

GENE EXPRESSION DURING OOGENESIS IN THE MOZAMBIQUE TILAPIA (*O. MOSSAMBICUS*)



Supervisor: Professor Doctor Adelino Canário
Co-supervisor: Doctor Laurence Deloffre

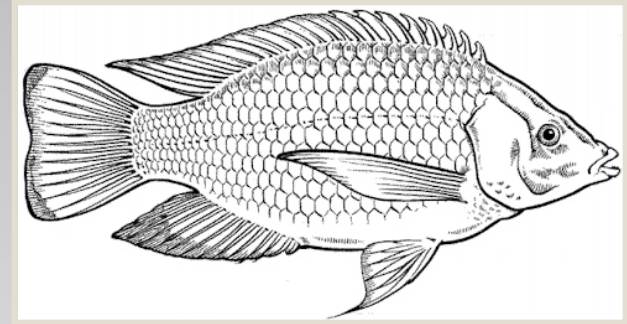
Nuno Manuel Cardoso Domingues

Faro, 26th October 2007

Introduction – Tilapia biology and value

TILAPIA:

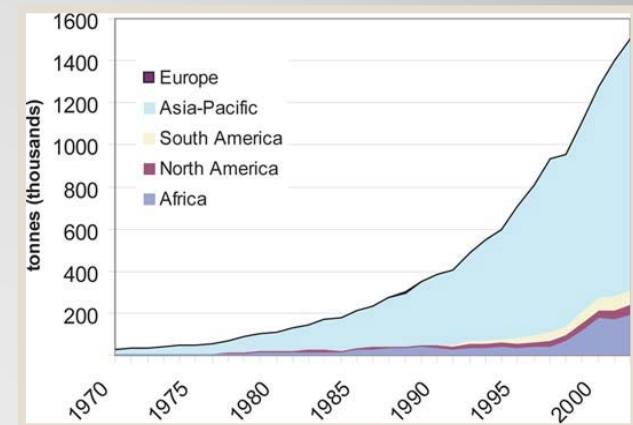
- ➔ **Ecological adaptations and tolerances;**
- ➔ **Tendency to overpopulate;**
- ➔ **Asynchronous ovaries.**



(Gunther Schmida in Mackenzie, et al. 2001)

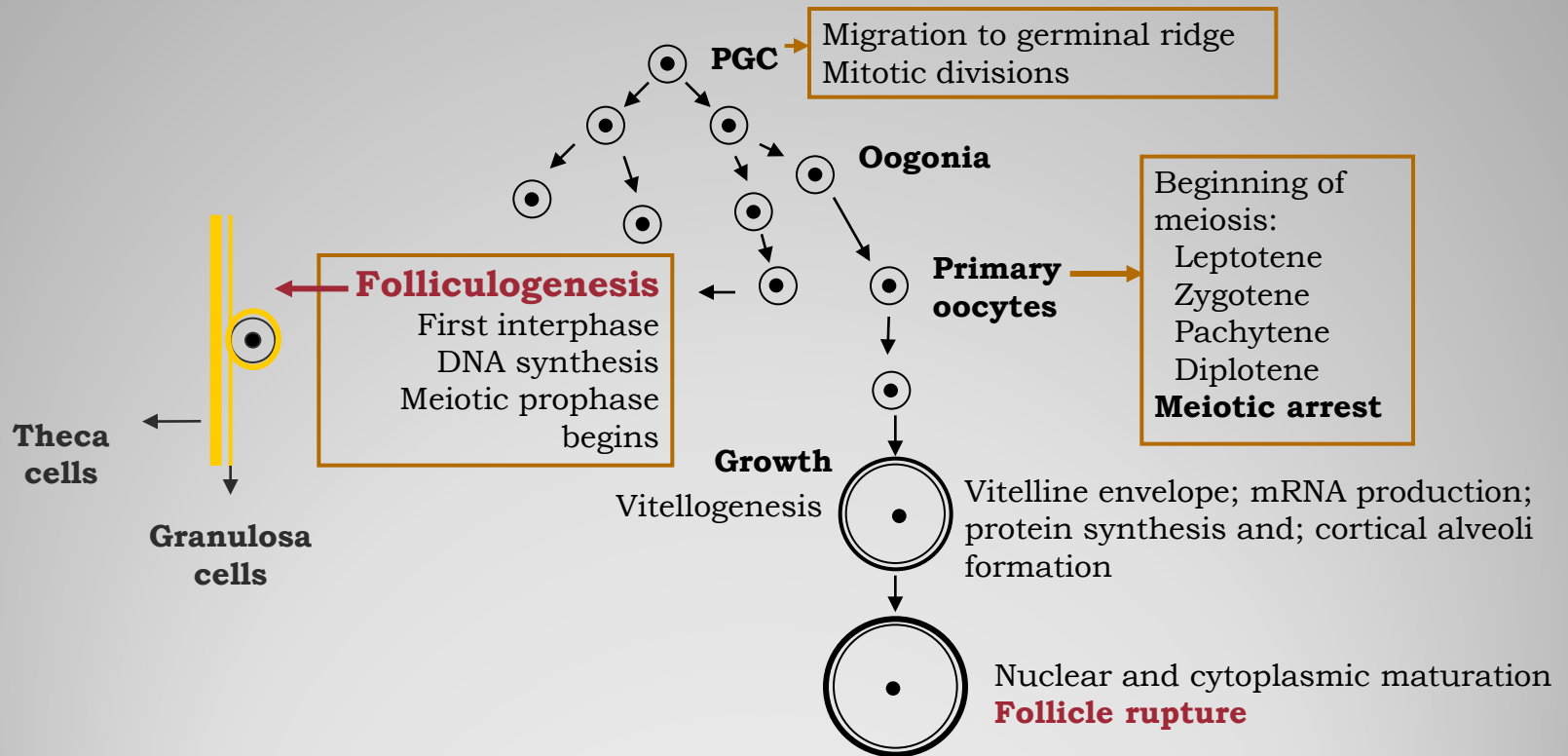
VALUE:

- ➔ **Interdisciplinary applications;**
- ➔ **Main area of interest aquaculture.**

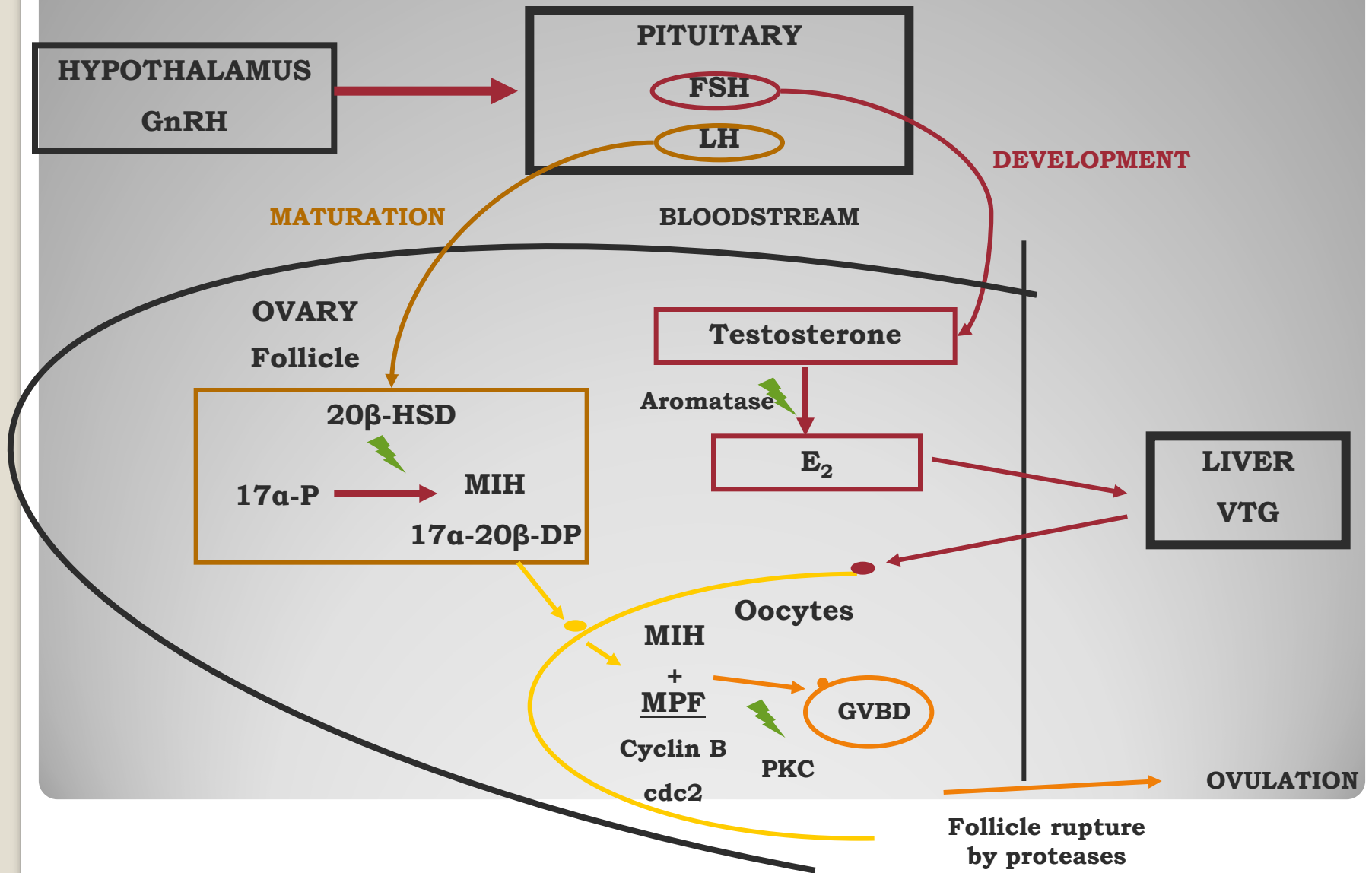


(De Silva et al. 2004).

Introduction – Ovary recrudescence in teleosts and oogenesis



Introduction – Hypothalamus-pituitary-gonadal axis in teleosts



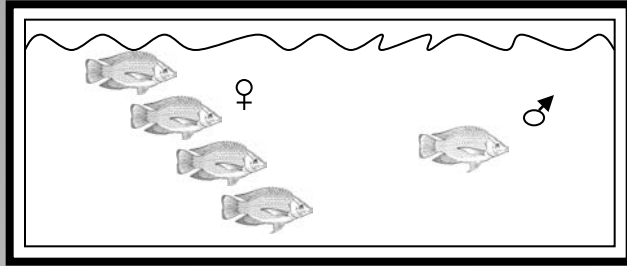
Objectives

➔ mRNA expression analysis of **putative factors** regulating the development of the ovary of *Oreochromis mossambicus*;

- BMP-R IB, RBMX, Aly protein, CPI-17, SART, PPMP protein, XP2, Clone 26

➔ Full sequences of these genes would be obtained by means of RACE-PCR.

