



# aquaculture europe

VOL. 40 (1) MARCH 2015

## Advances in Meagre

Year 1 of the  
EU DIVERSIFY  
project



Welcome to  
Rotterdam!

aquaculture  
europe 15

## Shellfish culture developments in Washington



eas



# ADVANCING AQUACULTURE AROUND THE WORLD

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## AQUACULTURE EUROPE

EAS is a non-profit  
society that aims at  
promoting contacts  
among all involved in  
aquaculture. EAS was

founded in 1976. Aquaculture Europe is the members'  
magazine of EAS.

### Secretariat

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I am pleased to present my greetings to all members of the European Aquaculture Society on behalf of the newly-elected Board of Directors. I very much look forward to working with the new board, which is a mixture of those who have been contributing to the EAS for some time and also with newly elected members from different parts of Europe.

The AE 2014 event at San Sebastian last October focusing on “Adding Value” was a great success. We owe this success to the old board and the extremely active local organisers. This event has left the new board with good news and also challenges to keep up the spirit and enthusiasm for the years to come.

The forthcoming event in Rotterdam, The Netherlands between the 20<sup>th</sup> and 23<sup>rd</sup> October 2015 shall deal with “Aquaculture, Nature and Society” where we do hope to continue to maintain the spirits of EAS in maintaining the close interaction between science and society in the large area of aquaculture.

EAS has also a major role in contributing towards EU policies in the area of aquaculture in its various facets. We shall as the board continue to play this role in an active manner. As representatives of the various stakeholders of European aquaculture, we have this commitment to support and strengthen our links with other regional aquaculture associations. It is also our mission to strengthen and support the students’ group of EAS and to further develop or reactivate our thematic groups.

I am honoured to represent all of you as President of the board of the EAS and shall strive to serve you well to keep up the spirit of EAS in all its activities.

