



# **Genetic basis of summer mortality in juvenile cupped oysters**

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**LPI Brest**

# **Summer mortality in juvenile Pacific oysters**

- Reported for many years and in several countries  
(Koganezawa, 1975 ; Glude, 1975 ; Gouletquer et al., 1998, Cheney et al., 2000...)
- “MOREST”
  - a multidisciplinary program to better understand and face summer mortality
    - pathogens,
    - environmental factors,
    - pollutants...

Genetics objectives : Determine if selective breeding programs could improve survival

# This talk :

## ■ First generation (G1) :

- survival in the field
- variance components for survival
- genetic parameters of survival



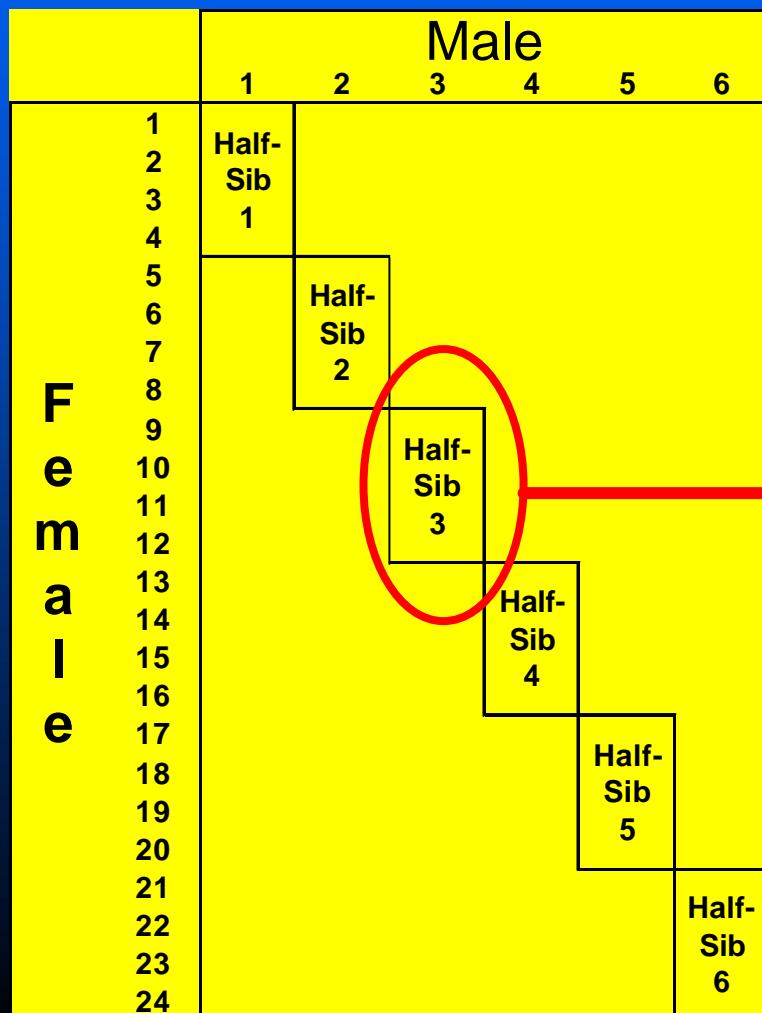
## ■ Second generation (G2) :

- divergent selection :
  - » trial in laboratory
  - » trial in the field : response to selection
- inbred lines :
  - » trials in the field and laboratory



# First generation (G1): material and methods

- 3 sets of 24 nested half-sib crosses giving 72 full-sib families



Half-sib family 3

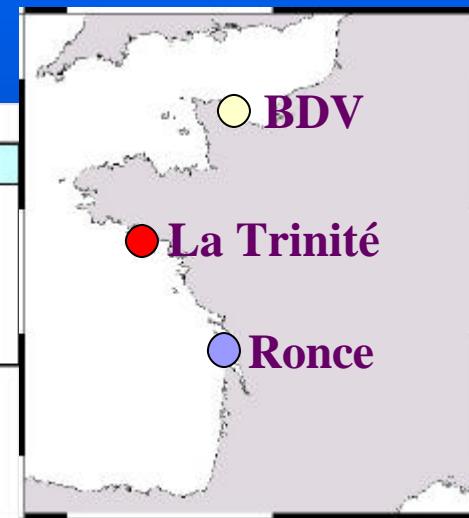
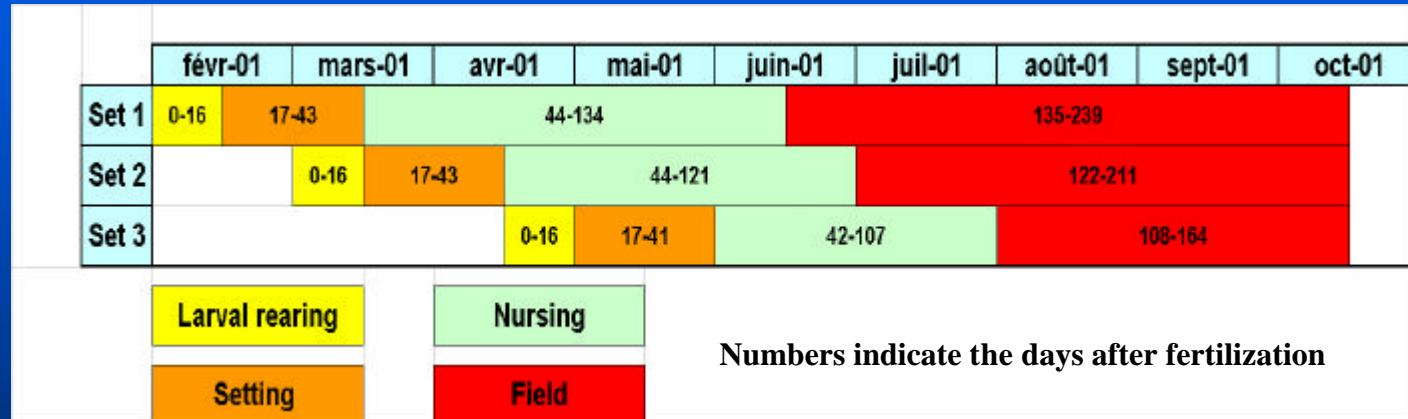
- Family 3-9
  - Family 3-10
  - Family 3-11
  - Family 3-12
- 4 full-sib families

Determination of genetic parameters  
(ANOVA - REML using SAS system)