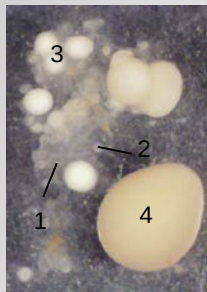
Material and methods – Biological material



- ➔ 200 L fresh water;
- ➔ 27°C;
- ➔ 2 hour photoperiod;
- ➔ constant oxygenation;
- ➔ sand on the bottom;
- ➔ fed daily with commercial cichlid ration.

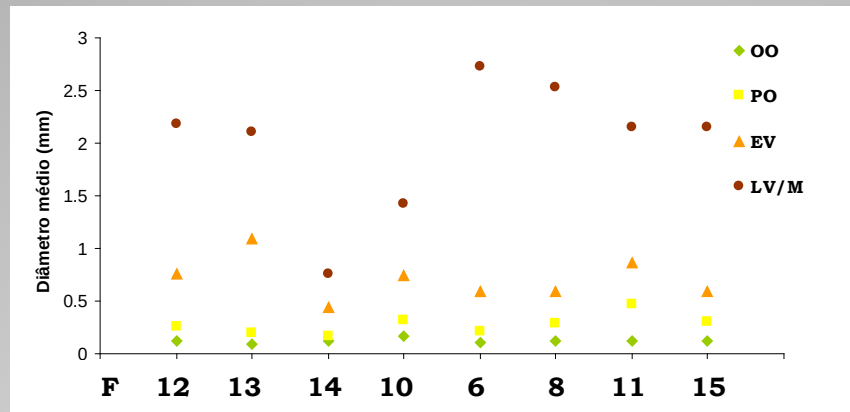
1) Lengths and weights registered (GSIs determined);

2) Oocytes separated under a binocular microscope in 0,9% NaCl:



- 1) oogonia (OO);
- 2) primary oocytes (PO);
- 3) early vitellogenic oocytes (EV);
- 4) late vitellogenic oocytes or mature oocytes (LV/M).

