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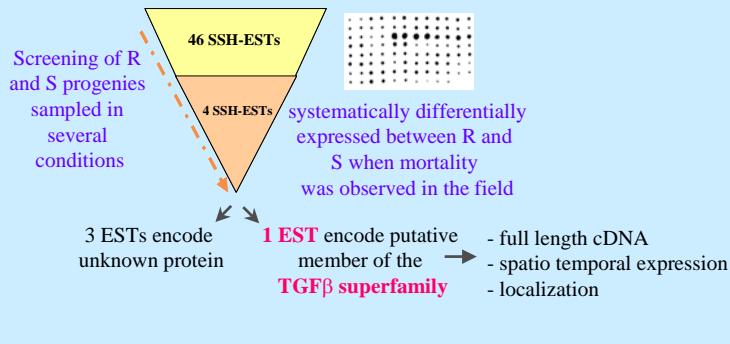
Introduction

To study the complex and non identified host causes of summer mortality in *C. gigas*, divergent selection criteria based upon summer survival have been applied to produce resistant (R) and susceptible (S) oyster families (Samain et al., 2007). A previous analysis by suppression-subtractive hybridization has characterized 46 differentially-expressed cDNAs between R and S families (Huvet et al., 2004). The present study reports the screening, using nylon array, of the 46 oyster SSH cDNAs from R and S selected families. Among the few differentially expressed genes characterized between R and S samples, one encodes a Transforming-Growth-Factor- β -related (TGF β) protein. Members of the TGF β superfamily are critical growth factors regulating a variety of important processes. We report the characterization and the spatio-temporal expression of *oyster-gonadal-TGF β -like* (*og-TGF β -like*), a TGF β superfamily member specifically expressed in the gonad of the cupped oyster, *C. gigas*.

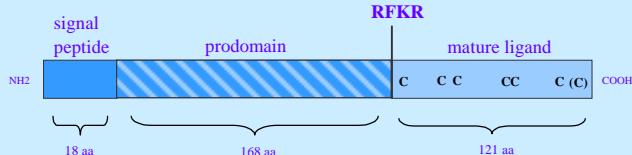
Results

1) cDNA macroarray analysis

cDNA macroarray hybridized by R and S samples collected in 4 experiments to focus on the genetic effect
(4 and 16 month-old oysters of 2^d and 3^d generations, collected in two sites, and in a nursery)



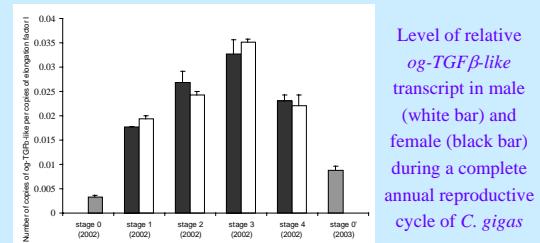
2) Isolation of *og-TGF β -like* cDNA



Nucleotide sequence encoding a preproprotein of 307 amino acids
- consensus cleavage site type « RXXR »
- 6 conserved cysteine (lacks the seventh characteristic cysteine)
- percentage of identity with other related ligands very low
→ derived member of the TGF β superfamily

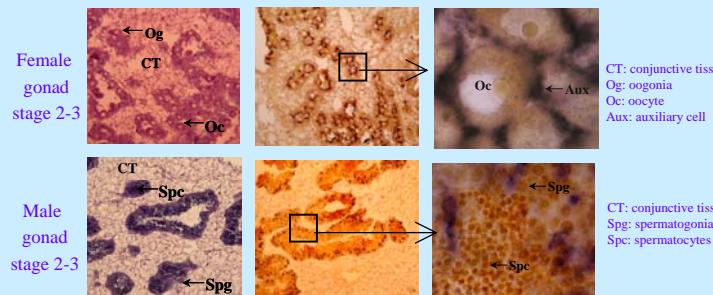
3) Spatio-temporal expression of *og-TGF β -like*

- No significant relative level of transcript detected in any early oyster developmental stage
- High relative level of transcript detected exclusively in the gonad, and not in other tissues, of both male and female
- Relative quantity of transcript increased continuously during the development of gonadic tubules in both sexes (figure)
- maximum level during active gametogenesis (stage 3)



4) Localization of *og-TGF β -like* by *in situ* hybridization

og-TGF β -like mRNA detected exclusively in cells between oocytes



og-TGF β -like mRNA detected exclusively in cells between spermatogonia

Conclusion

- ✓ *og-TGF β -like* has only 6 conserved cysteine as observed for teleost gonadal soma-derived growth factor (Sawatari et al., 2006)
- ✓ tissue-specific expression, exclusively in the gonad of both sexes and not during developmental stages nor in other tissues
- ✓ strongly expressed when germ cells are fully mature
- ✓ localized in somatic cells surrounding the germ cells, supposed to play a regulatory role in the differentiation and maturation of germ cells

→ derived member of the TGF β superfamily involved in gonadal development

- ✓ summer mortality is suggested to interplay with reproduction
- ✓ significant differences in reproduction observed between R and S
- ✓ some TGF β members (BMP15, GDF9) play crucial regulatory roles in ovarian follicle or testis development

Is that different expression of *og-TGF β -like* significant to the higher rate of summer mortality of S progeny ?