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- 1 The ontogeny of greater amberjack digestive system under different rearing
- 2 conditions: a histological and enzymatic approach

- 4 J.A. Pérez<sup>1\*</sup>, I.E. Papadakis<sup>2\*</sup>, N. Papandroulakis<sup>2</sup>, L. Cruces<sup>1</sup>, E. Cotou<sup>3</sup>, E. Gisbert<sup>4</sup>, A.
- 5 Lorenzo<sup>1</sup>, C.C. Mylonas<sup>2</sup>, C. Rodríguez<sup>1</sup>
- 6 \*These authors equally contributed to the study

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- <sup>1</sup> Departamento de Biología Animal, Edafología y Geología. Universidad de La Laguna.
- 9 38206 La Laguna, Tenerife, Spain
- 10 <sup>2</sup> Institute of Marine Biology, Biotechnology and Aquaculture. Hellenic Center for
- 11 Marine Research, P.O. Box 2214, Iraklion, Crete 71003, Greece
- <sup>3</sup> Institute of Marine Biology, Biotechnology and Aquaculture, Hellenic Center for
- 13 Marine Research, Agios Kosmas, Hellinikon 16777, Athens Greece
- <sup>4</sup> Institut de Recerca i Tecnologia Agroalimentaries, Centre de Sant Carles de la Ràpita
- 15 (IRTA-SCR), Crta. Poble Nou km 5.5, 43540 Sant Carles de la Ràpita, Spain

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- 21 Corresponding author: J.A. Pérez. Departamento de Biología Animal, Edafología y
- Geología. Facultad de Ciencias, Sección Biología. Universidad de La Laguna. 38206 La
- Laguna, Tenerife, Spain. **E-mail:** janperez@ull.edu.es; **Phone number**: +34 922318340

## **Abstract**

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26 An overall synchronization of morphological and physiological ontogenetic events of the digestive system occurred in greater amberjack (Seriola dumerili) larvae reared 27 28 under intensive (INT) or semi-intensive (MES) conditions for 30 days. The first differentiations of the digestive channel took place at 3-4 days post hatch (dph) (3.6-3.7) 29 mm TL). Differentiation of the endocrine and exocrine pancreas begun at 4-5 dph (4.0-30 31 4.1 mm TL), coinciding with a decrease of carbohydrase activity from egg to the onset 32 of exogenous feeding and the maintenance of bile salt-activated lipase and total alkaline proteases. Between 6 and 10 dph (4.1-4.5 mm TL), pepsin remained undetected and 33 pancreatic enzymes raised their activities compared to the lecitotrophic stage (0-5 dph). 34 35 The first gastric glands and pepsin activity were evident at 17-20 dph (5.5-6.2 mm TL) 36 in both fish groups. The appearance of pyloric caeca had a 5-day delay in the INTcompared to the MES-larvae (28 vs 23 dph). Althought both rearing protocols did not 37 globally differ in terms of somatic growth and maturation of the digestive function, 38 39 oxidative stress appeared to be less severe in larvae reared in MES, which may be 40 attributed to a lower stress and more stable culture conditions with respect to INT.

- 42 Key words: greater amberjack larvae, ontogeny, digestive system, digestive enzymes,
- 43 oxidative stress.

## 44 Introduction

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The greater amberjack Seriola dumerili is a marine warm-water large teleost fish with rapid growth and excellent flesh quality, which is receiving increasing research attention in the last years due to its great potential for aquaculture diversification (Rodríguez-Barreto et al. 2012, 2017; Mylonas et al. 2016; Zupa et al. 2017a,b; Jerez et al. 2018; Monge-Ortiz et al. 2018; Fakriadis et al. 2019; Sarih et al. 2019). Although knowledge about the optimal feeding regime for larval rearing is essential for a successful production of fry, available information on greater amberjack is still incomplete (Papandroulakis et al. 2005; Hamasaki et al. 2009; Mylonas et al. 2016). In fact, larval rearing is considered one of the major bottlenecks for the flourishing culture of this species, due to the low survival rates obtained during this period, which seriously compromise the availability of juveniles for the on-growing stage. Therefore, there is a great need to evolve culture techniques for mass seed production of the greater amberjack. Although there is scattered information about greater amberjack larviculture and larval development, studies evaluating different rearing technologies through physiological parameters, including the morphophysiological development of the digestive function and oxidative stress condition, are missing. Moreover, a holistic understanding of the feeding ecology and digestive functions of greater amberjack larvae is crucial for the design of specific dietary regimes, as well as for the adaptation of rearing protocols to meet larval requirements. This knowledge will allow the best presentation of prey and microdiets as well as synchronizing the larval stage of development with rearing processes in order to increase the productivity of this species during the first life stages. The digestive system enables fish to capture, ingest, digest and finally absorb

nutrients from the food, which are transported across the intestinal epithelium to the

circulatory system and then to the whole organism (Rønnestad et al. 2013). During the first stages of development until its transformation into a juvenile, numerous changes take place in the digestive system of fish larvae in terms of its morphology and functionality (Przybył et al. 2006; Papadakis et al. 2009, 2013; Gisbert et al. 2018). Over this period, activity of the digestive enzymes is affected by a number of different factors, and their levels are closely related to the state of maturation of the secreting digestive tissues, which show important variations between species, water temperature and rearing conditions (Lazo et al. 2011; Koven et al. 2019). Therefore, the knowledge of the digestive competence of a fish is essential in order to understand the digestive physiology of larvae and to adjust the feeding protocols to dietary qualitative and quantitative characteristics (Campoverde et al. 2017; Gisbert et al. 2018) contributing to the optimization of diets (Zambonino-Infante et al. 2008; Campoverde et al. 2017) and to the proper understanding of functions and limitations in the processing capacity of the digestive system. In this sense, the combined analysis of the ontogeny of the main digestive tract structures and related digestive enzymes during larval development is of special relevance for proper characterization of the plasticity of digestive processes to deliver nutrients to the rapidly growing larval tissues (Rønnestad et al. 2013) under changeable feeding and environmental conditions. Early life stages of marine fish are particularly sensitive to environmental stressors and rearing conditions, due to the lack or low functional capacity of some organ systems and to the high rates of metabolism needed to fuel growth and development (Pimentel et al. 2015). Thus, in order to better elucidate the possible impact of rearing protocols on larval condition and physiological function, oxidative stress biomarkers are broadly considered as reliable indicators (Izquierdo et al. 2013; Saleh et al. 2014; Garrido et al.

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2017; Suzuki et al. 2018). Oxidative stress is a progressive imbalance between reactive

oxygen species (ROS) production and the antioxidant defense system (*i.e.* prooxidant/antioxidant imbalance). The production of ROS is controlled by an efficient endogenous antioxidant capacity, characterized by a set of antioxidant enzymes, which can together detoxify ROS (Abele & Puntarulo 2004) and reduce their negative effects at cellular level that finally impact on the health and fitness of the organism (Malanga *et al.* 2004). The antioxidant defense system, which is comprised of endogenous enzymes such as superoxide dismutase (SOD), glutathione peroxidase (GPx), glutathione reductase (GR) and glutathione S-transferase (GST), is designed to maintain the lowest potential level of ROS in cells and is recognized as an essential component of the organism's response to maintain homeostasis (Castex *et al.* 2010; Adeyemi 2014). Moreover, non-enzymatic defenses such as reduced glutathione (GSH) levels are also employed to provide such protection (Harman 1972).

