

# Ecological Modelling

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

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<b>Corresponding Author:</b>	Teresa Grace Schwemmer, Ph.D. Story Brook University School of Marine and Atmospheric Sciences South Kingstown, RI UNITED STATES
<b>First Author:</b>	Teresa G. Schwemmer, Ph.D.
<b>Order of Authors:</b>	Teresa G. Schwemmer, Ph.D.
	Roger M. Nisbet, Ph.D.
	Janet A. Nye, Ph.D.
<b>Abstract:</b>	<p>Highlights</p> <ul style="list-style-type: none"><li>• Bioenergetic mechanisms of Atlantic silverside hypoxia responses were investigated.</li><li>• Hypoxia effects were modeled with a simplified Dynamic Energy Budget model.</li><li>• We connected physiology with energetic processes to identify potential mechanisms.</li><li>• Conversion efficiency and mortality parameters best explained hypoxia effects.</li><li>• This mechanism could impact energy flow across generations and trophic levels.</li></ul>
<b>Suggested Reviewers:</b>	Benjamin Martin University of Amsterdam Department of Theoretical and Computation Ecology b.t.martin@uva.nl Expertise in DEB models of fish, hypoxia and metabolism research, has used a DEBkiss model.
	Tjalling Jager DEBtox Research tjalling@debtox.nl Extensive expertise and experience with DEB modeling, developed the DEBkiss model, and has used it to model environmental stressor effects.
	Laure Pecquerie Université de Brest laure.pecquerie@ird.fr Expertise in DEB models of fish, fish metabolism and growth.
	Nina Marn Ruđer Bošković Institute nina.marn@irb.hr Expertise in DEB theory, physiology and energetics.
<b>Opposed Reviewers:</b>	

Dr. Teresa G. Schwemmer, Ph.D.  
Stony Brook University  
Stony Brook, NY 11794-5000  
teresa.schwemmer@gmail.com

March 29, 2024

Dr. Hsiao-Hsuan Wang, Ph.D.  
Editor-in-Chief  
*Ecological Modelling*

Dear Dr. Wang:

I am writing to submit our manuscript entitled “Attributing hypoxia responses of early life *Menidia menidia* to energetic mechanisms with Dynamic Energy Budget theory” for your consideration as a Research Article in *Ecological Modelling*, in the special issue “Metabolic organization across scales of space and time”.

We used a simplified Dynamic Energy Budget model (DEBkiss) to understand the mechanisms responsible for responses of Atlantic silverside (*Menidia menidia*) to hypoxia, an ecological stressor that is intensifying globally. This study expands on previous experimental work on Atlantic silversides that documented reduced early life growth, hatching success, and survival when reared under environmentally relevant levels of hypoxia. To connect these whole-organism responses to the energy budget, we employed the concept of synthesizing units (SU) to derive an oxygen-dependent correction factor that treats hypoxia as an inhibiting or damaging agent – a novel application of the SU concept. We identified conversion efficiency for growth and stage-specific mortality rates as the most likely DEB processes responsible for the experimentally observed responses to hypoxia.

This work will be of great interest to the readers of *Ecological Modelling* and this special issue because it takes a novel approach to modeling impacts of the real-world problem of intensifying hypoxia with a widely applicable DEB model. This model is a valuable tool to supplement ecological stressor experiments, particularly in early life stages when laboratory procedures are logistically constrained. Our results suggest that damage to the SUs, likely by anaerobic byproducts such as lactate, reduced the efficiency of growth and development, which could lead to lower recruitment and fecundity in realistic hypoxia scenarios. Readers will be interested in the implications of our findings using a model at the organismal scale of organization for population and ecosystem dynamics, as well as our suggestions for future work to refine the model using suborganismal data linked to physiological modes of action.

All authors have seen and approved the manuscript for submission to *Ecological Modelling* and have agreed to abide by the Editorial Policy. The authors confirm that this manuscript is original research that has not been previously published and is not under consideration for publication elsewhere. A previous version of this work is published only in my Ph.D. thesis and subsets of

our findings have been presented at the following conferences: the 2023 Dynamic Energy Budget Symposium in Baton Rouge, LA; the 2022 Larval Fish Conference in San Diego, CA; the 2021 Dynamic Energy Budget Symposium (virtual); and the 2021 American Fisheries Society meeting in Baltimore, MD.

Thank you for your time and consideration. I look forward to your response.

Sincerely,

Teresa G. Schwemmer, Ph.D.  
Corresponding Author

Highlights for Schwemmer et al.

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

### **Highlights**

- Bioenergetic mechanisms of Atlantic silverside hypoxia responses were investigated.
- Hypoxia effects were modeled with a simplified Dynamic Energy Budget model.
- We connected physiology with energetic processes to identify potential mechanisms.
- Conversion efficiency and mortality parameters best explained hypoxia effects.
- This mechanism could impact energy flow across generations and trophic levels.

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

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4     Teresa G. Schwemmer<sup>a,1</sup>, Roger M. Nisbet<sup>b</sup>, and Janet A. Nye<sup>c</sup>

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6     <sup>a</sup>School of Marine and Atmospheric Sciences, Stony Brook University, Stony Brook, NY 11794,  
7     U.S.A., teresa.schwemmer@gmail.com

8     <sup>b</sup>Department of Ecology, Evolution and Marine Biology, University of California Santa Barbara,  
9     Santa Barbara, CA 93106, U.S.A., rogernisbet@ucsb.edu

10    <sup>c</sup>Institute of Marine Sciences, University of North Carolina at Chapel Hill, Morehead City, NC  
11    28557, U.S.A., jnye@nyelab.org

12  
13    <sup>1</sup>Corresponding author, present affiliation: Teresa G. Schwemmer, Mid-Atlantic Coastal  
14    Acidification Network, Newark, DE 19716, U.S.A., teresa.schwemmer@gmail.com

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## Highlights

- Bioenergetic mechanisms of Atlantic silverside hypoxia responses were investigated.
- Hypoxia effects were modeled with a simplified Dynamic Energy Budget model.
- We connected physiology with energetic processes to identify potential mechanisms.
- Conversion efficiency and mortality parameters best explained hypoxia effects.
- This mechanism could impact energy flow across generations and trophic levels.

## Abstract

Ocean deoxygenation is intensifying worldwide due to warming and eutrophication, particularly in estuaries and coastal waters. Although the Atlantic silverside (*Menidia menidia*) is tolerant of the fluctuating environmental conditions in its estuarine habitat, 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 of Synthesizing Units, 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 the hypoxia effects to reduce the conversion efficiency of assimilates to structure accounted for some of the hypoxia-related changes in all three state variables. However, the best fit was achieved by simultaneously reducing the conversion efficiency and increasing both mortality parameters. In contrast, changing the parameter for maintenance rate with hypoxia provided little to no improvement of fit to the data. 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). In temperate estuaries, 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 episodic hypoxia (Farrell and Brauner, 2009; Zhu et al., 2013; Baumann, 2019), these are not necessarily adequate for tolerance of longer duration events. Fishes that spawn in the spring and summer may be particularly vulnerable because they are exposed to hypoxia during the 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 and/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 1/15<sup>th</sup> the ATP yield of aerobic respiration. Hypoxic exposure may lead to physiological responses such as depressed metabolism (Richards, 2009; Schwemmer, 2023),

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4 102 limited growth, increased ventilation, and changes to hematocrit, hemoglobin, and erythrocyte  
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6 103 quantities and characteristics (Taylor and Miller, 2001; Stierhoff et al., 2009; Bianchini and  
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8 104 Wright, 2013). Metabolism has also been shown to increase after temporary hypoxia as fish  
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10 105 remove lactate accumulated from anaerobic respiration (Heath and Pritchard, 1965).