# G1 : material and methods

- $3 \times 15 = 45$  families tested in 3 sites

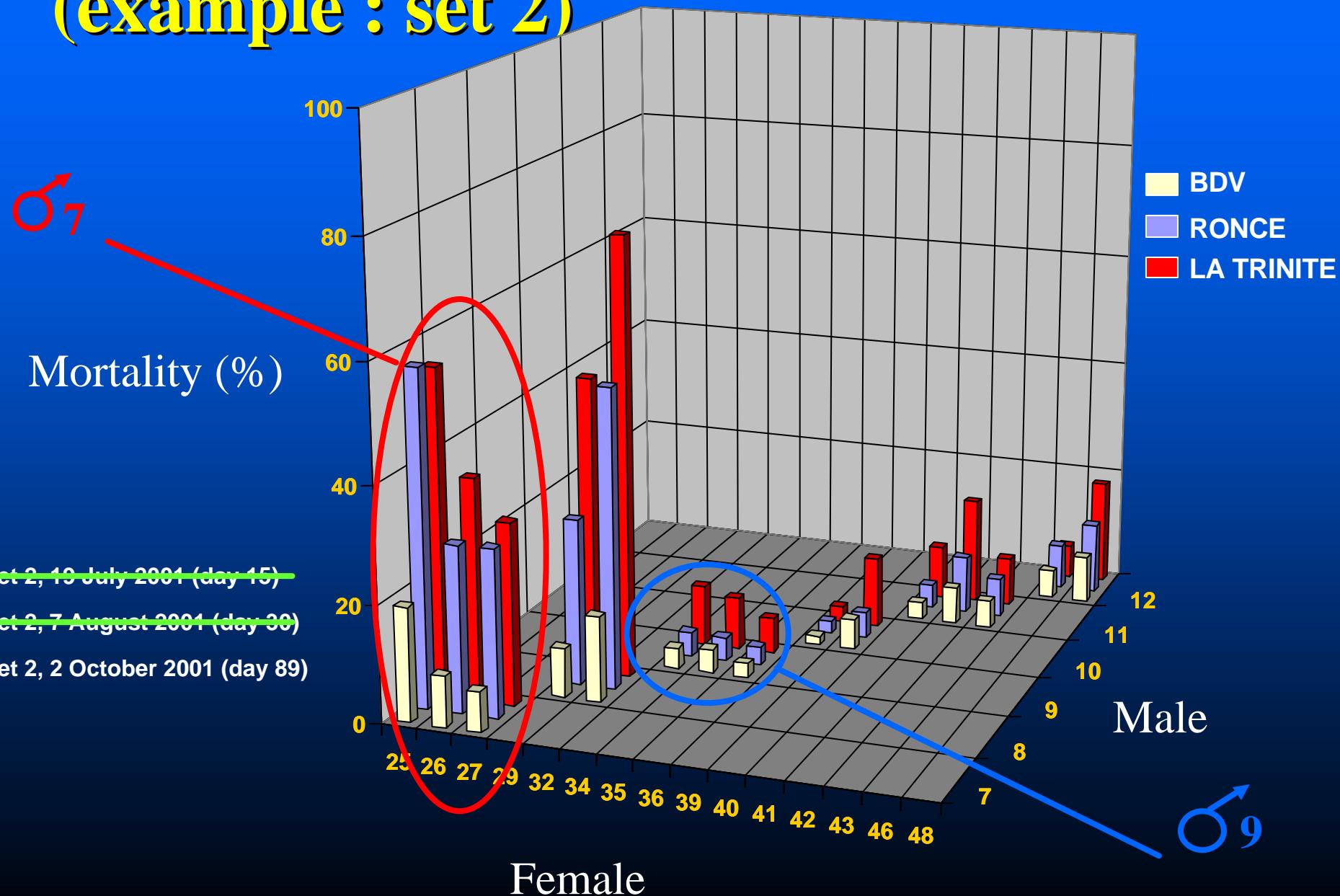
## Time schedule : 3 successive sets



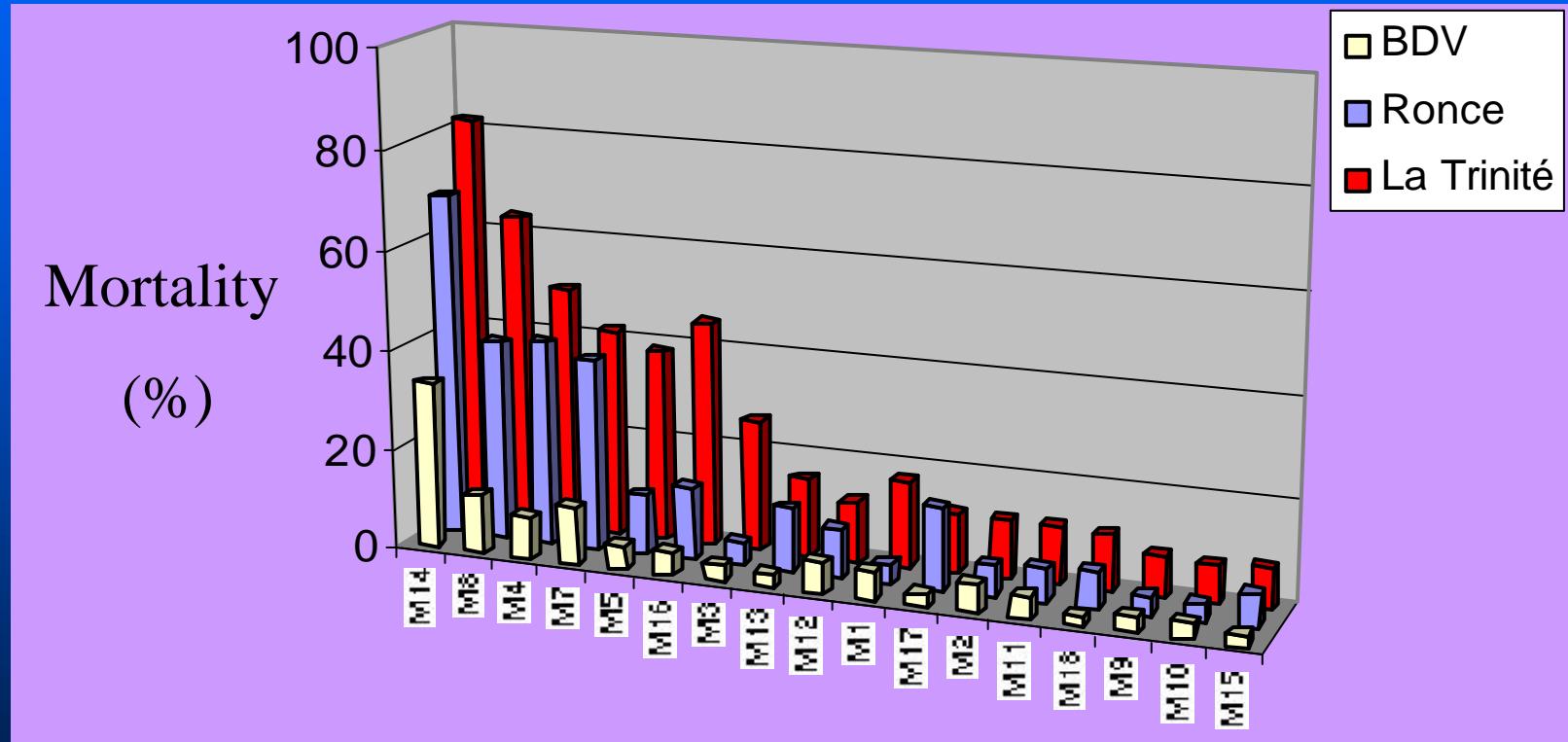
- Recorded traits :