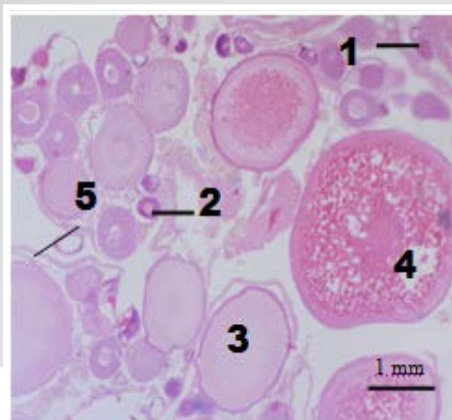
Material and Methods (continued)



Sizes significantly different from each others ($P \leq 0,001$)

Histology

Tissues stained with haematoxylin-eosin:



- 1) Oogonia;
- 2) Primary oocytes;
- 3) Early vitellogenic oocytes;
- 4) Late-vitellogenic oocyte;
- 5) Ovarian follicle.

Material and Methods – RNA extraction and cDNA synthesis

RNA extraction
(TRI Reagent)



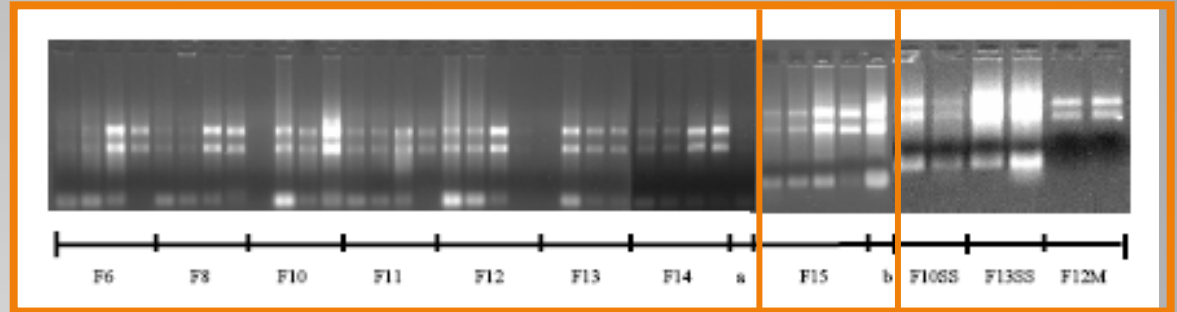
cDNA synthesis
with M-MLV RT



RT-PCR

Sexual dimorphic
expression

Patterns through
oocyte growth



DNase treatment



Primers:

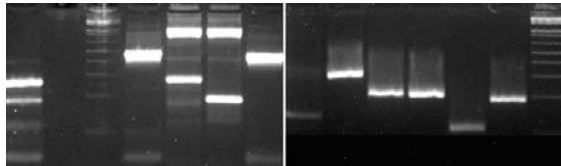
GC $\approx$ 50%

Annealing temperature FW-RV $\neq$ $< 5^{\circ}\text{C}$

No hairpin effect or primer dimers.

Material and Methods – RT-PCR

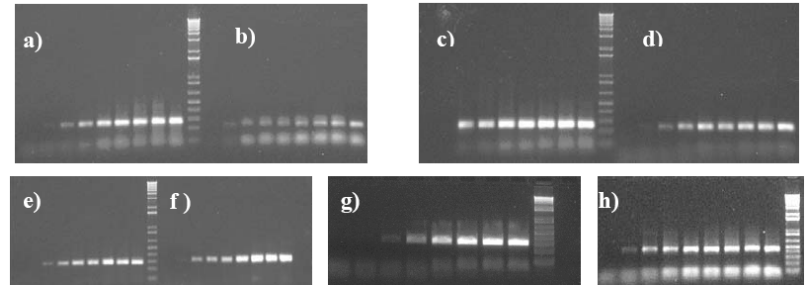
Optimization procedures



Temperature

MgCl₂

Nr of cycles



➔ RT-PCR in BIO RAD myCycler

94°C 4 minutes;

Y times {
- 94°C 25 s;
- **X** °C 25 s;
- 72°C 35 s.