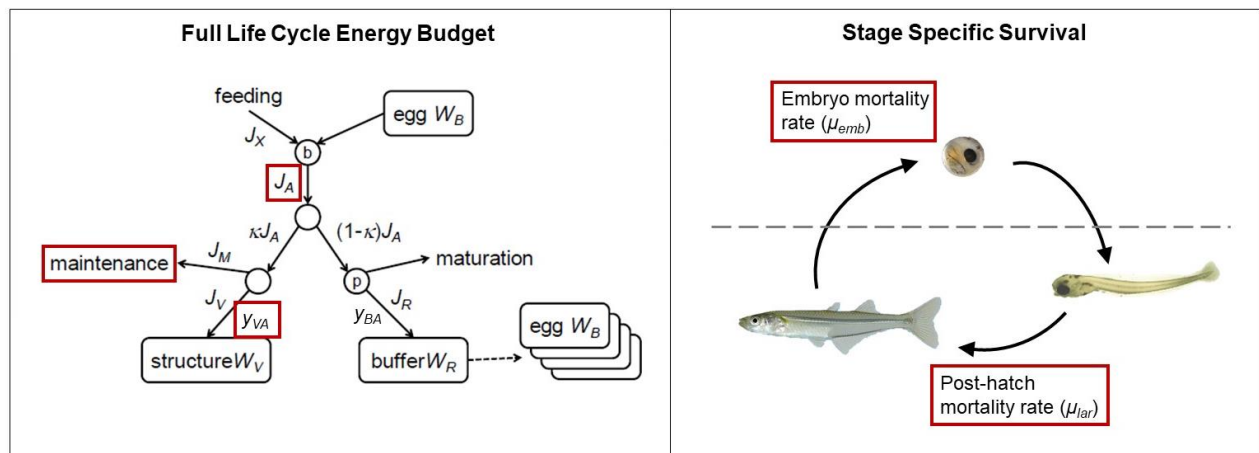
11  
12 106 Hypoxia often has interactive effects with other stressors such as temperature (Brandt et  
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14 107 al., 2009; McBryan et al., 2013; Earhart et al., 2022) and high carbon dioxide (CO<sub>2</sub>; Hancock and  
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16 108 Place, 2016; Miller et al., 2016; Morrell and Gobler, 2020). The Atlantic silverside (*Menidia*  
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18 109 *menidia*) is an estuarine forage fish that has frequently been used as a model species to  
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20 110 understand interactive effects of high CO<sub>2</sub> and hypoxia (DePasquale et al., 2015; Miller et al.,  
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22 111 2016; Murray and Baumann, 2018; Schwemmer et al., 2020). Rearing *M. menidia* offspring in  
23  
24 112 static low DO significantly delayed hatching, reduced survival to hatching and larval survival,  
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26 113 and reduced embryo and larval growth (Cross et al., 2019). However, the negative effects of both  
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28 114 hypoxia and high CO<sub>2</sub> were mitigated when offspring were exposed to diel-fluctuating  
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30 115 treatments rather than static (Cross et al., 2019). While diel fluctuations are a realistic  
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32 116 representation of current changes in community photosynthesis and respiration between day and  
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34 117 night, environmental change in coming years could extend hypoxic duration to reduce periods of  
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36 118 relief. Warming reduces oxygen solubility while increasing metabolic rates of organisms that  
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38 119 draw down oxygen when densely aggregated. At the same time, higher summer temperatures and  
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40 120 freshwater input in some regions will intensify stratification that separates low-oxygen water  
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42 121 from surface oxygen diffusion (Rabalais et al., 2009; Howarth et al., 2011). Currently *M.*  
43  
44 122 *menidia* is tolerant enough that population declines are not a concern, but without knowledge of  
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46 123 the mechanisms of early life impacts it is hard to anticipate whether this will change under  
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48 124 intensifying deoxygenation or with additional stressors (Baumann, 2019).

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50 125 Dynamic Energy Budget (DEB) theory is a bioenergetic framework designed to bridge  
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52 126 multiple levels of biological organization in assessing stressor effects and their mechanisms in a  
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54 127 vast variety of species (Kooijman, 2010a; AmP, 2023). This approach follows energy allocation,  
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56 128 represented in suborganismal metabolic fluxes, and how it leads to life history outcomes such as  
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58 129 growth rate, reproductive output, and survival, using physical and biological concepts that are  
59  
60 130 generalizable to most species (Jusup et al., 2017). It accounts for differences in the energy budget  
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62 131 at each stage to allow modeling of life stage transition timing and stage-specific responses to  
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64 132 stressors (Kooijman, 2010a). DEB theory is often used to connect experimental observations of  
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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 typically operate at this level, but this scaling requires additional steps and remains a challenge (but see Nisbet et al., 1989; Martin et al., 2013; Gear et al., 2020; Tai et al., 2021). 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 sometimes be beneficial (Evans et al., 2013), simple parameter-sparse models 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 explicit representation of reserve and assumes that assimilates are immediately allocated to structure, maintenance, and reproduction (Jager et al., 2013). This 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 makes it ideal for adaptation to many species of ecological or commercial value, even when the existing studies were not originally intended for this use.



**Figure 1. Conceptual diagram of the DEBkiss model highlighting parameters of interest for hypoxia effects.** The DEBkiss model (diagram adapted from Jager et al., 2013) used in this study includes 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 how the stage-specific survival modification is applied to embryos, larvae, and adults of *M. menidia*.

Romoli et al. (2024) present a detailed comparison of the advantages and limitations for ecological risk assessment of a model based on DEBkiss versus a model based on Kooijman's "standard" DEB model. The comparison focuses on application in ecotoxicology but the reasoning carries over to other forms of environmental stress, including this study on hypoxia. They highlight several "modeling choices" that should influence the choice of approach, including: (i) insufficient information in data sets; (ii) capturing differences between data sets for the same species; and (iii) auxiliary hypotheses. Consideration of each of these led us to choose DEBkiss for our work after much unsuccessful effort attempting to interpret parameter estimates with the standard model (using the Add-my-Pet software; AmPtool, 2022; Marques et al., 2018).

We used DEBkiss to test the hypothesis that changes in animal performance under hypoxia can be explained by changes in 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 the 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 (Muller et al 2019). SUs are generalized enzymes that produce products such as body structure or support maintenance requirements from incoming fluxes of substrate, i.e. food or egg buffer (Kooijman, 1998; Kooijman, 2010a). With single substrates for each life stage the SU formalism is equivalent to standard Michaelis-Menten kinetics, but the SU interpretation allowed us to exploit the subtleties in describing inhibition set out by 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 mechanism.

## 2. Methods

### 2.1. DEBkiss Model Description

The material flows are shown in Fig. 1. The yolk in an egg is treated as a buffer of “food” for the developing embryo that initially has an infinitesimally small structural biomass. There is no reserve compartment between food assimilation and its utilization. Hatching occurs when the egg buffer is fully depleted. After hatching, juveniles feed and assimilated food is allocated to growth and maintenance versus maturity or reproduction in accordance with the  $\kappa$ -rule.