- survival and growth 2 and 4 weeks after placement in the field
  - survival and growth at the end of the growing season
- 3 bags / family / site / sample = 1215 bags of 150 oysters**

# G1 : mortality in the field (example : set 2)

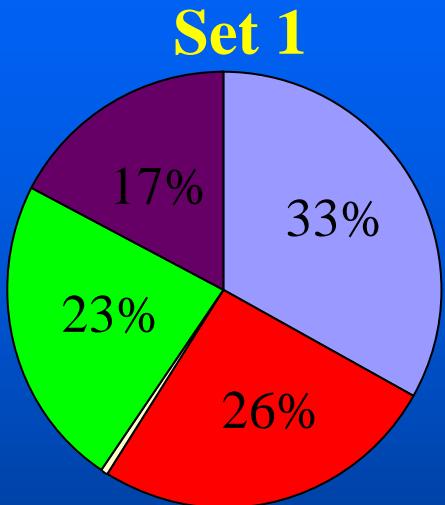


# G1: mortality in the field - October 2001 mean values per male and per site



- Differences among sites
- Differences among males → Important genetic basis for survival

# G1 : variance components for survival



$$h^2 = 0.23 \pm 0.35$$

$$h^2 = 0.85 \pm 0.46$$

$$h^2 = 1.21 \pm 0.57$$

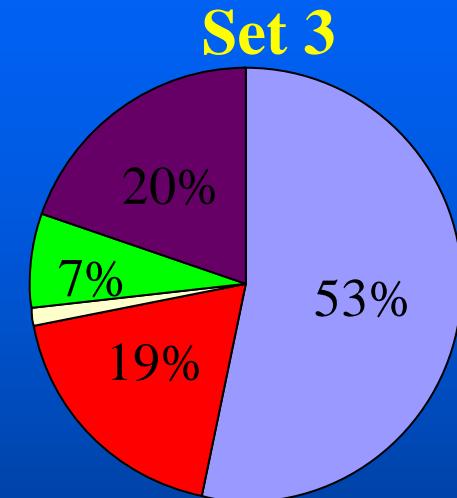
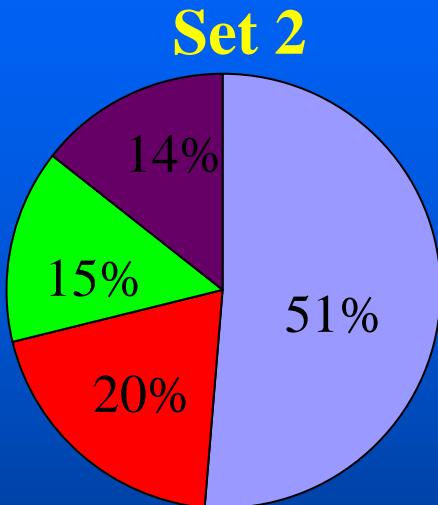
Family p<0.01

Site p<0.01

Replicate ns

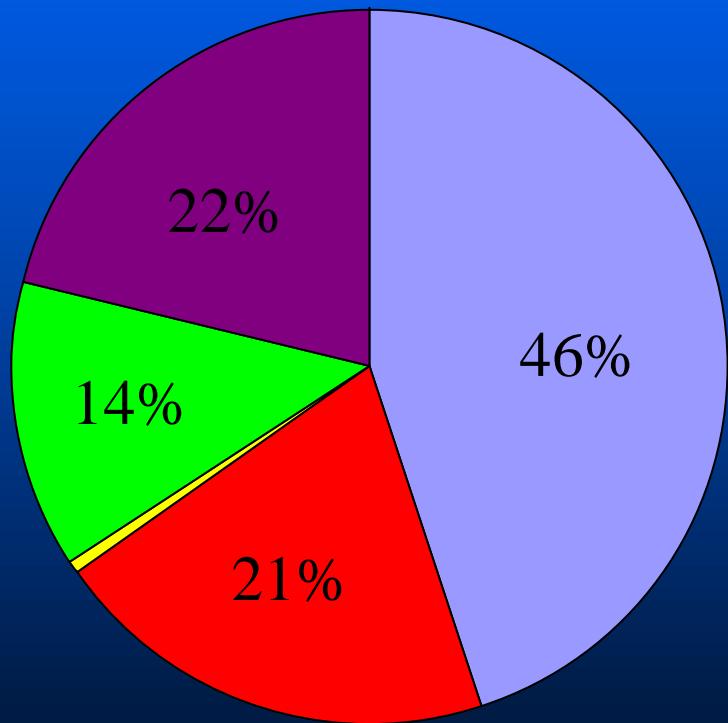
Family\*Site p<0.01

error



( $h^2$ = narrow sense heritability)

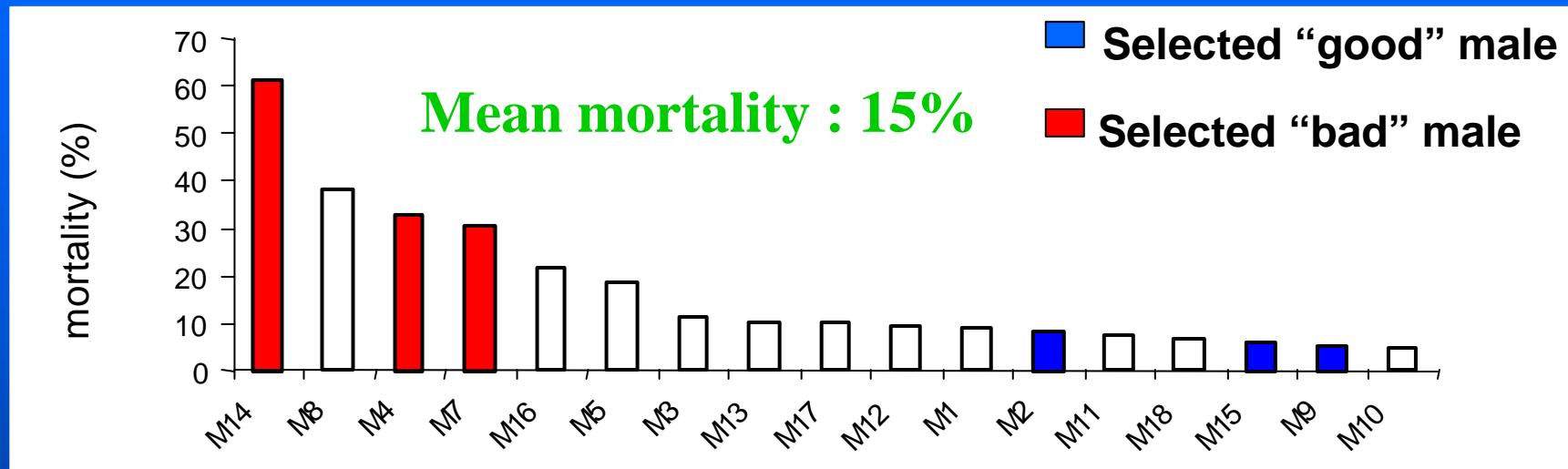
# G1 : variance components for survival overall sets - October 2001



