**Attributing hypoxia responses of early life *Menidia menidia* to energetic mechanisms with Dynamic Energy Budget theory**

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**Highlights**

* The bioenergetic mechanisms responsible for whole-organism hypoxia effects on Atlantic silversides were investigated using a simplified Dynamic Energy Budget model (DEBkiss).
* We connected physiological information with energetic processes to identify assimilation, conversion efficiency of assimilates to growth, maintenance rate, and mortality as potential mechanisms.
* The most likely mechanisms of observed experimental hypoxia effects on early life stages of silversides are conversion efficiency, embryonic mortality, and larval mortality.
* We used widely applicable biological theory to gain a better understanding of hypoxia effects on a model species, creating a tool that can be used to study effects of multiple stressors and population dynamics in future work.

**Abstract**

Ocean deoxygenation is intensifying worldwide due to warming and eutrophication, particularly in estuaries and coastal waters. Although the Atlantic silverside (*Menidia menidia*) has proven robust to the fluctuating environmental conditions in its estuarine environment, chronic hypoxia impairs hatching, growth, and survival in the early life stages. We used a simplified version of a Dynamic Energy Budget model (DEBkiss) to test the hypothesis that experimentally observed changes in animal performance can be explained by one or more of the rate processes in the model. We sought to identify the DEBkiss parameters that, when adjusted with a correction factor based on inhibition or damage to the Synthesizing Unit, provided the best fit to hypoxia effects in the three state variables of total length, egg buffer mass, and survival over time. Because hypoxia reduces survival in embryos and newly hatched larvae, we added a survival state variable controlled by pre- and post-hatching mortality parameters. Applying damage production to reduce the conversion efficiency of assimilates to structure accounted for some of the hypoxia-related changes in all three state variables. However, simultaneously reducing the conversion efficiency and increasing both mortality parameters provided the best fit. In contrast, increasing the parameter for maintenance rate with hypoxia had little impact on early life growth and egg buffer depletion, and no impact on survival rates. Reduced conversion efficiency under hypoxia would suggest that less of the energy invested by parents and consumed through predation is converted into biomass in *M. menidia* offspring, with implications for size at age that could threaten recruitment and alter the flow of energy through the food web.

**Keywords**

Dynamic Energy Budget; DEBkiss; early life stages; Atlantic silverside; hypoxia; stressors

1. **Introduction**

Hypoxia is common in coastal and estuarine waters and is expected to intensify with global warming (Diaz and Rosenberg, 2008; Breitburg et al., 2018). Between anthropogenic influence on nearshore waters and the natural dynamics of shallow, partially enclosed water bodies, hypoxia often co-occurs with other stressors such as high temperature, ocean acidification, and pollutants (Gruber, 2011). Along the Northeast United States coast, stratification and productivity associated with high temperatures in spring and summer cause hypoxic and eutrophic zones to form and great fluctuations in dissolved oxygen (DO) on diel to monthly time scales (O’Donnell et al., 2004; Baumann and Smith, 2018; Testa et al., 2018). While fish species that currently live in such areas tend to have mechanisms to cope with periods of hypoxia (Farrell and Brauner, 2009; Zhu et al., 2013; Baumann, 2019), these do not necessarily confer tolerance of longer durations. Fishes that spawn in the spring and summer face the additional threat of experiencing hypoxia during the particularly sensitive early life stages. Embryos and young larvae rely largely on diffusion for oxygen uptake and lack well-developed mechanisms, such as high surface area gills, to meet oxygen demands in low DO water (Rombough, 1988). While later stage fishes and even some early larvae can swim to avoid hypoxic habitats (Niklitschek and Secor, 2005; Chapman and McKenzie, 2009), embryos cannot utilize this response. Mortality can result directly from severe hypoxia or indirectly from reduced growth increasing susceptibility to predation. Even fish that survive may incur sublethal effects with lasting, lifelong consequences for growth, development, and reproduction (Stierhoff et al., 2006; Vanderplancke et al., 2015; Zambonino-Infante et al., 2017). Modeling the energetic mechanisms of responses to hypoxia using unified principles on model species can help connect physiology and life history to population-level changes and serve as a valuable alternative or supplement to time- and labor-intensive laboratory experiments on other species, particularly with very small embryos and larvae.

Hypoxia is known to inhibit growth and survival in early life fishes (Rombough, 1988; Cross et al., 2019; Del Rio et al., 2019), as oxygen is required for the processes that maintain homeostasis and convert food for growth and activity. Anaerobic energy production fuels these processes with about 1/15th the ATP yield of aerobic respiration. Hypoxic exposure leads to physiological responses such as depressed metabolism (Richards, 2009; Schwemmer, 2023), limited growth, increased ventilation, and changes to hematocrit, hemoglobin, and erythrocyte quantities and characteristics (Taylor and Miller, 2001; Stierhoff et al., 2009a; Bianchini and Wright, 2013). Metabolism has also been shown to increase after temporary hypoxia as fish remove lactate accumulated from anaerobic respiration (Heath and Pritchard, 1965).

Hypoxia often has interactive effects with other stressors such as temperature (Brandt et al., 2009; McBryan et al., 2013; Earhart et al., 2022) and high carbon dioxide (CO2; Hancock and Place, 2016; Miller et al., 2016; Morrell and Gobler, 2020). The Atlantic silverside (*Menidia menidia*) is an estuarine forage fish that has frequently been used as a model species to understand interactive effects of high CO2 and hypoxia (DePasquale et al., 2015; Miller et al., 2016; Murray and Baumann, 2018; Schwemmer et al., 2020). Rearing *M. menidia* offspring in static low DO significantly delayed hatching, reduced survival to hatching and larval survival, and reduced embryo and larval growth (Cross et al., 2019). However, the negative effects of both hypoxia and high CO2 were mitigated when offspring were exposed to diel-fluctuating treatments rather than static (Cross et al., 2019). While diel fluctuations are a realistic representation of changes in community photosynthesis and respiration between day and night, environmental change in coming years could extend hypoxic duration to reduce periods of relief. Warming reduces oxygen solubility while increasing metabolic rates of organisms that draw down oxygen when densely aggregated. At the same time, higher summer temperatures and freshwater input in some regions will intensify stratification that separates low-oxygen water from surface oxygen diffusion (Rabalais et al., 2009; Howarth et al., 2011). Currently *M. menidia* is tolerant enough that population declines are not a concern, but without knowledge of the mechanisms of early life impacts it is hard to anticipate whether this will change under intensifying deoxygenation or with additional stressors (Baumann, 2019).

Dynamic Energy Budget (DEB) modeling is a bioenergetic framework designed to bridge multiple levels of biological organization in assessing stressor effects and their mechanisms in a vast variety of species (Kooijman, 2010; AmP, 2023). This approach follows energy allocation, in the form of suborganismal metabolic fluxes, and how it leads to life history outcomes such as growth rate, reproductive output, and survival, using physical and biological concepts that are generalizable to most species (Jusup et al., 2017). It accounts for differences in the energy budget at each stage to allow modeling of life stage transition timing and stage-specific responses to stressors (Kooijman, 2010). DEB theory is often used to connect experimental observations of multiple stressor effects to both the underlying energetic mechanisms (Kooijman, 2018) and life history outcomes that feed into population dynamics (Nisbet et al., 2000; Martin et al., 2013; Smallegange et al., 2017). It is important to connect suborganismal and organismal responses to population implications because targeted conservation actions operate at this level, but this scaling requires additional steps and remains a challenge (but see Nisbet et al., 1989; Grear et al., 2020; Tai et al., 2021). Care must be taken as linear relationships between individual- and population-level impacts can not be assumed (Chambers and Trippel., 1997; Galic et al., 2018). The ability to bridge levels of biological organization from the molecular to population level makes DEB theory an excellent tool for enhancing the utility of experimental hypoxia data for conservation and management (Lavaud et al., 2021).

Depending on the application and types of data available, simplified versions of the standard DEB model can be used (e.g. Kooijman and Metz, 1984; Jager, 2018; Martin et al., 2017). Although complexity can be beneficial (Evans et al., 2013), simpler models with fewer parameters are often preferable for their predictive power and ability to be applied, tested, and interpreted widely (Holling, 1966; May, 2001; Jusup et al., 2017). The DEBkiss framework (Figure 1) is a moderately simplified variation on the standard DEB model for animals that eliminates the concept of reserve, a pool of assimilates that are allocated to structure, maintenance, and reproduction in the standard DEB model (Jager et al., 2013). This framework reduces the data requirements, the role of compound parameters, and, depending on the data, the total number of parameters to be estimated (Jager et al., 2013). The simplicity of DEBkiss and its easily understandable equations make it ideal for adaptation to many species of ecological or commercial value, even when the existing studies were not originally intended for this use, because it uses commonly measured data, such as growth and survival rates.

We used a DEBkiss model to test the hypothesis that changes in animal performance under hypoxia can be explained by one or more of the rate processes in the model, and to identify the bioenergetic mechanisms underlying experimental hatching, growth, and survival effects of hypoxia in early life stages of *M. menidia* observed in Cross et al. (2019). First, we fit a DEBkiss model to full-life data on total length, reproductive output, hatch timing, and survival and estimated or calculated parameters under fully oxygenated conditions. Second, we used the concept of Synthesizing Units (SU) that are inhibited or damaged by hypoxia to directly or indirectly change key parameters in the DEBkiss model. SUs are generalized enzymes that produce products such as body structure or maintenance from incoming fluxes of substrate, i.e. food or egg buffer (Kooijman, 1998; Kooijman, 2010; Muller et al., 2019). We used a correction factor based on inhibition or damage to the SU to fit the model to early-life data for four DO treatments. We evaluated which parameter or combination of parameters, when adjusted with the correction factor, was able to best account for the full set of hypoxia responses observed in experiments and thus allow inference of a mechanism.

1. **Methods**
   1. *DEBkiss Model Description*

To model the stage-specific energy budget of *M. menidia* in a way that would allow us to explain early-life hypoxia effects with bioenergetic processes, we used DEBkiss, a simplified and widely applicable DEB model (Jager et al., 2013; Jager, 2018). DEBkiss uses fewer parameters than the standard DEB model, which reduces data requirements and the risk of overfitting. It also has no reserve compartment between food assimilation and allocation, and for embryos this means that the egg buffer is assimilated into body structure and for maintenance, with hatching occurring when the egg buffer is fully depleted, instead of following reserve dynamics of the standard DEB model (Jager et al., 2013). The lack of reserve makes DEBkiss well-suited for animals with a small ultimate body size because reserve plays a smaller role in such species under DEB theory (Nisbet et al., 2000), but DEBkiss has been successfully applied to larger animals as well (e.g. Desforges et al., 2017).

The DEBkiss assumptions and equations we used are from Jager (2018). The parameters are defined in Table 1 and the variables, differential equations, and conversions are defined in Table 2. The flux of food or, for embryos, the egg buffer (*WB*) is immediately converted to assimilates which are allocated to a somatic fraction (*κ*) and a reproductive fraction (1-*κ*; Figure 1); these fractions are constant throughout the life cycle. The assimilation flux (*JA*) is the product of the scaled measure of resource availability (*f*), the volumetric surface area (*L2*), and the parameter maximum area-specific assimilation rate (*JaAm*) where *f* = 1 for embryos and for post-hatching fish fed *ad libitum*. Egg buffer mass decreases at a rate inversely proportional to assimilation flux (Table 2). Within the somatic branch, which does not change with life stage, a flux to maintenance (*JM*) is prioritized while the remainder goes to the flux for structure (*JV*) with a conversion efficiency *yVA*. The maintenance flux is the product of volume and the parameter for the volume-specific cost for maintenance (*JvM*; Table 2).

For juveniles, the non-somatic fraction of assimilates is spent on maturation, or increasing complexity through gonad development. While the standard DEB formulation uses a state variable for maturity that triggers changes between life stages, DEBkiss instead uses a constant size at puberty to specify when reproduction is initiated (Kooijman, 2010; Jager et al., 2013). Once the mass at puberty is reached (*WVp*), reproductive flux (*JR*) toward egg production begins in adults with a conversion efficiency *yBA*. Although *M. menidia* have a distinct larval and juvenile stage, we treated both as the juvenile stage because the relevant aspects of their energy budget for DEBkiss are assumed to be identical. DEBkiss also uses an optional flux to maturity maintenance (*JJ*) that comes from the 1-*κ* fraction of assimilates (Jager, 2018), which we chose to use in our model.

Because the model equations use dry weight for body size and our growth data are in total length, we calculated a shape correction coefficient (*δM*) and dry weight density (*dV*) to allow the model to convert between the two (Table 2). We calculated these constants using data on *M. menidia* length and egg volume (Klahre, 1997) and a total length to dry weight conversion (H. Baumann, personal communication):

(1)

To address the assumption of DEBkiss that eggs hatch when buffer is depleted, regardless of body size or developmental progress (Jager et al., 2013), we added a survival state variable (*S*). In addition to allowing an alternative outcome to hatching, this allowed us to examine survival as a consequence of hypoxia effects on the energy budget. We fit mortality parameters for embryos and post-hatch fish (*μemb* and *μlar*) to data for survival to hatching and larval/juvenile survival (Figure 1; Table 2). In our implementation of survival, the only DEB process influencing survival is egg buffer depletion, which determines the time to hatch and thus when the embryo mortality rate switches to the post-hatch mortality rate. This means survival is indirectly affected by the assimilation rate and conversion efficiency of assimilates into structure.

* 1. *Data*

We calculated and estimated DEBkiss parameters in normoxic conditions (Section 2.3) and modeled hypoxia effects (Section 2.5) based on four types of data (state variables): total length over time, egg buffer mass over time (and through this, time to hatching), cumulative egg production over time, and proportion surviving since fertilization over time. Data for total length were sourced from four studies. Length at hatching and 15 days post-hatching (dph) came from a study that reared *M. menidia* offspring in different static oxygen levels across two experiments (Cross et al., 2019). This provided data for parameter estimation at control oxygen levels in Section 2.3 and modeling three reduced oxygen treatments used in Section 2.5 (Table 2). We sourced additional length data for the full life span from control levels of experiments that exposed *M. menidia* offspring to ambient and elevated CO2 levels (Murray and Baumann, 2018; Murray and Baumann, 2020; Concannon et al., 2021). All total length data were obtained from fish maintained in static laboratory conditions at 24°C.

Data for the state variables of egg buffer mass (i.e. time to hatching, when egg buffer mass is zero), as well as survival at hatching and at 15 dph, were obtained from Cross et al. (2019). As with the total length data, the control data from these experiments were used to estimate parameters under normoxia (Section 2.3) while the other oxygen treatments were used in Section 2.5. We also obtained normoxic survival data from a study on the effects of temperature and CO2 on *M. menidia* early life survival, using only the 24°C and control CO2 data (Murray and Baumann, 2018). Four additional data points for long-term survival in laboratory conditions at 17°C were obtained from a study that exposed *M. menidia* offspring until 122 dph to two CO2 levels, of which we only used data from the control level (Murray et al., 2017). Lastly, the data for cumulative egg production over time, used to estimate parameters under normoxia (Section 2.3), were also obtained from control groups in Concannon et al. (2021), a study in which wild-caught juveniles were held in the laboratory at 20°C in different CO2 treatments and strip-spawned once they reached reproductive maturity.

* 1. *Parameter estimation*

We estimated four parameters by fitting them to full-lifespan data (*JaAm*, *yVA*, *μemb*, and *μ­lar*), calculated four parameters from existing data (*JvM*, *WB0*, *LVp*, and *f*) and fixed at suggested values parameters for which we had insufficient data to calculate or estimate (*yAV*, *yBA*, *κ*, and *fB*; Jager, 2018). The primary parameters and their calculated or estimated values are found in Table 1. Fitting was done in Matlab with the packages BYOM v.6.4 (Jager, 2022) and DEBkiss v.2.3a (Jager, 2021). BYOM uses a Nelder-Mead simplex search to optimize the parameters for a set of ordinary differential equations by minimizing negative log-likelihood (NLL). The DEBkiss package works under BYOM to bring in the DEBkiss model parameters, variables, and equations so that the parameters can be estimated based on their effect on the DEBkiss equations and the equations derived from them. The differential equations give the predicted data for each type of observed data (length, egg production, egg buffer mass, and survival over time) the difference between which is used to calculate NLL.

BYOM allows users to turn fitting on and off for each parameter, and with fitting turned off for all parameters it runs a simulation that calculates predicted data values over time for each state variable using the initial parameter values. Before estimating any parameters with the optimization described above, we ran simulations with fitting turned off using a set of recommended parameters (Jager, 2018) and parameters obtained from existing data on *M. menidia*. We visually assessed fit and checked NLL as we adjusted parameters to obtain a reasonable set of initial parameters before estimating any. Testing a range of parameters and obtaining realistic initial parameters helps avoid detecting local minima with the optimization. This also helped us reduce the number of parameters being estimated to avoid overfitting and so that there were not multiple correlated parameters free at once, because we were able to obtain a reasonable fit using suggested values for *yAV*, *yBA*, and *κ* that are thought to be widely applicable across species (Lika et al., 2011; Jager, 2018). The suggested value for *yVA­* of 0.8 from the literature (Lika et al., 2011; Jager, 2018) did not allow a realistic fit to the length data, but the length, reproduction, and egg buffer depletion data allowed it to be estimated with the BYOM optimization. Ultimate length was used to fit *JaAm* to a reasonable value while fixing all other parameters and before estimating *yVA*, because both parameters affect growth and egg buffer depletion in the model and therefore cannot be estimated simultaneously. Finally, we fixed all parameters except *μemb* and *μlar* to estimate these parameters, again using the visually best-fitting parameters from the simulations as initial values. The full-life predicted and observed data are shown in Figure 2.

The length and reproductive data allowed us to calculate length at puberty (*LVp*), which in the DEB literature is defined as the length at which egg production begins. We obtained *WB0* from *M. menidia* egg dry weight data (Klahre, 1997) and calculated *δM* and *dV* from total length, egg diameter, and egg mass data (Cross et al., 2019; Klahre, 1997; Concannon et al., 2021). To calculate volume-specific maintenance costs (*JvM*), we used data on the rate of decrease in larval dry weight over a period of starvation in the congeneric species *M. beryllina* (Letcher and Bengtson, 1993). Borrowing from closely related species is a common practice in bioenergetic modeling when the species has similar habitat, life history, and physiology (Sibly et al., 2013). *M. menidia* and *M. beryllina* have overlapping habitats and similar life history, egg sizes, and body sizes, although *M. beryllina* reaches a smaller ultimate length (Middaugh, 1981; Bengtson, 1984; Middaugh and Hemmer, 1992). All *M. menidia* experiments used in this study fed fish *ad libitum* in all treatment levels, so *f* was set to 1. For studies that exposed fish to different CO2 levels, we only used data from control groups to avoid potential CO2 effects in the data.

* 1. *Relating DEB processes to physiology*

We aimed to identify the DEBkiss parameters responsible for observed whole-organism effects of rearing *M. menidia* in hypoxia by applying a correction factor to modify one or more parameters with decreasing oxygen based on inhibition or damage to the SU by the physiological consequences of hypoxia exposure. SUs are generalized enzymes, including those that convert oxygen and food into assimilates and assimilates into body structure following kinetics similar to those of a Michaelis-Menten-Briggs-Haldane enzyme (Kooijman, 2010; Muller et al., 2011; Muller et al., 2019). Although oxygen can be a limiting substrate in SUs, previous work suggests that *M. menidia* embryos only become metabolically oxygen-limited below a critical level of 2.04 mg L-1 (Schwemmer, 2023), while it remains oxygen-independent at the treatments for which we have data (2.7, 3.1, 4.2, and 7.7 mg L-1; Schwemmer et al., 2020). We therefore considered a single-substrate growth SU in which food or egg buffer was the substrate and hypoxia effects on growth, hatching, and survival occur through inhibition of assimilation or damage that reduces the conversion efficiency of assimilates to growth. Inhibiting agents reversibly bind to SUs, preventing them from accepting substrates to proceed with their reaction. Damage, in contrast, induces dysfunction that is irreversible upon removal of the damaging agent; however, damaged SUs can be repaired or replaced (Muller et al., 2019). Although hypoxia itself cannot bind to an SU itself, it can induce the production of compounds that could in turn bind to SUs. We used existing information on the physiological responses of fish early life stages to hypoxia to identify the following DEBkiss parameters to which to apply the hypoxia-based correction factor: maximum assimilation rate, conversion efficiency of assimilates into structure (growth), maximum somatic maintenance rate, embryo mortality rate, and post-hatch mortality rate.

Assimilation is the transformation of food and oxygen into compounds that will go to structure, maintenance, or reproduction (Kooijman, 2010; Jager, 2018). Reduced food consumption, and thus limitation of substrate in the SU, is a primary mechanism by which the fish energy budget is thought to be impacted by hypoxia (Chabot and Dutil, 1999; Thomas et al., 2019). However, feeding effects cannot explain the observed hypoxia impacts on *M. menidia* hatch survival, timing, and size (Cross et al., 2019) because embryos do not yet ingest food. Instead, binding of inhibiting agents is a viable explanatory process by which hypoxia impacts the SU. Several genes controlling cell division and protein synthesis are regulated by hypoxia (Ton et al., 2003), such as insulin-like growth factor binding protein 1 (IGFBP-1), a protein controlled by hypoxia-inducible factor 1 (Hif-1) that is shown to reduce growth and delay development in fish embryos exposed to hypoxia (Kajimura et al., 2005; Kajimura et al., 2006; Sun et al., 2011; Tian et al., 2014). This factor is thought to trade off growth to reduce oxygen demand and help fish tolerate hypoxia. By preventing insulin-like growth factors from binding to their receptors, IGFBP-1 inhibits signaling for cell division and differentiation and energy can be diverted to processes necessary for survival (Kajimura et al., 2005). In the DEBkiss model such forms of inhibition to the SU under hypoxia would be represented by reduced assimilation rates (Muller et al., 2019). Changes to assimilation in response to hypoxia have been recorded in other species, but the direction of that effect is species-dependent (reviewed in Thomas et al., 2019).

The conversion efficiency of assimilates to structure controls growth and hatch timing because it is the fraction of assimilates that are converted into structure rather than burned on overhead costs of growth (Jager, 2018). When oxygen is low enough that anaerobic metabolism must be used, this reduces conversion efficiency so that less growth results from the same amount of yolk or food (Thomas et al., 2019). Damage to the SU may be responsible for reductions in conversion efficiency, perhaps through lactate accumulation and consequential declines in internal pH. Although capability for anaerobic glycolysis in embryos and yolk sac larvae appears to vary widely across species (Wieser, 1995; Finn 1995; Rombough, 1988), smaller fishes reach harmful levels of anaerobic end-products much faster than larger fishes due to their higher mass-specific metabolic rates (Nilsson and Östlund-Nilsson, 2008). We hypothesized that this contributed to a smaller hatch size and slower growth post-hatch.

Maintenance in DEBkiss is the energy allocated to any processes that support the integrity and functioning of the structural body (Jager, 2018), including homeostasis, damage repair, and activity. Demand for more protein turnover and cell repair can increase the volume-specific maintenance rate following damage (Bouma et al., 1994; Kooijman, 2010) and indeed maintenance has been shown to increase with damage to structural proteins (Maury et al., 2019). In addition to damage repair, maintenance rate could be elevated by the activity required for some of the behavioral responses fish exhibit under hypoxia (Thomas et al., 2019). *M. menidia* exposed to hypoxia swim to the surface to use aquatic surface respiration, taking advantage of the diffusion of oxygen from the air (Miller et al., 2016). This behavior is impossible in embryos but has been observed in larvae (Cross et al., 2019). Fishes also expend energy on faster ventilation and heartbeat when ambient DO is low to increase oxygen uptake (Kramer, 1987; Maxime et al., 2000) and remove accumulated CO2 and lactate (Perry et al., 2009; Heath and Pritchard, 1965), but these capabilities may be limited until development has progressed further.

Although mortality is not a process directly represented by an SU, it could indirectly be impacted by hypoxia effects on SUs through failure to meet developmental milestones – particularly for hatching – or damage that exceeds repair rates. In the parameter estimation using data from normoxic conditions (Section 2.3), our estimated survival parameter for embryo mortality was greater than that of larvae (Table 1). If assimilation rate or conversion efficiency of *M. menidia* decreases under hypoxia, the resulting slower egg buffer depletion would delay hatching, extending individuals’ time in the stage with greater mortality and thus accounting for reduced hatch survival under hypoxia. We therefore hypothesized that if either assimilation rate or conversion efficiency is modified by the hypoxia-based correction factor, additionally modifying the embryo mortality parameter would consequently not be necessary to account for hypoxia effects on all four state variables. However, this would not be the case for the post-hatch mortality parameter because none of the processes in the DEBkiss model indirectly affect mortality after hatching, so changing either the assimilation or conversion efficiency parameter in combination with the post-hatch mortality parameter may be necessary to fully replicate the observed changes to growth, hatch timing, and survival under hypoxia.

* 1. *Hypoxia Effects*

We tested the hypothesis that changes in *M. menidia* early life growth, hatch timing, and survival under reduced oxygen (Cross et al., 2019) can be explained by one or more DEBkiss processes (Figure 1). To summarize the experimental data on static hypoxia effects we are attempting to explain by altering these parameters, the mean values of data for each oxygen treatment are listed in Table 3. We used the parameter values from the model fit to full life data and altered one or more parameters at a time with oxygen-dependent correction factors, then fit the model to data for only the first 136 days by estimating a parameter that controls the correction factor’s relationship with DO. We only used early life data to fit the hypoxia-altered parameters because we did not have late-life or reproductive data for multiple oxygen treatments against which to validate observed changes. It did not make sense to include later life data in the calculations of NLL that influence the parameter estimates or to speculate about how well the predicted data match what we might expect to happen later in life if we not only lack late-life hypoxia data but also do not expect full-life hypoxia to occur in nature.

We derived a correction factor using the framework developed by Muller et al. (2019), in which inhibitors can act on SU dynamics in five different ways. Out of these, noncompetitive inhibition is best suited to this study because of data availability for *M. menidia*. In noncompetitive inhibition the arrival rate of substrate does not affect the binding and dissociation of inhibitors and therefore requires little information about the rate of food uptake (Muller et al., 2019). In this form of inhibition, the rate of assimilation by the SU is:

(2)

where *ji* is the arrival flux of the inhibitor and *ki* is the dissociation parameter. The effect of this relationship in our model is that assimilation declines as the arrival rate of hypoxia-related inhibitors increases. We set *ji* to depend on DO treatment above a DO threshold, DOc, below which *ji* is infinitely large, which would bring the rate of the process it is inhibiting to zero:

(3)

*Z* is a parameter that influences the shape of the relationship between *ji* and DO. We defined the correction factor *c* as the inhibition term (in parentheses in Equation 2) and replace *ji* with the function from Equation 3 for DO > DOc to derive the correction factor *c*:

(4)

As only the product of the parameters *ki* and *Z* appear in the formula and we have no need to estimate them separately, they can be combined into one parameter as *Z*. Simplifying Equation 4 and adding in the case for which DO ≤ DOc gives us the following correction factor:

(5)

The relationship between *c* and DO for three different sample values of *Z*, the parameter to be estimated, is shown in Figure 3. A larger *Z* value keeps *c* higher as oxygen decreases before a more abrupt drop, while a smaller *Z* gives a more constant decline in *c* with hypoxia. The value of *c* cannot exceed 1. DOc was fixed at a biologically relevant level of 2.04 mg L-1, which is the critical oxygen level below which embryonic routine metabolism becomes highly oxygen-dependent (Schwemmer, 2023). Attempts to estimate DOc and *Z* simultaneously showed that leaving DOc free did not improve the ability of the correction factor to fit the hypoxia data. This correction factor was multiplied by *JaAm* and *yVA* because these parameters were hypothesized to decrease under hypoxia. Although *yVA* is not a flux or process that can be inhibited, it can be impacted by damage, the irreversible destruction of functionality of an SU (Muller et al., 2019). However, if damage production is much slower than the maximum production rate of the SU, the model for noncompetitive damage is equivalent to that of noncompetitive inhibition (Muller et al., 2019). The parameters for maintenance and mortality were divided by *c* because they were hypothesized to increase, rather than decrease, with damage production and inhibition.

To find the best value of *Z* for each DEBkiss parameter or combination of parameters, we added *Z* as a model parameter and estimated it using the BYOM optimization to minimize NLL. We weighted the data equally across treatments to correct for differences in sample size across treatments and prevent one treatment group from disproportionately affecting the estimation of Z, so that all weights within each treatment added up to the same number. We did not apply the correction factor to *JaAm* and *yVA* simultaneously because they are multiplied together to obtain *JV* and their individual contributions to the growth and egg buffer depletion can not be fully separated. We only compared the fit of models in which *c* was applied to parameter(s) that resulted in all three early life datasets – total length, egg buffer mass, and survival – being affected by hypoxia. As a result, either *JaAm* or *yVA* is in each candidate model, because *Jv­M*, *μemb*, and *μlar* do not affect egg buffer depletion. To identify the most likely version of the model (which parameter or combination of parameters best explain the hypoxia effects on the state variables), we estimated *Z* for each of these scenarios and calculated Akaike’s Information Criterion for small sample sizes (AICc). We compared the AICc between each model using the difference between AICc values (ΔAICc) and the relative likelihood of each model using Akaike weights:

, (6)

where *wi*(AICc) is the Akaike weight of each model *i*, Δ*i*AICc is the difference between each model *i* and the model with the lowest AICc (AICcmin), and the denominator calculates the sum of relative likelihoods for every model starting at the first model *k* (Wagenmakers and Farrell, 2004). We used ΔAICc and ratios of Akaike weights to determine which combination of parameters best fit the data when inhibition or damage was applied and, therefore, which DEB processes best explain the hypoxia effects observed in experiments (Table 4).

1. **Results**
   1. *DEBkiss Model*

We obtained realistic fits to all datasets (Figure 2). The only exception is late-life survival, for which the mortality was too high beyond the larval stage but could not be better fit due to lack of full-life survival data. However, this did not impair our ability to model the effects of hypoxia on early life survival. Estimating *yVA* returned a value much lower than 0.8, which is the value suggested by Jager (2018) and has been applied in DEBkiss models of other species (e.g. Jager et al., 2018; Hamda et al., 2019). However, our value of *yVA* =0.365 is close to the maximum growth efficiency of 0.375 measured in the closely related *M. beryllina* (Letcher and Bengtson, 1993). This gave a realistic fit to the length data and allowed a detailed and very close fit to egg buffer mass over time (hatch timing). The observed and predicted data for full life span and early life are plotted in Figure 2.

* 1. *Hypoxia Effects*

Applying the oxygen-dependent correction factor to the parameter combinations listed above reproduced the direction of experimentally observed hypoxia effects, e.g. increasing *JaAm* reduced total length, increased time until egg buffer mass reaches 0, and reduced survival. The best model of experimental hypoxia effects on *M. menidia* early life stages simultaneously had *yVA* multiplied by *c*, and *μemb* and *μlar* divided by *c* (Figure 4, Table 4, Figure S1). Although applying damage to *yVA* alone affected all three state variables, concurrently increasing both mortality parameters improved the fit to the data (Table 4). It also had the lowest AICc of all candidate models, with an AICc of 794.03 (AICcmin). Adding a correction factor to *JvM* in addition to these three parameters yielded a slightly higher AICc of 795.97. The ratio of Akaike weights shows that the model with *c* applied to *yVA*, *μemb*, and *μlar*, is 2.67 times as likely as the one with *c* concurrently applied to *JvM* (Table 4). Applying a damage effect to maintenance was therefore not considered to have improved the fit. After estimating *Z* we calculated the values of *yVA*, *μemb*, and *μlar* when their respective correction factors are applied for each DO level (Table 5).

Interestingly, although *JaAm* affects the variables similarly to *yVA*, the ratio of Akaike weights showed that the best fitting model is about 3000 times as likely as the version applying inhibition to *JaAm*, *μemb*, and *μlar* (Table 4). Reducing *JaAm* with hypoxia using correction factor *c* resulted in a visually good fit to the data across oxygen levels and variables. Combining the adjusted *JaAm* with correction factors to increase both mortality rates improved the fit as well, but this model fit less well than the version that applied *c* to *yVA*, *μemb*, and *μlar*, with an AIC value of 810.21 in the former model compared to 794.03 in the latter.

The estimated best value of *Z*, the exponential coefficient in the correction factor *c*, enables us to calculate that *yVA* at the lowest oxygen level is 55% of its value with no hypoxia stress. Reducing conversion efficiencyalone produced small differences in survival at hatching because it prolongs the time spent in the embryo stage, which has a greater mortality rate than post-hatching in our model. Dividing both the pre- and post-hatching mortality rates by *c* more closely predicted the reduced survival rates in the low DO treatments, resulting in a best fitting model that explained observed hypoxia effects well by altering conversion efficiency, embryo mortality, and post-hatch mortality.

1. **Discussion**

By combining experimental data with unified principles for energetic allocation that are broadly applicable across species, we identified the conversion efficiency of assimilates into structure as the most likely process by which low oxygen levels affect early life stages of *M. menidia*. In comparing combinations of DEBkiss parameters that influence the ecological endpoints (total length, hatch timing, and survival), we discovered that applying correction factors based on damage production to the growth SU to reduce conversion efficiency (*yVA*) and increase pre- and post-hatching mortality rates (*μemb* and *μlar*) best predicted the experimental effects of hypoxia on larval length, time to hatching, and early life survival. Through this model we have found evidence that the mechanism largely responsible for the observed hypoxia impacts on growth, hatch timing, and survival is the efficiency by which assimilated food or egg yolk is converted into structure.

Reducing assimilation with hypoxia rather than conversion efficiency yielded a worse fit despite the two parameters’ similar contributions to the DEBkiss model in that both parameters are used to calculate predicted growth and egg buffer depletion. In our implementation of hypoxia effects on SUs, these two parameters are assumed to be impacted by different molecular mechanisms but the same noncompetitive inhibition-based correction factor, due to the assumption that the maximum rate at which the SUs form product is much greater than the damage production rate (Muller et al., 2019). Reducing either assimilation or conversion efficiency would extend developmental time, which is consistent with previous work showing yolk absorption slows under hypoxia (Polymeropoulos et al., 2017). As maintenance costs must continue to be paid, this would increase the energy expended to produce each unit of structure (Kamler, 2008). However, the mechanism for reduced conversion efficiency is assumed to be damage to the synthesizing unit, perhaps from buildup of anaerobic byproducts, and repairing the SU incurs additional maintenance costs compared to inhibition which requires no repair (Bouma et al., 1994; Kooijman, 2010; Muller et al., 2019). Conversion efficiency can also be reduced by the far less efficient production of ATP through anaerobic respiration combined with slower rates of tissue differentiation. The experimental DO levels are greater than the critical oxygen levels for oxygen-independent routine metabolism (*P*crit) of 2.04 mg L-1 and 1.56 mg L-1 for embryos and 5dph larvae, respectively (Schwemmer, 2023). *P*crit has been assumed by some to be the oxygen level at which anaerobic metabolism is triggered, but there is abundant evidence that some level of anaerobic metabolism can occur well above *P*crit (Nonnotte et al., 1993; Maxime et al., 2000; Wood et al., 2018). Additional activity such as swimming bursts can drive up the need for anaerobiosis (Di Santo et al., 2017). Our evidence that conversion efficiency is reduced by hypoxia-induced damage suggests that anaerobic metabolism is a mechanism of hypoxia effects in *M. menidia* early life stages even at oxygen levels above *P*crit.

A limitation of our model is that it does not allow us to test for the possibility that both conversion efficiency and assimilation rate are contributing to the observed hypoxia effects. It is not possible to simultaneously estimate both parameters as their product is used to calculate growth in the model; we can adjust one or the other with the correction factor and get similar effects on the flux for growth with no way of determining which is correct. If conversion efficiency were the only parameter varying across hypoxia treatments, one might expect the offspring to fully deplete the egg buffer and hatch at the same time, but with hatch size increasing with DO level. However, adjusting conversion efficiency with hypoxia does account for the observed significant differences in hatch timing between DO treatments in *M. menidia* larvae (Cross et al., 2019) because *yVA* reduces the body size at a given time, indirectly reducing the assimilation flux due to smaller body volume. It is nonetheless possible that *JaAm* is responsible for an unknown portion of the hypoxia effects.

Although both conversion efficiency and assimilation can explain hypoxia effects on total length and egg buffer mass over time, reducing them only produced a small decrease in survival relative to the data. Applying *c* to both mortality rates better captured the great reductions in survival at both hatching and 15 dph with hypoxia. In the experiments, the lowest oxygen level (2.7 mg L-1) had a mean hatch survival of 30.2% while the mean survival in the other three treatments was over 70% (Cross et al., 2019). By 15 dph fish from all three low oxygen treatments had lower survival than those from the normoxic treatment (Cross et al., 2019; Table 3). Including hypoxia effects for both pre- and post-hatching mortality rates allowed the model to predict these stage-specific differences in hypoxia effects more closely and improve the fit based on ΔAICc (Table 4). However, an intrinsic mortality rate isn’t as explicitly indicative of underlying energetic processes as the other DEB parameters are. The additional mortality that was not accounted for by *yVA* may have been related to unrepaired damage from buildup of toxic compounds during anaerobic metabolism (Richards, 2011). The mortality could also have resulted from failing to meet energetic demands with either aerobic or anaerobic metabolism (Richards, 2009) and, specifically in embryos, failure to reach a viable level of complexity before the yolk is depleted (Jager et al., 2013). Measurement of anaerobic byproducts such as lactate and morphometric assessment of dead embryos and larvae could help to identify the mechanisms underlying the mortality rates in future work. Although survival does not approach 0% during the larval stage in our best fitting model (Figure 4), all experimental replicates of the 2.7 mg L-1 DO treatment had 0% survival by 15 dph, making larvae apparently more sensitive than embryos (Cross et al., 2019). The authors of the study attribute this to a possibly lower ability to suppress metabolism in larvae compared to embryos. While the increased mobility of larvae may allow aquatic surface respiration (Miller et al., 2016; Cross et al., 2019) and escape from hypoxia in a patchy and stratified estuarine environment, activity comes with elevated maintenance costs in addition to those required to begin feeding almost immediately after hatching (Middaugh and Lempesis, 1976). This may also be a crucial time to repair damage to the SU (Muller et al., 2019), and the combination of these additional maintenance demands may be too great to meet without restoration of normoxia. Though beyond the scope of this work, a model that captures stage-specific differences in maintenance costs and links them explicitly to survival may better capture the mechanism of high mortality in larvae.

Adding a correction factor to the volume-specific maintenance rate in addition to *yVA*, *μemb*, and *μlar* did not substantially improve the fit according to AICc, suggesting that increasing maintenance costs is not a bioenergetic mechanism underlying hypoxia response in early life stages. This is consistent with laboratory measurements showing no effect of these hypoxia levels on embryonic or larval metabolic rates (Schwemmer et al., 2020). In our model, egg buffer depletion is insensitive to changes in volume-specific maintenance costs, requiring a quadrupling to see a noticeable delay in hatching. Changing maintenance has much greater effects on length later in life while failing to explain differences in length at the time of hatching. Repairing damage and increasing ventilation and swimming activity could both increase maintenance costs (Thomas et al., 2019), but at the embryo stage very little activity is possible. A common response to hypoxia in some fish embryos is premature hatching (Kamler, 2008) which could allow swimming escape responses that increase maintenance costs and in theory reduce growth, but studies on chorion removal have shown that the increased mobility can improve growth despite hypoxia exposure (Ciuhandu et al., 2005; Ninness et al., 2006). Some studies on fish responses to hypoxia suggest maintenance may drop temporarily due to the reduced capacity for aerobic metabolism at low DO levels, then subsequently be temporarily elevated after oxygen is restored because of recovery demands such as paying oxygen debt and removing or repairing damage from anaerobic byproducts (Heath and Pritchard, 1965; Claireaux and Chabot, 2016; Thomas et al., 2019). Such fluctuations in maintenance were not discernible in the time scale of our model, but future work should attempt to model the *M. menidia* early life energy budget during recovery from hypoxia.

Understanding the mechanisms of reduced growth and survival under hypoxia through DEB theory can be useful for predicting life history effects, and although modeling population growth rates was not within the scope of this study, our results have implications for processes that influence population dynamics. The predicted data resulting from fitting to early life data with a hypoxia-based correction factor predicts reductions in long-term growth and survival that would certainly be detrimental to population growth under extended periods of low oxygen. Under this model, even restoring normoxia after 15 days would result in smaller size at age and survival rates than the groups exposed to 7.7 mg L-1, and damage to the SU is not reversed upon return to normoxia, but rather requires energy to repair (Muller et al., 2019). However, compensation of growth may be possible after exposure to hypoxia (Wei et al., 2008) and other stressors (Russell and Wootton, 1992; Nicieza and Metcalfe, 1997; Ali et al., 2003). Delayed hatching and slower growth can lead to enhanced vulnerability to predation (Chambers and Leggett, 1987; Takasuka et al., 2007), further reducing survival rates beyond those observed in controlled laboratory conditions, although this is not always the case (Lankford et al., 2001; Robert et al., 2023). An important assumption of our model is that several of the parameters have the same value across life stages (e.g. *JaAm*, *JvM*, *yVA*) and similarly that values of the hypoxia correction factors are the same regardless of life stage. We lacked reproductive data to look at hypoxia effects on the proportion of total energy allocated to reproduction (1-*κ*), which is an additional component of DEB useful in connecting organismal effects to populations. Future experimentation could provide the high-resolution information for all life stages that is needed to extend this DEB model to predict population growth, which would be useful for resource management applications (Kooijman et al., 2020; Lavaud et al., 2021).

The oxygen levels in the estuaries inhabited by *M. menidia* undergo great diel and seasonal fluctuations (Baumann et al., 2015). The effects of fluctuating DO cannot be resolved in the time scales used by our DEBkiss model, so we assumed constant DO levels. As a result, the model is more useful in identifying mechanisms than in quantitatively predicting how *M. menidia* will respond to realistic hypoxia scenarios, as lifelong constant hypoxia is unrealistic. A study comparing the early life responses of this species to static and fluctuating hypoxia treatments suggested that fluctuations provided temporary relief that improved growth and survival (Cross et al., 2019), therefore our model would likely overestimate the negative impacts of hypoxia compared to outcomes in nature. It is also unrealistic for only a single environmental factor, in this case hypoxia, to influence the energy budget. This model can be adapted in future work to examine the interaction between hypoxia and acidification, which often co-occur in nature (Wallace and Gobler, 2021). Previous experiments on *M. menidia* embryos showed that high CO2 increases oxygen dependence of metabolism under both chronic (Schwemmer et al., 2020) and acute hypoxia (Schwemmer, 2023). Adding oxygen as a second substrate in the SU would allow a DEB model to incorporate the oxygen limitation that is evidently introduced at higher DO levels under acidification.

Our best fitting model overestimated time to hatching at 7.7 mg L-1 DO and overestimated survival at age for the 2.7 and 4.2 mg L-1 treatments, which suggests there either may be a different nonlinear correction factor function that better fits the relationship between DO and the DEBkiss parameters or that there were additional factors contributing to these differences that the model does not account for. For example, we do not have data on gonad development or reproductive output later in life after rearing *M. menidia* in hypoxia, which would allow us investigate if *κ* is an affected parameter. Hypoxia can reduce gonadosomatic index and gonad development in fish, suggesting that the reproductive branch of the energy budget might require additional energy to be redirected from the somatic branch (Wu et al., 2002; Thomas et al., 2006; Landry et al., 2007). Despite the potential for improvements with more data, the model was able to replicate the direction of effects and even account for some hypoxia effects in all three state variables simultaneously by changing only one parameter, either conversion efficiency or assimilation. Further, it provided these reasonable fits using an SU model based in well-studied and widely applicable Michaelis-Menten-Briggs-Haldane enzyme kinetics (Muller et al., 2019) rather than a more specialized or complex correction factor.

1. **Conclusions**

With this simple and widely applicable DEBkiss model we were able to attribute much of the hypoxia-related variability in total length, egg buffer mass, and survival over time to changes in DEB processes. The evidence for the mechanisms is inferred from a combination of experimentally observed responses and unified principles that apply to virtually all animal species (Jager et al., 2013; Kooijman, 2010). Similar approaches have applied correction factors to DEB parameters to model other species’ responses to hypoxia (Lavaud et al., 2019; Aguirre-Velarde et al., 2019) and other stressors such as seawater acidification (Jager et al., 2016; Pousse et al., 2022) and pollutants (Muller et al., 2010; Desforges et al., 2017). The success of this approach with a wide variety of stressors makes it an ideal supplement to multistressor experiments, which are limited by logistical constraints. Modeling stressor effects with DEBkiss parameters can yield additional information about energetic mechanisms of responses and, with careful attention to the assumptions being made, may be useful in extrapolating stressor effects to additional levels or combinations of stressors that would have been impractical to test experimentally or to species with certain shared physiology or life history traits (Goussen et al., 2020; Boult and Evans, 2021). The patterns modeled in this study should not be interpreted as a direct prediction of what will happen to wild *M. menidia* populations as coastal hypoxia intensifies. Lifelong constant oxygen levels do not occur and are not expected to occur in the future; rather, fluctuating oxygen levels will provide opportunities for recovery and may confer tolerance of temporary stress (Cross et al., 2019; Baumann, 2019). Instead, this approach demonstrates the value of identifying DEB parameters responsible for whole-organism effects of hypoxia to understand underlying energetic processes that are often time, labor, and cost-intensive to measure empirically, particularly in the early life stages, when biomass available for sampling is small and developmental changes are rapid. Through doing so we were able to highlight the conversion of assimilates to structure as a primary, but not sole, mechanism by which hypoxia reduces size, delays hatching, and increases mortality in an ecologically important forage fish.

**Declaration of Competing Interest**

The authors do not declare any competing interests.

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**Author Contributions**

Conceptualization – T.G.S., R.M.N., J.A.N.; Data curation – T.G.S.; Methodology – R.M.N., T.G.S.; Formal analysis – T.G.S.; Funding acquisition – T.G.S., J.A.N.; Visualization – T.G.S., Writing, original draft – T.G.S., Writing, reviewing and editing – T.G.S., R.M.N., J.A.N.

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**Data and Code**

I will add a list of DOIs for the data sources and my code.

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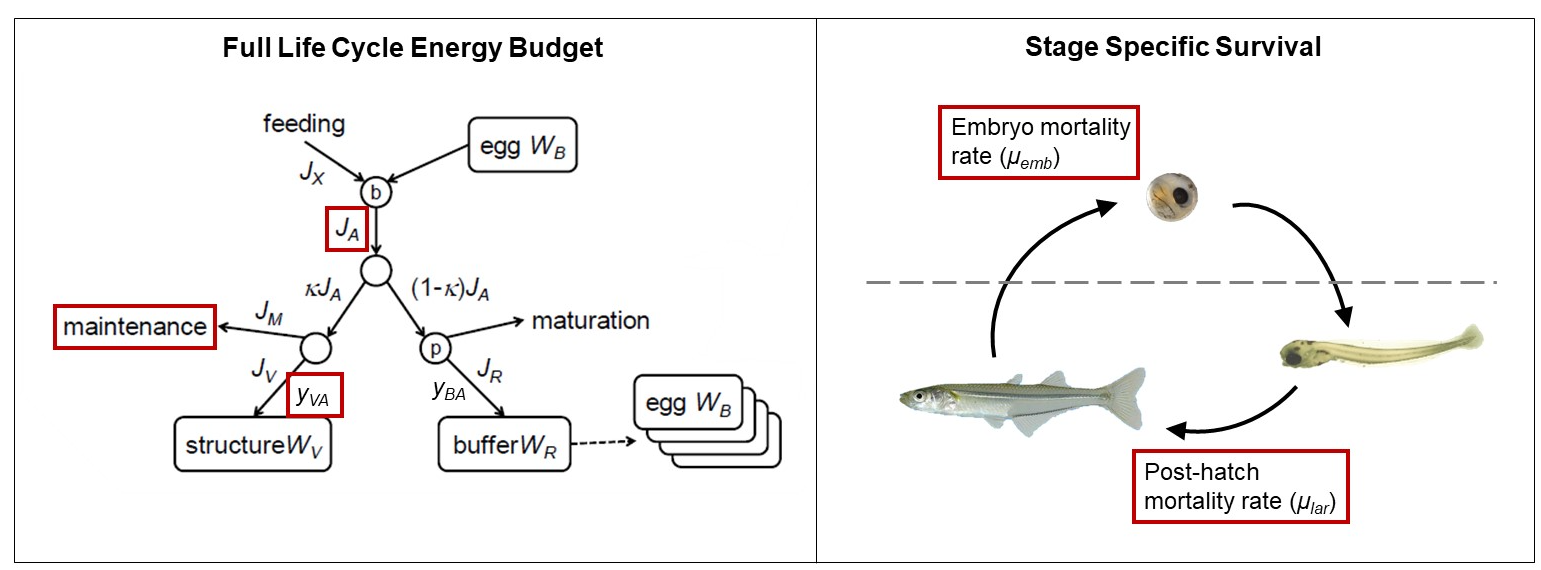
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**Figures**



**Figure 1.** The DEBkiss model (diagram adapted from Jager et al., 2013) with stage-specific survival parameters. The hypothesized parameters for hypoxia stress mechanisms are highlighted in red boxes. The left panel shows the energy budget for the full life cycle and the right panel shows the stage-specific survival modification.

A graph of a function

Description automatically generated with medium confidence

**Figure 2.** Predicted (lines) and observed data (dots) for the DEBkiss model of *M. menidia*. The state variables are (A) total length (mm) over time (days), (B) cumulative reproduction (eggs) over time (days), (C) egg buffer mass (mg) over time (days), and (D) survival over time (days). Predicted data lines were calculated with the parameter values listed in Table 1.

A graph of oxygen

Description automatically generated

**Figure 3.** The effect of DO on correction factor *c* at three different values of the exponential parameter *Z*. Actual estimated *Z* values are listed in Table 4, and the three *Z* values used in this figure are sample values to show how *Z* affects the relationship between DO and *c*.

**A diagram of a graph

Description automatically generated with medium confidence**

**Figure 4.** Best fit of DEBkiss model to experimental data from four DO levels, showing early life data only. The best fitting model was selected based on the requirement that all three response variables’ predicted values are affected by the hypoxia correction factor and based on lowest AICc. (A) is total length (mm) over time (days), (B) is egg buffer mass (mg) over time (days), and (C) is survival over time (days), with means rather than all data plotted for survival for ease of viewing. Full datasets used to estimate the correction factor parameter *Z* are plotted in Figure S1.

**Tables**

**Table 1.** DEBkiss parameters, their abbreviations, and their fixed or estimated values from fitting to full life data. Units are given with the value unless the parameter is a unitless ratio.

|  |  |  |  |
| --- | --- | --- | --- |
| **Parameter** | **Symbol** | **Fixed or estimated** | **Value** |
| Max. area-specific assimilation rate | *JaAm* | Estimated | 0.333 mg mm-2 d-1 |
| Max. volume-specific maintenance rate | *JvM* | Fixed | 0.0214 mg mm-3 d-1 |
| Initial egg buffer mass | *WB0* | Fixed | 0.15 mg |
| Total length at puberty | *LVp* | Fixed | 102 mm |
| Yield of assimilates on volume | *yAV* | Fixed | 0.8 |
| Yield of egg buffer on assimilates | *yBA* | Fixed | 0.95 |
| Yield of structure on assimilates | *yVA* | Estimated | 0.365 |
| Fraction of assimilates allocated to soma | *κ* | Fixed | 0.8 |
| Scaled food level | *f* | Fixed | 1 |
| Scaled food level for embryo | *fB* | Fixed | 1 |
| Mortality rate for embryos | *μemb* | Estimated | 0.0639 |
| Mortality rate for larvae | *μlar* | Estimated | 0.0294 |

**Table 2.** Fluxes, state variables, and differential equations in the DEBkiss model.

|  |  |  |  |
| --- | --- | --- | --- |
| **Flux** | **Symbol** | **Equation** | **Units** |
| Assimilation flux | *JA* |  | mg day-1 |
| Maintenance flux | *JM* |  | mg day-1 |
| Flux to structural growth | *JV* |  | mg day-1 |
| Flux to reproduction buffer | *JR* |  | mg day-1 |
| Flux to maturity | *JJ* |  | mg day-1 |
|  | | | |
| **State Variable** | **Symbol** | **Equation** | **Units** |
| Structural dry mass over time | *WV* |  | mg day-1 |
| Continuous reproduction rate | *R* |  | eggs day-1 |
| Egg buffer (yolk) mass | *WB* |  | mg day-1 |
| Survival | *S* |  | unitless  (range 0-1) |
|  | | | |
| **Other variables and conversions** | **Symbol** | **Equation** | **Units** |
| Total physical length | *LM* |  | mm |
| Volumetric length | *L* |  | mm (cubic root of volume) |
| Shape coefficient | *δM* |  | unitless |
| Dry weight density of structure | *dV* |  | mg mm-3 |
| Dry mass at puberty | *WVp* |  | mg |
| Volume-specific maturity maintenance costs | *JvJ* | *-* | mg mm-3 day-1 |
| Structural volume at puberty | *LVp3* | - | mm-3 |
| Scaled measure of resource availability | *f* | - | unitless  (range 0-1) |

**Table 3.** The mean survival to hatching, hatch time (at which egg buffer is zero), length at hatching, length at 15 dph, and survival to 15 dph from the different oxygen treatments in Cross et al. (2019). The control DO level means (7.7 mg l-1) also include data from Murray and Baumann (2018).

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
|  | **7.7 mg L-1** | **4.2 mg L-1** | **3.1 mg L-1** | **2.7 mg L-1** |
| Survival to hatching | 74.3% | 70.6% | 85.8% | 30.2% |
| Hatch time (egg buffer mass = 0) | 6 days | 7 days | 8 days | 9 days |
| Length at hatching | 5.3 mm | 4.6 mm | 4.4 mm | 4.1 mm |
| Larval length at 15 dph | 15.8 mm | 12.2 mm | 9.2 mm | - |
| Larval survival to 15 dph | 44.0% | 22.2% | 20.9% | 0% |

**Table 4.** The estimated *Z* value, AICc, ΔAICc, and Akaike weights when the correction factors were applied to each parameter or combination of parameters. ΔAICc and Akaike weights were calculated with AICcmin = 794.03 for the *yVA* + *μemb* + *μlar* model.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| **Parameter(s) affected by hypoxia correction factor** | **Estimated *Z* [95% CI]** | **AICc** | **ΔAICc** | **Akaike weight** |
| *JaAm* | 3.019 [2.512-3.612] | 856.06 | 62.03 | 2.5e-14 |
| *yVA* | 1.818 [1.601-2.342] | 848.65 | 54.62 | 1.0e-12 |
| *JaAm* + *JvM* | 3.105 [2.651-3.726] | 855.00 | 60.97 | 4.2e-14 |
| *yVA* + *JvM* | 1.985 [1.688-2.774] | 850.64 | 56.61 | 3.7e-13 |
| *JaAm* + *μemb* | 2.804 [1.605-3.287] | 823.24 | 29.21 | 3.3e-7 |
| *yVA* + *μemb* | 1.801 [1.570-2.167] | 808.12 | 14.09 | 6.3e-4 |
| *JaAm* + *μlar* | 2.930 [2.165-3.428] | 838.17 | 44.14 | 1.9e-10 |
| *yVA* + *μlar* | 1.767 [1.536-2.111] | 821.30 | 27.27 | 8.7e-7 |
| *JaAm* + *μemb* + *μlar* | 2.819 [1.920-3.286] | 810.21 | 16.18 | 2.2e-4 |
| *yVA* + *μemb* + *μlar* | 1.827 [1.620-2.269] | 794.03 | 0 | 0.72 |
| *JaAm* + *JvM* + *μemb* + *μlar* | 2.913 [2.288, 3.387] | 809.96 | 15.93 | 2.5e-4 |
| *yVA* + *JvM* + *μemb* + *μlar* | 1.981 [1.700, 2.456] | 795.97 | 1.94 | 0.27 |

**Table 5.** The value of the DEBkiss parameters that best reproduce the hypoxia effects observed experimentally, calculated (along with 95% confidence intervals in brackets) for each DO treatment level using the correction factor *c* and the estimated value of *Z* = 1.827.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
|  | **Product of correction factor and initial parameter value** | | | |
| **7.7 mg L-1** | **4.2 mg L-1** | **3.1 mg L-1** | **2.7 mg L-1** |
| ***yVA*** | 0.333  [0.329, 0.339] | 0.291  [0.284, 0.303] | 0.240  [0.230, 0.257] | 0.199  [0.188, 0.218] |
| ***μemb*** | 0.0701  [0.0689, 0.0709] | 0.0801  [0.0770, 0.0822] | 0.0970  [0.0906, 0.101] | 0.117  [0.107, 0.124] |
| ***μlar*** | 0.0322  [0.0317, 0.0326] | 0.0369  [0.0354, 0.0378] | 0.0446  [0.0417, 0.0466] | 0.0539  [0.0492, 0.0571] |



**Supplemental Figure**

A diagram of a curve

Description automatically generated with medium confidence

**Figure S1.** Best fit of DEBkiss model to all experimental data from four DO levels. The best fitting model was selected based on lowest AICc. (A) is total length (mm) over time (days), (B) is egg buffer mass (mg) over time (days), and (C) is survival over time (days).