Effects of Ocean Acidification over successive generations decrease larval resilience to Ocean Acidification & Warming but juvenile European sea bass could benefit from higher temperatures in the NE Atlantic

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Key words: Dicentrarchus labrax, ocean acidification, ocean warming, metabolic rates, larval growth, juvenile growth, teleost

Summary statement: We found that OA did not affect growth, RMR and SMR, while OW increased these traits. OAW decreased larval size at metamorphosis. We conclude that recruitment to nursery areas might decrease under OAW but juveniles might benefit from increased performance at higher temperatures in Atlantic waters.

Running title: OA decreases sea bass larval resilience to OAW, juveniles might benefit from OW

1. Abstract

European sea bass (*Dicentrarchus labrax*) is a large, economically important fish species with a long generation time whose long-term resilience to ocean acidification (OA) and warming (OW) is not clear. We incubated sea bass from Brittany (France) for two generations (>5 years in total) under ambient and predicted OA conditions (PCO_2: 650 and 1700 µatm) crossed with ambient and predicted ocean OW

conditions in F1 (temperature: 15-18°C and 20-23°C) to investigate the effects of climate change on larval and juvenile growth and metabolic rate.

We found that in F1, OA as single stressor at ambient temperature did not affect larval or juvenile growth and OW increased developmental time and growth rates, but OAW decreased larval size at metamorphosis. Larval routine and juvenile standard metabolic rates were significantly lower in cold compared to warm conditioned fish and also lower in F0 compared to F1 fish. We did not find any effect of OA as a single stressor on metabolic rates. Juvenile $P_{O_{2crit}}$ was not affected by OA or OAW in both generations.

We discuss the potential underlying mechanisms resulting in the resilience of F0 and F1 larvae and juveniles to OA and in the beneficial effects of OW on F1 larval growth and metabolic rate, but on the other hand in the vulnerability of F1, but not F0 larvae to OAW. With regard to the ecological perspective, we conclude that recruitment of larvae and early juveniles to nursery areas might decrease under OAW conditions but individuals reaching juvenile phase might benefit from increased performance at higher temperatures.

2. List of abbreviations

Δ1000 – Acidification condition (ambient PCO_2 + 1000 µatm)

A – Ambient PCO_2 condition

BL – Body length

C – Cold life conditioned group

CI – Complex I of the ETS (NADH dehydrogenase)

CII – Complex II of the ETS (succinate dehydrogenase)

dph – Days post hatch

dd – Degree days

DM – Dry mass

ETS – Electron transport system

IPCC – Intergovernmental Panel on Climate Change

MS-222 – Tricaine methanesulfonate

OA – Ocean acidification

OAW – Ocean acidification and warming

OW – Ocean warming

PCO_2 – Partial pressure of CO_2

PO_2 – Partial pressure of O_2

PO_{2crit} – Critical oxygen concentration

RCP – Representative concentration pathway

RMR – Routine metabolic rate

SDA – Specific dynamic action

SMR – Standard metabolic rate

W – Warm life conditioned group

WM – Wet mass

3. Introduction

Climate change is leading to increasing ocean surface temperatures (ocean warming – OW), as well as decreasing ocean pH (ocean acidification – OA). OW as a single stressor on fish metabolism has been investigated intensively since the 1980s in a variety of fish species and life stages and directly influences their metabolism and therefore their growth (Johnson & Katavic, 1986; Peck, 2002; Pörtner, et al., 2007), reproduction success (see review Llopiz, et al., 2014), as well as distribution range and abundance (Turner, et al., 2009; Pörtner, 2006). OW can increase growth rates of larval and juvenile fish (McMahon et al., 2020a; Baumann, 2019; Chauton, et al., 2015), within their thermal window. Although studies on larvae are less numerous than those on adults and juveniles, it has become obvious that larvae are less resilient to OW than adults and juveniles (Dahlke, et al., 2020a).

Initially, fish had been thought to be less vulnerable to OA due to well-developed acid-base regulation systems (Heuer & Grosell, 2014), yet their capacity to cope with OA and ocean acidification and warming (OAW) as co-occurring stressors has been investigated intensively during the last decade with species and life stage specific results (Cattano, et al., 2017): OA levels between 700 and 1600 µatm CO_2 can lead to increased larval growth (mahi-mahi, Bignami, et al., 2014; clownfish, Munday, et al., 2009), but decreased larval swimming performance (mahi-mahi, Bignami, et al., 2014; dolphinfish, Pimentel, et al., 2014) and larval metabolic rates (dolphinfish, Pimentel, et al., 2014). OA also induced severe to lethal tissue damage (cod larvae, Frommel, et al., 2011), decreased swimming performance, maximum metabolic rate and aerobic scope (Australasian snapper juveniles, McMahon et al., 2020b) and increased larval otolith size, with possible implications for hearing sensitivity (cobia and mahi-mahi, Bignami, et al., 2013; 2014, respectively). In other species, growth was decreased by OA (inland silverside juveniles, Baumann, et al., 2012), or not affected (Atlantic halibut juveniles, Gräns, et al., 2014; cobia larvae, Bignami, et al., 2013, Australasian snapper larvae, McMahon et al., 2020a). In some species OA even improved performance, e.g. OA increased survival (Australasian snapper larvae, McMahon et al., 2020a). Dahlke et al. (2020b) showed that Atlantic cod embryos demonstrated poor acid base regulation capacities before and during gastrulation, connected to increased mortality under OA and OAW. On the

contrary, acid base regulation capacities after gastrula were similar to that of adult cod. If both stressors were combined, the effects became more unidirectional and were synergistic in most fish species, e.g. OAW increased growth and survival in larval and juvenile sea bass in their Atlantic populations, but decreased physiological performance (Pope, et al., 2014). The cumulative consequences of these changes are to be determined.

An important factor for projecting whether a species will be able to keep their distribution range under changing conditions, is their potential and capacity to acclimate and adapt over generations. Few studies have so far reared fish for more than one generation or examined transgenerational effects of fish in the context of OAW, with trait- and species-specific capacities to adapt to future conditions. For example, in cinnamon anemone fish (*Amphiprion melanopus*) the negative effect of OA on escape responses was reduced in some traits if parents were exposed to OA (Allan, et al., 2014), whereas in spiny damselfish (*Acanthochromis polyacanthus*), negative effects on olfactory responses were not reduced after parental exposure to OA (Welch, et al., 2014). In addition to the low number of studies on transgenerational effects, they usually used small fish, with short generation times and applied only one stressor, either OW or OA. Little is known about the combined effect of several stressors on economically important larger-sized fish with longer generation times and thus multi-stressor, transgenerational studies on such fish are necessary to project future distribution of fish.

Consequently, in our study we used European sea bass *Dicentrarchus labrax* as a larger, long-lived model species. Sea bass is an economically important species in industrial and recreational fishing as well as in aquaculture (160 000 t in 2015, Bjørndal & Guillen, 2018). Sea bass can reach an age of up to 24 years in the Atlantic population (Irish waters, Kennedy and Fitzmaurice, 1972). Generally rather resilient towards environmental fluctuations, effects of OW and OA have been reported for several seabass life stages: OW increased growth rates in larval sea bass, although at the expense of decreased swimming performance (Atlantic population, 15 to 20°C, Cominassi, et al., 2019). Exposure to OA throughout larval development increased mineralization and reduced skeletal deformities (Atlantic population, 19°C and 15 and 20°C, respectively, Crespel, et al., 2017; Cominassi, et al., 2019). In combination, OAW did not have additional effects on larval growth, swimming ability and development than those already observed separately (Atlantic population, Cominassi, et al., 2019). Juvenile sea bass are highly tolerant to temperature (Dalla Via, et al., 1998; Claireaux & Lagardère, 1999) and show some degree of tolerance to OA as a single stressor at the mitochondrial level (Atlantic population, Howald, et al., 2019). OA and OW acted antagonistically: OW as a single stressor increased growth and digestive efficiency, while OA did not affect these traits. Both stressors combined resulted in reduced growth and digestive efficiency compared to the impact of OW alone. Low food ratios enhanced this effect resulting in an even more pronounced growth and digestive efficiency reduction than under OAW alone (Atlantic population, Cominassi, et al., 2020).

This study aimed to investigate the effect of OAW as well as the effect of OA over two successive generations (F0 and F1) on larval and juvenile growth and metabolism. Therefore, we incubated sea bass from an Atlantic population for two generations (>5 years in total) under current and predicted OA conditions (PCO_2: 650 and 1700 µatm) and applied a warming condition on larvae and juveniles of the F1 generation (ambient, 15-18°C, and ∆5°C, 20-23°C). To study the effect of OA (F0, F1), OW (F1), and OAW (F1) on sea bass, we investigated growth (F0, F1) through ontogeny as a proxy for whole organism fitness. In addition we measured routine metabolic rates (RMR, F1) of larvae, as well as standard metabolic rates (SMR, F0, F1) and critical oxygen concentration (PO_{2crit}, F0, F1) of juvenile sea bass, to unravel the underlying mechanisms resulting in possible growth differences. In F0, no effect of OA on larval and juvenile growth or juvenile SMR and PO_{2crit} were found (Crespel, et al., 2017; 2019). Those traits were compared in F0 and F1 fish to determine the effects due to parental acclimation to different OA levels. Our hypotheses were: (1) OW will lead to increased growth and metabolic rates in F1 larvae and juveniles. (2) OA alone will not have significant effects on larval and juvenile growth and metabolism in F1, as sea bass seem to be quite tolerant to OA and no detrimental effects were found in F0. (3) In combination, OA will lead to synergistic OAW effects, reflected in lower growth in larvae and juveniles.

4. Materials and Methods

The present work was performed within the facilities of the Ifremer-Centre de Bretagne (agreement number: B29-212-05). Experiments were conducted according to the ethics and guidelines of the French law and legislated by the local ethics committee (Comité d'Ethique Finistérien en Experimentation Animal, CEFEA, registering code C2EA-74) (Authorizations APAFIS 4341.03, #201620211505680.V3 and APAFIS 14203-2018032209421223 for F0 and F1, respectively).

4.1. Animals and experimental conditions

Sea bass were reared from early larval stage onwards in two OA treatments in F0 and four OAW treatments in F1. A flow chart summarizing temperature and PCO_2 conditions as well as replicate tank number, tank volume and number of individuals per tank is shown in Figure 1, the timeline of the rearing of the fish is shown in Figure S 1. F0 fish were reared in two OA scenarios, following the predictions of the Intergovernmental Panel on Climate Change (IPCC, 2021) for the next 130 years: today's ambient situation in coastal waters of Brittany and the Bay of Brest (A, approx. 650 µatm (cf. Pope, et al., 2014; Duteil, et al., 2016)) and a scenario according to SSP5-8.5, projecting a ΔPCO_2 of 1000 µatm (∆1000, approx. 1700 µatm). Adults from these two treatments were used in the reproduction experiments to generate F1. Sea bass of F1 were reared under the same OA conditions as their respective parents. Additionally two different temperatures were applied on each OA condition in F1 to create a cold and a warm life condition scenario or four OAW conditions (C-A, C-Δ1000, W-A and W-Δ1000), respectively.

As larvae and post-larval juveniles would display different growth rates under the different life condition scenarios, we adopted the concept of degree days (dph · T(°C)) as a basis for comparison between them. This concept allows to compare them at their physiological age rather than their chronological age and has been shown to be an effective way of normalizing growth at different temperatures (Peck et al., 2012).

Larval rearing was performed in a temperature controlled room and water temperatures were fixed to 19°C in F0, and 15 and 20°C in F1-C and F1-W, respectively. In juveniles and adults, water temperatures of F0 and F1-C sea bass were adjusted to ambient temperatures in the Bay of Brest during summer (up to 19°C), but were kept constant at 15 and 12°C for juveniles and adults, respectively, when ambient temperature decreased below these values. F1-W was always 5°C warmer than the F1-C treatment.

During larval rearing, the photoperiod was set to 24h darkness during the first week and 16h light and 8h darkness (12h each in F1, respectively) per day afterwards. Light intensity increased progressively during the larval rearing period from total darkness to about 100 lux (Table S 1). To work in the larval rearing facilities, headlamps were used (set to lowest light intensity). In the juvenile and adult rearing facilities photoperiod followed natural conditions (adjustment once a week).