The present study aims to describe the morphological and physiological ontogeny of the digestive system and the response of the antioxidant defences of greater amberjack larvae cultured under two different rearing conditions, intensive and semi-intensive. The intensive rearing conditions and the feeding protocol used is an adaptation from the established methods in Meditteranean hatcheries whereas the semi-intensive methods that allow the use of wild plankton seems to be more appropriate for the culture of this species (Papandrolakis *et al.* 2005). Although there exist several studies describing larval development in other carangid species (Carton 2005; Stuart *et al.* 2011; Martínez-Montaño *et al.* 2016; Plaza *et al.* 2017 among others), there is missing information about greater amberjack. In addition, species-specific conclusions can not be drawn from the above-mentioned studies in order to be applied on greater amberjack larviculture, since each species presents its own developmental patterns and requirements that need to be individually defined. Our results may facilitate the

implementation of an adequate feeding strategy adapted to the digestive capacity and nutritional needs of greater amberjack during early development, while also addressing options to reduce cannibalism and size dispersion which are of primary importance to boost larval survival and growth. This information will be of great value for synchronizing the larval stage of development with rearing practices; thus, improving actual larval rearing protocols for greater amberjack.

#### Materials and methods

127 Larval rearing

Larval rearing trials were performed in the facilities of the Institute of Marine Biology,

Biotechnology and Aquaculture at the Hellenic Center for Marine Research (HCMR,

Crete, Greece). Greater amberjack larvae were cultured under two different rearing

conditions, the semi-intensive (MES) and the intensive (INT) systems. Eggs used for

this study were obtained from induced spawning of wild breeders kept in Argosaronikos

SA cage farm (ARGO). After collection, eggs were shipped by air to the hatchery

facilities of the HCMR in polystyrene boxes (~12 hours trip).

Semi-intensive system (MES)

A total of one hundred and ten thousand eggs were stocked in a 40 m<sup>3</sup> indoor tank (2.75 eggs  $1^{-1}$ ) filled with filtered (5 µm) natural seawater (salinity 37 psu) treated with UV, which was also the water used for subsequent renewal. During the larval rearing, seawater temperature was maintained at  $24.0 \pm 0.7^{\circ}$ C and the pH fluctuated from 8.0 to 8.2. Dissolved oxygen varied from 5.8 to 6.8 mg  $1^{-1}$ , whereas the rate of water renewal was increased progressively from the initial 15% to 35% of total water volume per day at 17 days post hatching (dph), to 100% at 22 dph and then to 200% at 30 dph. Aeration

was provided at 5 points along the perimeter and in the center of the tank. A surface skimmer was operational during the appropriate period (5 to 13 dph) to keep the surface free from lipids, a requisite for good swim bladder inflation. The photoperiod was adjusted to constant light from mouth opening to 20 dph and then reduced to 18L:06D for the remaining experimental period. Light intensity varied according to the weather conditions between 500 lux on cloudy days to 1,000 lux on sunny days. During the night when prolonged photophase was applied, light intensity was about 250 lux. The rearing technology employed here is a semi-intensive technique for production of greater amberjack larvae (Papandroulakis *et al.* 2005).

*Intensive rearing in closed water recirculation system (INT)* 

Thirty-six thousand eggs were placed in 0.5 m<sup>3</sup> cylindro-conical tanks (72 eggs 1<sup>-1</sup>) connected to a closed water recirculating system coupled to a biological filter. Tanks were filled with borehole 35 psu-water kept at 24.0 ± 0.5°C, pH ranged from 8.0 to 8.2 and the dissolved oxygen was maintained between 6.8 and 7.2 mg 1<sup>-1</sup>. Water circulation was achieved through a biological filter during embryogenesis, egg hatching and the autotrophic larval stage with aeration provided at 150-250 ml min<sup>-1</sup>. After first feeding, water circulation was obtained for each tank by means of an airlift pump in order to maintain stable the rearing environment. The water in the biological filter was used for renewal in the larval rearing tanks at a rate of 3% daily until 15 dph, then increased gradually to 50% until 25 dph. A skimmer was installed at the appropriate period (5-15 dph) to keep the surface free from lipids. The photophase was 24L:00D from mouth opening until 20 dph, and 18L:06D for the remaining experimental period. Light intensity varied between 200-800 lux during the day, and was ~200 lux at night.

#### Feeding protocols

The duration and type of diet for each rearing protocol during the trial is presented in Figure 1. Microalgae (*Chlorella sp*) and rotifers (*Brachionus sp*) enriched with DHA Protein Selco (INVE S.A., Belgium) were daily added in the rearing tanks from 3-4 dph to 23 dph. Rotifers' concentration was kept at 2-3 individuals ml<sup>-1</sup> in the MES and at 4-5 individuals ml<sup>-1</sup> in the INT. Instar I *Artemia* AF nauplii (12 to 14 dph) and *Artemia* EG instar II nauplii (14 to 30 dph) enriched with A1 DHA Selco (INVE S.A.) were offered to the larvae at a starting concentration of 0.05 to 0.35 nauplii ml<sup>-1</sup>. Enrichment in all cases was performed according to manufacturer's instructions. In both rearing systems, microdiets were added progressively according to fish size from 16 dph (MES) and 21 dph (INT) (NRD 2/4, grain size 200–300 μm; NRD 3/5 grain size 300–500 μm, INVE S.A., Derdenmonde, Belgium). In the semi-intensive system the feeding was supplemented with potential wild prey developed in the tank (mainly harpacticoida copepods) and also, after 20 dph, with eggs (live or frozen) of gilthead sea bream (*Sparus aurata*) and newly hatched larvae. The egg/larvae addition was based on observations during rearing of other pelagic species.

#### Sampling procedures

As no tank replication was available for the MES treatment, larvae within the 40 m<sup>3</sup> tank were collected from three different and distant tank areas in order to allow for larval spatial and temporal heterogenity to attain meaningful significance levels (Gamble 1990). Once captured, fish larvae were sacrificed with an overdose of anesthetic phenoxy ethanol (Sigma-Aldrich, Darmstadt, Germany). For the histological study, random samples of greater amberjack eggs and larvae (n = 10) were collected on the following days: 1 day before hatching, 0 (hatching), 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 12,

13, 15, 17, 20, 23, 25, 28 and 30 dph. The total length (TL) of the larvae was determined under graduated stereoscope. Fish samples were preserved in a buffered fixative containing 4% formaldehyde and 1% gluteraldehyde for at least 24 hours (McDowell & Trump 1976). For comparisons of main digestive pancreatic and gastric enzyme activities, eggs and larvae were collected and pooled according to their age-size (3 replicates per experimental condition) at different ontogenic periods: 0-5, 6-10, 11-15 and 21-30 dph (1500-2000, 380-450, 50-80 and 5-9 larvae, respectively), rinsed in distilled water to remove external salt and immediately frozen at -80 °C until analysis. Finally, for the study of the oxidative stress status,  $125 \pm 25$  mg wet weight of larvae per stage were collected at 7 dph, 18 dph (flexion), 23 dph and 30 dph. The samples at 23 dph included big (B) and small (S) size larvae (TL 8.52  $\pm$  0.65 mm and 5.95  $\pm$  0.49 mm, respectively).

All animal experiments were approved by the Ethics Committee of the Institute of Marine Research and the relevant veterinary authorities (Ref Number 255332) and were conducted in certified laboratories (EL91-BIOexp-04) in accordance with legal regulations (EU Directive 2010/63).

## Histological analyses

Before embedding in methacrylate resin (Technovit 7100®, Heraeus Kulzer, Germany), larvae were dehydrated in gradually increasing ethanol solutions (70-96%). Serial sections of 3 µm were obtained with a microtome (Leica, RM 2245, Germany). Sections were stained with Methylene Blue (Sigma-Aldrich)/Azure II (Sigma-Aldrich)/Basic Fuchsin (Polysciences Inc., Warrington, PA, USA) according to Bennett *et al.* (1976). In order to describe the ontogeny of the digestive system and stomach content analysis, all

the sections were examined using a compound microscope (Nikon Eclipse 50i, Melville, NY, USA).

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min per ml of homogenate at 37 °C.