- Set (ns)
- Family(set) (p<0.01)
- Site (p<0.01)
- Replicate (ns)
- Set\*Site (ns)
- Site\*Family(Set) (p<0.01)
- error

$$h^2 = 0.81 \pm 0.29$$

# Second generation : divergent selection 2002



Mortality : 43%

S = 28 %

Low selected group : 12 families

Male	4	7	14			
Family	F4-15	F4-16	F7-25	F7-26	F14-54	F14-55
4	F4-15		13	14	17	18
	F4-16		15	16	19	20
7	F7-25				21	22
	F7-26				23	24
14	F14-54					
	F14-55					



Mortality : 4%

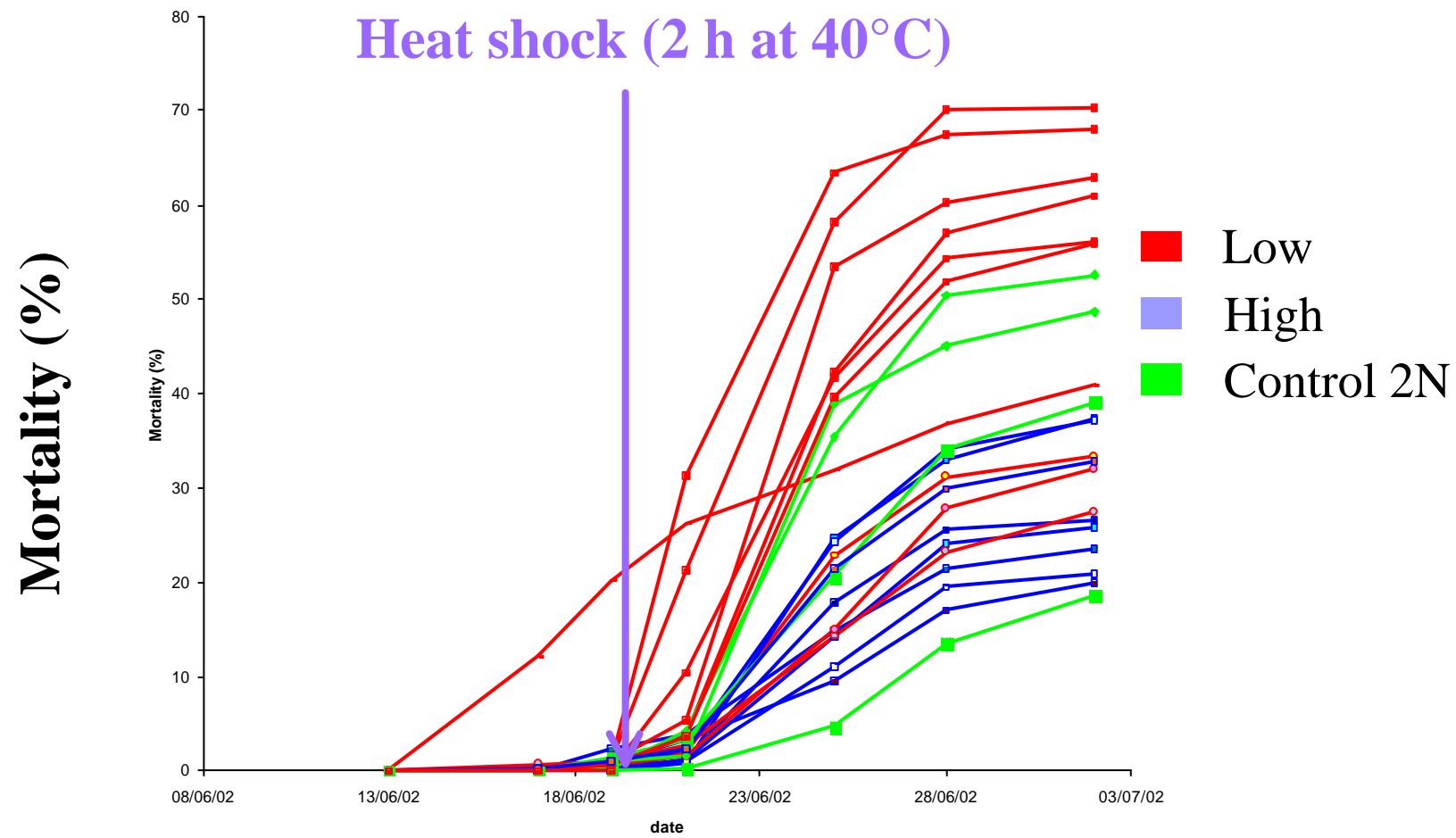
S = 11 %

High selected group : 12 families

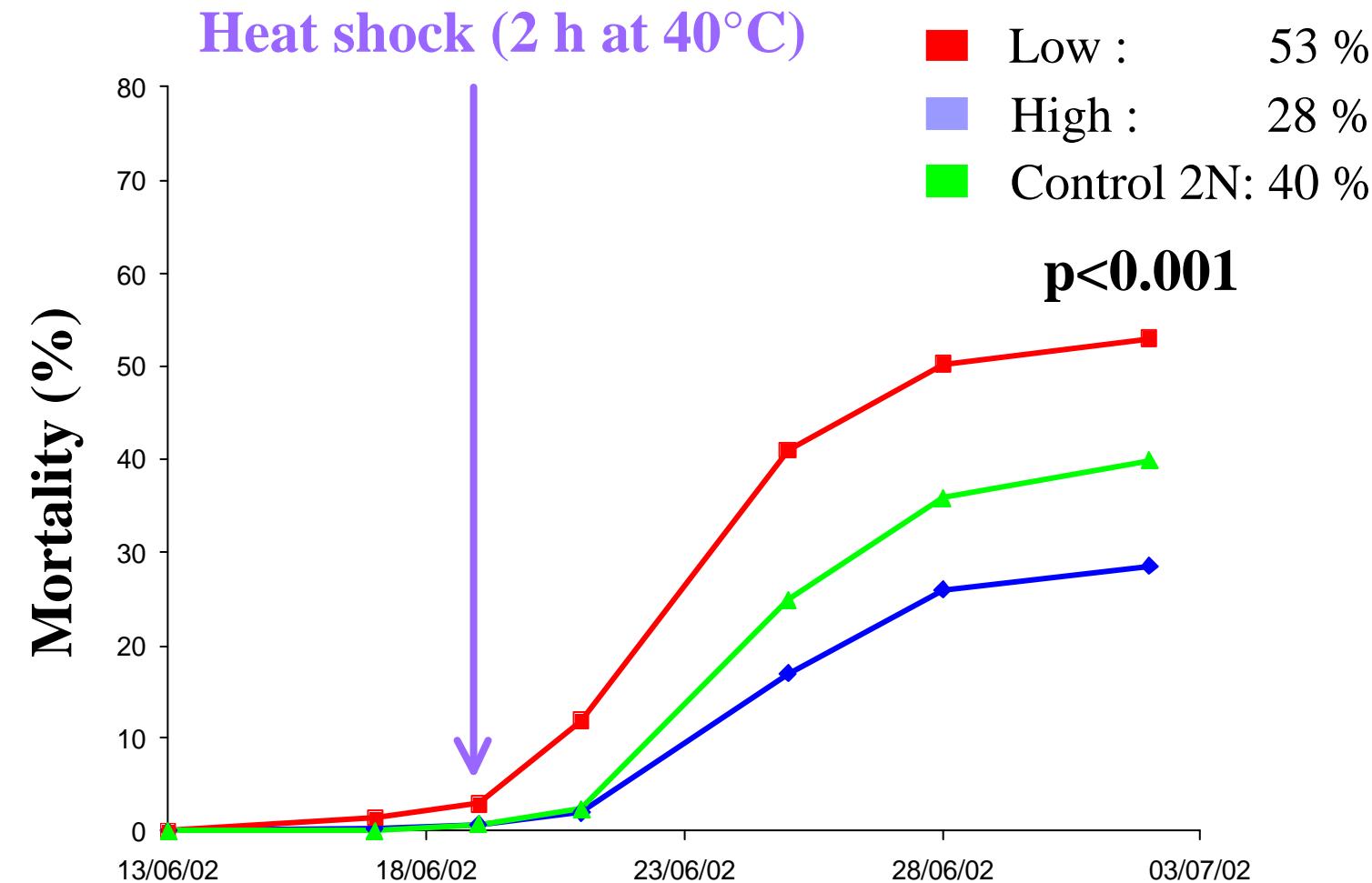
Male	2	9	15			
Family	F2-5	F2-8	F9-35	F9-36	F15-57	F15-58
2	F2-5				1	2
	F2-8				3	4
9	F9-35					5
	F9-36					6
15	F15-57					9
	F15-58					10
						11
						12