4.1.1. F0 generation

4.1.1.1. Larval rearing

F0 larval rearing and origin is described in detail in Crespel et al. (2017; 2019), briefly, larvae were obtained from the aquaculture facility Aquastream (Ploemeur-Lorient, France) at 2 dph (October 2013). F0 larvae were randomly distributed among the two OA conditions described above. Larvae were reared in nine black 38 L tanks initially stocked with *ca*. 2200 larvae $tank^{-1}$ in triplicates for all conditions. Larvae were fed *ad libitum* via continuous delivery of Artemia nauplii until 28 dph. Afterwards, commercial dry pellets (Neo Start, LeGouessant, France) were fed for the rest of the larval period.

4.1.1.2. Juvenile rearing

Juvenile rearing was described in detail in Crespel et al. (2019). Briefly, the early juveniles were counted per tank and transferred from larval to juvenile rearing facilities at approx. 820 degree-days (dd) (45 dph). Juveniles of one condition were combined and kept in square shaped 450L tanks (n=1500 fish per condition). At 8 months (about 250 dph), juveniles were PIT tagged (marked with passive integrated transponders). Juveniles were fed daily with commercial fish food (Neo Start), which was adjusted in size and amount, as recommended by the supplier (Le Gouessant, Lamballe, France). Food ratios were adjusted after each sampling for growth, approx. every 30 days or 3-4 weeks in F0 and F1, respectively (see below), using the formulae provided by Le Gouessant. Daily food ratios were supplied to the tanks by automatic feeders during day time.

4.1.1.3. Adult rearing

During the reproductive season 2017 (fish were 3.5 years old), sex steroid plasma concentration was measured regularly in all adult F0 fish. The individuals with the highest concentrations were kept in round black tanks with a volume of 3 m^3 and a depth of 1.3 m. Each of the two tanks (one for each condition) was stocked with 22 males and 11 females, resulting in fish density of 11.6 kg m^{-3} and 11.0 kg m^{-3} in A and Δ1000, respectively. Mass and length were regularly measured and commercial fish food was adjusted accordingly. Fish were fed Vitalis CAL (Skretting, Norway) during reproduction season and Vitalis REPRO (Skretting, Norway) during the rest of the year. Vitalis REPRO was supplied to the tanks with automatic feeders during daytime. Vitalis CAL was supplied to the tank manually in three to four rations during week days.

4.1.2. F1 generation

Embryos were obtained by artificial reproduction of F0 fish. Succinctly, once the water temperature reached 13°C and the first naturally spawned eggs were observed in the egg collectors, females were injected with gonadotropin-releasing hormone (GnRH, 10 µg kg^{-1}) to accelerate oocyte maturation (23.03.2018). After three days (26.03.2018) eggs and milt were stripped from ripe females and males, respectively, and artificial fertilization was performed following the protocol of Parazo et al. (1998). Briefly, eggs (10 ml l^{-1}) were mixed with sea water and milt (0.05 ml milt L^{-1} seawater). Ten females (1.56 ± 0.24 kg) were crossed with 18 males (1.07 ± 0.16 kg) and 11 females (1.28 ± 0.30 kg) were crossed with 19 males (0.99 ± 0.19 kg) in the A and Δ1000 groups, respectively. Fertilized eggs were incubated in 40 L tanks (without replicates) at 15°C and at the same PCO_2 conditions as respective F0. Hatching occurred after four days (30.03.2018).

4.1.2.1. Larval rearing

Two days after hatch (02.04.2018), larvae were distributed into twelve black 35 L tanks. Triplicate tanks were allocated to each of the four OAW treatments with *ca*. 4500 and 4200 larvae $tank^{-1}$ in A and Δ1000 tanks, resulting in a total of *ca.* 13500 and 12800 larvae $condition^{-1}$ in A and Δ1000, respectively. The temperature of the tanks allocated to warm life condition was increased stepwise by 1 °C day^{-1} during the following five days. Starting at 7 days post-hatch (dph) (mouth opening), larvae were fed with live artemia, hatched from High HUFA Premium artemia cysts (Catvis, AE 's-Hertogenbosch, Netherlands). Artemia were fed to the larvae 24h after rearing cysts in sea water. Larvae were fed *ad libitum* with artemia during the day, excess artemia left the tank via the waste water outflow. Larval mortality was 26-96 %, without any pattern for OAW condition (Table S 2). High mortality of sea bass larvae, especially during early larval rearing are common in science and aquaculture (e.g. Nolting, et al., 1999; Suzer, et al., 2007; Villamizar, et al., 2009). We could not find any signs of infection neither in the tanks with high mortality, nor in the tanks with lower mortality rates. However, as larval mortality was unreasonably high (96%) within the first week in one of the replicate tanks of the W-A treatment, remaining larvae in

this tank were euthanized (sedation followed by an anaesthetic overdose) and not used for further analysis. Water surface was kept free of oily films using a protein skimmer. Water exchange was set to 25 l/h and stepwise increased to 40 l/h at the end of larval rearing.

4.1.2.2. Juvenile rearing

At approx. 950 dd, the early juveniles were counted per tank and transferred from larval to juvenile rearing (48 dph, 17.05.2018 and 63 dph, 01.06.2018 for W and C, respectively). For F1-W, only the Δ1000 fish were transferred to juvenile rearing facilities. Juveniles were randomly allocated to duplicate tanks per condition. Swim bladder test was done at 1680 dd (83 dph, 21.06.2018) and 1661 dd (104 dph, 12.07.2018) for F1-W and F1-C, respectively. Briefly, the fish were anaesthetized and introduced into a test container with a salinity of 65 psu (Marine SeaSalt, Tetra, Melle, Germany). In F1-W, all floating fish with a developed swim bladder were counted and kept in the rearing tanks, resulting in 355 fish per tank (710 fish in total). In F1-C, 410 fish per tank were randomly selected (820 fish per condition), to have similar stocking densities in W and C. Non-floating fish as well as excess F1-C fish were counted and euthanized (sedation followed by an anesthetic overdose). The juveniles were reared in round tanks with a volume of 0.67 m^3 and a depth of 0.65 m. During the first five days after moving to juvenile rearing, the juveniles were fed artemia nauplii and commercial fish food. Afterwards commercial fish food was fed as described above.

4.1.3. Experimental conditions

4.1.3.1. Sea water preparation

The sea water used in the aquaria was pumped in from the Bay of Brest from a depth of 20 m approximately 500 m from the coastline, passed through a sand filter (~500 µm), heated (tungsten, Plate Heat Exchanger, Vicarb, Sweden), degassed using a column packed with plastic rings, filtered using a 2 µm membrane and finally UV sterilized (PZ50, 75W, Ocene, France) assuring high water quality.

Water conditions for the rearing tanks were preadjusted to the desired OAW condition in header tanks. Sea water arrived in a reservoir next to the rearing facilities, after passing the tungsten heater, in F1, two different reservoirs were used to create the different temperature conditions. The temperature controlled water supplied the header tanks within the rearing facilities to adjust the water to the desired OA condition. Each header tank supplied water to all replicate tanks of the respective condition.

In F0 larvae and juveniles the water pH in the header tank was controlled by an automatic injection system connected to a pH electrode (pH Control, JBL, Germany), which injected either air (A) or CO_2 (Δ1000), to control water pH. For the Δ1000 F1 larvae the CO_2-bubbling was installed in the middle of the header tank and the water was mixed continuously with a pump. The CO_2-bubbling was adjusted by a flow control unit, when pH deviated from the desired value.

Older F0-A juveniles (> 2 years) and adults, as well as F1-A larvae and juveniles received water directly from the respective reservoir, without header tank. Additionally as water exchange rates became too high for the automatic injection system and the header tank, PVC columns were installed to control the pH in the rearing tanks. The temperature controlled water arrived at the top of the column and was pumped from the bottom of the column to the rearing tanks. The CO_2-bubbling was installed at the bottom of the column and was adjusted by a flow control unit, when pH deviated from the desired value.

4.1.3.2. Calculation of water chemistry

The Microsoft Excel macro CO2sys (Lewis & Wallace, 1998) was used to calculate seawater carbonate chemistry, the constants after Mehrbach et al. (1973, as cited in CO2sys) refit by Dickson and Millero (1987, as cited in CO2sys), were employed.

From October 2015 onwards (late juveniles of F0), total alkalinity was measured following the protocol of Anderson & Robinson (1946) and Strickland & Parsons (1972): 50 ml of filtered tank water (200 µm nylon mesh) were mixed with 15 ml HCl (0.01 M) and pH was measured immediately. Total alkalinity was then calculated with the following formula:

$$TA = \frac{V_{HCl} \cdot c_{HCl}}{V_{sample}} - \frac{\left(V_{HCl} + V_{sample}\right)}{V_{sample}} \cdot \frac{\{H^+\}}{\gamma_{H^+}} \left[\frac{mol}{l}\right]$$

With: TA – total alkalinity [mol · l^{-1}], V_{HCl} – volume HCl [l], c_{HCl} – concentration HCl [mol · l^{-1}], V_{sample} – volume of sample [l], H^+ – hydrogen activity (10^{-pH}), γ^{H+} – hydrogen activity coefficient (here $\gamma^{H+} = 0.758$).

4.1.3.3. Water quality control

Temperature and pH were checked each morning with a handheld WTW 330i or 3110 pH meter (Xylem Analytics Germany, Weilheim, Germany; with electrode: WTW Sentix 41, NIST scale) before feeding the fish. Until F0 juveniles reached 2 years, the pH meter and the automatic injection system were calibrated weekly with fresh buffers (Merk, Germany). Measured values never differed more than 2% from the target values. Afterwards the pH meter was calibrated daily with NIST certified WTW technical buffers pH 4.01 and pH 7.00 (Xylem Analytics Germany, Weilheim, Germany).

Total pH was determined twice during F0 larval rearing (start and end) and nine times during F0 juvenile rearing following Dickson et al. (2007) using m-cresol purple as indicator. Additionally, water samples were sent to LABOCEA (France) to measure total alkalinity by titration, as well as phosphate and silicate concentration by segmented flow analysis following Aminot et al. (2009).

In later F0 juveniles (> 2 years) and adults as well as F1 larvae and juveniles, total alkalinity was measured monthly or weekly in F0 and F1, respectively, following the protocol described above. Oxygen saturation (WTW Oxi 340, Xylem Analytics Germany, Weilheim, Germany) and salinity (WTW LF325, Xylem Analytics Germany, Weilheim, Germany) were measured together with total alkalinity (monthly in F0 and weekly in F1). The tanks were cleaned daily after pH-measurements. Water flow within the tanks

was adjusted once a week, so that oxygen saturation levels were kept >85%, with equal flow rates in all tanks of one temperature. All water parameters are summarized in Table 1 for F0 larvae and juveniles and Table 2 for F0 adults (two years before spawning) and F1 larvae and juveniles.

4.2. Growth

4.2.1. Larval growth

4.2.1.1. F0 larvae

Larval growth was measured as described in Crespel et al. (2017). Briefly, 10 larvae per tank were sampled each week, starting at 15 dph and ending at 45 dph, when 30 larvae per tank were sampled. For growth measurements, larvae were anaesthetized with phenoxyethanol (200 ppm) and their wet mass (WM), as well as body length (BL) were measured. BL in F0 larvae was measured with a caliper from the tip of the snout to the end of the notochord until flexion, afterwards fork length was considered as BL, see Figure S 2.

4.2.1.2. F1 larvae

In F1 larvae, individuals were sampled every 200 dd from 100 – 900 dd to follow growth throughout the larval phase. At each sampling point, 20 larvae per tank were anaesthetized with MS-222 (50 mg l^{-1}, Pharma Q) prior to feeding and directly photographed individually with a microscope (Leica M165C). The larvae were then frozen in liquid nitrogen and stored at -80°C until dry mass (DM) measurements. The software ImageJ (Schneider, et al., 2012) was used to determine BL of larvae, see Figure S 2 on the definition of BL.

4.2.2. Growth in juveniles

BL and WM were measured approx. every 30 days in F0 and every 3 – 4 weeks in F1 juveniles. Early juveniles were starved for one day prior to growth samplings. Later on, two days of starving were put into practice, to make sure that digestive tracts were empty. Juveniles were caught from their tanks and anaesthetized with MS-222 (Pharma Q). Concentration of anesthetic was adjusted to reach a loss of equilibrium within less than 5 minutes, typically 0.2 g l^{-1}. WM and BL were directly determined with a precision balance (Sartorius MC1 AC210P) and calipers. For all sampling, only the morning hours were used, to avoid diurnal artefacts in data.