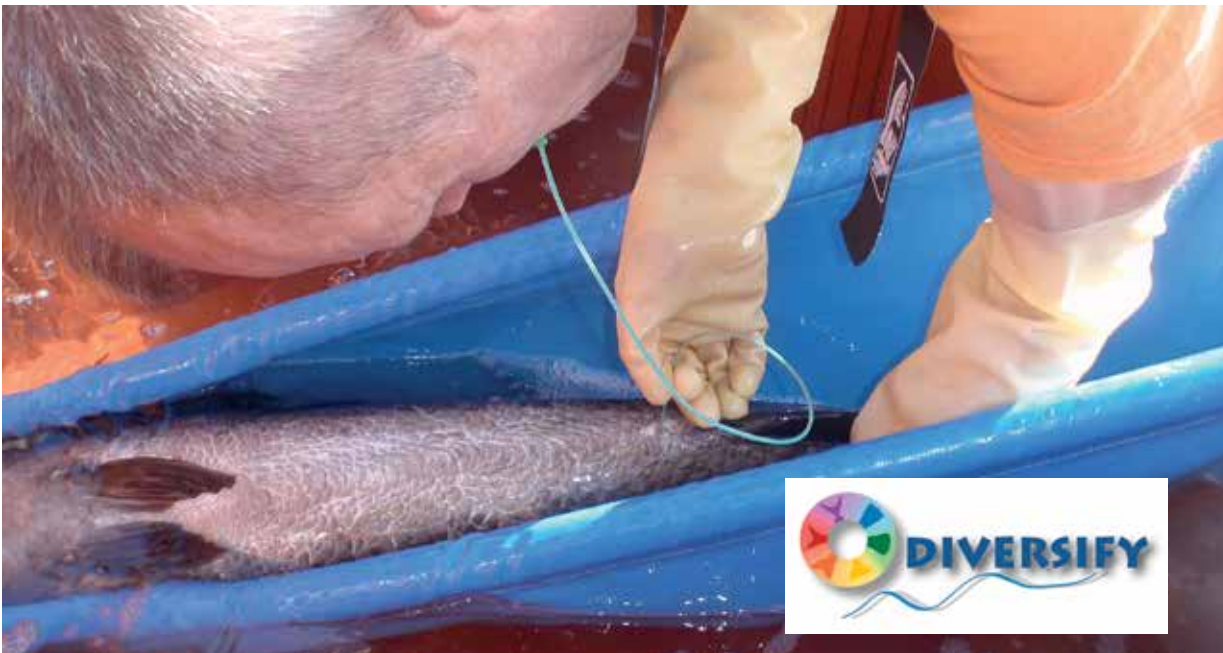
*Sachi*



The 2014-2016 EAS Board. From left to right: Elena Mente (GR), Diego Mendiola (ES), Kjell Maroni (NO), Bjorn Myrseth (NO), Sachi Kaushik (FR), Herve Migaud (UK), Margriet Drouillon (BE) and Hilde Toften (NO). Inset – Stefan Meyer (DE).



**MARK YOUR CALENDAR  
ROTTERDAM, OCT 20-23, 2015**



Taking an ovarian biopsy from meagre to determine stage of maturity.

# ADVANCES IN MEAGRE (*Argyrosomus regius*) RESEARCH DURING THE FIRST YEAR OF THE PROJECT "DIVERSIFY"

CONSTANTINOS MYLONAS AND ROCIO ROBLES

During the first year of the DIVERSIFY project (Dec 2013–Nov 2014), a variety of research activities have been undertaken with meagre, and a summary of the most relevant results is provided below.

## REPRODUCTION

An evaluation of the genetic variation of a large number of the available captive meagre broodstocks of 13 research institutes and SMEs from 7 European countries has been carried out by Fundacion Canaria Parque Cientifico Tecnológico de la Universidad de las Palmas de Gran Canaria (FCPCT, Spain, Dr. J.M. Afonso) using 2 multiplexes of 18 microsatellite markers. The examined broodstocks, as a whole, appeared to originate from three different populations and sufficient genetic variation exists to form a base population for a breeding program (Fig. 1). However, care will be needed in selecting families within each broodstock and an increase in the number of families is recommended, in order to avoid problems and ensure improvement of desirable traits.

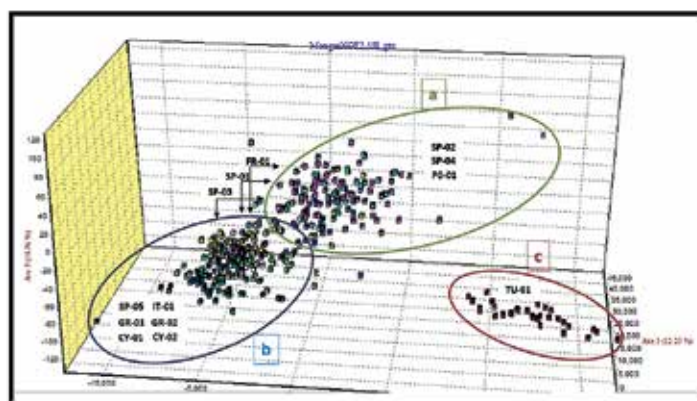


Fig 1.- Factorial Correspondence Analysis from 18 loci and 376 fish distributed in 13 Mediterranean meagre broodstocks maintained in captivity for research or aquaculture production, showed three different original populations.

continued on page 6

Updated findings of all of the species involved in the DIVERSIFY project will be presented in a session entitled “New and Emerging Finfish Species” at Aquaculture Europe 2015 in Rotterdam



continued from page 5

Paired crossings with six pairs of females and males were carried out in the Institute de Recerca I Tecnologia Agroalimentàries (IRTA, Spain, Dr. N. Duncan). Spawning was induced with GnRH $\alpha$  injections (15  $\mu\text{g Kg}^{-1}$  for females and 7.5  $\text{g kg}^{-1}$  for males) every 7-10 days. Breeders that did not spawn after 2-3 induced spawning attempts were replaced. A total of 41 different pairs were induced to spawn, of which 10 pairs produced >500,000 eggs, 16 pairs produced >250,000 eggs and 19 pairs produced >100,000 eggs that hatched (Fig. 2). Poor spawning results were not caused by maturity status, repeated spawning or inductions, and different individuals had clear differences in egg production and quality.