Hatchery control : 2N and 3N

# G2 divergente selection : mortality *in labo* 2002 :

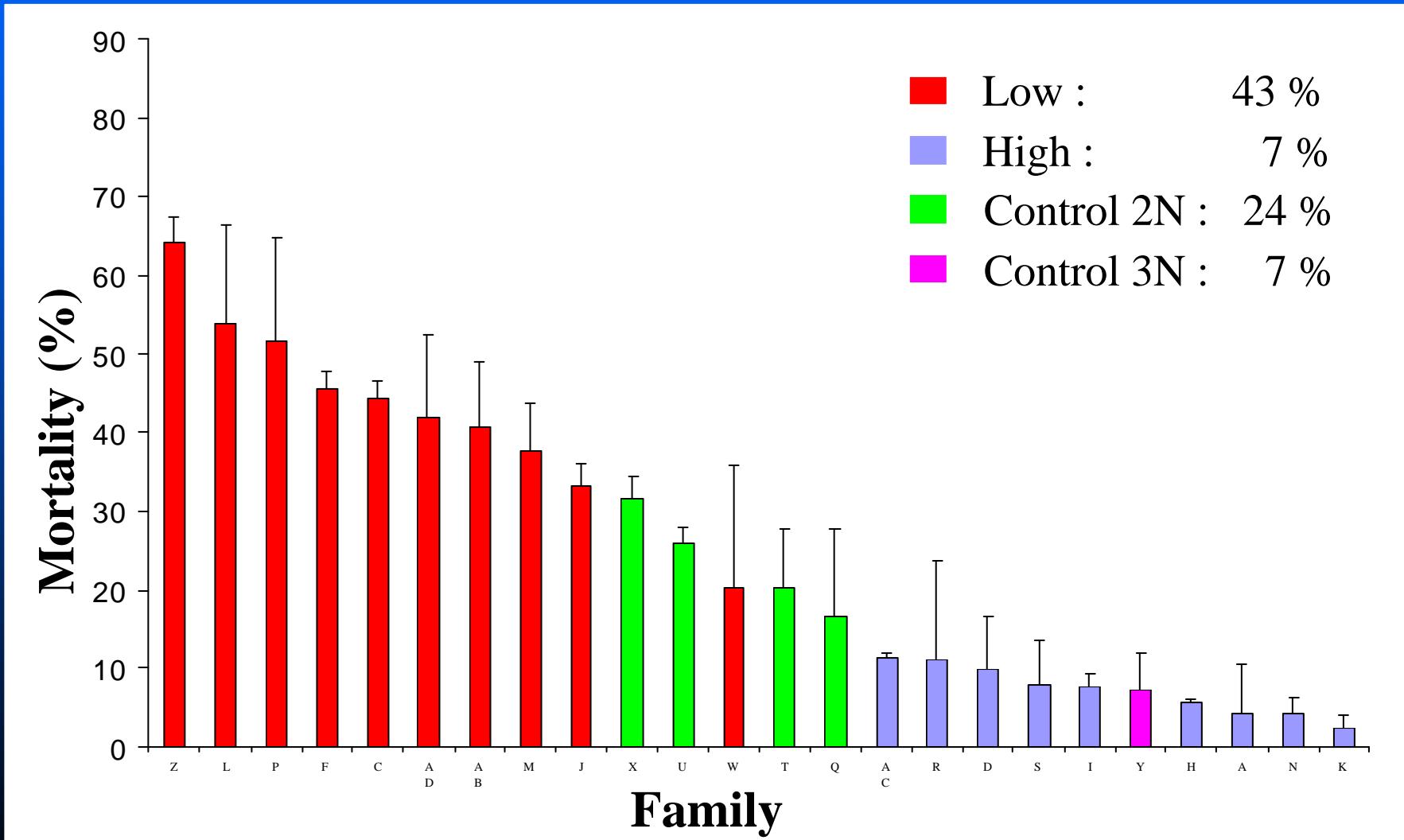


# G2 divergente selection : mortality *in labo* 2002 :

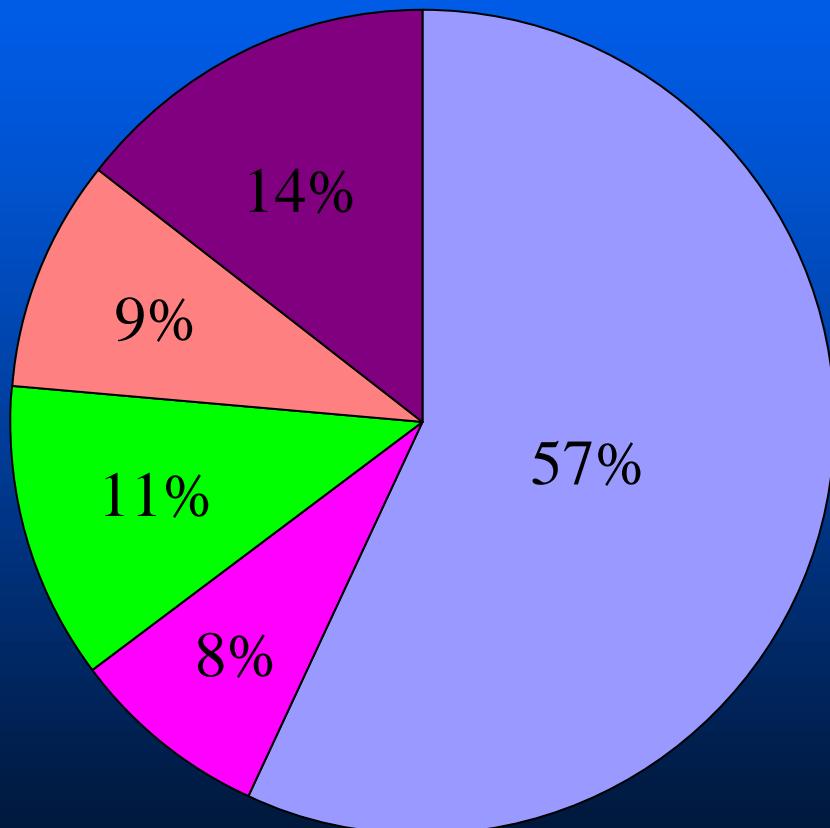


# G2 divergente selection : mortality in the field

Site : La Trinité (South Brittany)