The DEBkiss assumptions and equations 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, from the egg buffer ( $W_B$ ) is immediately converted to assimilates which are allocated to a somatic fraction ( $\kappa$ ) and a reproductive fraction ( $1-\kappa$ ; Figure 1). The assimilation flux ( $J_A$ ) is the product of the scaled measure of resource availability ( $f$ ), the volumetric surface area ( $L^2$ ), and the parameter maximum area-specific assimilation rate ( $J_{Am}^a$ ) where  $f=1$  for embryos and for post-hatching fish fed *ad libitum*. Within the somatic branch, a flux to maintenance ( $J_M$ ) is prioritized while the remainder goes to structural mass ( $J_V$ ) with a conversion efficiency  $y_{VA}$ . The maintenance flux is proportional to structure.

Parameter	Symbol	Fixed or estimated	Value
Max. area-specific assimilation rate	$J_{Am}^a$	Estimated	$0.333 \text{ mg mm}^{-2} \text{ d}^{-1}$
Max. volume-specific maintenance rate	$J_M^v$	Fixed	$0.0214 \text{ mg mm}^{-3} \text{ d}^{-1}$
Initial egg buffer mass	$W_{B0}$	Fixed	0.15 mg
Total length at puberty	$L_{vp}$	Fixed	102 mm
Yield of assimilates on volume	$y_{AV}$	Fixed	0.8
Yield of egg buffer on assimilates	$y_{BA}$	Fixed	0.95
Yield of structure on assimilates	$y_{VA}$	Estimated	0.365
Fraction of assimilates allocated to soma	$\kappa$	Fixed	0.8
Scaled food level	$f$	Fixed	1
Scaled food level for embryo	$f_B$	Fixed	1
Mortality rate for embryos	$\mu_{emb}$	Estimated	0.0639

Mortality rate for larvae	$\mu_{lar}$	Estimated	0.0294
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**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.

Flux	Symbol	Equation	Units
Assimilation flux	$J_A$	$J_A = f J_{Am}^a L^2$	mg day <sup>-1</sup>
Maintenance flux	$J_M$	$J_M = J_M^v L^3$	mg day <sup>-1</sup>
Flux to structural growth	$J_V$	$J_V = y_{VA} (\kappa J_A - J_M)$	mg day <sup>-1</sup>
Flux to reproduction buffer	$J_R$	$J_R = (1 - \kappa) J_A - J_J$ when $W_V \geq W_{Vp}$ $J_R = 0$ when $W_V < W_{Vp}$	mg day <sup>-1</sup>
Flux to maturity	$J_J$	$J_J = J_J^v L^3$ when $W_V < W_{Vp}$ $J_J = J_J^v L_{Vp}^3$ when $W_V \geq W_{Vp}$	mg day <sup>-1</sup>
State Variable	Symbol	Equation	Units
Structural dry mass over time	$W_V$	$\frac{dW_V}{dt} = J_V$	mg day <sup>-1</sup>
Continuous reproduction rate	$R$	$\frac{dR}{dt} = \frac{y_{BA} J_R}{W_{B0}}$	eggs day <sup>-1</sup>
Egg buffer (yolk) mass	$W_B$	$\frac{dW_B}{dt} = -J_A$	mg day <sup>-1</sup>
Survival	$S$	$\frac{dS}{dt} = -\mu_{emb} S$ when $W_B > 0$ $\frac{dS}{dt} = -\mu_{lar} S$ when $W_B = 0$	unitless (range 0-1)
Other variables and conversions	Symbol	Equation	Units
Total physical length	$L^M$	$L^M = \frac{L}{\delta_M}$	mm
Volumetric length	$L$	$L = \sqrt[3]{L^3}$	mm
Shape coefficient	$\delta_M$	$\delta_M = \frac{L}{L^M}$	unitless
Dry weight density of structure	$d_V$	$d_V = \frac{W_V}{L^3}$	mg mm <sup>-3</sup>
Dry mass at puberty	$W_{Vp}$	$W_{Vp} = d_V * (L_{Vp} * \delta_M)^3$	mg
Volume-specific maturity maintenance costs	$J_J^v$	-	mg mm <sup>-3</sup> day <sup>-1</sup>
Structural volume at puberty	$L_{Vp}^3$	-	mm <sup>-3</sup>

Scaled measure of resource availability	$f$	-	unitless (range 0-1)
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**Table 2. Model definition.** Fluxes, state variables, and differential equations in the DEBkiss model.

For juveniles, the non-somatic fraction of assimilates is spent on maturation, or increasing complexity. 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, 2010b; Jager et al., 2013), so the maturity variable plays no role in the current work. Once the mass at puberty is reached ( $W_{VP}$ ), reproductive flux ( $J_R$ ) toward egg production begins in adults with a conversion efficiency  $y_{BA}$ . 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.

Because our growth data are in total length, we used a shape correction coefficient ( $\delta_M$ ) and dry weight density ( $d_V$ ) to connect length with the model state variables (Table 2). We calculated these constants using data on *M. menidia* length and egg volume (Klahre, 1997) and a total length ( $L^M$ ) to dry weight ( $W_V$ ) conversion (H. Baumann, personal communication):

$$W_V = e^{(2.997 \cdot \ln(L^M) - 6.7)} \quad (1)$$

After calculating  $W_V$  from  $L^M = 5.3$  mm at hatching (Cross et al., 2019), we obtained a dry weight at hatching of 0.18 mg. Assuming there is negligible change in weight or volume during hatching, we used the volume of an egg immediately before hatching,  $L^3 = 0.45$  mm<sup>3</sup>, to calculate  $d_V$  using:

$$d_V = \frac{W_V}{L^3} \quad (2)$$

This gave us  $d_V = 0.4$  mg mm<sup>-3</sup>. We similarly used the egg volume to calculate volumetric length of an embryo as  $L = 0.77$  mm, which gives us a  $\delta_M$  of 0.145 using the following equation:

$$\delta_M = \frac{L}{L^M} \quad (3)$$

However, this value led the model to underestimate total length later in the life span, suggesting the  $\delta_M$  value was too high for this long and slender fish. We manually adjusted  $\delta_M$  to a final value of 0.107 which provided a reasonable fit to length data and a better starting point for parameter estimation.

We added a survival state variable ( $S$ ) which, in addition to allowing an alternative outcome to hatching, enabled us to model survival as a consequence of hypoxia effects on the energy budget. We fit mortality parameters for embryos and post-hatch fish ( $\mu_{emb}$  and  $\mu_{lar}$ ) 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.

## 2.2. 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: total length over time, egg buffer mass over time (extrapolated from knowledge of egg mass and age at hatching), cumulative egg production over time, and proportion surviving since fertilization over time. As described in the introduction, the data available for this model led us to use DEBkiss over the “standard” DEB model based on the factors highlighted by Romoli et al. (2024). We had insufficient data, had to integrate information from multiple studies of the same (and similar) species, and had to hypothesize plausible values for a few parameters.

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 described in Section 2.3 and modeling three reduced oxygen treatments (Section 2.5 and 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 CO<sub>2</sub> 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 (via 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). The control data from these experiments were used to estimate parameters under normoxia (Section 2.3). We also obtained normoxic survival data from a study on the effects of temperature and CO<sub>2</sub> on *M. menidia* early life survival, using only the 24°C and control CO<sub>2</sub> 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 CO<sub>2</sub> 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 CO<sub>2</sub> treatments and strip-spawned once they reached reproductive maturity.