An additional experiment was also carried out at the Hellenic Center for Marine Research (HCMR, Greece, Dr. C.C. Mylonas) with four pairs of breeders to determine how many successful spawns can be produced in response to consecutive weekly injections of GnRH $\alpha$ . Up to 17 consecutive spawns were obtained with high quality eggs that had >80% hatching success and larval survival to 5 days post hatch. These two trials demonstrated that paired spawning of high quality eggs is possible, and the method could be used in breeding selection programs.

### LARVAL CULTURE

A weaning assay was carried out in IRTA (Dr. A. Estevez) to advance the time for weaning in meagre. Larvae were weaned either at age 12, 15 and 20 (the usual age) days post hatch (dph) using half the amounts of enriched *Artemia* metanauplii and a commercial weaning diet (Gemma Micro, Skretting). Growth (Fig. 3), survival rate (Fig. 4), fatty acid composition as well as digestive system development (histology and enzymes) were analysed. A high incidence of cannibalism was detected from day 12 dph onwards, resulting in very low survival (2-3.3%). The experiments will be repeated in 2015 and several new approaches

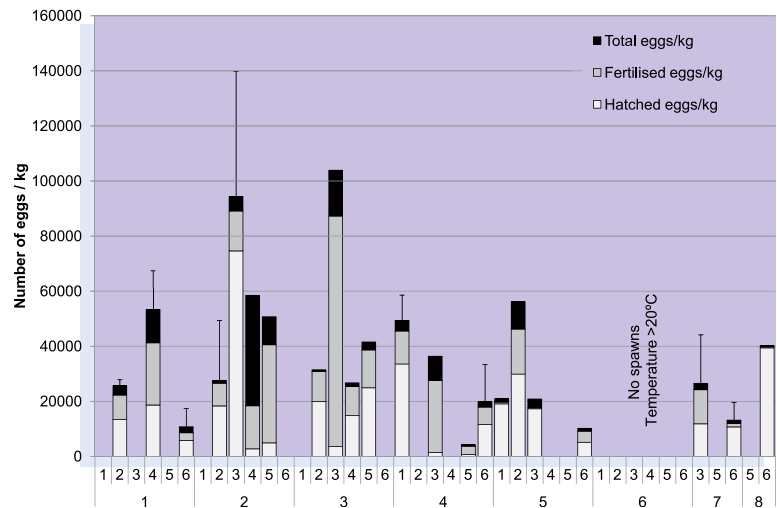
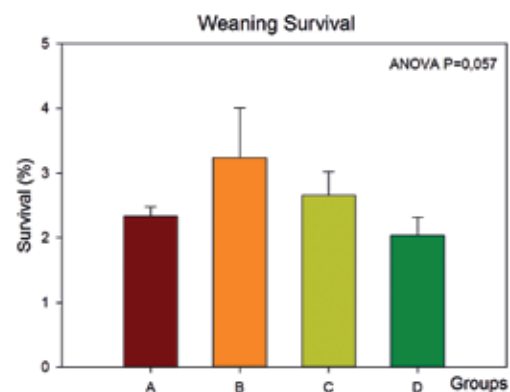
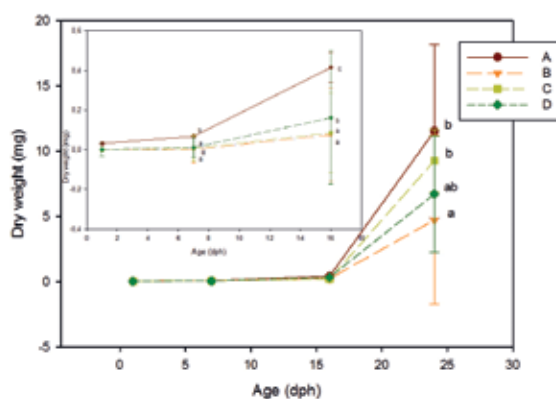