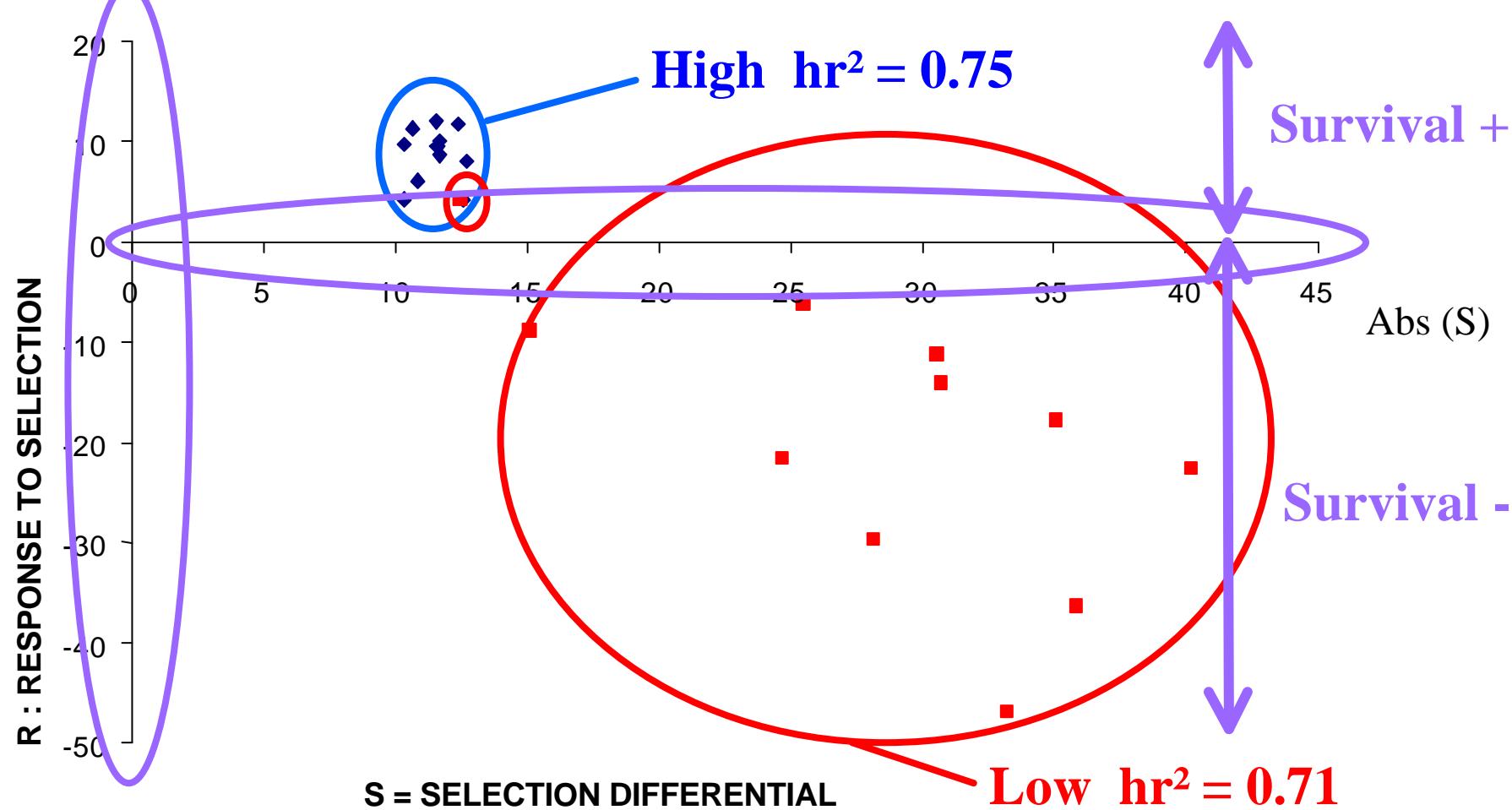


## ■ G2 divergent selection : variance components for survival - October 2002

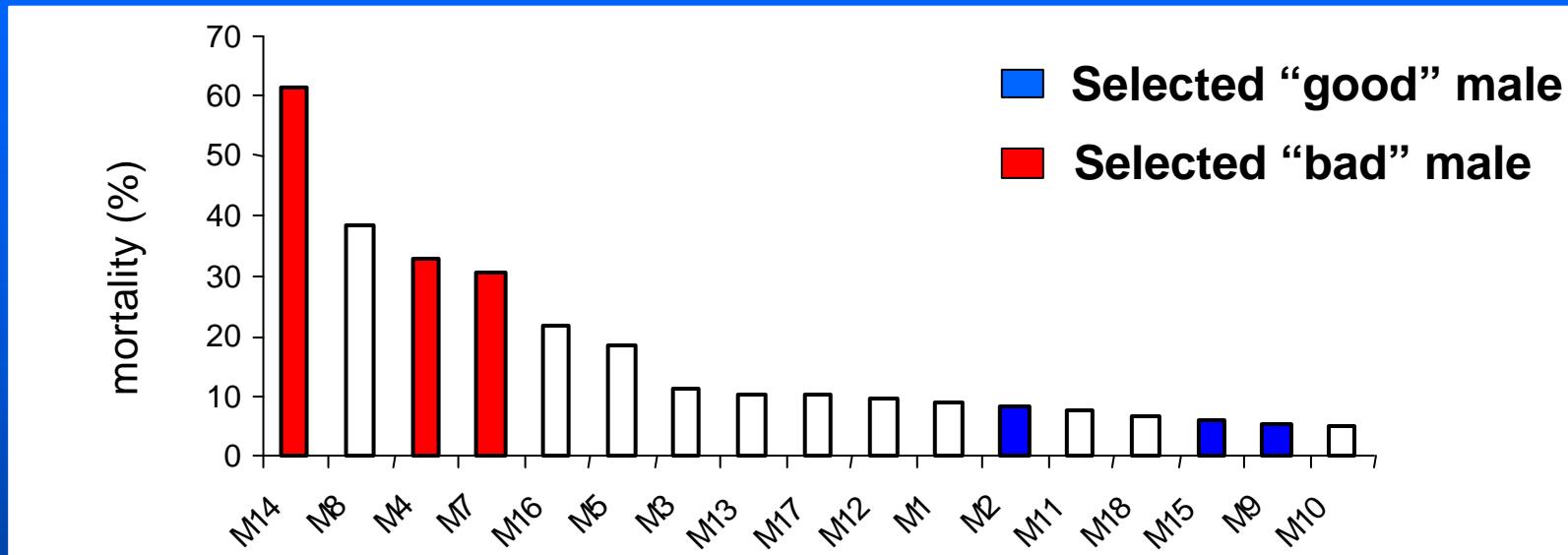


- **Site** p<0.01
- **Group** p<0.01
- **Family(group)** p<0.01
- **Replicate** ns
- **Site\*Group** p<0.01
- **Site\*Family(group)** p<0.01
- **error**