4.2.3. Data handling

For F1 larvae and juveniles, mean specific growth rates (SGR [% day^{-1}]) of each tank were calculated after Sutcliffe (1970) with the following formula:

$$SGR = 100 \cdot (e^{g} - 1)$$

The instantaneous growth coefficient (g) was calculated as followed:

335 $$g = \frac{\ln S_1 - \ln S_0}{\Delta t}$$

336 With: S_0 and S_1 – initial and final size (BL, WM or DM) and Δt – time between the two measurements
337 [days]. Initial and final sizes were calculated for three quantiles (0.05, 0.5 and 0.95) for each tank ("ecdf"
338 function in R).
339 Q_{10} was calculated with the following formula:

340 $$Q_{10} = \left(\frac{SGR_W}{SGR_C}\right)^{\left(\frac{10}{T_W - T_C}\right)}$$

341 With: SGR – specific growth rate, T – temperature, W and C as subscripts for W and C condition.

342 **4.3. Respirometry**

343 **4.3.1. F1 larvae**

344 Larval respiration measurements were conducted from approx. 350 to 950 dd in all conditions (18 – 47
345 dph and 25 – 63 dph in W and C, respectively).
346 Larval respiration was measured in an intermittent flow system. The setup consisted of up to eight 4 ml
347 micro respiration chambers with a glass ring (Unisense A/S, Aarhus, Denmark), equipped with a glass
348 coated magnetic stirrer (Loligo® Systems, Viborg, Denmark) and a stainless steel mesh (Loligo® Systems,
349 Viborg, Denmark), to separate the stirrer from the larva. The magnetic stirrers were connected to one
350 stirrer controller (Rank Brothers Ltd., Cambridge, England). The chamber was closed with a custom-
351 made glass lid with three metal ports: two with a diameter of 0.8 mm for water inflow and outflow
352 during the flushing, and one with 1.2 mm to insert the oxygen sensor into the chamber. Oxygen
353 concentration within the chamber was measured with oxygen microsensors connected to a FireSting
354 oxygen meter (PyroScience GmbH, Aachen, Germany). The respiration chambers were placed within a
355 rack without shielding between the individual chambers. The rack holding the respiration chambers was
356 fully submerged in a water reservoir, which received flow through water from the respective header
357 tanks of the larval rearing. Water conditions within the water reservoir were kept at 15.5±1.5 °C and
358 21.2±1.0°C for W and C larvae, combined with the OA condition of the origin tank of the respective
359 larvae. The reservoir was a black container, which shielded the respiration setup from external
360 disturbances. During the flushing periods, water from the reservoir was pumped into the respiration
361 chambers using computer-controlled flush pumps (Miniature DC pump, Loligo® Systems, Viborg,
362 Denmark), relays and software (AquaResp, Copenhagen, Denmark). Four chambers were connected to
363 one flush pump and controlled by one computer. Oxygen microsensors were calibrated to 0% saturation
364 (nitrogen purged seawater) and 100% saturation (fully aerated seawater) prior to each measurement.
365 Respiration measurements were done in the larval rearing facilities with the same light conditions as for
366 larval rearing. Larvae were fasted at least three hours prior to respiration measurements to minimize

the effect of specific dynamic action (SDA) on metabolic rate. Preliminary tests with measurements overnight proved that oxygen consumption during the 12 h after the 3-h fasting period was similar, suggesting no contribution of SDA and thus that the 3h-fasting was sufficient for our setup. Larvae were individually placed in the respiration chambers. Oxygen partial pressure was measured every second for approx. four hours. Cycles were composed of 420 s flush, followed by 60 s wait time (time after flush pump stopped to wait for stable drop in oxygen concentration) and 600 to 180 s measurement time (13-20 cycles per larvae). Measurement time was decreased with increasing larval size. Oxygen concentration was restored to normoxia during the flush time of each cycle and was usually kept above 75% air saturation. Background respiration was measured for 30 min (one slope) after 11 and 18 measurements in F1-C and F1-W larvae, respectively. The mean bacterial respiration was calculated for each temperature treatment and subtracted from total respiration of all larvae of this temperature to obtain oxygen consumption of the larva. Background respiration was typically 0.5 - 6 % of total respiration. Only declines in oxygen concentration displaying $R^2 > 0.80$ were used for analysis. After the measurement, larvae were checked if alive, anaesthetized with MS-222 (50 mg l^{-1} Pharma Q), photographed individually and frozen in liquid nitrogen. Length and DM of the larvae was obtained as described above (see Table S 3). After each experiment, the respiration system was rinsed with fresh water and let dry. For disinfection, respiration chambers, the tubing of the flush pump and the oxygen sensors were additionally rinsed with ethanol, which was allowed to sit in the chambers and the tubing for at least 30 min followed by rinsing with distilled water.

4.3.2. Juveniles

4.3.2.1. Set up F0 juveniles

Measurements on the 15 months old F0 juveniles (F0-old) were described in Crespel et al. (2019), measurements on the 5 months old F0 juveniles (F0-young) were done similarly and if different, the information for F0-young are given in brackets. Briefly, F0 juvenile respiration was measured individually in one of four (eight) intermittent flow respirometry chambers with a volume of 2.1 l (60 ml), which were submerged in a tank which received flow-through seawater at 15 ± 0.25 °C and the respective acidification condition. The water was recirculated within the chamber with a peristaltic pump with gas-tight tubing. The oxygen probe (FireSting oxygen meter, PyroScience GmbH, Aachen, Germany or multichannel oxygen meter, PreSens Precision Sensing GmbH, Regensburg, Germany) was placed within the recirculation loop. Oxygen sensors were calibrated to 0% saturation (sodium sulfite, saturated) and 100% saturation (fully aerated seawater) prior to each experiment. The flush pumps were controlled by relays and software (AquaResp, Copenhagen, Denmark). The setup was placed behind a curtain to avoid disturbances. Background respiration was measured after each experiment and estimated for the whole

experiment by linear regression assuming zero background respiration at the beginning of the run as the entire system was disinfected with household bleach between each trial.

4.3.2.2. Set up F1 juveniles

F1 juvenile respiration was measured in an intermittent flow system. The setup consisted of up to eight 450 ml custom-made respiration chambers. The chambers were made from Lock&Lock glass containers with plastic lid. Four rubber ports were placed into the lid: two for water inflow and outflow during flushing cycles and two to connect the chamber to a mixing pump (Miniature DC pump, Loligo® Systems, Viborg, Denmark). Oxygen concentration was measured with robust oxygen probes placed within the circulation loop and connected to a FireSting oxygen meter (PyroScience GmbH, Aachen, Germany) or to a multichannel oxygen meter (PreSens Precision Sensing GmbH, Regensburg, Germany). The respiration chambers were fully submerged in a flow-through water reservoir. Water conditions within the water reservoir were kept at 14.9±1.0 °C and 22.3±1.8°C for C and W larvae, combined with the OA condition of the origin tank of the respective juvenile. During the flushing periods, water from the reservoir was pumped into the respiration chambers using computer-controlled flush pumps (EHEIM GmbH & Co. KG, Deizisau, Germany), relays and software (AquaResp, Copenhagen, Denmark). Four chambers were connected to one flush pump and controlled by one computer, running either the FireSting or the PreSens oxygen meter. The setup was covered with black foil to avoid disturbances. Oxygen sensors were calibrated to 0% saturation (nitrogen purged seawater) and 100% saturation (fully aerated seawater) prior to each experiment. Background respiration was measured for 30 min (one slope) after each measurement and the run was discarded, if background respiration was >10 %. After each experiment the whole system excluding the oxygen sensors was disinfected with household bleach or Virkon® (Antec International Limited, Suffolk, United Kingdom) and rinsed with freshwater afterwards.

4.3.2.3. Measurement protocol

Respiration measurements of F0 juveniles were done on approx. 5 (119 – 165 dph) and 15 months (454 – 495 dph) old juveniles. F1 juvenile respiration measurements were conducted from 2900 to 3900 dd (137-178 dph, 5 months) and 4700-5100 dd (291-318 dph, 10 months) for F1-W and F1-C, respectively. F1-C fish were older than F1-W fish at the measurement time in order to have comparable fish sizes (see Table S 4).

Juvenile sea bass were fasted for 48-72h prior to respiration measurements to minimize the effect of residual SDA (Dupont-Prinet, et al., 2010). Juveniles were randomly taken from their tank and placed individually in the respiration chambers. The whole setup was shielded from external disturbances with curtains or black foil, but the individual respiration chambers were not shielded from each other. F0 juveniles were chased until exhaustion prior to introduction to the chambers (MMR data partly

published in Crespel et al., 2019). Each experiment lasted for about 70 hours in F0 and 65 hours in F1. Oxygen partial pressure was measured 1/s and was usually kept above 80%, until start of critical oxygen concentration (PO_{2crit}) trial (see below). Each cycle was composed of 360 s (F0) and 540 s (F1) flush time, during which oxygen concentration was restored to normoxia (until PO_{2crit} trial), followed by 30 s wait and 210 s (F0) and 180 s (F1) measurement time. In F0 only the measurements taken after the fish fully recovered from chasing stress were used to calculate SMR, usually after 10 hours. In F1, the first 5 hours of each experiment were not used for analysis of SMR, to account for acclimation of the fish to the respirometer and recovery from handling stress, resulting in approx. 390 and 310 cycles in F0 and F1 juveniles, respectively. Analyses were performed only on declines in oxygen concentration displaying $R^2 > 0.85$ and $R^2 > 0.90$ in F0 and F1, respectively. On the third morning, a PO_{2crit} trial was done on F0-old and F1 juveniles, see below. After finishing the trial or the respiration measurement for F0-young, fish were removed from the chamber. F0 juveniles were weighed and measured in BL prior to the experiment, F1 juveniles after the experiment. F0-old juveniles were identified by their PIT tag and returned to their origin tank after the experiment. F0-young juveniles and F1 juveniles were killed by a cut through the spine after the experiment.

4.3.2.4. Critical oxygen concentration trial

On the third morning, oxygen concentration in the tank surrounding the chambers was continuously decreased, in F0-old by passing the water through a gas equilibration column supplied with nitrogen gas before pumping it to the tank. In F1 the decrease in oxygen concentration was done by bubbling nitrogen directly into the surrounding water bath. The decrease lasted over a period of four to six hours to determine PO_{2crit}. When the fish lost equilibrium in the oxygen depleted chambers, they were removed from their chamber and treated as described above.

4.3.3. Data handling

In F0 juveniles the metabolic rate (MR, in mg O_2 h^{-1} kg WW^{-1} in F0) was calculated by the Aquaresp software. In F1 oxygen concentration was converted from % air saturation to nmol l^{-1} and mmol l^{-1} in larvae and juveniles, respectively ("conv_O2" function of "respirometry" package, (Birk, 2020)). MRs were calculated from the raw data with the following formulas:

$$MR = Slope \cdot V_{Resp}$$

With: Slope – oxygen decline in the respiration chamber during one measurement cycle ([nmol O_2 l^{-1} h^{-1}] and [mmol O_2 l^{-1} h^{-1}] for larvae and juveniles, respectively), V_{resp} – Volume of respirometer [l].

RMR of F1 larvae was calculated as the mean MR throughout the measuring period (approx. 4h). SMR of F0 juveniles was calculated following the protocol of Chabot et al. (2016) as described in Crespel et al. (2019). SMR of F1 juveniles was calculated in R with the "calcSMR" function of "fishMO2" package (Chabot, 2020), derived from this protocol. Briefly, the best SMR was chosen as described in Chabot et

al. (2016) as either the SMR deriving from the mean of the lowest normal distribution (MLND) method (SMR_{MLND}) or the SMR deriving from the quantile method with p=0.2 (SMR_{quant}). SMR_{MLND} was used when the coefficient of variation (CV) was < 7% or < 5.4%, in F0 and F1, respectively, otherwise SMR_{quant} was applied. Both RMR and SMR were divided by fish mass (resulting in RMR_{Raw} and SMR_{Raw}) and then corrected for allometric scaling with the following formulas:

$$RMR = RMR_{Raw} \cdot \left(\frac{DM}{DM_{mean}}\right)^{1-coeff_{Larvae}}$$

$$SMR = SMR_{Raw} \cdot \left(\frac{WM}{WM_{mean}}\right)^{1-coeff_{Juv}}$$

With: RMR_{Raw} and SMR_{Raw} – RMR [nmol O_2 µg DM^{-1} h^{-1}] and SMR [mmol O_2 kg WW^{-1} h^{-1}] calculated as described in the text, DM – larval dry mass [µg], WM – juvenile wet mass [kg], DM_{mean} and WM_{mean} – Mean DM and WM of all larvae and juveniles, respectively, $coeff_{Larvae}$ and $coeff_{Juv}$ – allometric scaling coefficient for larvae (0.89) and juveniles (0.99), respectively. The allometric scaling coefficients used were the slopes of linear regressions of MR over mass in the whole larval (F1) and juvenile (F0 and F1 together) dataset. Q_{10} was calculated with the same formula as used for SGR (see section 4.2.3).