Fig 2.- Mean ( $\pm$  SD) daily fecundity of meagre in response to multiple (8) GnRH $\alpha$  injections. Total number of eggs was multiplied by percentage fertilization and hatch to determine the number of fertilized eggs and larvae produced.

will be taken, including increasing the photoperiod to give more chances of the fish to eat the weaning diet or increase the initial stocking density.

### NUTRITION

A trial was conducted by FCPCT (Dr. L. Robaina) to investigate the requirements of meagre larvae for n-3 HUFA in relation with vitamin E (vit E) and vitamin C (vit C). After feeding the larvae with combinations of different levels of n-3 HUFA (0.5 and 3.5%) and vit E and vit C (150 vit E + 180 vit C, 300 vit E + 180 vit C and 300 vit E + 360 vit C) from day 14 to 28 dph, results showed a clear improvement in growth when dietary n-3 HUFA levels were 3.5%, whereas the effects of vit E or vit C and the interaction between both nutrients and the n-3 HUFA levels were not significant. Regarding biochemical composition, larval contents of n-3 HUFA reflected clearly dietary levels (table 1), being significantly higher in larvae fed fish oil, and elevation of dietary n-3 HUFA and vit E + vit C tended to increase larval lipid contents. Study of larval foregut histological characteristics showed that larvae fed 0.5% HUFA presented very pigmented enterocytes with centered nucleoli and very little lipid vacuoles, while larvae fed higher levels of dietary HUFA, such as in the 3.5/150/180 combination, showed larger



Figs. 3 and 4: Growth (dry weight in mg) and survival (%) of larvae of the different groups after weaning.





Table 1.- Culture performance and morphometric parameters of meagre larvae (initial total length 4.07±0.26 mm and dry weight 0.058±0.01 mg) fed early weaning diets containing several n-3 HUFA, vitamin E and vitamin C levels from 14 dph to 28 dph.

|                              | Diet        |             |             |             |             |             |
|------------------------------|-------------|-------------|-------------|-------------|-------------|-------------|
|                              | 0.5/150/180 | 0.5/300/180 | 0.5/300/360 | 3.5/150/180 | 3.5/300/180 | 3.5/300/360 |
| <b>Total length (24 dah)</b> | 4.8±0.44b   | 5.0±0.39a   | 4.9±0.40ab  | 5.0±0.45a   | 5.0±0.48a   | 5.1±0.38a   |
| <b>Total length (28 dah)</b> | 5.2±0.46ab  | 5.2±0.43ab  | 5.1±0.51ab  | 5.3±0.44a   | 5.0±0.31b   | 5.3±0.59a   |
| <b>Dry weight (24 dah)</b>   | 0.19±0.04c  | 0.21±0.02bc | 0.20±0.03bc | 0.21±0.02bc | 0.22±0.02ab | 0.24±0.03a  |
| <b>Dry weight (28 dah)</b>   | 0.23±0.02   | 0.21±0.04   | 0.21±0.03   | 0.27±0.05   | 0.23±0.05   | 0.24±0.04   |
| <b>Survival (%)</b>          | 12.1±4.9    | 8.0±5.2     | 15.1±4.1    | 14.2±8.3    | 16.7±3.5    | 15.2±7.7    |

\* Values (mean ± standard deviation) with the same letters are not significantly different; ANOVA. P<sub>Length</sub> <0.01; P<sub>Weight</sub> <0.05.