# Response to selection for survival in the field :



# G2 : Inbred lines



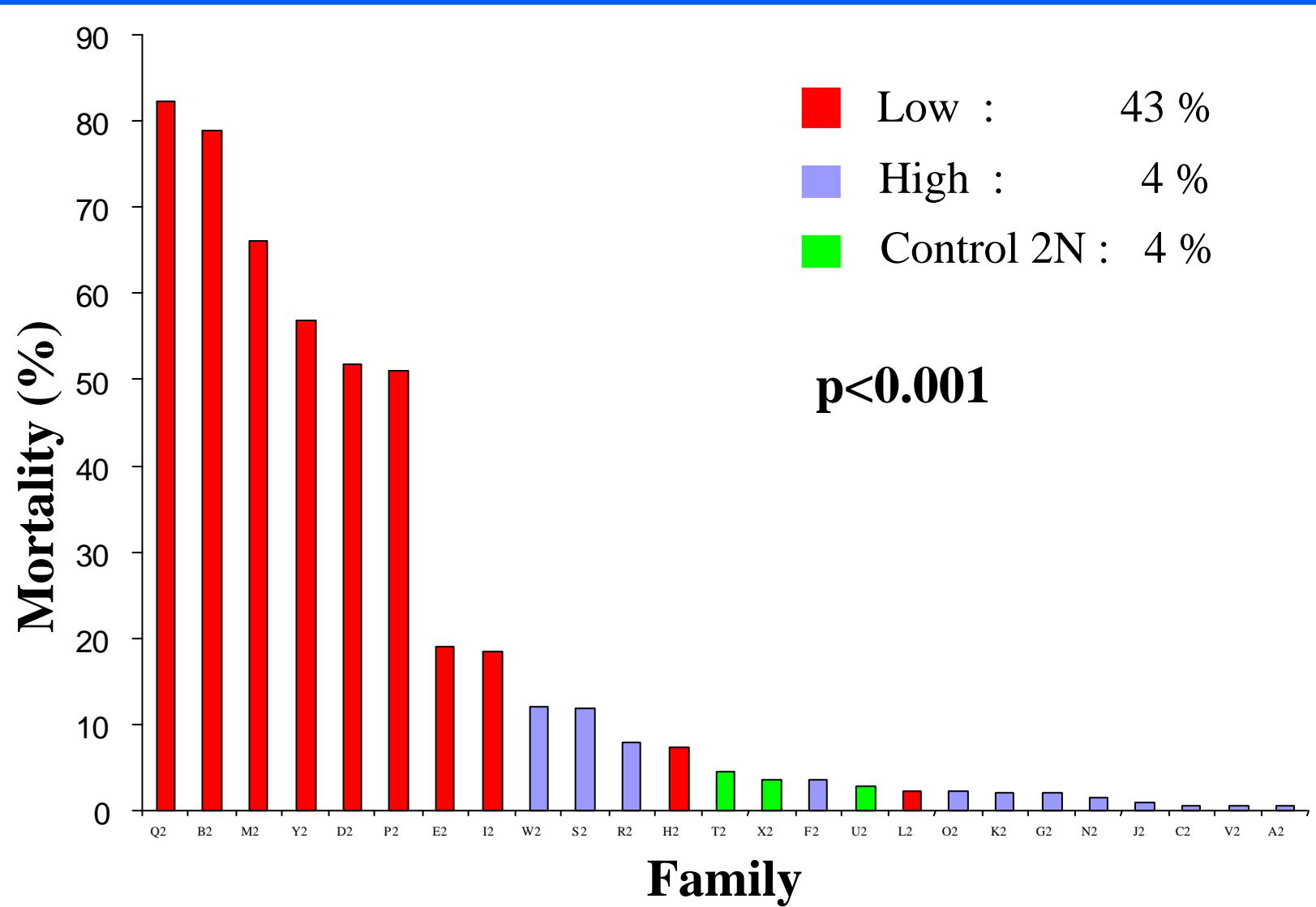
Hatchery  
Control :  
2N

Male	4						7			14				
	Family		F4-15	F4-16	F7-25	F7-26	F14-54	F14-55						
4	F4-15		13	14										
	F4-16		15	16										
7	F7-25				17	18								
	F7-26				19	20								
14	F14-54						21	22						
	F14-55						23	24						