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Determination of digestive enzyme activities

Determinations of pancreatic (α-amylase, bile salt-activated lipase, total alkaline proteases) and gastric (pepsin) enzyme activities were based on methods previously performed and described by Gisbert et al. (2009) and processed as recommended by Solovyev & Gisbert (2016) in order to prevent sample deterioration. Briefly, samples were completely homogenized (Ultra-Turrax T8, IKA©-Werke, Germany) in 5 volumes (v/w) of ice-cold Milli-Q water, centrifuged at 3,300 x g for 3 min at 4°C, and 1 mlaliquots of supernatant kept at -80°C until their analysis for enzyme quantification. Larval digestive capacities during early life stages before stomach development and acidic digestion was evaluated by measuring the activity of total alkaline proteases according to the method of García-Careño and Haard (1993). Alkaline proteases activity was determined at room temperature using azocasein (0.5%) as substrate in Tris-HCl 50 nmol 1<sup>-1</sup> (pH 9) for 10 min. Reaction was stopped with 20% TCA (trichloroacetic acid) and samples were centrifuged at 10,000 x g for 5 min and absorbance of the supernatant was measured at  $\lambda = 366$  nm. One unit (U) of alkaline proteases activity was defined as 1 µmol of azo dye released per minute and per ml of homogenate. Alpha-amylase (E.C. 3.2.1.1) was quantified according to Métais & Bieth (1968) using 0.3% soluble starch dissolved in Na<sub>2</sub>HPO<sub>4</sub> buffer (pH 7.4) as substrate. The reaction was stopped with 1 N HCl and, after the addition of 2 ml of N/3000 iodine solution (Merck, Darmstadt, Germany) the absorbance was read at  $\lambda = 580$  nm. Alpha-

amylase activity (U) was defined as the amount of starch (mg) hydrolysed during 30

Bile salt-activated lipase (BAL, E.C. 3.1.1) activity was assayed for 30 min at 30°C using p-nitrophenyl myristate as substrate dissolved in 0.25 mM Tris-HCl, pH 9.0, 0.25 mM 2-methoxyethanol and 5 mM sodium cholate buffer. The reaction was stopped with a mixture of acetone: n-heptane (5:2), the extract centrifuged at 6,000 x g and the increase in absorbance of the supernatant determined at  $\lambda = 405$  nm. BAL activity (U) corresponded to the  $\mu$ mol of substrate hydrolyzed per min per ml of enzyme extract (Iijima et~al. 1998).

Finally, pepsin (E.C. 3.4.23.1) was quantified by mixing the enzymatic extract with the substrate (2% hemoglobin solution in 1 N HCl at pH 2.0) and incubated for 10 min at 37 °C. The reaction was stopped with 5% TCA and the extract centrifuged at 4,000 x g for 6 min at 4 °C. The absorbance of the supernatant was read at  $\lambda = 280$  nm. One unit of activity (U) was defined as 1  $\mu$ mol of tyrosine liberated per min and ml of homogenate (Worthington Biochemical Corporation, 1972).

Enzymatic activities are expressed as the total activity defined as units per larva (U/larva) and specific activity as units per mg protein (U/mg protein). Soluble protein of crude enzyme extracts was quantified by means of the Bradford's method (Bradford 1976) using bovine serum albumin as standard. All the assays were made in triplicate from each pool of larvae and absorbance read using a spectrophotometer (Beckman Coulter DU800, Fullerton, CA).

### Determination of oxidative stress status

To assess the oxidative stress of the developing greater amberjack larvae, the concentration of total GSH and the enzymes' activities of total SOD, GPx, GR and GST were determined. Supernatants were prepared and analyzed for enzyme activities in a manner similar to that described in Hamre *et al.* (2014) with minor modifications. In

brief, 25 mg (for GSH) and 100 mg (for SOD, GPx, GR and GST) of frozen samples from each biological sample were placed in Eppendorf tubes (1.5 ml). A 12 x volume of ice-cold homogenization buffer was added in each sample, homogenized with a pellet pestle (cordless motor, Sigma-Aldrich) and centrifuged at 3,000 x g for 10 min or at 10,000 x g for 20 min, for GSH or for all the other enzymes, respectively. Homogenization buffer for GSH was a 5% (w/v) metaphosphoric acid and 0.6% sulfosalicylic acid (w/v) mixture, while for all the other enzymes a 0.1 M phosphate buffer pH 7.4 containing 0.15 M KCl, 1 mM dithiothreitol (DTT), 1 mM EDTA and 0.1 M phenyl-methylsulfonylfluoride (PMSF) was used. Supernatants were collected and stored at -80 °C until analysis. The GSH was analyzed according to Rahman *et al.* (2006) and the GPx, GR and GST according to McFarland *et al.* (1999). The SOD activity was analyzed with a commercial kit (706002, Cayman Chemical Co., MI) according to the manufacturer's instructions. Total protein concentrations were measured with a Coomassie brilliant blue reagent (Sigma-Aldrich) according to Bradford (1976).

Statistical analysis

Data are presented as mean ± SEM unless otherwise stated. For the description of growth performance as total length (TL) as a function of time, an exponential equation was used. Prior to the statistical analysis of the activities of digestive and antioxidant enzymes the data were checked for normality with the one-sample Kolmogorov–Smirnov test and for homogeneity of variances with the Levene's test. A one way analysis of variance (ANOVA) followed by a Tukey post hoc test or a Games-Howell test were performed to determine significant differences among developmental stages (unless otherwise stated). A Student's *t*-test was applied for comparisons between both

- rearing systems (MES vs INT). The level of significance was established at P < 0.05.
- All statistical comparisons were conducted using SPSS for Windows 21.0 (IBM-SPSS
- 295 Inc., Chicago, Il, USA).

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#### Results

- 298 *Growth performance*
- No differences in growth of greater amberjack larvae were observed between both
- 300 culture systems during the experimental period (P > 0.05). Growth performance
- adjusted exponentially to the equation:  $y = 3.223 e^{0.033x}$ ,  $R^2 = 0.9436$ , for the MES and y
- $= 3.3065 e^{0.0294x}$ , R<sup>2</sup> = 0.9703, for the INT rearing systems (Figure 1).

- 304 *Ontogeny of the digestive system*
- 305 Overall, there was a synchronization of the ontogenetic events occurring in greater
- amberjack larvae from both rearing protocols during the first life stages (Figure 2).
- 307 *Period 1: 0-5 dph (MES, 3.5 4.0 mm TL; INT, 3.5 4.0 mm TL)*
- From hatching (3.5  $\pm$  0.04 mm TL) until 2 dph (3.7  $\pm$  0.05 mm TL), the digestive tract
- appeared as a closed straight tube located dorsally to the yolk sac (Figure 3a) and it
- 310 consisted of a single-layer epithelium of cuboidal and columnar cells. The liver
- developed rapidly. The early hepatic cells appeared at 2-3 dph (3.8  $\pm$  0.1 mm TL) and
- 312 were initially located behind the yolk sac under the anterior intestine and later
- 313 surrounding the anterior part of the intestine (Figure 3b). The pancreas appeared as an
- undifferentiated tissue at 2 dph, but differentiation in endocrine and exocrine regions
- begun between 4-5 dph  $(4.0 \pm 0.1 \text{ mm TL})$  (Figure 3c).
- Regardless of the rearing protocol, the first differentiation events of the digestive
- system took place at around 3-4 dph. During this period, the mouth and the anus opened