and more developed enterocytes containing lipid vacuoles around the nucleus, reflecting the higher lipid absorption activity. These results suggest that there is a high requirement of this species for n-3 HUFA to promote growth, and vit E and vit C to prevent fatty acid oxidation during larval stages. Thus, weaning diets for meagre larvae must be supplemented with increased n-3 HUFA, vit E and vit C in order to be improved. Selected diets were used to conduct studies on resistance to handling stress, stress bio-markers such as gene expression of HSPs (FCPCT), specific fish behaviour, evaluation of metabolic cost after sub-lethal stress, video analysis of activity, escape responses and sensory acuity (Danmarks Tekniske Universitet, Denmark, Dr. I. Lund) and digestive enzyme (protease, amylase and lipase) and gut ATPase activities (University of La Laguna, Spain, Dr. C. Rodriguez).

In the following years, the essential fatty acid requirements will be examined in grow out diets (Skretting Aquaculture Research Center, Norway, Dr. R. Fontanillas) for meagre by feeding six levels of docosahexaenoic, eicosapentaenoic and arachidonic acids (FCPCT). During the last three months of 2014, information on the nutritional requirements of meagre have been collected and a basal diet formulation for grow out has been defined.

### ONGROWING

Size variability in juvenile meagre during pre-grow out makes regular grading essential to avoid cannibalism, and grades of smaller fish may be related to poor performance when transferred to sea cages. Experiments were carried out by IRTA using meagre juveniles of a mixture of 5 known families, to simulate the commercial hatchery situation and in order to study differences in growth rate. Juveniles were separated into three size grades, and were stocked into tanks at the same initial density and fed the same commercial diet. After 4 months the distribution of all the size grades across the different tanks / grades was compared and 70% of the population was observed to be in the size range of 15-30 g (Fig 5).

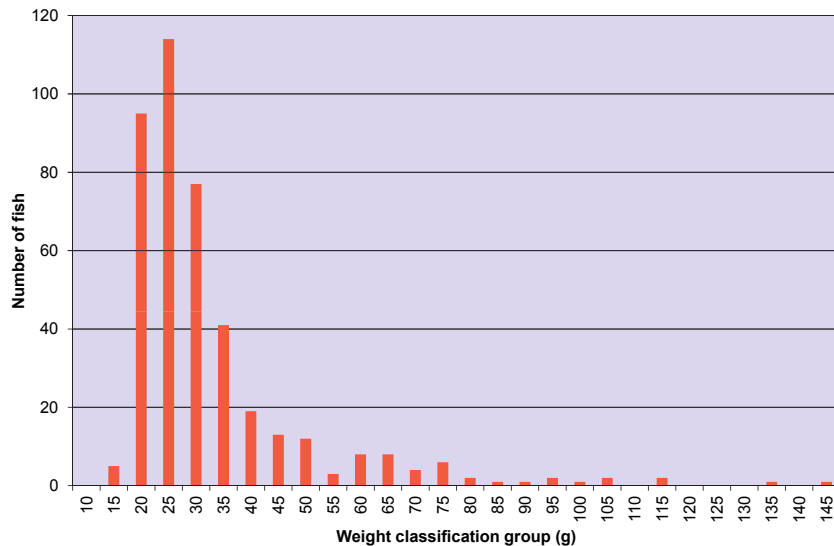


Fig 5.- Frequency distribution of meagre in each 5-g size classification. The weight shown is the upper value of the classification, for example classification 15 g contains fish from 10.1 to 15 g.

The population was skewed to larger fish with 30% of the population in the range of 30-145 g and this wide dispersion of sizes made management difficult. The normally distributed 70% of the population was graded into three grades of 73 large fish (25-30 g), 89 medium fish (20-25 g) and 86 small fish (15-20 g) and growth was monitored.