PO_{2crit} was calculated with the "calcO2crit" functions of "fishMO2" package (Chabot, 2020), or according to Claireaux and Chabot (2016).

4.4. Statistical analysis

All statistics were performed with R (R Core Team, 2020). All data were tested for outliers (Nalimov test), normality (Shapiro-Wilk's test) and homogeneity (Levene's test). None of the datasets met the assumptions for ANOVA, therefore all data were fitted to linear mixed effects models (LME models, "lme" function of the "nlme" package, Pinheiro et al., 2017). Rearing tank was included as a random effect in all models. For the respirometry experiments, respirometer was also included as random effect. In case of heterogeneity of data, variance structures were included in the random part of the model. The best variance structure was chosen according to lowest Akaike information criteria (AIC) values. After fitting fixed and random effects, a backwards model selection process was applied to determine the significant and fixed variables and interactions. If significant effects were detected in the linear mixed effect models, posthoc Tukey tests were performed with the "lsmeans" function ("lsmeans" package, Lenth, 2016). Significance for all statistical tests was set at $p < 0.05$. All graphs are produced from the lsmeans-data with the "ggplot2" package (Wickham, 2016). All data are shown as lsmeans ± s.e.m. (standard error of the mean).

4.4.1. Growth data

Larval BL (F0 and F1 larvae)

Larval BL at mouth opening was only measured in F1 larvae. As these were reared in a full factorial design, temperature condition, PCO_2 concentration and their interactions were included as fixed effects in the model. Across generations, the dataset for larval BL at metamorphosis and over time was imbalanced, therefore it was not possible to test the effect of temperature, PCO_2 condition, generation and their interaction separately, instead treatment was used as fixed variable in the model for larval BL at metamorphosis, resulting in six groups: F0-A, F0-Δ1000, F1-C-A, F1-C-Δ1000, F1-W-A and F1-W-Δ1000. For larval BL over time, treatment, age and the interaction between group and age were included as fixed effects in the model.

Larval DM (F1 larvae)

Larval DM was only measured in F1 larvae, therefore temperature condition, PCO_2 concentration, age and their interactions were included as fixed effects in the model for log-transformed larval DM over time. Larval DM at mouth opening and metamorphosis was analyzed with temperature condition, PCO_2 concentration and their interactions as fixed effects.

Juvenile BL and WM over time (F1 juveniles)

As F0 and F1 juveniles had different temperature life histories as well as rearing conditions, their growth rates over time were not directly compared. Due to an imbalanced dataset in F1 juveniles, it was not possible to test the effect of temperature, PCO_2 condition and their interaction separately. Instead, as for larval BL, treatment was used as fixed variable, resulting in three groups: F1-C-A, F1-C-Δ1000 and F1-W-Δ1000. Treatment, age and the interaction between treatment and age were included as fixed effects in the models for juvenile BL and log-transformed juvenile WM over time.

Juvenile BL and WM at 3000dd (F0 and F1 juveniles)

Juvenile BL and WM were compared at 3000 dd across generations. Due to the imbalanced dataset, treatment was again used as fixed effect. For juvenile BL and WM, treatment included the following five groups: F0-A, F0-Δ1000, F1-C-A, F1-C-Δ1000 and F1-W-Δ1000.

4.4.2. Respirometry

Larval RMR (F1 larvae)

As larvae were reared in a full factorial design, temperature condition, PCO_2 concentrations and their interactions were included as fixed effects in the model.

Juvenile SMR and PO_{2crit} (F0 and F1 juveniles)

Due to an imbalanced dataset for juvenile respirometry, it was not possible to test the effect of temperature, PCO_2 condition, generation, age and their interaction separately, instead treatment was used as fixed variable, resulting in seven groups for SMR: F0-A-young, F0-Δ1000-young, F0-A-old, F0-

533 Δ1000-old, F1-C-A, F0-C-Δ1000 and F1-W-Δ1000 and five groups for PO_{2crit}: F0-A-old, F0-Δ1000-old, F1-
534 C-A, F0-C-Δ1000 and F1-W-Δ1000.

535 **5. Results**

536 **5.1. Growth**

537 Neither temperature nor PCO_2 treatment had a significant effect on larval size at mouth opening stage
538 in F1 larvae (Figure 2A and D, Table 4). During the following larval development, higher temperatures
539 significantly increased growth if larvae were compared at the same age (dph): F1-C larvae were smaller
540 than F0 and F1-W larvae at higher temperature (Figure 3A and B, Table 4). SGR ranged from 7.85 to 9.75
541 % day^{-1} for larval DM and 11.67 to 14.76 % day^{-1} for larval BL (Table 3). The higher growth rates in F1-W
542 larvae resulted in Q_{10} of 1.67-2.12 and 1.81-2.35 for DM and BL (Table 3). PCO_2 had no effect on growth
543 of F0 and F1-C larvae, but reduced growth significantly in F1-W larvae (Table 4). Due to the longer larval
544 duration in colder temperatures (900 dd equals 45 dph at 20°C and 60 dph at 15°C), F1-C larvae were of
545 comparable size to F1-W-A and F0 larvae at metamorphosis. In contrast, F1-W-Δ1000 larvae were
546 significantly smaller at metamorphosis than any other group of larvae (Figure 2B and E, Table 4).
547 In juveniles, the overall positive effect of temperature on growth persisted, with F1-W juveniles
548 displaying significantly higher growth rates than F1-C juveniles (Figure 3C and D, Table 4). SGR ranged
549 from 2.88-5.16 % day^{-1} for juvenile WM and 0.84-1.55 % day^{-1} for juvenile BL, the higher growth rates
550 resulted in Q_{10} of 2.41-2.72 and 2.31-2.52 for WM and BL, respectively, in F1-Δ1000 juveniles. If
551 compared at the age of 3000 dd (165, 140 and 181 dph for F0, F1-W and F1-C juveniles, respectively),
552 the difference in size was inverted compared to metamorphosis, F1-W-Δ1000 juveniles were now
553 significantly larger than any other group (Figure 2C and F, Table 4). PCO_2 did not have any significant
554 effect on growth of F0 or F1-C juveniles. The effect of PCO_2on F1-W juveniles was not determined due
555 to the missing F1-W-A treatment.

556 **5.2. Metabolic rates**

557 Metabolic rate estimations were done on larvae with mean size ranging from approx. 1.5 to 3.0 mg DM
558 and 11.5 to 14 mm BL with no significant differences in size between treatments (BL and DM, Table S 3).
559 For juveniles, mean size ranged from approx. 3 to 62 g WM and 9 to 20 cm BL (Table S 4), with no
560 significant differences in size (BL and WM) or condition factor between acidification treatments of the
561 same age and generation (ANOVA, $P>0.05$ for F0-old; LME, $P>0.05$ for F0-young and F1-C) nor between
562 F1-C and F1-W (LME, $P>0.05$). The positive effect of temperature on growth was mirrored in larval RMR
563 in F1: RMR was significantly lower in F1-C compared to F1-W. But in contrast to growth no effect of PCO_2
564 treatment or an interaction of temperature and PCO_2 treatment on larval RMR was observed (Figure 4A,
565 Table 4). A Q_{10} of 2.24 and 2.51 was calculated for larval RMR for F1-A and F1-Δ1000 larvae, respectively.

Similarly, juvenile SMR was significantly lower in F1-C compared to F1-W juveniles (Figure 4B, Table 4), with Q_{10} of 1.61 for F1-Δ1000 juveniles. The comparison between the two generations showed that the SMR in the F0 juveniles did not change significantly between 5- and 15-months old juveniles, but F0-SMR estimates were significantly lower than those in F1 juveniles (Table 4). Comparable to larval RMR, there was no significant effect of PCO_2 in juvenile SMR at each thermal treatment. Although the LME model states a significant effect of treatment on the critical oxygen concentration PO_{2crit} (Figure 4C, Table 4), posthoc tests revealed only a significant difference between F0-Δ1000 and F1-C-A (P<0.04), all other groups were not significantly different from each other.

6. Discussion

Long-term experiments exploring the potential of fish to adapt to OAW are still scarce, especially in larger, temperate species with long generation times. In this long-term experiment, we observed that OW as single driver increased growth rates and RMR in the warm F1 larval sea bass, but due to the decreased larval phase duration at warmer temperatures, F1-C-A and F1-W-A larvae had similar size at metamorphosis. OA as single driver had no effects on F1 larval and juvenile growth nor on metabolism at ambient (cold) temperature. Under OAW, F1-W-Δ1000 larvae were significantly smaller at metamorphosis than any other group, while maintaining similar RMR as F1-W-A larvae. As they grew into juveniles, F1-W-Δ1000 fish were bigger than F1-C fish at 3000 dd and had the highest SMR. Unfortunately, the F1-W-A group could not be kept until juvenile phase. Although F0 and F1-W larvae were both raised at increased temperatures, we observed that the detrimental effects of OAW occurred only in F1-W-Δ1000 and not in F0-Δ1000. We also observed that juvenile SMR was lower in F0 than in F1-C and F1-W, with no effect of OA in F0 and F1-C. Juvenile PO_{2crit} was not affected by OA or OAW in both generations.

6.1. Effects of OW on European sea bass growth and metabolism

F1-C larvae were reared at 15°C, reflecting ambient temperature towards middle to end of the spawning season in the Bay of Brest. We applied a warming scenario of + 5°C on F1-W larvae, which reflects typical rearing temperatures in aquaculture, as well as natural temperatures towards middle to end of the spawning season in the Mediterranean (Ayala, et al., 2003). This thermal treatment (20°C) is well below the upper thermal limits for seabass larvae from the Bay of Brest (27°C, Moyano, et al., 2017). OW as a single driver at ambient PCO_2 significantly increased growth rates and decreased the time to reach metamorphosis in F1-W-A larvae in comparison to F1-C-A larvae. Due to the longer larval phase duration, size at metamorphosis was comparable between F1-C-A larvae and F1-W-A larvae. Faster growth at higher temperatures and similar size at metamorphosis despite different temperatures has been shown in other studies for sea bass from Mediterranean and Atlantic populations (Ayala, et al., 2001; 2003).

OW also increased RMR in F1-W-A larvae compared to F1-C-A larvae. The increase in RMR was similar to the increase in SGR, reflected by similar Q_{10} (1.96, 2.22 and 2.24 for SGR of dry mass and body length (DM and BL, 0.5 Quantile) and RMR, respectively). This reflects the expected Q_{10} increase of 2-3 for biological processes and confirmed our hypothesis that OW will lead to increased growth and RMR in larval sea bass of this particular population. We did not determine the effects of OW as a single driver on growth and metabolism in F1 juveniles, due to the absence of F1-W-A.

6.2. Effects of OA on European sea bass growth and metabolism

OA as single driver within the cold temperature condition did neither affect growth and metabolism (RMR; SMR), nor PO_{2crit} in F1 European sea bass larvae or juveniles. In the wild, sea bass eggs are spawned in stable open ocean conditions and larvae develop during the drifting towards the coast, therefore larvae were thought to be less resilient to OA than juveniles and adults. This has already been proven not to be the case for sea bass in scenarios up to SSP5-8.5 and similar (Pope, et al., 2014; F0 in Crespel, et al., 2017) and was further confirmed by this study, as larval growth and RMR were not affected by OA within the cold temperature group. As juvenile sea bass inhabit coastal areas and have been shown to be tolerant to a broad range of environmental factors, including temperature and salinity (Dalla Via, et al., 1998; Claireaux & Lagardère, 1999), their tolerance to OA was expected and could be confirmed in this study – no effects of OA within the cold temperature group on growth, SMR and PO_{2crit} were observed. Our study also supports the hypothesis of Montgomery et al. (2019) that an observed 20% decrease in PO_{2crit} under acute increase of PCO_2 (3 to 5 fold increase in PCO_2 within approx. 6 hours) in European sea bass will vanish after long-term acclimation to OA.