➔ 2 or 3% agarose gel by electrophoresis;

➔ Quantification (Biorad Quantity One);

➔ Statistical analysis (ANOVA, ANOVA on ranks and Pearson correlation).

1x Taq Buffer

MgCl₂ (3 mM)

Taq DNA polymerase (0,5 units)

dNTPs (0,2 mM each)

Primers (0,4 pmol/μl or 0,8 pmol/μl)

cDNA

H₂O (Sigma)

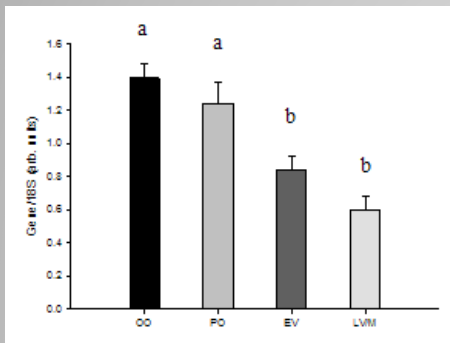
25 μl

Results – Control genes

18S – reference gene

Accumulation in oocytes? ➡ Vasa, FoxL2 and CYP19a tested.

Vasa



➡ PGC formation and migration to germinal ridge;

➡ Decrease in vitellogenic stages.

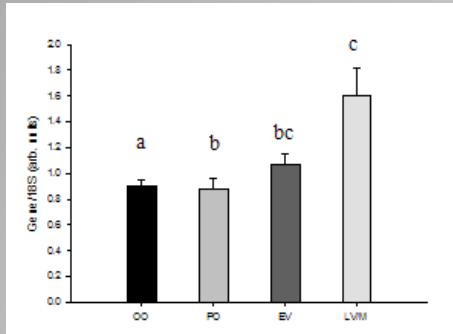
➡ Dispersion through the cytoplasm?
(Kobayashi et al. 2000; Ye et al. 2007)

Observed in:

- ➡ **Nile tilapia** (Kobayashi et al. 2000);
- ➡ **Zebrafish** (Braat et al. 1999b);
- ➡ **Rainbow trout** (Ye et al. 2007).

Results – FoxL2 and CYP19a

FoxL2 (transcription factor)



➞ Ovarian development and function;

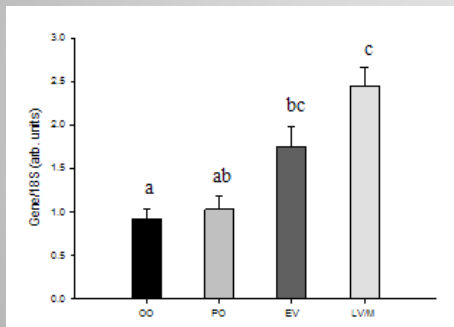
➞ Differentiation of PGC to oogonia.

FoxL2 binds to CYP19a
ACAAATA in promoter

Positively
correlated

Activation of CYP19a

CYP19a (ovarian aromatase gene)



➞ Higher in vitellogenic stages ➞ E₂ synthesis.

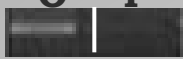
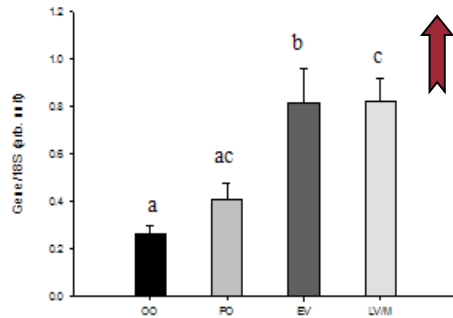
Results agree with:

- Wang et al. 2007;
- Yoshiura et al. 2003;
- Pannetier et al. 2006.