A random sample of 50 fish from each group was weighed and measured (length) every 3 weeks. The large fish have grown from 27.2±1.5 g to 113±21.0 g, medium fish have grown from 22.7±12.2 g to 94.2±19.8 g and small fish have grown from 17.9±1.8 g to 71.6±31.31 g (Fig. 6). On all sample dates there have been significant differences (ANOVA, P<0.05) between the grades, and the fish in each group have grown significantly during the study (ANOVA, P<0.05). The different size grades appeared to have very similar growth potential. The trial finished on 11th December 2014 and the fish will be characterised genetically for parentage assignment (HCMR, Dr. C. Tsigenopoulos) to establish if differences in growth were a consequence of genetic origin.

The effect of cage depth on meagre grow out was studied by HCMR. The trial started in May 2014 using cages of 180- m<sup>3</sup> (6x6x5 m, Shallow) and 290- m<sup>3</sup> (6x6x8 m, Deep) at the Souda Bay pilot cage farm (Dr. N. Papandroulakis) in duplicates, and juveniles

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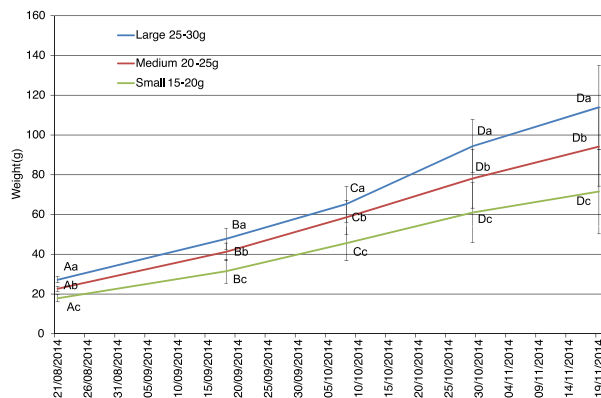


Fig 6.- Mean ( $\pm$  SD) wet weight of meagre classified to three grades as large (initially 25-30 g), medium (initially 20-25 g) and small (initially 15-20 g). Capital letters represent significant differences ( $P < 0.05$ ) between sample dates for the same size grade. Lower case letters represent significant differences ( $P < 0.05$ ) between size grades on the same sample date.

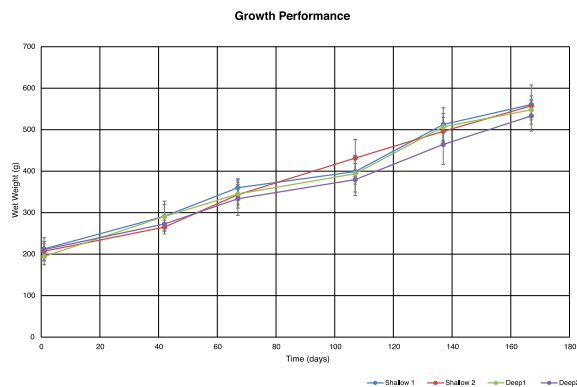


Fig 7.- Mean ( $\pm$ SD) growth performance of meagre at the Souda Bay pilot cage farm (HCMR).

obtained from the HCMR hatchery. Eggs were from a single spawning and larval rearing was performed at the Mesocosm hatchery. Juveniles of 2 g were transferred at the cage facility and they were reared under similar conditions until the beginning of the trial. Then, 4 groups were created, two of ~5,150 fish for the 180-m<sup>3</sup> cages and two of ~8,240 fish for the 290-m<sup>3</sup> ones. The wet weight at the beginning of the trial was 200 $\pm$ 20 g. During the experiment, growth performance was estimated with monthly samples (Fig. 7).

Every second month, blood samples were taken for haematological (hematocrite, hemoglobin), biochemical (osmotic pressure, glucose, lactic acid, free fatty acids), immunological (lysozyme, myeloperoxidase serum) and hormonal (cortisol) evaluation. The samples are currently being analyzed.