- and the separation of the digestive tract into distinc regions occurred (Figure 3d). The
- 319 ileorectal valve that separates the midgut from the hindgut also appeared at this stage
- 320 (Figure 3e).
- Regarding digestive enzymes, α-amylase, BAL and total alkaline protease activities
- were detected at this stage, even prior to hatching and to the onset of exogenous feeding
- 323 (3 dph;  $3.8 \pm 0.1$  mm TL). The specific activity of  $\alpha$ -amylase (U/mg protein)
- significantly decreased from the egg to 5 dph larvae (Figure 4; ANOVA, P < 0.05),
- while that of BAL and total alkaline proteases remained stable (ANOVA, P > 0.05).
- 326 Pepsin activity was not detected during this developmental stage.
- 327 *Period 2: 6-10 dph (MES, 4.0 4.2 mm TL; INT, 3.9 4.3 mm TL)*
- Within this period, both feeding regimes were based mainly on rotifers. At 6 dph (3.9-
- 329 4.0 mm TL), folding of the oesophageal mucosa occurred (Figure 3f), whereas the
- 330 pyloric and cardiac sphincters at the intestine indicated the area where the stomach will
- start developing (Figure 3g). Supranuclear vacuoles were present in the larval hindgut at
- 8 dph in the MES (4.1  $\pm$  0.1 mm TL) and at 10 dph in the INT (4.3  $\pm$  0.1 mm TL)
- 333 (Figure 3h), being visible up to 30 dph in both rearing systems.
- In addition, pepsin remained undetected, whereas BAL activity was maintained
- unvariable and that of amylase and alkaline proteases increased with respect to the
- previous period when data are expressed as U/mg protein (ANOVA, P < 0.05) but
- remained constant when reported as U/larva (Figure 4; ANOVA, P > 0.05).
- 338 *Period 3: 11-15 dph (MES, 4.5 5.0 mm TL; INT, 4.5 5.3 mm TL)*
- The first taste buds were formed along the buccopharyngeal epithelium at 12 dph (4.5)
- mm TL) (Figure 5a), whereas goblet cells appeared at the oesophagus at 15 dph (INT =
- $5.3 \pm 0.2$  mm; MES =  $5.0 \pm 0.2$  mm), increasing their number over time (Figure 5b). At

- this age, the first pharyngeal teeth also appeared at the posterior part of the buccopharynx area (Figure 5c).
- During this period, the activity of pancreatic enzymes ( $\alpha$ -amylase, BAL and total alkaline proteases) remained stable compared to the previous stage (ANOVA, P > 0.05), whereas pepsin was firstly detected in both rearing systems. When comparing the activity of larval digestive enzymes between treatments at 12 dph,  $\alpha$ -amylase was higher in intensive- than in semi-intensive-reared larvae (Figure 6; t-test, P < 0.05), BAL and total alkaline proteases presented the opposite trend (P < 0.05), and pepsin did not show any significant variation (Figure 6; t-test, P < 0.05).
- 351 *Period 4: 20-30 dph (MES, 6.5 8.7 mm TL; INT, 6.0 8.2 mm TL)*
- 352 The first gastric glands at the pyloric portion of the stomach were evident at 20 dph (6.0
- $\pm$  0.2 mm TL) in the INT group and at 17 dph (5.7  $\pm$  0.4 mm TL) in the MES group
- 354 (Figure 5d).
- Supranuclear bodies in the midgut appeared between 20-23 and 20-25 dph in INT
- and MES-larvae, respectively (Figure 2). The first goblet cells were evident in the
- midgut at 23 dph (7.2  $\pm$  0.7 mm TL) in the MES group and at 25 dph (7.8  $\pm$  1.0 mm TL)
- in the INT one (Figure 5e). Moreover, the appearance of pyloric caeca had a 5-day delay
- in larvae reared under intensive conditions compared to those of the semi-intensive
- group (28 vs 23 dph) where mean fish length were 8.5  $\pm$  1.2 and 7.2  $\pm$  0.7 mm TL,
- 361 respectively (Figure 5f).
- As it is shown in Figure 4, pancreatic enzyme activities were similar to those of the previous stage (ANOVA, P > 0.05) when given as U/mg protein, but dramatically
- 364 increased when reported per individual larvae. Regardless of the units considered,
- pepsin significantly increased in comparison to previous stages (ANOVA, P < 0.05). At
- 366 30 dph, the activity of pepsin was *ca*. 2.5 times higher in internsive-reared larvae than in

MES-larvae (P < 0.05) whereas alkaline proteases was higher in the INT group only when expressed as U/mg protein (Figure 6).

Oxidative stress

Regarding the antioxidant system and the levels of the oxidative stress biomarkers, GPx, SOD, and GST activities tended to significantly decreased in every developmental stage compàred to 7 dph (Figure 7; ANOVA, P < 0.05). In addition, significantly lower activities of GPx, SOD and GST were also evident in MES-larvae compared to INT-larvae in most developmental stages (Figure 7; t-test, P < 0.05). On the other hand, GSH content was significantly higher at 18 dph than at 7 dph for both rearing systems (Figure 7; t-test, P < 0.05), following a trend to a progressive reduction of the enzyme content with larval age. A gradual decrease of GR activities was also recorded with age which was however not significant between culture conditions, with the exception of those measured at 7 dph which were lower for the MES group (Figure 7; t-test, P < 0.05).

#### **Discussion**

Ontogeny of greater amberjack digestive system

The ontogeny of the digestive system of greater amberjack can be considered as a rapid process, similarly to other carangid species like *S. lalandi* (Chen *et al.* 2006) and *S. rivoliana* (Teles *et al.*, 2017). The development of the digestive capacity is controlled by endogenous factors and generally, it is genetically programmed, which results in similar developmental patterns between teleost fish larvae (Rønnestad *et al.* 2013). However, the time of appearance of the digestive structures and their functionality can be influenced by a number of factors of which temperature and feeding regime are among the most critical issues (Kamler 2002). As developmental rates are correlated with the

larval TL, rearing protocols have also to be synchronized with larval size. Generally, in marine fish species, the feeding protocol has to include firstly rotifers, whereas *Artemia* nauplii have to be offered at larger sizes. However, these general principles have to be tailored for each fish species in order to adapt rearing conditions to larval ontogeny. In this sense, the histological description of the main digestive organs, the ontogenetic profile of digestive enzymes, and their dietary adaptation may be used as reliable indicators of larval development, food acceptance, digestive capacity and of their further larval performance (Ueberschär 1993; Rønnestad *et al.* 2013). It is of special relevance to merge data in the same study from two types of approaches, *i.e.* morphological (histological analysis) and functional (assessment of the activity of digestive enzymes) development information on the digestive system; as in some cases the presence of a morphologically distinct organ (*i.e.* stomach) does not match with its functionality (Solovyev *et al.* 2016).

The pattern of development of digestive enzyme activities in *S. dumerili* is similar to that reported to other temperate and warm water marine species (Rønnestad *et al.* 2013), and especially to that of *S. lalandi* (Chen *et al.* 2006). In brief, the activity of the main carbohydrases, proteolytic and lipolytic enzymes were detected before and just after hatching. Regardless of detecting BAL activity during the embryonic and at hatching stages in greater amberjack, it does not mean that embryos and/or newly hatched larvae digest lipids contained in their yolk sac reserves by means of this lipolytic enzyme produced by the exocrine pancreas. In fact, these results indicate that the spectrophotometric method for assessing this enzyme, whose activity is enhanced by means of bile salts (sodium cholate), is not specific (Nolasco-Soria *et al.* 2018) and it may also detect lipases hydrolyzing tryglicerides and wax esters contained in the yolk (Heming & Buddington 1988). Yolk protein serves two primary functions: it provides

amino acids for tissue growth and supplies energy via catabolic processes (Heming & Buddington 1988). However, the detection of alkaline protease activities before hatching may be mainly attributed to the presence of chorionase, an alkaline proteolytic enzyme involved in hatching (Hagenmaier 1974; Yamagami 1988) rather than to yolk protein digestion. These results would be in agreement to those reported by Segner et al. (1989), who stated that trypsin, the main alkaline pancreatic protease, activity was not found in the yolk syncytium of fish larvae. In this sense, yolk amino acids and proteins may be obtained by non-selective bulk endocytosis (Heming & Buddington 1988) and receptor-mediated pinocytosis (Rønnestad & Fihn 1993), respectively. Amylolytic activity have been reported by several authors in the yolk of fish (Gawlicka et al. 2000; Naz 2009, present study) and poultry embryos (Ikeno & Ikeno 1991), although this activity may not be correlated to pancreatic  $\alpha$ -amylase, since the exocrine pancreas was not fully differentiated during the embryo stage. Amylolytic activity during embryonic development and during the lecitotrophic stage may be attributed to carbohydrate (glycoproteins) utilization contained in yolk reserves as source of energy and structural components (Cetta & Capuzzo 1982; Whyte et al. 1993; Kamler 2002).