### 2.3. Parameter estimation under normoxia

We estimated four parameters by fitting them to full-lifespan data listed in Section 2.2 ( $J^a_{Am}$ ,  $y_{VA}$ ,  $\mu_{emb}$ , and  $\mu_{lar}$ ), calculated four parameters from data ( $J^a_M$ ,  $W_{B0}$ ,  $L_{Vp}$ , and  $f$ ), and fixed at suggested values parameters for which we had insufficient data to calculate or estimate ( $y_{AV}$ ,  $y_{BA}$ ,  $\kappa$ , and  $f_B$ ; Jager, 2018). The primary parameters and their calculated or estimated values are found in Table 1. Fitting was done in Matlab with the platform BYOM v.6.4 and the package DEBkiss v.2.3a (<https://www.debtox.info/byom.html>). 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 estimate model parameters based on their effect on the DEBkiss equations and the auxiliary equations. The differential equations predict length, egg production, egg buffer mass, and survival over time with the differences from observations used to calculate NLL.

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 we calculated from data on *M. menidia*, as described below. We visually assessed fit and noted the NLL calculated from each simulation 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. Furthermore, we were able to obtain a reasonable fit using suggested values for  $y_{AV}$ ,  $y_{BA}$ , and  $\kappa$  for unstressed fish that are thought to be widely applicable across species (Lika et al., 2011; Jager, 2018). The suggested value for  $y_{VA}$  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 re-estimated with the BYOM optimization. Ultimate length was used to fit  $J^a_{Am}$  to a reasonable value while fixing all other parameters before estimating  $y_{VA}$ , because both parameters affect growth and egg buffer depletion in the model and therefore cannot be estimated simultaneously. Finally, we used the BYOM optimization to estimate  $\mu_{emb}$  and  $\mu_{lar}$ .

The length and reproductive data allowed us to calculate “length at puberty” ( $L_{VP}$ ), defined as the length at which egg production begins. We obtained  $W_{B0}$  from *M. menidia* egg dry weight data (Klahre, 1997) and calculated  $\delta_M$  and  $d_V$  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 ( $J^V_M$ ), 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 CO<sub>2</sub> levels, we only used data from control groups to avoid potential CO<sub>2</sub> effects in the data.

#### 2.4. 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 of or damage to a SU. 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<sup>-1</sup> (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<sup>-1</sup>; Schwemmer et al., 2020). We therefore considered a single-substrate growth SU in which food or egg buffer was the substrate. The mathematical characterization of inhibition and damage is in Section 2.5.

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). The idea is that hypoxia induces the production of compounds that in turn bind to SUs. We used existing information on the physiological responses of fish early life stages to hypoxia to identify the following candidate 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. Hypoxia effects on growth and hatching time can occur either through inhibition of assimilation or through damage that reduces the conversion efficiency of assimilates to growth. Hypoxia impact on survival may be caused directly through damage or by inhibition of damage repair processes. Hypoxia impact on somatic maintenance rate may be most plausibly represented as damage. We now discuss how these abstract concepts, damage and inhibition, may relate to observations on hypoxia.

Assimilation is the transformation of food (or yolk) and oxygen into compounds that will go to structure, maintenance, or reproduction (Kooijman, 2010a; Jager, 2018). Reduced food consumption or reduced conversion of food into utilizable compounds, and thus limitation of input of substrate to 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 with findings reinterpreted for DEBkiss).

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 has been 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 for other oxygen-demanding processes 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, though the link to survival is not represented explicitly.

The conversion efficiency of assimilates to structure determines growth and hatch timing because it represents 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 also be responsible for reductions in conversion efficiency through lactate accumulation and consequential declines in internal pH. Even at oxygen levels above the critical level at which oxygen consumption declines, anaerobic glycolysis may increase (Nonnotte et al., 1993; Maxime et al., 2000; Wood et al., 2018). 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, 2010a) 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 CO<sub>2</sub> and lactate (Perry et al., 2009; Heath and Pritchard, 1965), but these capabilities may be limited until development has progressed further.

The maintenance flux in DEBkiss is represented in units of assimilated carbon required to meet the energy demand from maintenance. It therefore is only indirectly related to respiration rates measured as oxygen consumption. For example, an increase in the abstract maintenance parameter in the model could be caused by an increase in anaerobic processes without impacting oxygen consumption. The *measured* oxygen consumption rates of *M. menidia* early life stages did not significantly increase under experimental chronic hypoxia, but great variability in metabolic rates among individuals combined with the short respirometry periods used (<1 hour)

may make small increases related to damage repair or activity difficult to detect (Schwemmer et al., 2020).

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 directly through increase in damage production or inhibition of damage 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.

## 2.5. 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 inhibition or damage linked to 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.

Variable	7.7 mg L <sup>-1</sup>	4.2 mg L <sup>-1</sup>	3.1 mg L <sup>-1</sup>	2.7 mg L <sup>-1</sup>
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 3. Summary of experimental data for each DO level.** 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 DO treatments in Cross et al. (2019). The control DO level means (7.7 mg l<sup>-1</sup>) also include data from Murray and Baumann (2018).

We derived a correction factor for *inhibition* 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 well-suited to this study because of the limitations 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:

$$J_A = f J_{Am}^a L^2 \left( \frac{1}{1 + \frac{j_i}{k_i}} \right) \quad (2)$$

where  $j_i$  is the arrival flux of the inhibitor and  $k_i$  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  $j_i$  to depend on DO treatment above a DO threshold,  $DO_c$ , below which  $j_i$  is infinitely large, which would bring the rate of the process it is inhibiting to zero:

$$j_i = \begin{cases} \infty & \text{if } DO \leq DO_c \\ \frac{1}{Z(DO - DO_c)} & \text{if } DO > DO_c \end{cases} \quad (3)$$

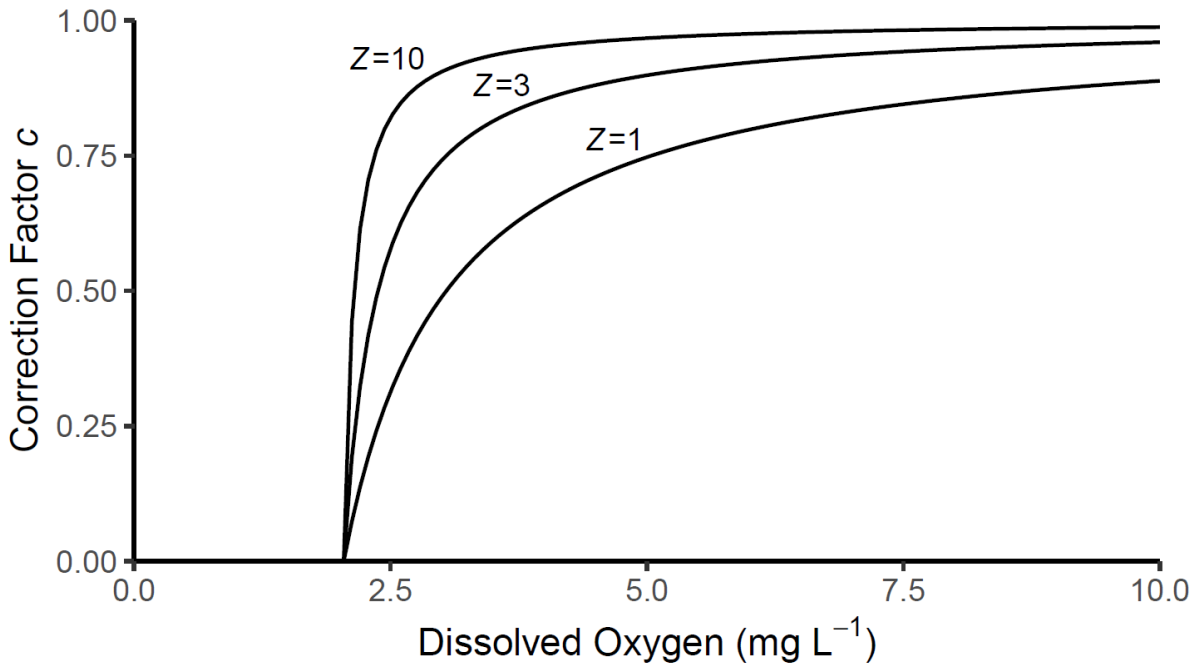
$Z$  is a parameter that influences the shape of the relationship between  $j_i$  and DO. We defined the correction factor  $c$  as the inhibition term (in parentheses in Equation 2) and replace  $j_i$  with the function from Equation 3 for  $DO > DO_c$  to derive the correction factor  $c$ :