The vertical distribution in cages was monitored using an echo integrator. Although a technical problem has not allowed the monitor during the first month of the trial, an upgraded system (CageEye 1.3, Lindem Data Acquisition AS, Norway) was installed in June 2014 and the trial was implemented as planned without further alterations. The analysis of the data is not completed yet, but an interesting observation has

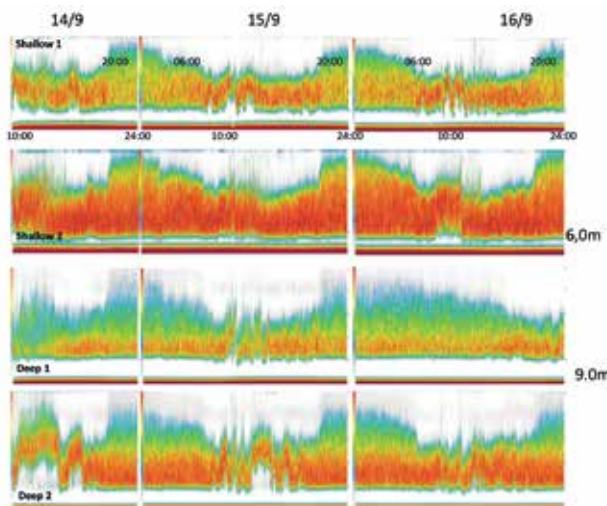


Fig 8.- Vertical distribution of meagre in the experimental cages for a period of 3 days.



Monitoring growth of meagre juveniles

been made already. The vertical distribution of meagre shown for a period of 3 days (Fig. 8), demonstrates clearly that the fish are located mostly at the lower half of the cage for a period of ~12 hours, while the rest of the period are distributed almost homogeneously in the whole available volume of the cage. This observation is independent of the cage depth and it is correlated with the light and dark period of the day. To our knowledge this is the first time that such behavior has been observed.

### HEALTH

Meagre were sampled for collecting data on specific growth rate and to collect chronological samples for the immune ontogeny study. Duplicate sets of samples were collected at each time point (Fig. 9); one set was fixed in formalin for histological analysis, and a second set was collected in RNAlater for extraction of RNA to be used in gene expression analysis. As fish became more developed and organ tissues were recognized easily, individual tissue samples were collected in formalin and RNAlater. Tissues collected included spleen, head kidney, gills and intestine. Samples for immune gene expression analysis are being stored at -80°C. The original plan was to collect animals that were of a medium size, as well as animals from the larger end of the growth spectrum to see how differential growth may lead to premature immune maturation. We eliminated this idea due to a reduction in the overall size of the population. The original population was diminished greatly due to cannibalism during the growout.

A search of the online database GenBank was performed to identify and collect existing sequences





for genes of interest from extant marine teleost species for the study of the immune system. The sequences collected were used for the preparation alignments for designing degenerate/consensus primers for amplification from cDNA of meagre tissues. Samples for the preparation of RNA and subsequent synthesis of cDNA for preparation of these gene expression assays (table 2) has already been done during the growout period of fish being used in the earlier experiment. All of this process for isolation of gene sequences and development of the specific gene expression assays were initiated in the first quarter of 2015. ■



**Co-funded by the Seventh Framework Programme of the European Union**

This 5-year-long project (2013-2018) has received funding from the European Union’s Seventh Framework Programme for research, technological development and demonstration (KBBE-2013-07 single stage, GA 603121, DIVERSIFY). The consortium includes 38 partners from 12 European countries –including 9 SMEs, 3 Large Enterprises, 5 professional associations and 1 Consumer NGO- and is coordinated by the Hellenic Center for Marine Research, Greece. Further information may be obtained from the project site at “www.diversifyfish.eu”.