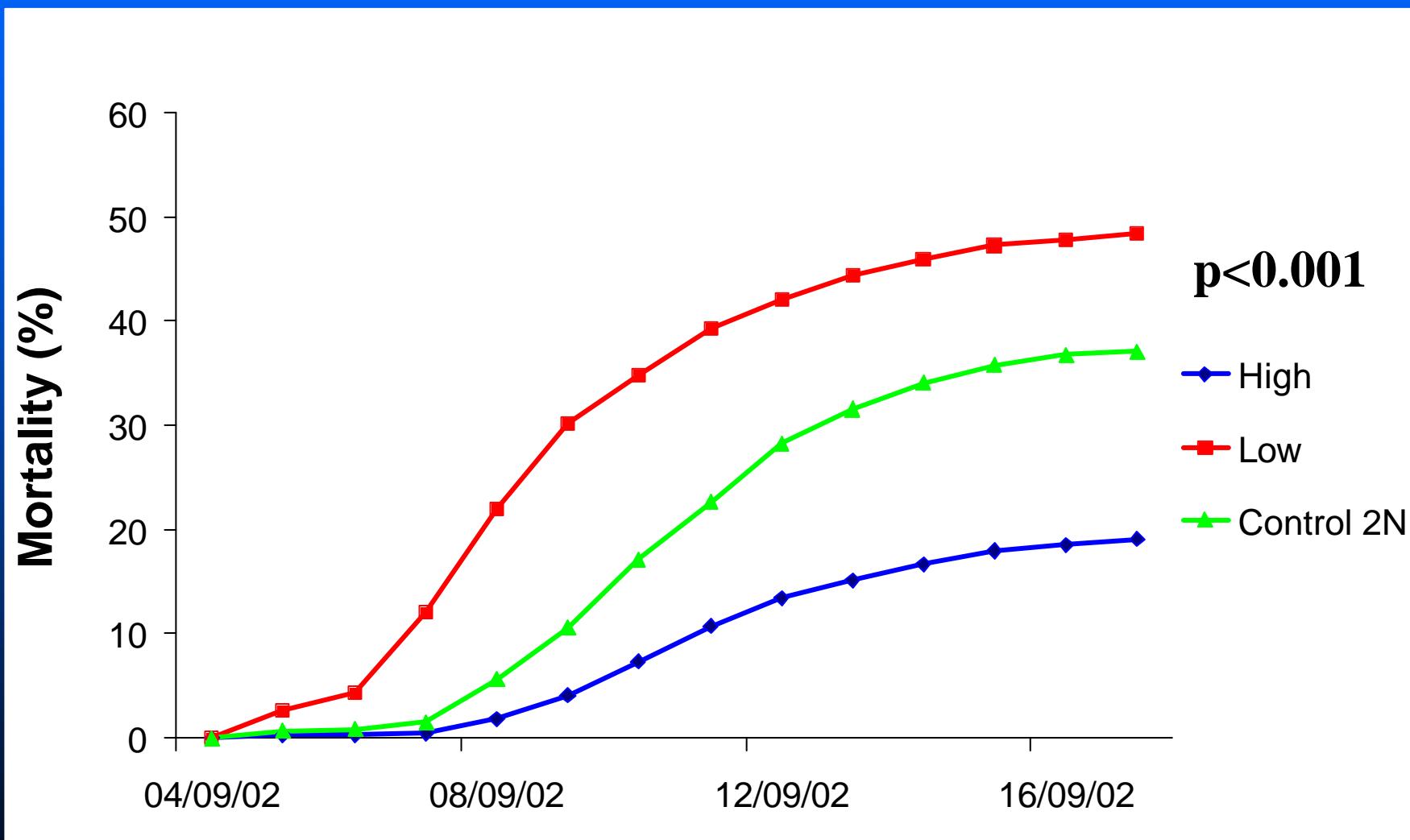
Male	2						9			15				
	Family		F2-5	F2-8	F9-35	F9-36	F15-57	F15-58						
2	F2-5				1	2								
	F2-8				3	4								
9	F9-35						5	6						
	F9-36						7	8						
15	F15-57								9	10				
	F15-58								11	12				

# G2 Inbred lines : mortality in the field

Site : Ronce (Marennes-Oléron Bay)



# G2 inbred lines : mortality *in laboratory*



# Conclusion :

- Differences of survival among half-sib families (G1)
- High response to selection and confirmation of the strong heritabilities for survival between the G1 and G2

*Selective breeding programs could improve survival of juveniles oysters*

- No negative effect on the growth

# Perspectives :



- 1- To reconstitute the divergent selection to confirm the results of the second generation
- 2- Development of inbred lines for the families that showed high and low levels of survival
- 3- Cryopreservation of gametes for the selected families
- 4- Availability of selected families with high and low survival performances for the others disciplines of the framework « MOREST » (immunology, physiology, ecotoxicology...)

Specials thanks to :

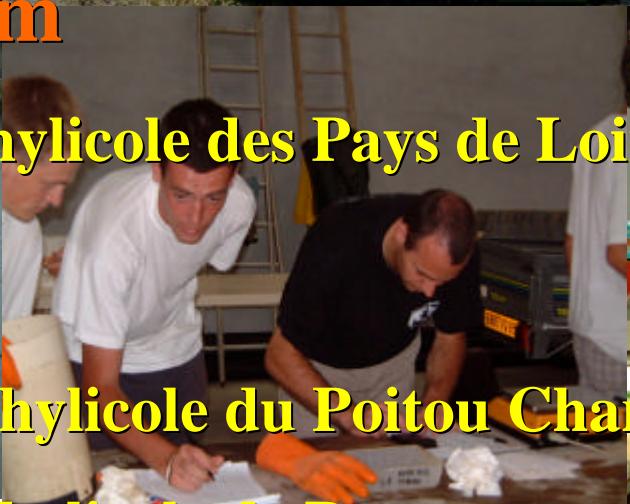
- Hatchery team

Laboratoire de Génétique et Pathologie



- Nursery team

Laboratoire conchylicole des Pays de Loire



- Field teams

Laboratoire Conchylicole du Poitou Charentes



Laboratoire Conchylicole de Bretagne

- BIG BOSS TEAM

Laboratoire de Physiologie des Invertébrés Marins