After hatching, the activity of the above-mentioned enzymes tended to increase in parallel to the development of the exocrine pancreas as it has been reported in other fish species (see review by Rønnestad *et al.* 2013). Similar to most fish species described so far, the increment in activity of pancreatic enzymes in *S. dumerili* would be correlated to the morphogenesis of the exocrine pancreas (4-5 dph;  $4.0 \pm 0.1$  mm TL) and the accumulation of zymogen granules in pancreocytes, and to the complete resorption of the yolk sac and transition to exogenous feeding, as our histological data indicated. These results are in agreement to those reported in another carangid species like the golden pompano *Trachinotus ovatus* (Ma *et al.* 2014) and *S. rivoliana* (Teles *et al.*,

2017). The decrease in activity of  $\alpha$ -amylase from hatching to the onset of exogenous feeding in comparison to values found in embryos may be genetically programmed rather than dietarily induced as it has been reported in most of carnivorous fish larvae, even though the magnitude of the above-mentioned ontogenic decrease in activity is species-specific (Govoni et al. 1986; Zambonino-Infante & Cahu 2001; Rønnestad et al. 2013). In addition, the higher activity of  $\alpha$ -amylase found in S. dumerilii larvae aged 12 dph reared under INT conditions may be attributed to a delay in the maturation of the digestive function (Zouiten et al. 2011), as it has been previously reported when data of α-amylase activity has been used as marker of the functionality of the digestive system and accessory glands (Zambonino-Infante & Cahu 2001, 2007). However, the abovementioned delay in digestive function maturation in S. dumerilii larvae was compensated within a few days for intensive-reared specimens as data from larvae sampled at 30 dph from both rearing systems indicated. After the onset of exogenous feeding, total activity of the three assayed pancreatic enzymes, when expressed as U/larvae, sharply increased until the end of the study at 30 dph (8.2-8.7 mm TL). Similarly to other species, total alkaline proteases and BAL were the main pancreatic enzymes during the first days after first feeding in S. dumerili, indicating that peptides and proteins, as well as lipids (triglycerides) are the principal sources of energy for sustaining larval growth and development. In particular, alkaline proteolytic enzymes, especially trypsin and chymotrypsin the main pancreatic alkaline proteases, are generally regarded as being particularly significant in the early life stages of fish larvae because of the absence of a functional stomach with its acid protease, pepsin (Rønnestad et al. 2013). Under present rearing standard conditions and in agreement with other studies assessing the changes in alkaline protease specific activities along larval development in carnivorous fast-growing species (Chen et al. 2006; Ma et al. 2014;

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Solovyev et al. 2016; among others), major changes in the activity of total alkaline proteases were found between the onset of exogenous feeding until the early juvenile stage at 30 dph. However, the large deviation values in alkaline protease specific activities found in different sampling points (6-10, 11-15 and 20-30 dph) denoted a heterogeneous larval population, which probably masked somehow the abovementioned activity peaks associated with pancreas differentiation and changes in food items. Under the current experimental conditions, alkaline proteolytic specific activity was slightly different among the rearing system (MES vs. INT) and tended to decrease at late larval stages (21-30 dph), concomitantly to the sharp increase in pepsin activity observed during this period. However, when expressed per larvae, the pancreatic protease activity was still high at day 30 post hatching. These results suggested that even though the stomach was functional at early ages (17 dph,  $5.7 \pm 0.4$  mm TL for MES), and peptic activity appeared and increased due to the presence of abundant gastric glands as histological data revealed, the complete transition to a juveniledigestion pattern based on acid digestion was not achieved in S. dumerili specimens aged 30 dph and measuring 8.2-8.7 mm in TL, as the presence of eosinophilic supranuclear bodies in the hindgut also suggested (Ma et al. 2005; Teles et al., 2017). It is really feasible that this process occurred at later stages (Zambonino-Infante & Cahu 2001), but current experimental design and sampling schedule did not allow to validate this hypothesis and further longer studies (>30 dph) are needed to elucidate this concern. Regardless of this fact, present results are in agreement to those reported for T. ovatus, where pepsin activity was detected at higher body sizes of 5 mm in TL and then progressively increased at older ages (Ma et al. 2014). When comparing pepsin activity between both rearing protocols, present results show that regardless of the late appearance of gastric glands in larvae from the INT group in comparison to those larvae

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reared in MES (Figure 2), pepsin activity was higher in *S. dumerili* from the INT group at 30 dph. Such differences in pepsin activity between both systems may be attributed to the feeding protocol and could be considered as dietarily induced rather than differences in the maturation of the stomach, since larvae from the INT were mainly fed on enriched *Artemia* and a compound diet, whereas those from the MES treatment were offered a more diverse diet (*Artemia*, copepods, compound diet, and eggs and fish larvae). Thus, differences in the level of protein, peptide and free amino acid contents between food items may explain such differences in acid protease activity (Zambonino-Infante & Cahu 2007).

BAL is an important enzyme for the hydrolysis of a wide range of lipids such as glycerophospholipids, cholesterol esters and lipid-soluble vitamins (Rønnestad et al. 2013). In S. dumerili, BAL total activity sharply increased with the transition to exogenous feeding and remained stable afterwards, whereas there were not clear differences in activity regarding the larval rearing method considered. In T. ovatus lipase showed two ontogenic activity peaks, one coinciding with first feeding as in the present study in S. dumerili, and a second one when golden pompano were fed with Artemia nauplii (Ma et al. 2014). In contrast, in S. dumerili and S. lalandi this peak in activity associated to a shift in the type of live prey (rofiter vs. Artemia nauplii) (11-15 dph) was not detected, which may be associated to the larval sampling procedure used in the present work where individuals were pooled by periods of 5 - 10 days, whereas in other similar studies larvae were sampled more often. Independently of the rearing system considered, histological data revealed that the percentage of area covered by lipid vacuoles in the liver decreased between 11 and 15 dph in comparison to younger ages. These results differed from those observed in S. rivoliana (Teles et al., 2017), differences that may be attributed to different larval rearing and Artemia enrichment

procedures. Under present experimental conditions, the above-mentioned decrease in the level of accumulation of fat stores in the hepatic parenchyma in *S. dumerili* larvae may be correlated to a change in food items (*Artemia* AF vs. EG nauplii) coupled with a higher energy demand to support the higher somatic larval growth observed during this period. In addition, the former results were also correlated with a decrease in the activity of GR, GSH, GPX and GST enzymes, which is in agreement with the above-mentioned hypothesis.

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The comparison between both larval rearing methods, MES vs. INT, revealed that there was a large conservation in the ontogenic differentiation of digestive structures during the first developmental stages, independently of the rearing procedure applied as it is shown in Figure 2. The only remarkable variations between both culture procedures were found in the final differentiation of the digestive system and in particular related to the appearance of gastric glands and pyloric caeca. These two digestive structures were detected in S. dumerili larvae at smaller sizes in larvae reared in MES (5.7 and 7.2 mm in TL, respectively) in comparison to those from the INT system (6.0 and 8.5 mm in TL, respectively). This pattern was also observed in European sea bass (Dicentrarchus labrax) (Zouiten et al. 2011), as well as in meagre (Argyrosomus regius) (Papadakis et al. 2013; Solovyev et al. 2016) when comparing these two rearing systems. Although there are very few studies evaluating the impact of these two different rearing procedures on larval development and quality, the former authors suggested that the proliferation of wild zooplankton on semi intensive tanks besides the presence of enriched live preys (rotifers and Artemia nauplii) had a key effect on promoting larval development, although this process was compensated within a few days for the intensive-reared larvae (Zouiten et al. 2011). Deviations from these values obtained under standard rearing conditions may be indicative of problems in larval quality, development and/or rearing conditions (Zambonino-Infante & Cahu 2001).

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Antioxidant endogenous system plays an important role in providing cell protection from oxidative stress during larval development (Dandapat et al. 2003). This period is highly demanding in energy and oxygen uptake, thus the influence of a poor nutritional status or other unfavourable conditions can enhance oxidative damage. There have been a number of studies on the antioxidant defence system in fish, particularly in relation to specific oxidative stress conditions and also in relation to age and development (Peters & Livingstone 1996; Dorval et al. 2003; Kalaimani et al. 2007; Skjærven et al. 2013; Hamre et al. 2014; Liravi et al. 2014). Generally, the endogenous antioxidant system includes some enzymes which catalyze the reaction of ROS degradation. Therefore, SOD protects against oxidative damage by catalyzing the reaction of dismutation of the superoxide anion to H<sub>2</sub>O<sub>2</sub>, GRx reduces both hydrogen peroxide and organic hydroperoxides, GR catalyses the reaction to form GSH (glutathione) and maintain a ratio GSH/GSSG (oxidized GSH) under oxidative stress, and GST detoxify some of the secondary ROS produced by reaction with cellular constituents (Rudneva 2013). However, there have not been relevant studies specifically investigating the activity of the antioxidant enzyme systems in developmental larval stages of the greater amberjack. The present study has shown that readily measurable specific activities of most important antioxidant enzymes were present in the first life stages of greater amberjack larvae (from 7 dph to 30 dph) in both rearing technologies. Overall, the results indicated that the activities of GPx, SOD, GR and GST tended to decrease at 18 dph, after the flexion stage, while this reduction was significantly lower in the semi-intensive than in the intensive rearing system. According to literature, a similar progressively reduction in antioxidant enzyme activities with regard, however, to different dph fishes-dependant stages of larval development have been found in Atlantic cod (Gadus morhua) for GPx, but not for SOD (Hamre et al. 2014), in turbot (Scophthalmus maximas) for SOD but not for GR and GPx (Peters & Livingstone 1996), in Dentex dentex for SOD (Mourente et al. 1999) and in silver carp (Hypophthalmichthys molitrix) for GPx and SOD (Liravi et al. 2014). Conversely, a progressively induction for GPx have been indicated in the fast-growing Asian seabass (Lates calcarifer) (Kalaimani et al. 2007) and for SOD in Salmo iridaeus (Aceto et al. 1994). Depletion in GSH content has been attributed either to higher level of ROS production that convert more reduced GSH to its oxidized form (GSSG) (Ou et al. 1996) or to a decreased activity of GR (Costagliola 1991). Thus, the depletion in GSH content observed at 7 dph greater amberjack could be explained by a higher level of ROS production as the activity of GR was significantly higher at this stage compared to all the other stages. The overall patterns of all antioxidant enzymes observed during the developmental stages of S. dumerili larvae might suggest specific compensatory mechanisms of antioxidant defense compensate ROS to production/removal (neutralize ROS) and to eliminate the damage of oxidative stress due to a high metabolic rate (Solé et al. 2004). Present results also revealed that oxidative stress appeared to be more severe in INT compared to MES rearing systems. Reduced antioxidant enzyme activities in semi-intensive systems could have been related to a lower oxidative stress and more stable rearing conditions than in the intensive system. Thus, we assumed that the decrease/increase of antioxidant enzymes activity indicated during larval stages might be related to the changes in the levels of ROS. It has been reported that oxidative stress in aquatic organisms is more profound during nutritional deficiency (Mourente et al. 1999; Hidalgo et al. 2002; Tocher et al. 2003; Morales et al. 2004), elevated temperature (Hwang & Lu 2002), hypoxia (Kolkovski et al. 2000) and exposure to xenobiotics (Pedrajas et al. 1995; Dandapat et

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al. 2003; Peña-Llopis et al. 2003; Rudneva & Zalevskaya 2004). Taken into account the two rearing protocols, we hypothesize that the change in the type of food given together with the food availability after the 12 dph stage and after the 20 dph stage (Figure 1) might have influenced the lower antioxidant enzymes activities obtained after 18 dph compared to the 7 dph stage and in the 23 dph stage compared to 7 and 18 dph stages. Besides, the increase of antioxidant and digestive enzymes activities at similar larval stages (i.e. 7 dph and 6-10 dph, respectively), as well as the progressive decrease in both antioxidant and digestive enzymes activities after the 7 dph stage revealed that both biomarkers followed the same trend regardless of the physiological process considered. Overall, various aspects of intensive aquaculture stimulate stress responses in fish larvae because artificial systems are frequently exposed to a range of 'unnatural' stressors, which are related to rearing practices (Zouiten et al. 2011). Thus, antioxidant enzymes play an important role in inactivation of ROS and thereby control oxidative stress as well as redox signaling. Both processes change across the life span of the organism and thus modulate its sensitivity and resistance against free radical damage.

#### **Conclusions**

This study is the first one describing the morphoanatomical and functional development of the digestive system in *S. dumerili* and assessing the impact of two different larval rearing systems on the larval digestive function. In addition, the integration of data on oxidative stress was used to evaluate ther impact of both rearing protocols in this novel species. Coupling histological and biochemical data we were able to show that *S. dumerili* larvae had a functional digestive system at the onset of exogenous feeding when the digestive process was basically alkaline, regardless of the rearing technology considered. The morphogenesis of the stomach and pyloric caeca were affected by the

rearing protocol, appearing earlier in fish cultured in semi-intensive in comparison to those reared in intensive conditions. However, the above-mentioned differences were compensated within a few days for the intensive-reared larvae as data on pepsin activity indicated. Although both tested rearing strategies did not differ in larval performance in terms of somatic growth and maturation of the digestive function, present results showed that oxidative stress appeared to be more severe in larvae reared in intensive than in semi-intensive systems, which may be attributed to more stable rearing conditions in the semi-intensive rearing technology.

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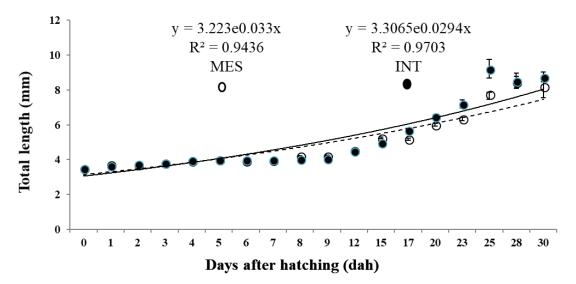
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# **Data Availability Statement**

The data that support the findings of this study are available from the corresponding author upon reasonable request.

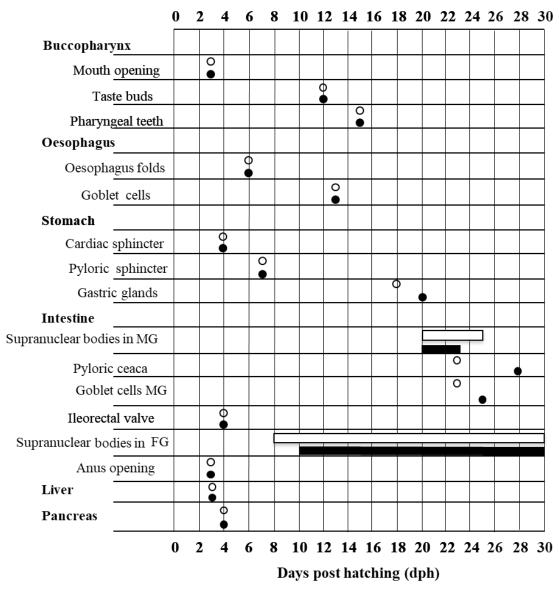
## Figure 1.



## **MES**

| 1,122                       |   |   |   |   |   |   |   |   |   |   |    |    |    |    |    |    |    |    |    |     |                           |      |    |     |    |    |     |           |      |      |      |
|-----------------------------|---|---|---|---|---|---|---|---|---|---|----|----|----|----|----|----|----|----|----|-----|---------------------------|------|----|-----|----|----|-----|-----------|------|------|------|
|                             | 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | 13 | 14 | 15 | 16 | 17 | 18 | 19  | 20                        | 21   | 22 | 23  | 24 | 25 | 26  | 27        | 28   | 29   | 30   |
| Phytoplankton               |   |   |   |   |   |   |   |   |   |   |    |    |    |    |    |    |    |    |    |     |                           |      |    |     |    |    |     |           |      |      |      |
| Rotifers                    |   |   |   |   |   |   |   |   |   |   |    |    |    |    |    |    |    |    |    |     |                           |      |    |     |    |    |     |           |      |      |      |
| Artemia nauplii AF          |   |   |   |   |   |   |   |   |   |   |    |    |    |    |    |    |    |    |    |     |                           |      |    |     |    |    |     |           |      |      |      |
| Artemia nauplii EG          |   |   |   |   |   |   |   |   |   |   |    |    |    |    |    |    |    |    |    |     |                           |      |    |     |    |    |     |           |      |      |      |
| Artificial food             |   |   |   |   |   |   |   |   |   |   |    |    |    |    |    |    |    |    |    |     |                           |      |    |     |    |    |     |           |      |      |      |
| Frozen eggs (Sparus aurata) |   |   |   |   |   |   |   |   |   |   |    |    |    |    |    |    |    |    |    |     | $\mathbb{H}^{\mathbb{N}}$ | ļij. |    | 175 | H  |    | Įij | 175       | ļi;i | ijij | Įij. |
| Fish larvae (Sparus aurata) |   |   |   |   |   |   |   |   |   |   |    |    |    |    |    |    |    |    |    |     | Ш                         |      | Ш  | Ш   | Ш  | Ш  | Ш   | Ш         | Ш    |      | Ш    |
| INT                         |   |   |   |   |   |   |   |   |   |   |    |    |    |    |    |    |    |    |    |     |                           |      |    |     |    |    |     |           |      |      |      |
|                             | 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | 13 | 14 | 15 | 16 | 17 | 18 | 19  | 20                        | 21   | 22 | 23  | 24 | 25 | 26  | 27        | 28   | 29   | 30   |
| Phytoplankton               |   |   |   |   |   |   |   |   |   |   |    |    |    |    |    |    |    |    |    |     |                           |      |    |     |    |    |     |           |      |      |      |
| Rotifers                    |   |   |   |   |   |   |   |   |   |   |    |    |    |    |    |    |    |    |    |     |                           |      |    |     |    |    |     |           |      |      |      |
| Artemia nauplii AF          |   |   |   |   |   |   |   |   |   |   |    |    |    |    |    |    |    |    |    |     |                           |      |    |     |    |    |     |           |      |      |      |
| Artemia nauplii EG          |   |   |   |   |   |   |   |   |   |   |    |    |    |    |    | 13 |    |    |    | 133 | $\odot$                   | 98   | 18 | 18  |    |    | ×   | $\otimes$ | 00   |      |      |
| Artificial food             |   |   |   |   |   |   |   |   |   |   |    |    |    |    |    |    |    |    |    |     |                           |      |    |     |    |    |     |           |      |      |      |

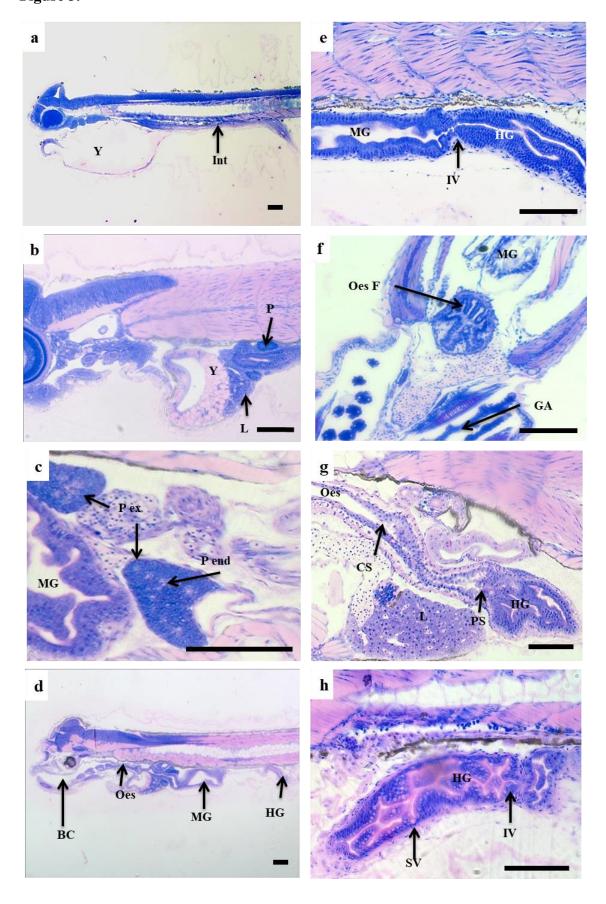
## **Figure 2.**

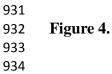


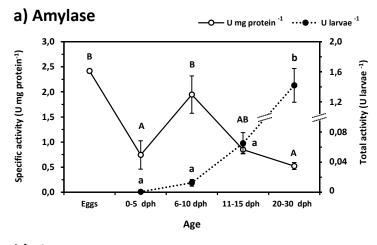
|     | 0    | 1    | 2    | 3    | 4    | 5    | 6    | 7    | 8    | 9    | 12   | 15   | 17   | 20   | 23   | 25   | 28   | 30   |
|-----|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|------|
| INT | 3.52 | 3.72 | 3.73 | 3.81 | 3.96 | 4.01 | 3.94 | 3.98 | 4.23 | 4.23 | 4.55 | 5.26 | 5.19 | 6.01 | 6.35 | 7.77 | 8.47 | 8.20 |
| MES | 3.52 | 3.67 | 3.75 | 3.83 | 3.99 | 4.05 | 4.03 | 4.02 | 4.05 | 4.09 | 4.53 | 5.00 | 5.69 | 6.49 | 7.2  | 9.23 | 8.54 | 8.74 |

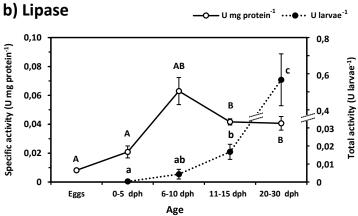
Total length in (mm)

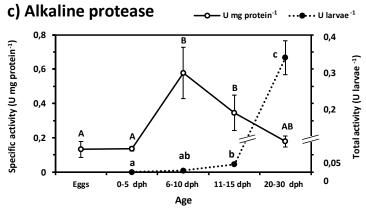
**Figure 3.**930

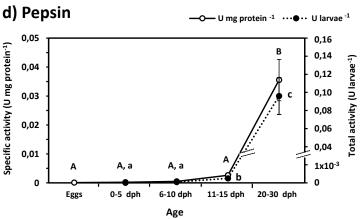




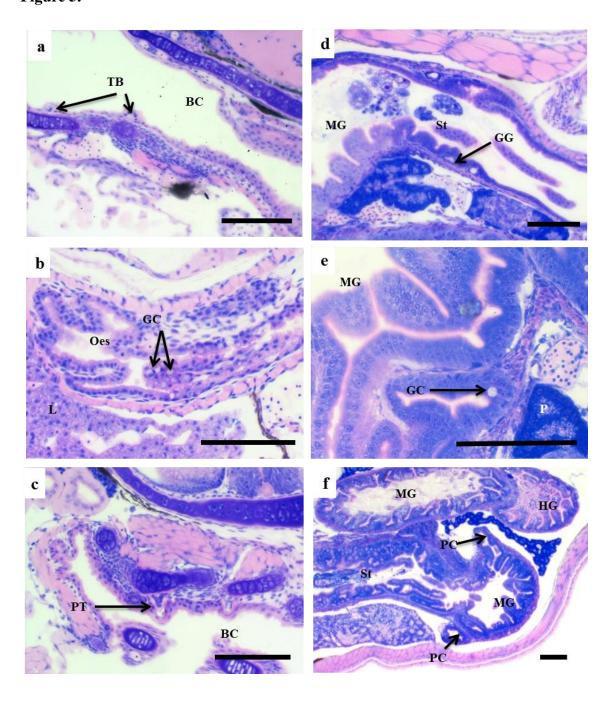








**Figure 5.** 



960 961 Figure 6. 962 a) Amylase a) Amylase Specific activity (U mg protein-1) **■ INT ■ MES ■ INT ■ MES** 1.8 1.0 Totalactivity (U larvae-1) 1.6 1.4 0.8 1.2 0.6 1.0 0.8 0.4 0.6 0.4 0.2 0.2 0.0 12 dph 30 dph 12 dph 30 dph Age Age b) Lipase b) Lipase 0.10 **■ INT ■ MES** ■ INT ■ MES 12 dph 30 dph 12 dph 30 dph Age Age c) Alkaline protease c) Alkaline protease **■ MES ■ INT** ■ INT ■ MES 0.5 Total activity (U larvae-1) 0.4 0.4 0.3 0.3 0.2 0.2 0.1 0.1 0 12 dph 30 dph 12 dph 30 dph Age Age d) Pepsin d) Pepsin 0.08 0.20 Specific activity (U mg protein<sup>-1</sup>) 0.00 0.00 0.00 0.00 0.00 0.01 ■ INT ■ MES **■ MES** ■ INT Total activity (U larvae<sup>-1</sup>) 0.10 20.0

43

12 dph

30 dph

Age

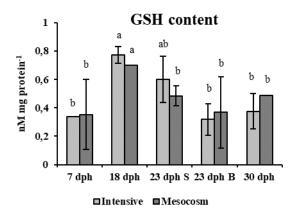
30 dph

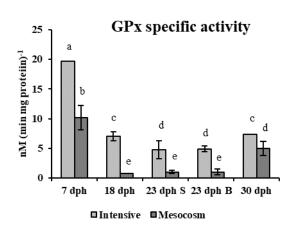
Age

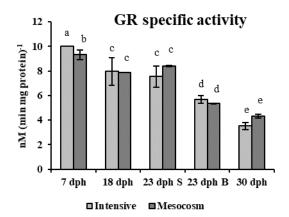
Specific activity (U mg protein-1)

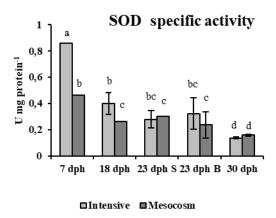
12 dph

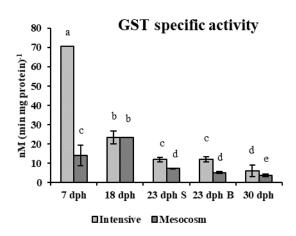
Figure 7.











- Figure 1. Growth performance of greater amberjack larvae (mean  $\pm$  SD of total length)
- 966 cultured in intensive (INT) and semi-intensive (MES) rearing systems. Below the graph,
- the rearing protocols used during the rearing procedure are presented, including type
- and duration of food items provided.
- 969 Figure 2. Schematic representation of the appearance (open solid circles indicate the
- 970 MES, black solid circles indicate the INT system) of the main developmental structures
- 971 examined in greater amberjack larval digestive system, as a function of days after
- 972 hatching (dph, horizontal axis). Horizontal bars (white MES and black INT) indicate the
- period that supranuclear bodies (vacuoles) were present in the anterior-median intestine
- 974 (mid gut, MG) and hindgut (HG). Below, mean values of the total length of greater
- amberiack larvae for each sampling day and rearing system are presented.
- 976 Figure 3. Microphotographs of histological sections from greater amberjack larvae at
- 977 different developmental stages. (a) At 1dph showing digestive tract as a closed straight
- 978 tube located dorsal to the yolk sac. (b) At 2 dph when the liver and pancreas appeared.
- 979 (c) At 4 dph when the pancreas differentiated to endocrine and exocrine parts. (d) At 4
- 980 dph showing the different characteristic areas of the digestive system. (e) At 4 dph when
- 981 the ileo-rectal valve appeared. (f) At 6 dph showing the formation of folds at the
- 982 oesophagus. (g) At 6 dph showing the formation of the stomach area. (h) At 8 dph
- showing the supranuclear vacuoles at the hindgut (MES). BC = buccopharynx; CS =
- cardiac sphincter; GA = gill arches; HG = hindgut;Int = intestine; IV = ileo-rectal valve;
- L = liver; MG = midgut; Oes = oesophagus; Oes F = oesophageal folds; P = pancreas; P
- end = endocrine pancreas; P ex = exocrine pancreas; PS = pyloric sphincter; SV = supranuclear vacuoles; Y = yolk. Bar represents 100 μm.
- 988 **Figure 4.** Specific activity (U mg protein<sup>-1</sup>, white circles) and total activity (U larvae<sup>-1</sup>,
- 989 black circles) of digestive enzymes during the ontogeny of the digestive tract of greater
- 990 amberjack larvae cultured under intensive conditions. Different letters indicate
- significant differences (P<0.05) between larval ages.
- 992 Figure 5. Microphotographs of histological sections of greater amberjack larvae at
- 993 different developmental stages. (a) At 12 dph when the taste buds appeared. (b) At 15
- 994 dph when the goblet cells appeared at the oesophagus. (c) At 15 dph showing the
- 995 pharyngeal teeth at the buccopharynx. (d) At 17-20 dph showing the gastric glands at
- 996 the stomach (MES vs INT). (e) At 23-25 dph showing the goblet cells at the midgut
- 997 (MES vs INT) and (f) at 23-28 dph showing the formation of the pyloric caeca (MES vs
- 998 INT). BC = buccopharynx; GC = goblet cells; GG = gastric glands; HG = hindgut; IV =
- ileo-rectal valve; L = liver; G = midgut, Oes = oesophagus; PT = pharyngeal teeth; PC =
- pyloric caeca; St = stomach; TB = taste buds. Bar represents 100  $\mu m$ .
- 1001 **Figure 6.** Comparison of specific activity (U mg protein<sup>-1</sup>) and total activity (U larvae<sup>-1</sup>)
- of digestive enzymes of greater amberiack larvae between intensive and semi-intensive
- rearing systems at 12 and 30 dph. Asterisks (\*) indicate significant differences (P<0.05)
- between INT and MES rearing systems for a particular enzyme and larval age.
- 1005 Figure 7. Changes in antioxidant defense systems (GSH content and GPx, GR, SOD and
- 1006 GST specific activities) of developmental larval stages of Serioladumerilicultured in
- intensive and semi-intensive rearing systems (S and B refer to small and large larvae,
- respectively). Different letters indicate significant differences (P<0.05) between larval
- 1009 ages.