$$c = \frac{1}{1 + \frac{1}{k_i Z (DO - DO_c)}} \quad (4)$$

As only the product of the parameters  $k_i$  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  $DO \leq DO_c$  gives us the following correction factor:

$$c = \begin{cases} 0 & \text{if } DO \leq DO_c \\ \frac{Z(DO - DO_c)}{1 + Z(DO - DO_c)} & \text{if } DO > DO_c \end{cases} \quad (5)$$

The relationship between  $c$  and  $DO$  for three different sample values of  $Z$ , the parameter to be estimated, is shown in Figure 2. 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.  $DO_c$  was fixed at a biologically relevant level of  $2.04 \text{ mg L}^{-1}$ , which is the critical oxygen level below which embryonic routine metabolism becomes highly oxygen-dependent (Schwemmer, 2023). Attempts to estimate  $DO_c$  and  $Z$  simultaneously showed that leaving  $DO_c$  free did not improve the ability of the correction factor to fit the hypoxia data.



**Figure 2. The correction factor  $c$  used to apply hypoxia effects to DEBkiss parameters.** The effect of  $DO$  on correction factor  $c$  is shown 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  $c$ .

Similar simplification of the reasoning by Muller et al. (2019) can be used to derive an analogous correction factor for *damage*. Assuming that the proportional change in the rate of damage production to the SU (e.g. via “damage inducing compounds”; Kooijman 2010a)),  $j_d$  has the same form as Equation (3). If damage production is quickly balanced by repair or mitigation, then fluxes that decrease through hypoxia will again be reduced by the factor given by Equation (5). This was recognized by Muller et al. (2019) who noted that if damage production is much slower than the maximum production rate of an SU, the formalism for noncompetitive damage is equivalent to that of noncompetitive inhibition (Muller et al., 2019). Further submodels relating damage to rates that may increase in response to hypoxia (e.g. maintenance and mortality) are needed to derive functional forms for appropriate conversion factors here. Absent information to support such submodels, we hypothesize that the increase was inversely proportional to  $c$  defined by Equation (5).

The correction factor  $c$  was multiplied by  $J_{Am}^a$  and  $y_{VA}$  because these parameters were hypothesized to decrease under hypoxia irrespective of the underlying cause (inhibition or damage). Reductions in the parameter  $y_{VA}$  through hypoxia are most plausibly interpreted as damage, the irreversible destruction of functionality of an SU. However, 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  $J_{Am}^a$  and  $y_{VA}$  simultaneously because they are multiplied together to obtain  $J_v$  and their individual contributions to the growth and egg buffer depletion cannot 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  $J^a_{Am}$  or  $y_{VA}$  is in each candidate model, because  $J^v_M$ ,  $\mu_{emb}$ , and  $\mu_{lar}$  do not affect egg buffer depletion.

To identify the most likely version of the model (which parameter or combination of parameters best explains 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 ( $\Delta AICc$ ) and the relative likelihood of each model using Akaike weights:

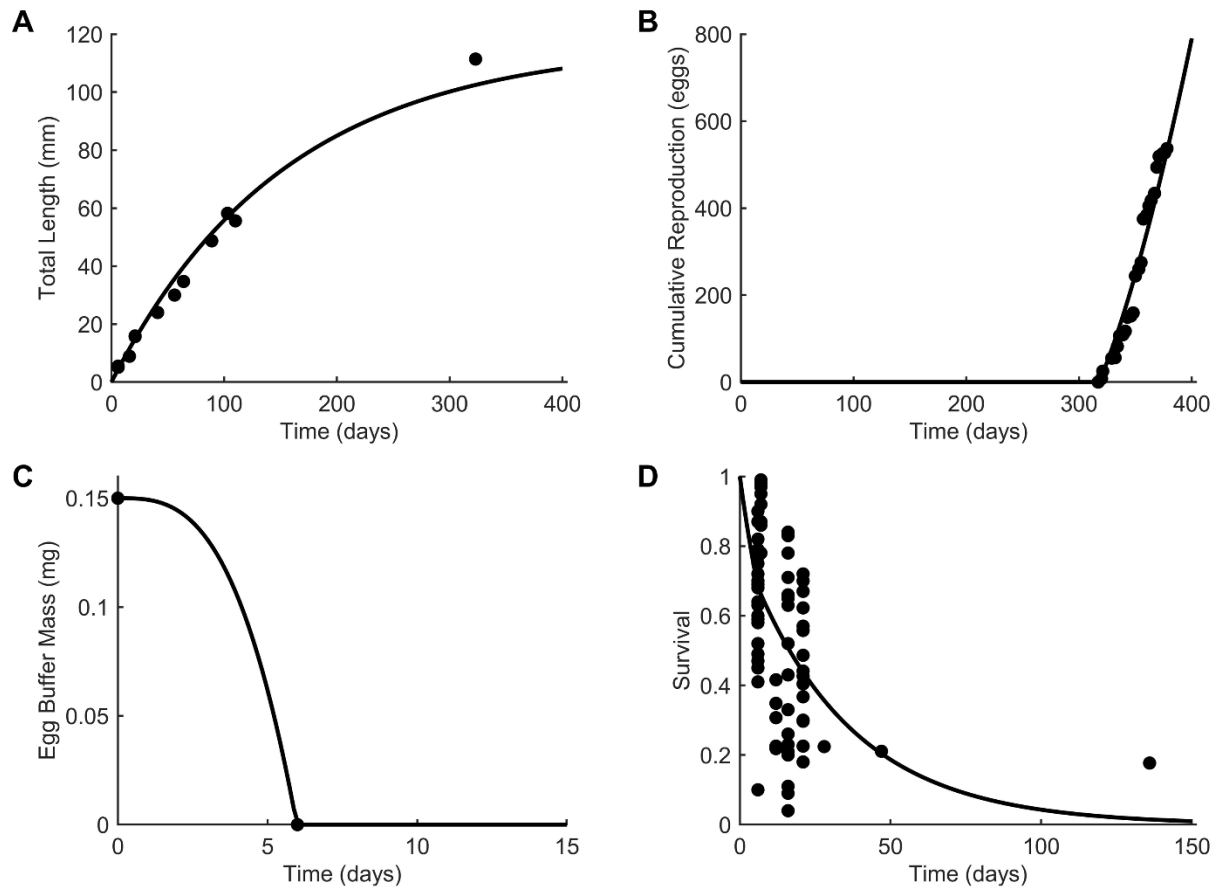
$$w_i(AICc) = e^{-0.5 \cdot \Delta_i AICc} / \sum_{k=1}^K e^{-0.5 \cdot \Delta_k AICc}, \quad (6)$$

where  $w_i(AICc)$  is the Akaike weight of each model  $i$ ,  $\Delta_i AICc$  is the difference between each model  $i$  and the model with the lowest AICc ( $AICc_{min}$ ), and the denominator calculates the sum of relative likelihoods for every model starting at the first model  $k$  (Wagenmakers and Farrell, 2004). We used  $\Delta AICc$  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).