Results – Tested genes

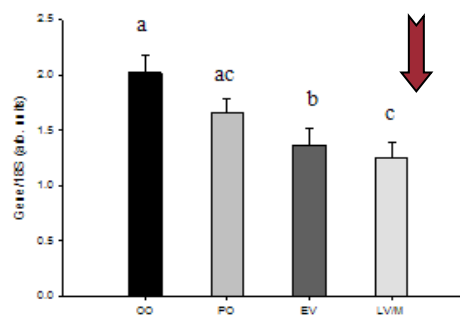
BMP-R IB

O T

RBMX

O T

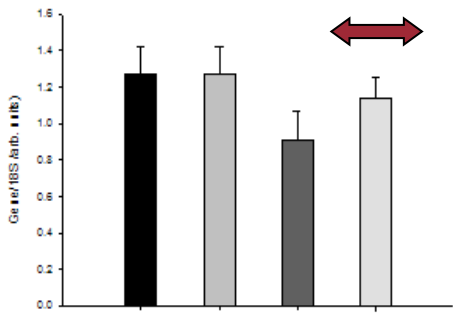



Major role in cellular differentiation

Major role in maturation

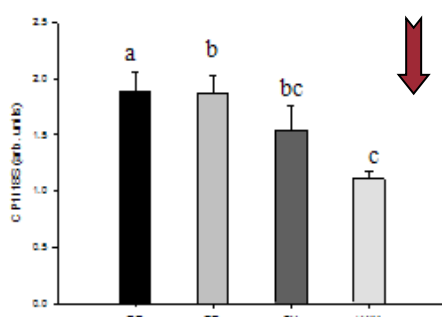
Aly protein

O T

CPI-17

O T

Basic cellular processes

18S

O T

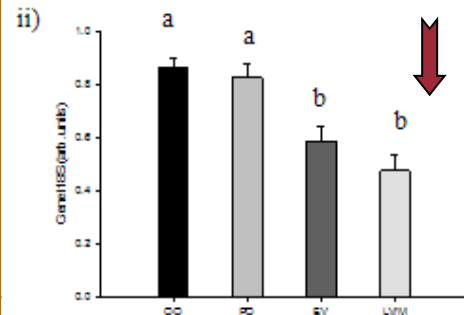
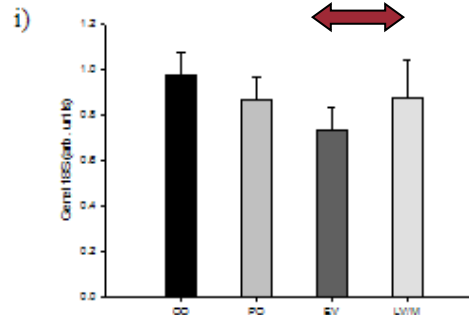


Results – Tested genes

SART



PPMP



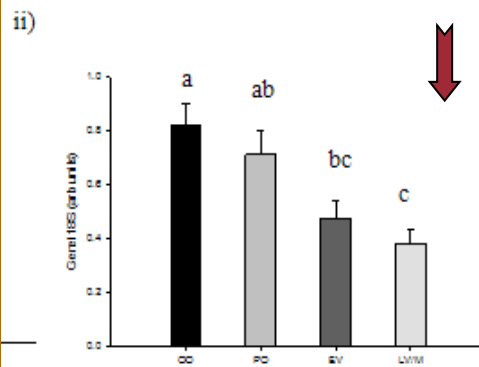
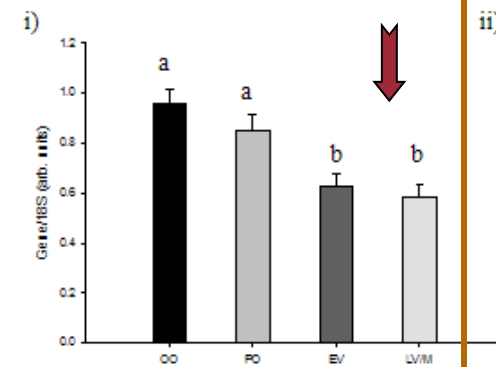
Major role in cellular differentiation

Major role in maturation

XP2



Clone 26



Basic cellular processes

18S



Discussion – BMP-R IB

➞ Expressed in mouse granulosa cells, theca cells (higher) and oocytes

At theca cells
BMP - 4
BMP - 7

At oocytes
BMP -6
BMP - 15

Autocrine /
paracrine actions

**FSH-BMP signalling
in mammals**

➞

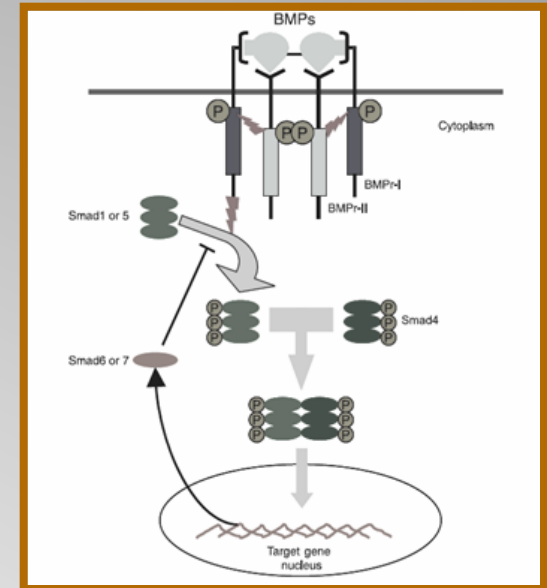
**BMPs increase FSH
signalling by Smads
cascade**