6.3. Combined effects of OA and OW on European sea bass growth and metabolism

However, the combined effects of OA and OW (OAW) changed the picture for larval resilience. While growth rates increased sufficiently in F1-W-A to reach the same size at metamorphosis than F1-C-A, F1-W-Δ1000 larvae were significantly smaller at metamorphosis than larvae from any other treatment, but maintained RMR as high as F1-W-A larvae. Q_{10} values revealed that temperature had a stronger effect on metabolic rate than on growth under OA: 1.67 and 1.95 for SGR of DM and BL (0.5 Quantile) and 2.51 for RMR, respectively. This suggests that F1-W-Δ1000 larvae either allocated the energy differently, such as using more energy for movement or different regulatory processes, or that their energy production and oxygen usage was not as efficient as in the other groups. Although it is possible that the higher RMRs are due to higher activity of the F1-W-Δ1000 larvae during the measurements, larvae were regularly observed during the trials and the inter-individual variability in movement did not seem related to treatment. Therefore it seems more plausible that larvae under OAW needed energy for different regulatory processes, probably combined with decreased energy production efficiency. In this sense, we

already found that OAW decreased the efficiency of complex II (CII) of the electron transport system (ETS) in cardiac mitochondria of juvenile sea bass in the W-Δ1000 treatment under acute temperature change (Howald, et al., 2019). Inhibition of CII by OA was also found in other studies on mammals and fish (Simpson, 1967; Wanders, et al., 1983; Strobel, et al., 2013). In Atlantic cod embryos, reduced activity of complex I (CI) of the ETS resulted in reduced mitochondrial phosphorylation capacity and subsequently in reduced oxygen consumption rates, while energy requirements were simultaneously increased (Dahlke, et al., 2017). Although CII was only affected in juvenile sea bass under acute temperature change, it is probable that larvae are more vulnerable than juveniles (Dahlke, et al., 2020a): similar to embryos (Leo, et al., 2018), they are less developed while at the same time investing all available energy into growth without reserving excess capacity for environmental regulation and are therefore already affected at their acclimation temperature if OA and OW are combined. This inability to cope with OAW has not been observed in European sea bass larvae before, contrastingly in former studies growth of larval European sea bass has been shown to be resilient to OA even at rearing temperature of 19°C (Pope, et al., 2014; F0 larvae in Crespel, et al., 2017). Potential explanations why these differences first occurred in F1 are likely related to their parents being reared under OA conditions, as well as effects due to different rearing protocols, which are both addressed below (section 6.4).

In contrast to larvae, F1-W-Δ1000 juveniles displayed a greater thermal plasticity and grew significantly faster than F1-C juveniles, resulting in larger fish at 3000 dd in the F1-W-Δ1000 than in F1-C-A and F1-C-Δ1000. High growth rates were supported by high SMR, which were also highest in F1-W-Δ1000 juveniles in comparison to F1-C-A and F1-C-Δ1000. As we did not incubate the F1-W-A treatment to juvenile phase, it is unclear whether the detrimental effects of OAW on growth and metabolism in larval European sea bass would have persisted into the juvenile phase. The increased growth rates and bigger size at 3000 dd in F1-W-Δ1000 juveniles in comparison to F1-C-A and F1-C-Δ1000 juveniles might either indicate that OA did not affect growth in juveniles or that growth under OW was so much accelerated in juveniles that F1-W-Δ1000 fish were able to catch up and grow to bigger sizes than F1-C fish masking the negative effects of OAW. The latter suggestion is supported by the findings in SMR and by the Q_{10} of SMR and SGR: in F1-Δ1000 juveniles, SMR was less affected by temperature (Q_{10} 1.61) than SGR (Q_{10} 2.63 and 2.45 for SGR of WM and BL (0.5 Quantile)). Q_{10} for SGR and SMR are well in the range found in other studies on European sea bass from the Atlantic (Q_{10} for SGR of WM ~ 2.4 (15-20°C, calculated from Gourtay, et al., 2018) and Q_{10} for SMR 2.09 (14-22°C, Montgomery, et al., 2021 preprint)) and from the Western Mediterranean populations (Q_{10} for SGR of WM and RMR of 2.40 and 1.70, respectively, 13-25°C, calculated from Person-Le Ruyet, et al., 2004). The authors of the latter study explained the different Q_{10} of RMR and SGR with increased growth rates due to increased feed intake. As the fish in our study were fed *ad libitum*, they were able to increase food intake to support high growth rates, too. The better capacity of juveniles to cope with and even profit from higher temperatures even under OAW

in comparison to larvae is probably due to the reproduction biology of European sea bass, as well as to the generally higher capacity for acid-base regulation in juveniles in comparison to larvae. Larvae are developing during spring in the open ocean resulting in stable and relatively cold temperatures (8-13°C for Atlantic specimen, Jennings & Pawson, 1992), with optimal larval growth temperatures of 15-17°C (Mediterranean specimens, Koumoundouros, et al., 2001; Ayala, et al., 2003). Juveniles on the other hand live in shallow coastal areas, resulting in higher temperatures during summer but also higher daily and seasonal variation (6-18°C for Atlantic specimens, Russel, et al., 1996) with optimal growth temperatures of 22-28°C (Mediterranean specimens, Lanari, et al., 2002; Person-Le Ruyet, et al., 2004). Consequently, in terms of growth and metabolism juvenile sea bass at the northern distribution range might benefit from higher temperatures, as already found in other studies (Howald, et al., 2019; Montgomery, et al., 2021 preprint) and do not seem to be severely affected by OA.

6.4. Effects of OA on European sea bass growth and metabolism over two successive generations

In addition to the effects of the single and combined stressors OA, OW and OAW on individual groups of fish, we also studied the effects of OA in two successive generations on the ability of sea bass larvae and juveniles to cope with upcoming conditions. This study is to our knowledge the first one to examine the effects of OA on European sea bass or other long-lived teleost in more than one generation. Interestingly the detrimental effect of OAW on larval growth was only observed in F1 and not in F0 larvae of European sea bass, despite their respective parental generation's identical thermal history, and thus appears to be an OA effect. This can be explained by several reasons: First, the provisioning of necessary resources when parents have already encountered the same conditions as the future offspring, e.g. via egg size and composition (Munday, 2014) could explain the observed trend in F1-W-Δ1000 larvae. Parental effects can lead in different directions and can last throughout larval phase: parental effects influence growth in stickleback under OW and OA (Shama, et al., 2014; Schade, et al., 2014) and explained differences in embryo mortality and hatching success in Atlantic cod under OW (Dahlke, et al., 2017). In our study, we did not measure egg size and quality nor did we incubate offspring of F0-A in cross factorial Δ1000 scenarios, so we cannot directly quantify parental or transgenerational effects. However, the size of F1 larvae at mouth opening, a landmark until which the larvae depend on yolk sac reserves, did not differ across treatments. Thus, using this landmark as indirect indicator, parental provisioning does not seem to explain differences in larval growth rates. Second, the incubation protocol differed between F0 and F1. While F0 larvae were first incubated under OA conditions at 2 dph, F1 sea bass were constantly reared under OA conditions from fertilization onwards, although warming was also applied from 2 dph onwards. It is possible that effects of OA during embryogenesis shaped the reaction of F1 larvae to OAW, e.g. via epigenetic signaling. As reviewed by Dahlke et al. (2020a), it seems that spawning adults and embryos are the most vulnerable life stages in fish, possessing the lowest tolerance

to OW, e.g. Atlantic cod embryos exposed to OAW showed reduced hatching success and oxygen consumption rates (Dahlke, et al., 2016) and OA decreased the Q_{10} of RMR in Atlantic silverside embryos (Schwemmer, et al., 2020). To summarize, the different reaction of F0 and F1 larvae to OAW could be due to parental effects or effects during embryogenesis and more research is necessary to determine the underlying mechanisms.

As the different temperature life histories and replication schemes (no replicate tanks in F0 juveniles) did not allow a direct comparison of growth rates between F0 and F1 juveniles, we compared size at the age of approx. 3000 dd (165, 140 and 181 dph for F0, F1-W and F1-C juveniles). Due to their high growth rates during juvenile phase, F1-W-Δ1000 fish were largest at 3000 dd, while F0 and F1-C fish were smaller (WM and BL) but similar to each other. This matched the result of SMR, which was not affected by OA and was higher in F1-W compared to F1-C fish. Surprisingly, SMR was also higher in F1-C compared to F0 fish. This difference might be explained by the different temperature life histories. While F0 fish had been raised at warmer temperatures and were acclimated to colder temperatures afterwards, F1-C fish had been reared at 15°C throughout their life, except for summer months, when temperatures reached up to 19°C. No detrimental effects on juvenile growth rate under OA were visible in the second generation of sea bass reared under OA conditions, reflected by similar SMR and SGR between A and Δ1000 condition. Due to the missing F1-W-A treatment, we cannot state if the detrimental effects of OAW observed in F1 larvae persisted to juvenile phase.

6.5. Ecological perspective

Larvae are not fully developed compared to later stages and are exposed to higher predation and starvation risks, as such they had been thought to be more vulnerable to environmental stressors such as OAW (as reviewed in Houde, 2009). In this context, OAW could impact larval survival and recruitment success via different mechanisms. If OAW leads to faster growth rates and increased metabolic rates (as seen in this study between F1-W and F1-C larvae), larvae need more food in shorter time to support these growth rates, therefore it is essential that they match adequate prey fields (prey abundance, size and quality). In our study the larvae were fed *ad libitum* at both temperatures, supporting increased energetic demands for the high growth and metabolic rates at higher temperatures. However, in the ocean it is possible that food availability is not sufficient to support accelerated growth under OW. Bochdansky et al. (2005) showed that fish larvae with higher growth and metabolic rates died earlier, when food was limited, but profited when fed at saturation level. In sea bass larvae, high growth rates were also only supported under high food ratios, but survival was not significantly decreased, even at one eighths of saturation ratio (Zambonino Infante, et al., 1996). This might indicate that sea bass will not grow as fast as in our study under future OW scenarios if food is scarce, but might still survive to juvenile stage.

Besides food-related aspects, OAW can also have a large impact on larval behavior and dispersal, which can later influence recruitment success. Sea bass spawn in the open ocean and larvae are drifted inshore (Jennings & Pawson, 1992). As with many temperate species, their swimming behavior and its effect on dispersal has not been studied as extensively as for coral reef fish that have well developed sensory abilities (hearing, olfaction, vision) and show directional swimming early on (as reviewed in Leis, 2018; Berenshtein, et al., 2021 preprint). To the best knowledge, it seems that early seabass larvae are more dependent on currents than on their swimming performance and that they are able choose a certain depth and therefore a certain current in the preferred direction (Jennings & Pawson, 1992). When being drifted closer to the coast, sea bass larvae wait for certain cues from nursery areas, which are present from June onwards (Jennings & Pawson, 1992).

OW accelerates development of sea bass larvae and therefore possibly alters the timing and spacing of dispersal. Studies have shown species-specific responses of fish behavior to OA, OW and OAW, e.g. OW increased activity level in larval kingfish but not boldness, while OA had no effect on these behavioral traits (Laubenstein, et al., 2019). Yet, OA decreased swimming duration and orientation in larval dolphinfish (Pimentel, et al., 2014) and reversed orientation towards settlement habitat cues in barramundi (Rossi, et al., 2015). To our knowledge, larval sea bass behavior has not been measured under OAW yet. Consequently due to the altered timing of larval development and in combination with the possibility of altered behavior and impacted senses, reaching nursery areas might be challenging for sea bass larvae under OAW, especially if (1) food is not abundant and (2) cues are weaker and/or different due to greater distance and/or earlier timing. Once the larvae entered the coastal areas and metamorphose, they are exposed to a more changing environment. Although this study could confirm that juvenile sea bass are less vulnerable to OAW than larval sea bass, food availability and behavior will determine, if the observed increased growth under OAW in F1 will occur in the wild, too. In a sister study on offspring of wild caught European sea bass, OAW reduced digestive enzyme activity under restricted food ratios resulting in severely reduced food conversion efficiency and reduced growth rates (Cominassi, et al., 2020). Additionally, OA decreased the distance which early juvenile sea bass needed to sense food or predator cues (Porteus, et al., 2018) and juvenile sea bass behavior was altered by OW resulting in decreased latency of escape response and mirror responsiveness (Manciocco, et al., 2015). Consequently, although faster larval (OW) and juvenile growth (OAW) as well as earlier metamorphosis (OW, OAW) is generally beneficial for larvae and early juveniles, many factors may modulate this effect and whether it will translate into higher larval survival, recruitment and increased growth rates in the wild. Further research should determine the effects of limited food under OAW on larval and juvenile growth and behaviour.