### 3. Results

#### 3.1. DEBkiss Model

We obtained realistic fits to the full life cycle data (Figure 3). 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 (Figure 3D). Silversides are an annual species so survival should be greater than 0% after 150 days. However, this did not impair our ability to model the effects of hypoxia on early life survival, which is most important given that the present study focuses on hypoxia in the early life stages. Estimating  $y_{VA}$  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  $y_{VA} = 0.365$  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 are plotted in Figure 3.



**Figure 3. Full life model fits to data for four state variables.** Predicted (lines) and observed data (dots) for the DEBkiss model of *M. menidia* are shown. 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.

### 3.2. Hypoxia Effects

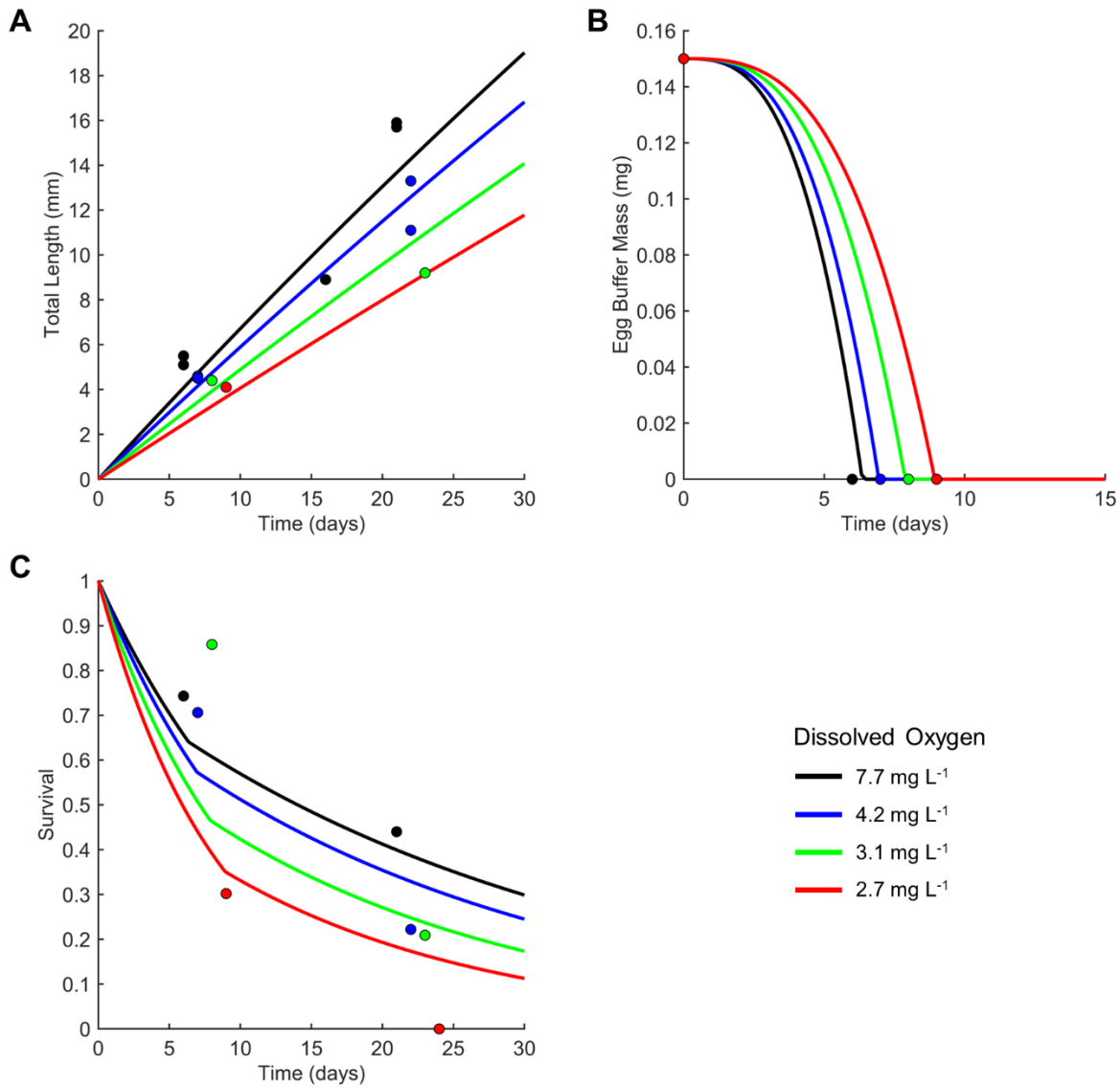
Applying the oxygen-dependent correction factor to the parameter combinations listed in Table 4 reproduced the direction of experimentally observed hypoxia effects, e.g. increasing  $J^a_{Am}$  reduced total length, increased time until egg buffer mass reaches 0, and reduced survival. The best model of experimental hypoxia effects on *M. menidia* early life stages simultaneously had  $y_{VA}$  multiplied by  $c$ , and  $\mu_{emb}$  and  $\mu_{lar}$  divided by  $c$  (Figure 4, Table 4, Figure S1). Although



applying damage to  $y_{VA}$  alone affected all three state variables, concurrently increasing both mortality parameters improved the fit to the data (Table 4). The model in which the correction factor was applied to  $y_{VA}$ ,  $\mu_{emb}$ , and  $\mu_{lar}$  also had the lowest AICc of all candidate models, with an AICc of 794.03 (AICc<sub>min</sub>). Adding a correction factor to  $J^v_M$  in simultaneously with these three parameters yielded a slightly higher AICc of 795.97 (Table 4). The ratio of Akaike weights shows that the model with  $c$  applied to  $y_{VA}$ ,  $\mu_{emb}$ , and  $\mu_{lar}$ , is 2.67 times as likely as the one with  $c$  concurrently applied to  $J^v_M$  (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  $y_{VA}$ ,  $\mu_{emb}$ , and  $\mu_{lar}$  when their respective correction factors are applied for each DO level (Table 5).