➞

**Potentialtion of
steroidogenic pathway**

➞

**Supress LH receptor
expression**



(Source: Granjeiro, et al. 2005).

**Correlated
positively with
FoxL2 and CYP19a**

Role in inhibition/delaying of luteinization and granulosa cell differentiation

Discussion – hnRNPGs

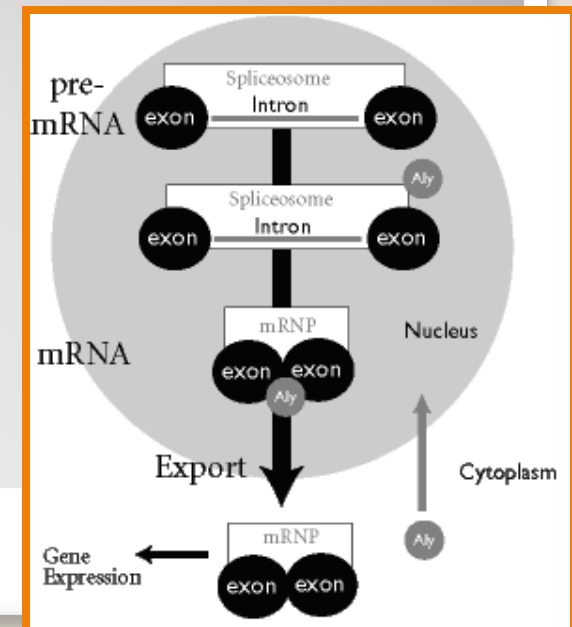
RBMX ↓

- ➔ Expression suggests involvement in basic development processes (brain of *Danio rerio*);
- ➔ Decrease through follicle differentiation;
- ➔ Required for mRNA splicing.

Positively
correlated

Aly protein ↔

- ➔ Exportation of mRNA from nucleus to cytoplasm and involved in splicing;
- ➔ Aly recycles to the nucleus (Zhou et al. 2000).



(Source: Zhou et al. 2000)

Discussion – CPI-17

```
graph TD; A[Discussion – CPI-17] --> B[Cysteine protease inhibitor – 17 KDa]; B --> C[Phosphorylated by PKC at Thr38]; C --> D[ACTIVATION]; D --> E((Inhibitor of PP1s)); E --> F([Regulation of cell division processes and transcription by PP1s  
Involvement of CPI-17?]); E --> G([PPs 1 and 2 in steroidogenesis at follicle cells  
CPI-17 expression decreases during vitellogenesis]);
```

Cysteine protease inhibitor – 17 KDa

Phosphorylated by PKC at Thr38

ACTIVATION

Regulation of cell division processes and transcription by PP1s

Involvement of CPI-17?

→ Inhibitor of PP1s

PPs 1 and 2 in steroidogenesis at follicle cells

CPI-17 expression decreases during vitellogenesis

Discussion (SART, PPMP, XP2 and Clone 26)

↔ SART-1 → Squamous cell carcinoma-associated reactive antigen for cytotoxic T cells;

↓ PPMP protein → no reference to this gene.

↓ XP2 → hypothetical new protein (also in *Danio rerio*);

↓ Clone 26 → no homolog sequence (might be UTR).

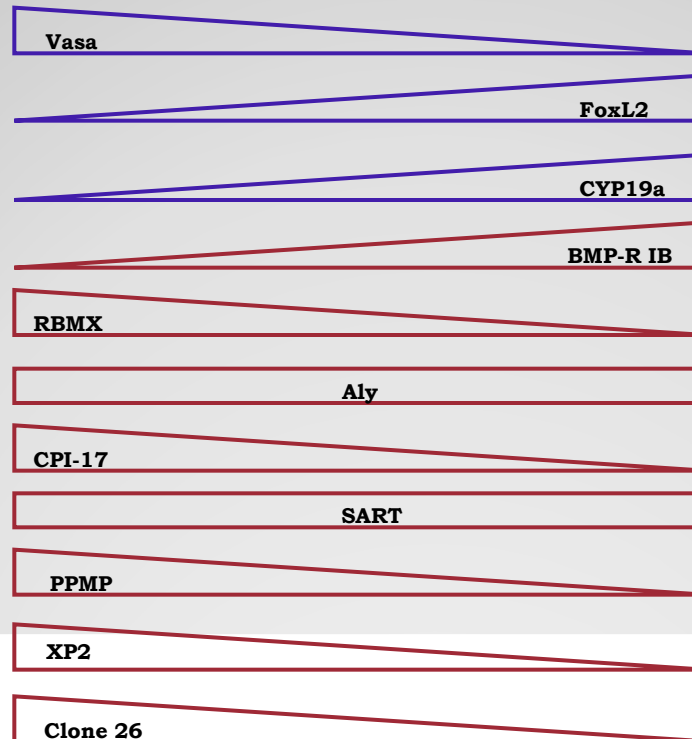
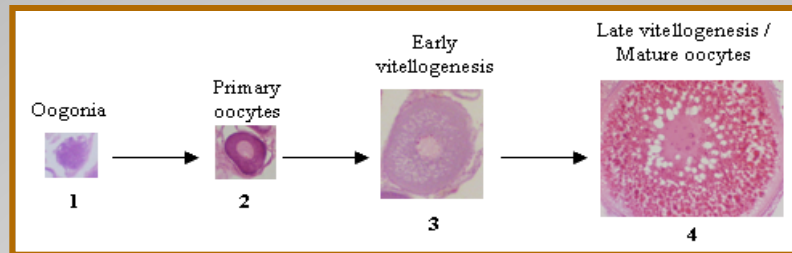
Further characterization required

Conclusions and Further Work

➡ 18S is a good reference gene proven by Vasa, FoxL2 and CYP9a results;

Conclusions and Further Work

➡ 18S is a good reference gene proven by Vasa, FoxL2 and CYP9a results;



To confirm reliability of rRNA 18S as reference

No proven identities or known functions in the ovary

Conclusions and Further Work

- ➡ **18S is a good reference gene proven by Vasa, FoxL2 and CYP9a results;**
- ➡ **Decreasing expression - role in cellular differentiation;**
 - **RBMX, PPMP, Clone 26, XP2 and CPI-17**
- ➡ **Increasing expression – important role in maturation;**
 - **BMP-R IB**
- ➡ **Constant expression – always necessary;**
 - **Aly protein and SART**
- ➡ **Further work required for characterization of unidentified genes.**

References

- Coward K & Bromage NR** 2000 Reproductive physiology of female tilapia broodstock. *Reviews in Fish Biology and Fisheries* 10 1-25.
- De Silva SS, Subasinghe RP, Bartley DM & Lowther A** 2004 Tilapias as alien aquatics in Asia and the Pacific: a review. In *FAO Fisheries Technical Paper*, p 65: FAO.
- Granjeiro JM, Oliveira RC, Bustos-Valenzuela JC, Sogayar MC & Taga R** 2005 Bone morphogenetic proteins: from structure to clinical use. *Braz J Med Biol Res* 38 1463-1473.
- Janz DM** 2000 Endocrine systems: gross functional anatomy. In *The laboratory fish - The Handbook of Experimental Animals*, pp 189-217. Ed GK Ostrander. Bath: Academic Press.
- Mackenzie R, Jackson P & Cotterell E** 2001 An operational strategy for Queensland freshwaters. In *Control of Exotic Pest Fishes*. Brisbane: Department of Primary Industries.
- Zhou Z, Luo MJ, Straesser K, Katahira J, Hurt E & Reed R** 2000 The protein Aly links pre-messenger-RNA splicing to nuclear export in metazoans. *Nature* 407 401-405.

Acknowledgements

Professor Doutor Adelino Canário

Doutora Laurence Deloffre

Laboratório de Endocrinologia Molecular Comparada (FCMA)

Laboratório de Biologia Molecular (FERN)

Olinda, Nádia, Rita, Ângela, Rute, João, Isabel, Patrícia e os outros

Joana, Tiago, Alejandro, Andrea

Pai, Mãe, João.

THE END