**Sampling Schedule**

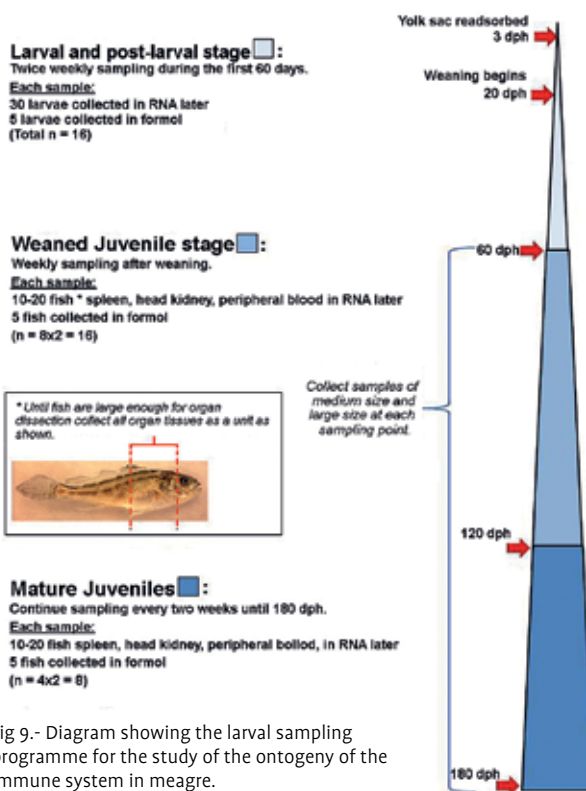


Fig 9.- Diagram showing the larval sampling programme for the study of the ontogeny of the immune system in meagre.

Table 2.- Genes targeted for characterization of the immune system of meagre. The unknown gene sequences should provide amplicon sizes approximating those shown, if there exists a high degree of conservation between species. These estimates are based upon data from existing sequences found in GenBank.

|                       | Target Gene                                    | Degenerate/Consensus Primers | Amplicon size |
|-----------------------|--|------------------------------|---------------|
| Endogeneous Controls  | EF1 (Elongation Factor)                        | X                            | 230           |
|                       | GAPDH (Glyceraldehyde Phosphate Dehydrogenase) | X                            | 239           |
|                       | 18S  | X                            | -             |
| Innate Immunity       | Piscidin1 (“Defensin”)                         | X                            | 110           |
|                       | Piscidin2 (“Defensin”)                         | -                            | -             |
|                       | Piscidin3 (“Defensin”)                         | -                            | -             |
|                       | Lysozyme                                       | X                            | 220           |
|                       | Metallothionein                                | X                            | 80            |
|                       | MX protein                                     | X                            | 570           |
|                       | NOD2 (Toll Like Receptor - TLR)                | X                            | 1390          |
| Adaptive Response     | RAG1 (Recombination Activating Gene)           |                              |               |
|                       | IgM  |                              |               |
|                       | IgT  |                              |               |
|                       | TcR (T-cell Receptor)                          |                              |               |
|                       | C3 (complement)                                | X                            | 1202          |
|                       | TNFa (Tumor Necrosis Factor)                   | X                            | 250           |
|                       | IFN alpha (interferon)                         |                              |               |
|                       | IFN gamma                                      |                              |               |
|                       | IL-1beta (Interleukin)                         |                              |               |
|                       | IL-2   |                              |               |
|                       | IL-4   |                              |               |
| IL-10                 |  |                              |               |
| IL-17                 |  |                              |               |
| IL-22                 |  |                              |               |
| Inflammatory Response | COX2 (cyclooxygenase 2)                        | X                            | 1500          |
|                       | MyD88 (myeloid differentiating factor)         | X                            | 130           |



We have the pleasure to welcome you to Aquaculture Europe 2015, to be held from October 20-23 in Rotterdam and with the theme “Aquaculture, Nature and Society.

Abstract submission is now open and we are proposing a very wide range of parallel sessions for you to submit to. Abstracts should be submitted before May 1.

The following articles give you some idea of recent developments in Dutch aquaculture that show the diversity and innovative approaches being made in this small country - one that has a long history in aquaculture research and farming systems for finfish and for shellfish.

The Netherlands also plays a significant role in seafood processing, traceability and international trade.