Parameter(s) affected by hypoxia correction factor	Estimated Z [95% CI]	AICc	$\Delta$ AICc	Akaike weight
$J^a_{Am}$	3.019 [2.512-3.612]	856.06	62.03	2.5e-14
$y_{VA}$	1.818 [1.601-2.342]	848.65	54.62	1.0e-12
$J^a_{Am} + J^v_M$	3.105 [2.651-3.726]	855.00	60.97	4.2e-14
$y_{VA} + J^v_M$	1.985 [1.688-2.774]	850.64	56.61	3.7e-13
$J^a_{Am} + \mu_{emb}$	2.804 [1.605-3.287]	823.24	29.21	3.3e-7
$y_{VA} + \mu_{emb}$	1.801 [1.570-2.167]	808.12	14.09	6.3e-4
$J^a_{Am} + \mu_{lar}$	2.930 [2.165-3.428]	838.17	44.14	1.9e-10
$y_{VA} + \mu_{lar}$	1.767 [1.536-2.111]	821.30	27.27	8.7e-7
$J^a_{Am} + \mu_{emb} + \mu_{lar}$	2.819 [1.920-3.286]	810.21	16.18	2.2e-4
<b><math>y_{VA} + \mu_{emb} + \mu_{lar}</math></b>	<b>1.827 [1.620-2.269]</b>	<b>794.03</b>	<b>0</b>	<b>0.72</b>
$J^a_{Am} + J^v_M + \mu_{emb} + \mu_{lar}$	2.913 [2.288, 3.387]	809.96	15.93	2.5e-4
$y_{VA} + J^v_M + \mu_{emb} + \mu_{lar}$	1.981 [1.700, 2.456]	795.97	1.94	0.27

**Table 4. Parameter Z estimates and model selection results.** The estimated Z value, AICc,  $\Delta$ AICc, and Akaike weights when the correction factors were applied to each parameter or combination of parameters.  $\Delta$ AICc and Akaike weights were calculated with AICc<sub>min</sub> = 794.03 for the  $y_{VA} + \mu_{emb} + \mu_{lar}$  model (bold).



**Figure 4. Best fit of DEBkiss model to experimental data from four DO levels.** The best fit of the predicted data (lines) to the observed data (dots) for four DO levels is shown, for early life data only. 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), 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.

Interestingly, although  $J^a_{Am}$  affects the variables similarly to  $y_{VA}$ , the ratio of Akaike weights showed that the best fitting model is about 3000 times as likely as the version applying

inhibition to  $J^a_{Am}$ ,  $\mu_{emb}$ , and  $\mu_{lar}$  (Table 4). Reducing  $J^a_{Am}$  with hypoxia using the correction factor resulted in a visually good fit to the data across oxygen levels and variables. Simultaneously applying  $c$  to  $J^a_{Am}$  and both mortality parameters improved the fit compared to only applying it to  $J^a_{Am}$ , but this model fit less well than the version that applied  $c$  to  $y_{VA}$ ,  $\mu_{emb}$ , and  $\mu_{lar}$ , 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  $y_{VA}$  at the lowest oxygen level is 55% of its value with no hypoxia stress. Reducing conversion efficiency alone 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.

	Product of correction factor and initial parameter value			
	7.7 mg L <sup>-1</sup>	4.2 mg L <sup>-1</sup>	3.1 mg L <sup>-1</sup>	2.7 mg L <sup>-1</sup>
$y_{VA}$	0.333 [0.329, 0.339]	0.291 [0.284, 0.303]	0.240 [0.230, 0.257]	0.199 [0.188, 0.218]
$\mu_{emb}$	0.0701 [0.0689, 0.0709]	0.0801 [0.0770, 0.0822]	0.0970 [0.0906, 0.101]	0.117 [0.107, 0.124]
$\mu_{lar}$	0.0322 [0.0317, 0.0326]	0.0369 [0.0354, 0.0378]	0.0446 [0.0417, 0.0466]	0.0539 [0.0492, 0.0571]

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

#### 4. 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 ( $y_{VA}$ ) and increase pre- and post-hatching mortality rates ( $\mu_{emb}$  and  $\mu_{lar}$ ) 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 with which assimilated food or egg yolk is converted into structure. The limitations of this inference are discussed later.

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). In *M. menidia*, however, 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 most plausibly interpreted as 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, 2010a; 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<sup>-1</sup> and 1.56 mg L<sup>-1</sup> 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

may be a mechanism of hypoxia effects in *M. menidia* early life stages even at oxygen levels above  $P_{crit}$ .

While  $y_{VA}$  is the best parameter to explain the hypoxia effects according to our model and AICc, it is nonetheless possible that  $J^a_{Am}$  is responsible for an unknown portion of the hypoxia effects. Because of near collinearity between  $J^a_{Am}$  and  $y_{VA}$ , our model does not allow us to test for the possibility that both parameters are simultaneously 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. We therefore cannot test for partial contribution of the two parameters to hypoxia effects or quantify their relative contributions. 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  $y_{VA}$  reduces the body size at a given time, indirectly reducing the assimilation flux due to smaller body volume.

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. Simultaneously 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<sup>-1</sup>) 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  $\Delta AICc$  (Table 4). The additional mortality that was not accounted for by  $y_{VA}$  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<sup>-1</sup> 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  $y_{VA}$ ,  $\mu_{emb}$ , and  $\mu_{lar}$  did not substantially improve the fit according to AICc, suggesting that increasing maintenance costs is not an important 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), but as noted earlier interpretation of respiration data is challenging and there was high individual variability in the data. 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 is 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 fish 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 control group, 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 in aquatic ectotherms 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 fish 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.  $J^a_{Am}$ ,  $J^v_M$ ,  $y_{VA}$ ) and similarly that values of the hypoxia correction factors are the same regardless of life stage. We lacked data on the effects of hypoxia on the proportion of total energy allocated to reproduction ( $1-\kappa$ ), which is an additional component of DEB useful in connecting organismal effects to populations. Future experimentation could provide the adult-stage information 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), given the ecological importance of forage fishes and the value of model species like *M. menidia*.

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 and this assumption may lead to overestimation of hypoxia effects. Studies comparing fish responses to static and fluctuating hypoxia treatments have suggested that fluctuations provided temporary relief and reduced sensitivity (Cross et al., 2019; Williams et al., 2019; Wang et al., 2021), although conflicting results also exist (Morrell and Gobler, 2020). It is also unrealistic for only a single environmental factor, in this case hypoxia, to influence the energy budget. Other studies have applied correction factors to DEB parameters to model other species' responses to hypoxia (Lavaud et al., 2019; Aguirre-Velarde et al., 2019), seawater acidification (Jager et al., 2016; Moreira et al., 2022; 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 magnitudes 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; Boulton and Evans, 2021). In the case of *M. menidia*, previous work showed that high CO<sub>2</sub> 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 induced by acidification.

Our best fitting model overestimated time to hatching at 7.7 mg L<sup>-1</sup> DO and overestimated survival at age for the 2.7 and 4.2 mg L<sup>-1</sup> 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  $\kappa$  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.

We end with a comment on the limitation of the “DEBtox” approach (Kooijman et al., 2009), a toxicology application of DEB from which DEBkiss stems, to identifying physiological modes of action in response to environmental stress. In Section 1 we cite the paper by Romoli et al. (2024) that highlighted the difficult modeling choices that are required. Here we chose to use DEBkiss coupled with several hypothesized responses to hypoxia. We selected the combination of DEB model and response hypothesis that best described given data (in an information theoretic sense using AICc), *conditional on the “correctness” of the model and of assumed values for some parameters*. Yet, for a case study in ecotoxicology, Romoli et al. showed that a different dominant physiological model of action was selected when using two different underlying DEB models that both give visually good fits to control data. Muller et al. (2010) demonstrated a related issue for a study of early life stage growth by identifying best fit submodel for larval growth of two closely related bivalve species exposed to mercury. Implausibly, the selected submodels were different to the extent that the best fit for one species was the worst for the other.