As the hypoxia tolerance of European sea bass juveniles was unaffected by OA, OW and OAW, they might cope well with upcoming hypoxia events in coastal areas. However, it is important to note here

that we measured $P_{O_{2crit}}$ only at SMR and thus may have estimated P_{O_2} effects too conservatively. Recent studies suggest that this $P_{O_{2crit}}$ at SMR might not be the most ecologically relevant estimate (see Seibel & Deutsch, 2020 and references therein): Long-term survival of individuals and their population would require that the fish are able to digest food, grow and reproduce, which would require more energy than provided by SMR. Consequently, depending on the duration and intensity of hypoxia events, individuals might be able to survive over short terms, but other fitness related traits such as growth might be affected in the long-term.

7. Conclusion

We confirmed our hypotheses that OW increases growth and metabolism in the European sea bass, and that larvae as well as juveniles are resilient to OA if it occurs as a single stressor. Yet, we could also confirm that OAW had detrimental effects on larval growth. Our results together with other findings on larval fish and European sea bass suggest that it is possible that under OAW fewer individuals will reach metamorphosis, e.g. due to limited food to support high growth rates, different dispersal to nursery areas by altered developmental timing, changed behavior or affected olfactory senses. However, those individuals that reach the juvenile phase might benefit from higher temperatures, due to increased performance.

8. Acknowledgements

We thank Mark Suquet for his support during the reproduction period. We acknowledge the technicians and researchers from the Laboratory of Adaptation, and Nutrition of Fish for their support in animal welfare during the weekends. We want to specially thank Patrick Quazuguel and Nicolas LeBayon for their help in rearing the fish from larvae to reproduction and beyond. Finally, we would like to thank two anonymous reviewers for their detailed and constructive comments on the original manuscript.

9. Competing interests

The authors declare no competing or financial interests.

10. Author contributions

Conceptualization: SH, AC, MM, GC, MP, FCM

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Formal analysis: SH, AC, MM, FCM

Funding acquisition: MP, GC, FCM

Investigation: SH, AC, LC, LLK

Methodology: SH, MM, GC, FCM

Project administration: MM, GC, MP, FCM

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Supervision: MM, GC, MP, FCM

Validation: SH, MM, FCM

Visualization: SH

Writing -original draft: SH

Writing - review & editing: SH, MM, AC, LC, GC, MP, FCM

11. Funding

This work was supported by the German Science Foundation (Deutsche Forschungsgemeinschaft, DFG) and part of the FITNESS project (DFG grants MA 4271/3-1 to F.C.M. and PE 1157/8-1 to M. Peck, University of Hamburg, Germany).

12. Data availability

Datasets of growth, metabolic rates and water conditions during rearing are available online from PANGAEA (www.pangaea.de)

13. References

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14. Figures and Tables

Table 1 Water parameters during larval and early juvenile phase of F0: Larval period until (45 dph, ~900 dd), early juveniles until 1.5 years. Means ± s.e.m. over all measurements per condition (triplicate tanks in larvae, single tanks in juveniles). Temperature (Temp.) and pH (NBS scale) were measured daily. pH (total scale), salinity, phosphate, silicate and total alkalinity (TA) were measured once at the beginning and once at the end of the larval phase and 9 times during juvenile phase; $pCO_2$was calculated with CO2sys; A–Ambient pCO_2, Δ1000 –ambient + 1000 µatm CO_2,L – Larvae, J – Juveniles, (see Crespel, et al., 2017; Crespel, et al., 2019).

Treatment	pH_{NBS} [-]	pH_{total} [-]	Temp. [°C]	Salinity [psu]	TA [µmol L^{-1}]	pCO_2 [µatm]	PO_4^{3-} [µmol L^{-1}]	SiO_4 [µmol L^{-1}]
L A	7.96±0.01	7.89±0.01	19.2±0.3	33.8±0.2	2294±3	589±10	0.57±0.01	8.94±0.06
L Δ1000	7.59±0.00	7.54±0.03	19.2±0.3	33.8±0.2	2306±9	1521±97	0.57±0.01	8.94±0.06
J A	8.05±0.01	7.94±0.03	15.3±0.1	34.3±0.2	2294±10	516±31	0.71±0.08	8.35±0.26
J Δ1000	7.61±0.01	7.53±0.02	15.3±0.1	34.3±0.2	2280±16	1489±42	0.71±0.08	8.35±0.26

Table 2 Water parameters in the 2 years before spawning of F0 (2016-2018) and during larval and juvenile phase of F1: Larval period until 17.05.2018 (48 dph, ~900 dd) and 01.06.2018 (63 dph, ~900 dd) for warm and cold life condition respectively, for the juveniles until 28.09.2018 (180 dph, ~4000 dd) and 12.02.2019 (319 dph, ~5100 dd) for warm and cold conditioned fish respectively. Means ± s.e. over all replicate tanks per condition. Temperature (Temp.), pH (free scale), salinity, oxygen and total alkalinity (TA) were measured weekly in F1 and monthly in F0; pCO_2 was calculated with CO2sys; sea water (SW) measurements were conducted in 2017 and 2018; A – Ambient pCO_2, Δ1000 – ambient + 1000 µatm CO_2, L – Larvae, J – Juveniles, C – cold life condition, W – warm life condition.

Treatment	pH_{free} [-]	Temp. [°C]	Salinity [psu]	O_2 [% airsat.]	TA [-]	pCO_2 [µatm]
F0 A	7.95±0.02	14.1±0.6	33.6±0.3	92.4±1.7	2406±49	670±40
F0 Δ1000	7.59±0.02	14.1±0.6	33.6±0.3	92.4±1.9	2411±46	1616±74
F1 L C A	8.06±0.01	15.3±0.1	31.8±0.1	94.3±1.0	2360±23	504±19
F1 L C Δ1000	7.53±0.01	15.5±0.1	31.8±0.1	94.3±0.8	2330±22	1872±74
F1 L W A	7.96±0.01	20.2±0.2	31.7±0.0	84.9±3.4	2311±32	656±22
F1 L W Δ1000	7.61±0.01	20.2±0.2	31.8±0.0	88.1±1.7	2321±32	1624±59
F1 J C A	7.94±0.01	16.1±0.2	33.0±0.1	92.4±0.5	2376±15	696±19
F1 J C Δ1000	7.60±0.01	16.3±0.2	33.0±0.1	94.3±0.5	2380±14	1603±32
F1 J W A	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.
F1 J W Δ1000	7.57±0.02	22.7±0.2	33.0±0.2	86.3±1.3	2323±16	1866±83
SW	8.07±0.01	15.0±0.5	34.6±0.3	101.0±0.8	2441±23	609±37

Table 3 Specific growth rates (SGR) and their respective Q_{10} of larval and juvenile mass and body length of European sea bass. SGR [% day^{-1}]and Q_{10} [-] are given for 0.05, 0.5 and 0.95 quantile of the cohort. Means ± s.e. over all replicate tanks per condition. A – Ambient PCO_2, Δ1000 – ambient + 1000 µatm CO_2, L – Larvae, J – Juveniles, C – cold life condition, W – warm life condition, n.a. – treatment was not available or not measured at this state

Treatment	n	0.05 Quantile	0.5 Quantile	0.95 Quantile
Larval dry mass				
SGR F1 C A	3	9.19±0.37	9.57±0.23	9.63±0.25
SGR F1 C Δ1000	1	7.85	9.26	9.75
SGR F1 W A	2	12.92±0.38	14.25±0.11	14.76±0.08
SGR F1 W Δ1000	3	11.67±0.29	12.92±0.26	13.73±0.20
Q_{10} A		1.84	1.96	2.12
Q_{10} Δ1000		1.81	1.67	1.80
Larval body length				
SGR F1 C A	3	2.27±0.08	2.41±0.04	2.40±0.06
SGR F1 C Δ1000	3	2.14±0.07	2.33±0.02	2.43±0.03
SGR F1 W A	2	3.09±0.15	3.37±0.06	3.50±0.02
SGR F1 W Δ1000	3	2.88±0.01	3.01±0.06	3.26±0.04
Q_{10} A		1.98	2.22	2.35
Q_{10} Δ1000		1.81	1.95	1.98
Juvenile wet mass				
SGR F1 C A	2	3.07±0.09	2.94±0.03	3.04±0.04
SGR F1 C Δ1000	2	2.96±0.05	2.88±0.01	2.92±0.04
SGR F1 W A		n.a.	n.a.	n.a.
SGR F1 W Δ1000	2	5.16±0.02	4.93±0.11	4.82±0.13
Q_{10} A		n.a.	n.a.	n.a.
Q_{10} Δ1000		2.72	2.63	2.41
Juvenile body length				
SGR F1 C A	2	0.91±0.02	0.87±0.00	0.87±0.00
SGR F1 C Δ1000	2	0.85±0.02	0.84±0.00	0.86±0.01
SGR F1 W A		n.a.	n.a.	n.a.
SGR F1 W Δ1000	2	1.55±0.01	1.49±0.02	1.46±0.06
Q_{10} A		n.a.	n.a.	n.a.
Q_{10} Δ1000		2.52	2.45	2.31

Table 4 F- and p-values of fixed effects from the linear mixed models on growth and metabolic rates of F0 and F1 larval and juvenile European sea bass. n.a. – treatment was not available or not measured at this state

	OAW Treatment		pCO_2 Treatment		Temperature		pCO_2:Temperature	
	F-value	**p-Value**	**F-value**	**p-Value**	**F-value**	**p-Value**	**F-value**	**p-Value**
Larval growth								
Dry mass								
at mouth opening	n.a.	n.a.	1.18	0.3	4.49	0.06	2.13	0.18
at metamorphosis	n.a.	n.a.	11.69	0.01	6.37	0.05	2.73	0.16
over time	n.a.	n.a.	17.27	0.0032	2.61	0.1447	8.01	0.0221
Body length								
at mouth opening	n.a.	n.a.	0.23	0.66	0.21	0.64	1.72	0.23
at metamorphosis	10.04	0.0008	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.
over time	275.09	<.0001	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.
Juvenile Growth								
Wet mass								
at 3000 dd	16.41	0.0222	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.
over time	240.515	0.0005	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.
Body length								
at 3000 dd	46.93	0.0049	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.
over time	1111.59	<.0001	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.
Metabolic rates								
RMR	n.a.	n.a.	0.01	0.94	29.62	<.0001	0.06	0.82
SMR	95.44	<.0001	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.
PO2crit	3.79	0.0064	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.

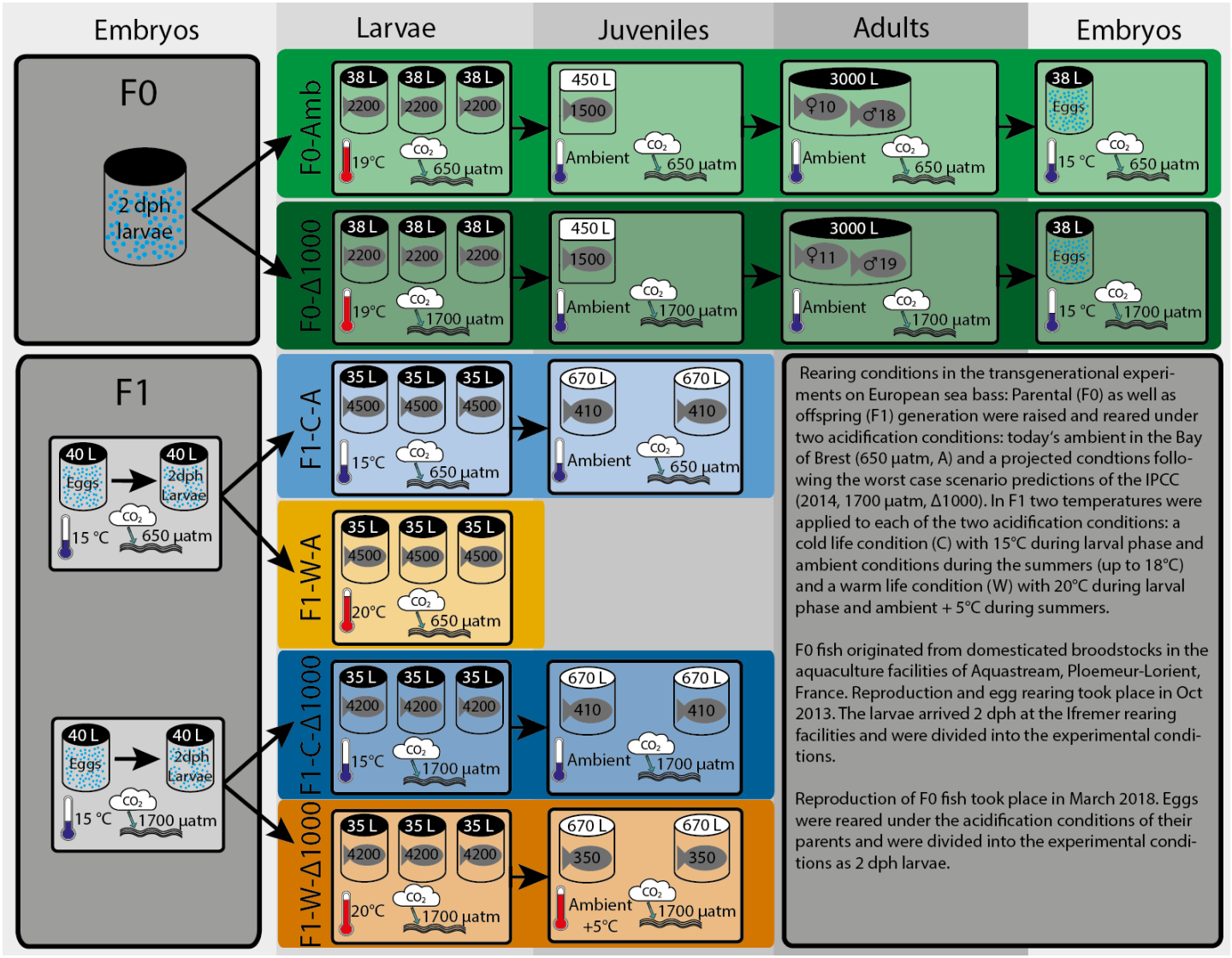

Figure 1 Schematic overview summarizing the rearing conditions of two generations of sea bass under different OAW scenarios.

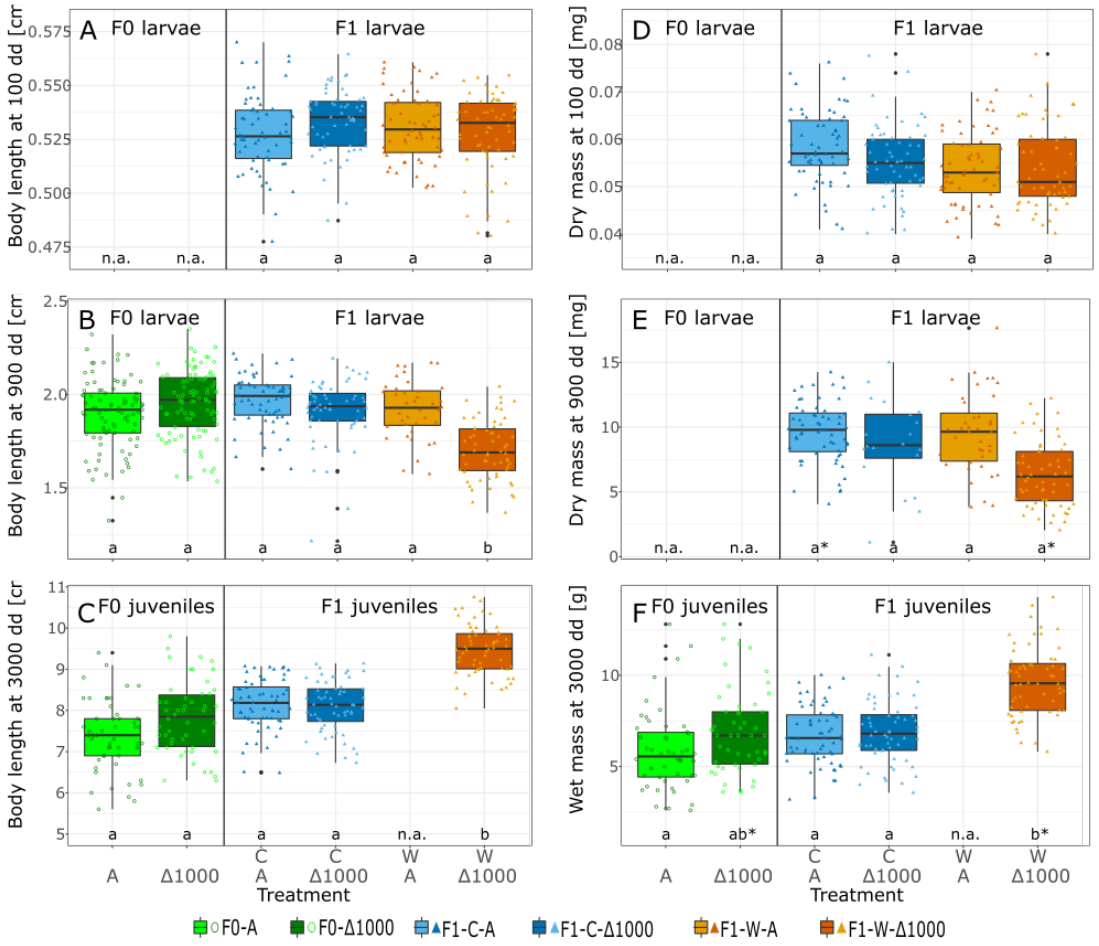

Figure 2 Body length and mass of European sea bass at approx. 100 dd (mouth opening, A and D, 7 dph), 900 dd (metamorphosis, B and E) and 3000 dd (C and F) in F0 and F1 fish. Overlying dots are the individual data points of each treatment, different letters indicate significant differences [linear mixed effects (LME), $P<0.05$], asterisks indicate statistical trends [LME, $P<0.1$], A – Ambient pCO_2, Δ1000 – ambient + 1000 µatm CO_2, C – cold life condition, W – warm life condition, n.a. – treatment was not available or not measured at this state, n=40-90.

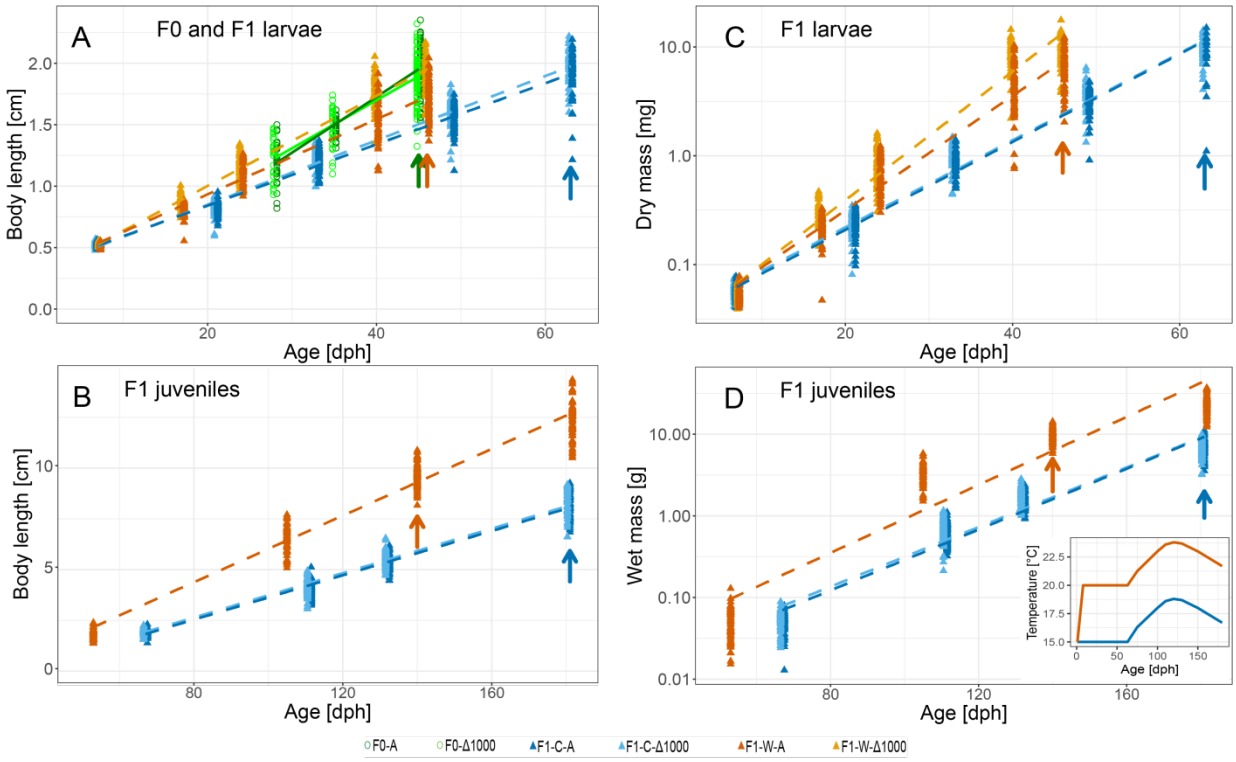

Figure 3 Growth of F0 and F1 larvae (A, C) and F1 juveniles (B, D) of European sea bass with linear regression lines. Shown are individual data points of body length (A, B), larval dry mass (C) and juvenile wet mass (D). F1-C larvae grew significantly slower than F1-W (A, C) and F0 larvae (A). F1-W-Δ1000 larvae grew significantly slower than F1-W-A (A, C) and F0-A larvae (A). F1-W-Δ1000 juveniles grew significantly faster than F1-C juveniles (B, D). No differences were observed between PCO_2 treatments in F0 larvae (A), F1-C larvae (A, C) and F1-C juveniles (B, D), respectively. All data were tested with LME models, F- and p-Values are summarized in Table 4. Arrows indicate the data points at metamorphosis (900dd, A, C) and at 3000 dd (B, D), data of different PCO_2 conditions of the same age are slightly moved for better visibility. Insert in D shows the temperature history of F1 larvae and juveniles. A – Ambient PCO_2, Δ1000 – ambient + 1000 µatm CO_2, C – cold life condition, W – warm life condition.

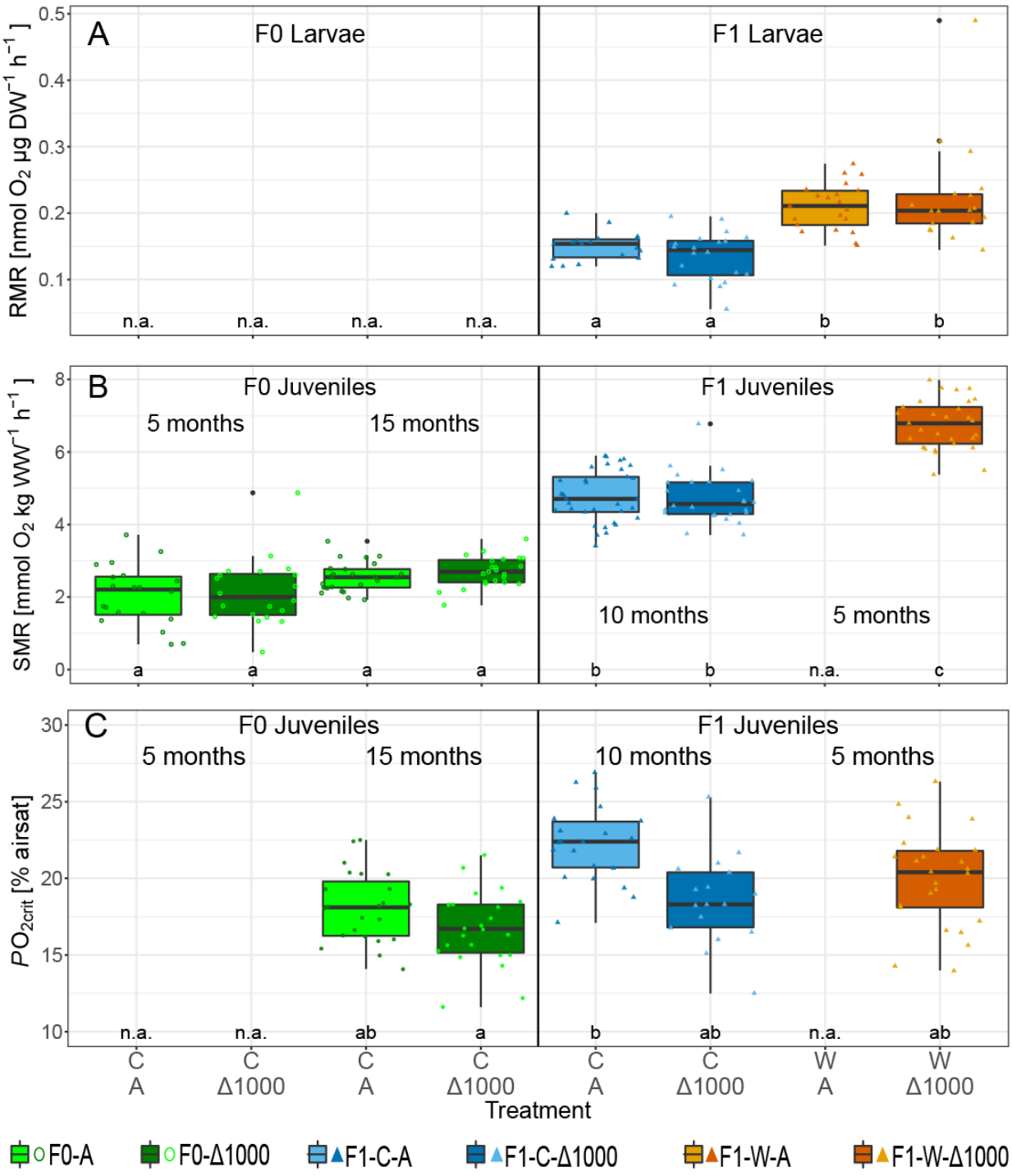

Figure 4 Routine (RMR, A) and standard metabolic rates (SMR, B) and critical oxygen concentration (PO_{2crit}, C) of F0 and F1 larvae and juveniles. Overlying dots are the individual data points of each treatment. Metabolic rates are corrected with allometric scaling factors (0.89 and 0.98 for larvae and juveniles, respectively). Different letters indicate significant differences [linear mixed effects (LME), $P<0.05$]; data of 15 months old F0 juveniles taken from Crespel et al. (2019). A – Ambient PCO_2, Δ1000 – ambient + 1000 µatm CO_2, C – cold life condition, W – warm life condition, n.a. – treatment was not available or not measured at this state, n=20-35.

15. Supplement

Table S 1 Light intensity during rearing phase of European sea bass larvae. Age is given in days post hatch (dph). Light intensity was changed at the indicated days and remained identical during the light phase until the next increase.

Age [dph]	2	8	11	20	30	32	36	46
Light intensity [lux]	0	0-1	1	7	10	31	59	96

Table S 2 Larval mortality in % in the different larvalrearing tanks (n=3). A – Ambient PCO_2 and Δ1000 – ambient + 1000 µatm CO_2, T – temperature, Rep 1-3 – replicate tank 1-3.

	A			Δ1000		
T [°C]	Rep 1	Rep 2	Rep 3	Rep 1	Rep 2	Rep 3
15	73.5	28.8	83.1	67.9	55.0	47.2
20	96.4	76.2	25.8	59.3	52.5	53.7

Table S 3 Biometrical data of larvae used for respiration measurements. Treatments: C – cold life condition (15°C), W – warm life condition (20°C), A – ambient PCO_2, Δ1000 – ambient PCO_2 + 1000 µatm, values are means ± s.e.m. Different letters indicate significant differences between groups (LME, $P<0.05$).

Treatment	n	Dry weight [mg]	Bodylength [mm]
C – A	18	2.87±0.51[a]	13.96±0.77[a]
C – Δ1000	20	2.95±0.46[a]	14.04±0.80[a]
W – A	21	2.51±0.43[a]	13.04±0.71[a]
W – Δ1000	18	1.70±0.53[a]	11.63±0.85[a]

Table S 4 Biometrical data of juveniles used for respiration measurements. Treatments: C – cold life condition (up to 18°C), W – warm life condition (up to 23°C), A – ambient PCO_2, Δ1000 – ambient PCO_2 + 1000 µatm, values are means ± s.e.m. Different letters indicate significant differences between groups (LME, P<0.05).

Generation	Treatment	Age [m]	n	Fish mass [g]	Forklength [mm]	Condition factor [-]
F0	C – A	6	20	5.06±0.24[a]	-	-
F0	C – Δ1000	6	20	5.85±0.27[a]	-	-
F0	C – A	18	24	81.80±2.60[b]	18.11±0.18[b]	1.37±0.02[b]
F0	C – Δ1000	18	24	81.40±3.22[b]	18.25±0.22[b]	1.33±0.03[b]
F1	C – A	10	33	15.00±0.69[c]	10.86±0.14[c]	1.14±0.02[c]
F1	C – Δ1000	10	26	13.05±0.56[c]	10.31±0.14[c]	1.17±0.01[c]
F1	W – Δ1000	5	29	15.73±1.01[c]	11.04±0.20[c]	1.12±0.02[c]

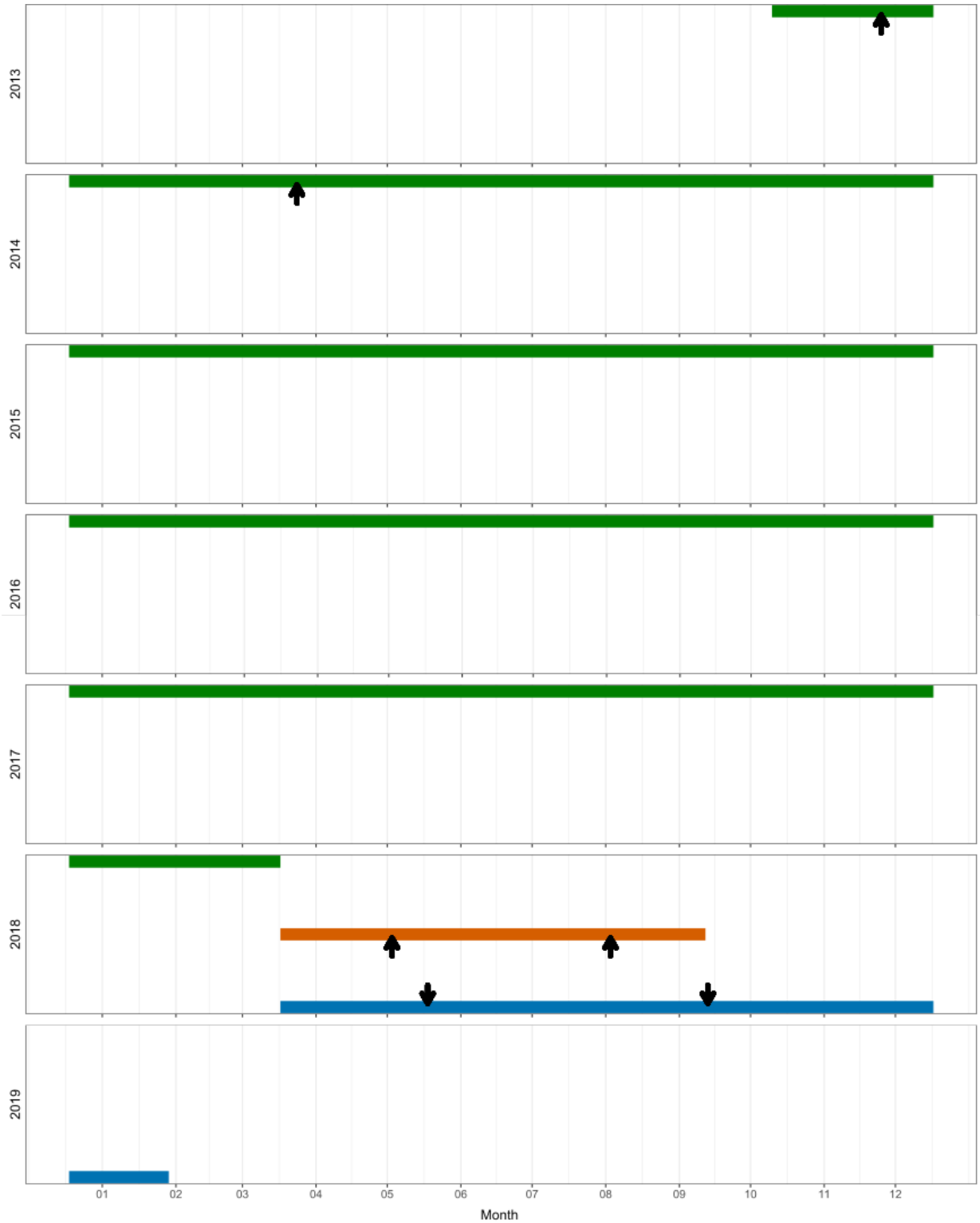

Figure S 1 Timeline of the rearing of the different treatment. Green (2013-2018): rearing of F0 fish; Orange (2018): rearing of F1-W fish; Blue (2018-2019): rearing of F1-C fish. Arrows indicate the time of metamorphosis from larvae to juveniles (first arrow per treatment) and when the fish reached the age of 3000 dd (second arrow per treatment). C- Cold life condition, W- Warm life condition.

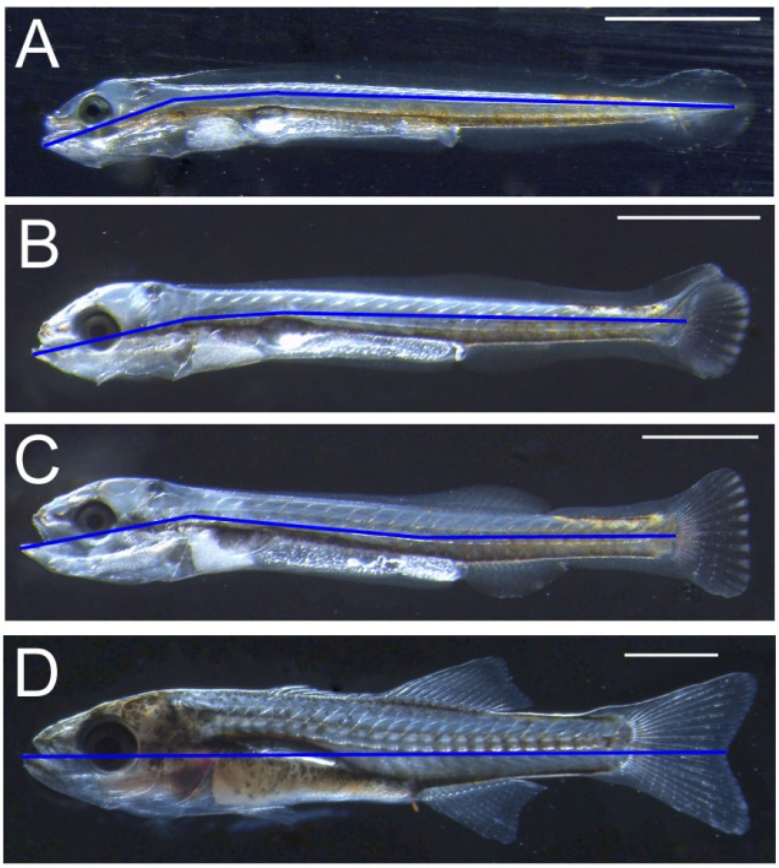

Figure S 2 Body length measurements in larvae at different developmental stages. A – pre flexion (about 300 dd), B – flexion (about 460 dd), D- post flexion (about 460 dd) and (post)metamorphosis (about 900 dd). Until post flexion the segmented line tool in the software ImageJ (Schneider, et al., 2012) was used to measure the length of the larva, afterwards the length of the larvae was measured as a straight line, as it would be done with calipers. The lines of the measurement are marked in blue.