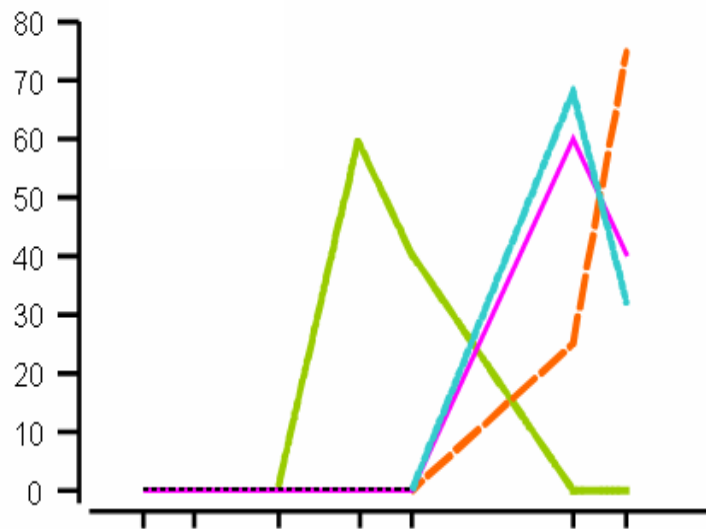
Dynamics of Female release



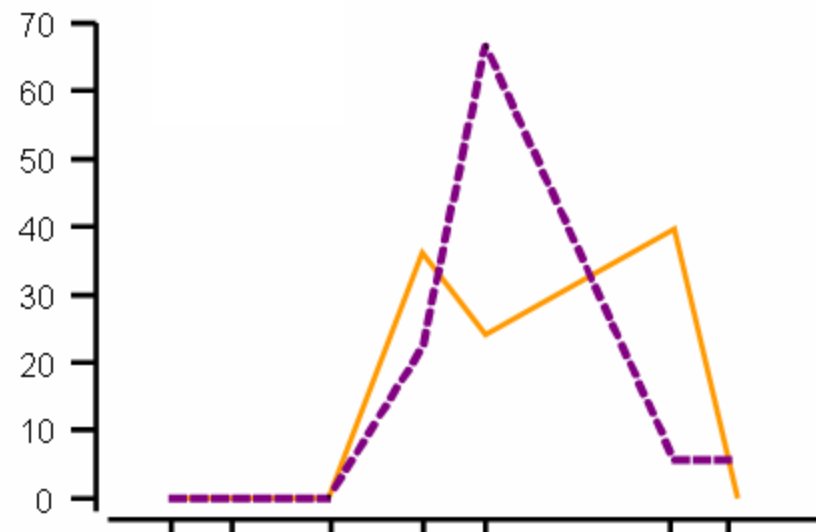
In 1 mass spawning



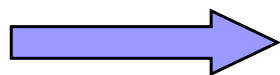
In 2 spaced mass spawning



In 2 successive mass spawning

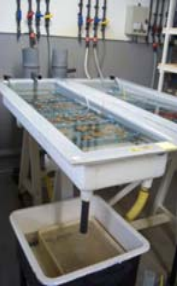
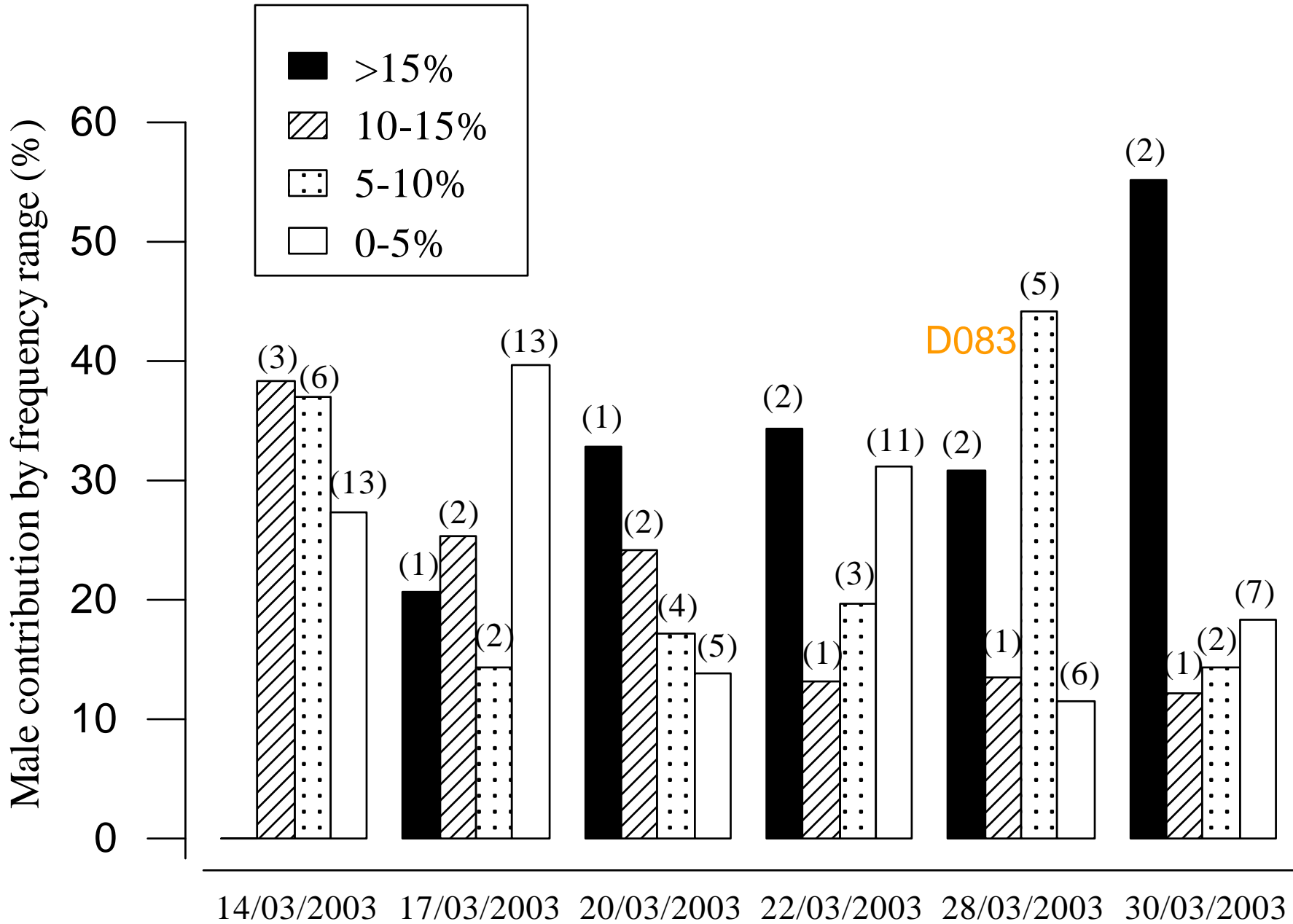


In 3 or 4 mass spawning

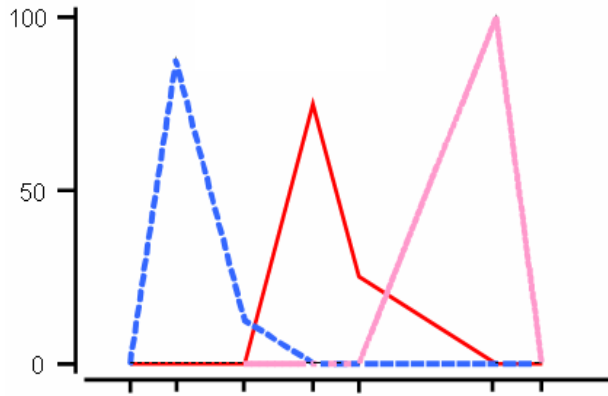
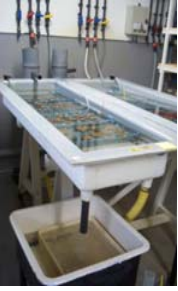


Variable extension in time of larval release and its dynamics

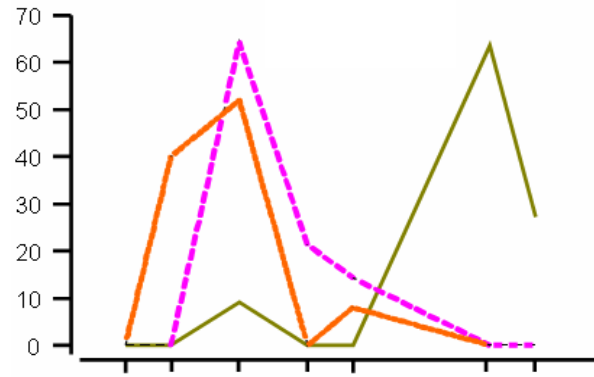
Male contribution



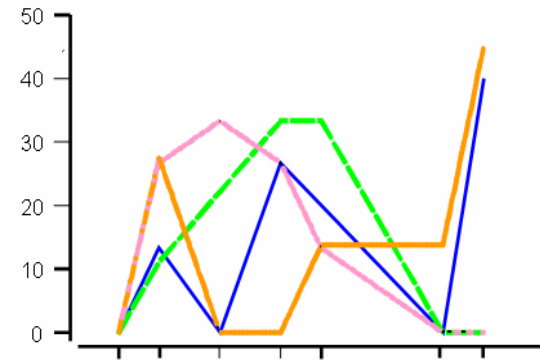
Dynamics of Male release



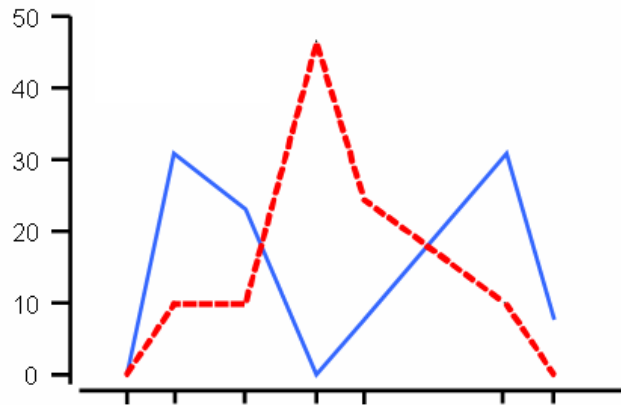
In 1 or 2 mass spawning



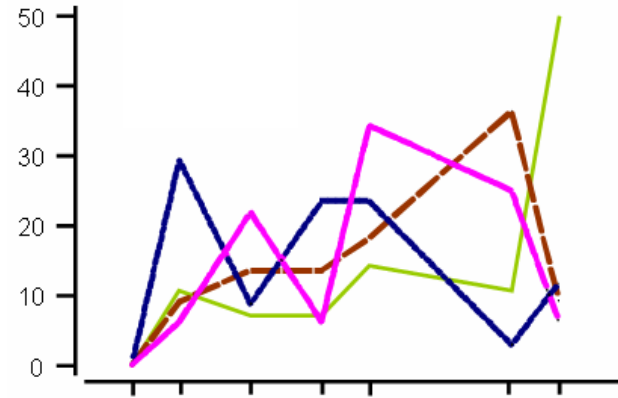
In 3 mass spawning



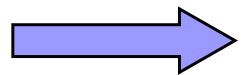
In 4 mass spawning



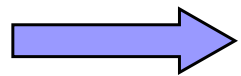
In 5 mass spawning



In 6 mass spawning

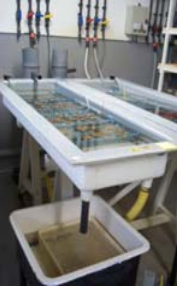


More complex pattern

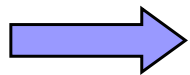


Sperm release can be extended in time

Conclusions



- **Better knowledge of reproductive biology of *O. edulis***
 - Ability to change of sex very quickly (maturation of 2 types of gametes simultaneous)
 - Spawning of one individual does not stimulate the others
 - High variability in the temporal dynamics of emission of gametes and larvae
- **Relative variance in reproductive success in males and females**
 - For most females, spawning events are constituted by the progeny of several males (especially in our hatchery experiment)
 - Most males fertilize several females (hatchery experiment)
 - Effective reproductive period is longer for males than for females (hatchery experiment)



Higher variance in reproductive success
in females than in males

Thank you for your attention