In the preceding discussion, we have offered a few suggestions for empirical work on whole organisms that would significantly help narrow down the DEB processes responsible for responses to hypoxia. However, it is likely that an additional, very promising way forward is to determine *suborganismal* processes co-occurring with the observed whole-organism responses. Transcriptomic data represent a particularly promising candidate (Murphy et al 2018). We recognized this qualitatively in Section 2.4 when invoking genes controlling cell division and protein synthesis that are regulated by hypoxia-inducible factors. Stevenson et al. (2023) demonstrated the power of transcriptomic data in a study of killifish embryos exposed to a toxicant. The molecular data helped to identify damage mechanisms that in turn led to changes in DEB parameters. There are many further exciting possibilities for integrating suborganismal (molecular) data with DEBtox modeling.

## 5. Conclusions

With this simple and widely applicable DEBkiss model we were able to attribute hypoxia-related variability in *M. menidia* growth, hatch timing, and survival to damage-induced reductions in conversion efficiency of assimilates into structure. Applying hypoxia corrections simultaneously to conversion efficiency and the mortality parameters for embryos and larvae provided the best fit, suggesting that hypoxia leads to both wasted energy and damage that cannot be sufficiently repaired in the early life stages. As lifelong, constant oxygen conditions are unrealistic in nature, the patterns modeled in this study should not be interpreted as a standalone prediction of what will happen to wild *M. menidia* populations as coastal hypoxia intensifies. Instead, this approach demonstrates the value of identifying energetic processes 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 support the utility of modeling inhibition and damage to synthesizing units and highlight conversion efficiency of food into growth as a primary mechanism by which hypoxia impacts an ecologically important forage fish and model species. Measuring suborganismal processes to identify physiological modes of action can refine this model so that it can better model this species' response to realistic hypoxia scenarios and, ultimately, how reductions in conversion efficiency could affect energy flow through food webs.

## 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., R.M.N.; Writing, reviewing and editing – T.G.S., R.M.N.,  
J.A.N.

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## **Data and Code**

The datasets used for modeling can be found on BCO-DMO: early life total length, survival, and  
hatching: doi: 10.1575/1912/bco-dmo.742200; early life total length with oxygen treatments: doi:  
10.1575/1912/bco-dmo.777130.1; hatching and survival with oxygen treatments: doi:  
10.1575/1912/bco-dmo.777117.1; total length of adults: doi: 10.26008/1912/bco-dmo.845906.1;  
total length of larvae and juveniles: doi: 10.1575/1912/bco-dmo.652124; egg production: doi:  
10.26008/1912/bco-dmo.845633.1. The BYOM and DEBkiss packages can be found at  
<https://www.debtox.info/byom.html>. The code for inputting data, parameter estimation, and  
plotting, for both the normoxic model and the model with hypoxia effects, can be found at  
[github.com/tschwemmer/MenidiaDEB](https://github.com/tschwemmer/MenidiaDEB).

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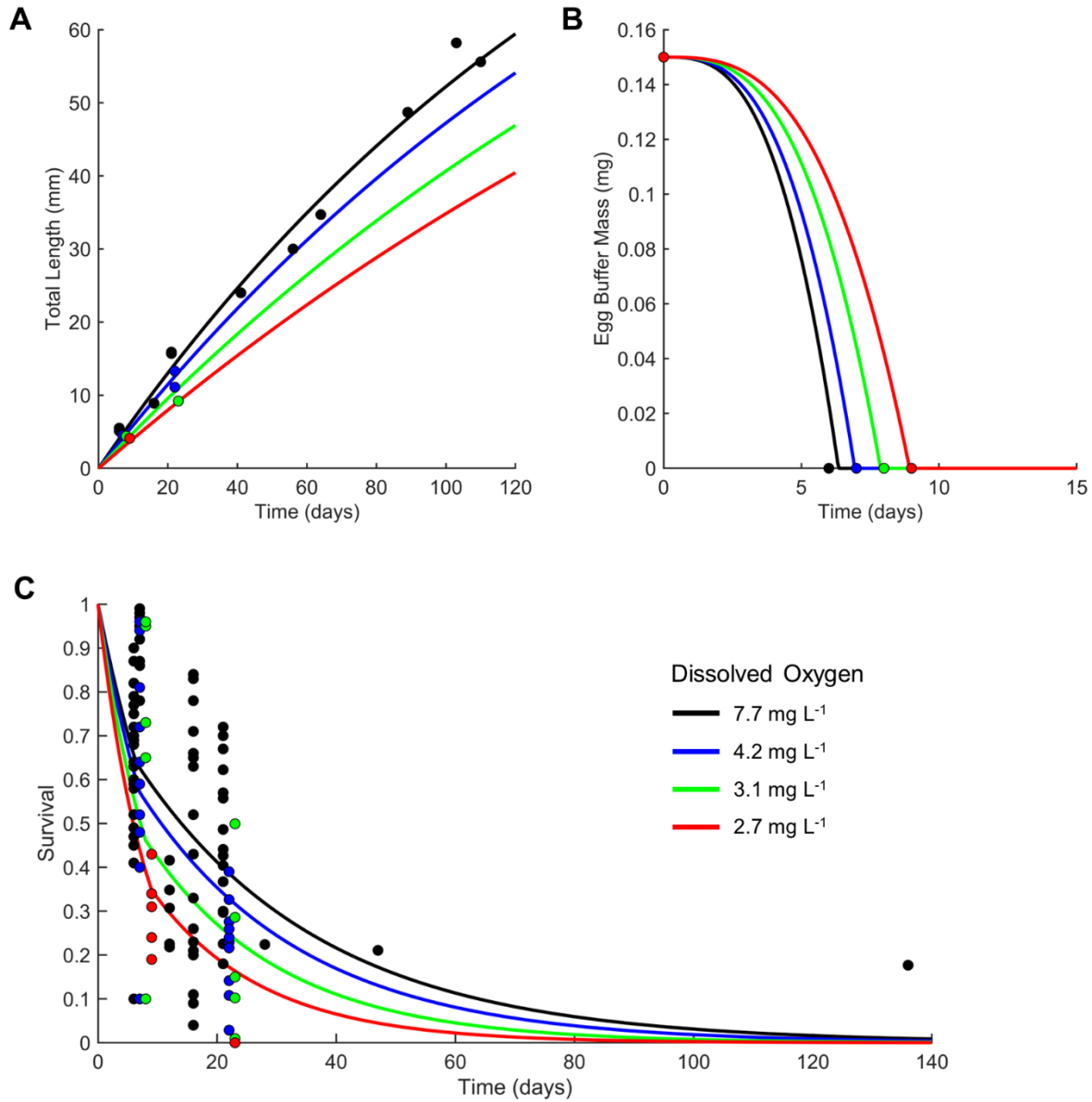
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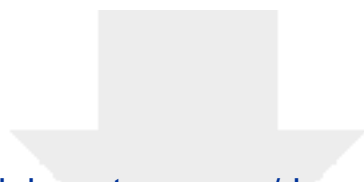
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## Supplementary Figure



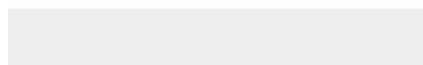
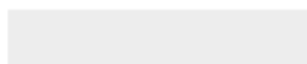
**Figure S1. Best fit of DEBkiss model to all experimental data from four DO levels.** The model was fitted to early life data (embryos, larvae, and juveniles) and the best fitting model was selected based on lowest AICc. (A) is total length (mm) over time (days), (B) is egg buffer mass (mg) over time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





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**Declaration of interests**

☒The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

☐The authors declare the following financial interests/personal relationships which may be considered as potential competing interests: