1 Intergenerational plasticity aligns with temperature-dependent selection on

2 offspring metabolic rates

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9 Abstract

- 10 Metabolic rates are linked to key life history traits that are thought to set the pace of life and 11 affect fitness, yet the role that parents may have in shaping the metabolism of their offspring 12 to enhance survival remains unclear. Here, we investigated the effect of temperature (24 °C or 30 °C) and feeding frequency experienced by parent zebrafish (Danio rerio) on offspring 13 14 phenotypes and early survival at different developmental temperatures (24 °C or 30 °C). We 15 found that embryo size was larger, but survival lower, in offspring from the parental low food 16 treatment. Parents exposed to the warmer temperature and lower food treatment also 17 produced offspring with lower standard metabolic rates – aligning with selection on embryo 18 metabolic rates. Lower metabolic rates were correlated with reduced developmental and 19 growth rates, suggesting selection for a slow pace of life. Our results show that 20 intergenerational effects on offspring size and metabolic rate can be adaptive when parent 21 and offspring temperatures are matched: the direction of selection on embryo size and 22 metabolism aligned with intergenerational plasticity towards lower metabolism at higher 23 temperatures, particularly in offspring from low condition parents. These findings highlight 24 the importance of anticipatory parental effects, but only when parental and offspring 25 environments match.
- 26

27 Keywords

28 Development, energy, maternal effects, metabolism, parental investment, reproductive

29 investment

30 Introduction

31 Selection on life history strategies can drive the evolution of metabolic rate, which represents 32 the energetic cost of living [1,2]. Metabolic rates expressed during early life are associated 33 with key life history traits: individuals with faster minimal metabolic rates have faster 34 developmental and growth rates, earlier onset of reproduction, and shorter lifespan, than 35 slow metabolic phenotypes [3,4]. The majority of ectotherms undergo embryonic 36 development in eggs, with a finite amount of energy reserves available to sustain cell 37 division, differentiation, and maintenance costs until post-hatching feeding [5]. Hence, 38 variation in metabolic rates will also determine how quickly energy reserves are depleted for 39 these species, with important consequences for survival [6]. It might be expected therefore, 40 that selection should act to suppress minimal rates of metabolism to conserve energy, yet 41 variation in metabolism is ubiquitous - varying by up to three-fold, even after accounting for 42 embryo size and developmental temperature [7]. Furthermore, selection for a fast pace-of-43 life may mediate the expression of higher metabolic rates [8]] that can be beneficial in high 44 competition environments [9]. Investigating the interplay between metabolic rates and 45 survival – and the environmental dependence of this relationship – is crucial for 46 understanding the potential adaptive capacity of variation in metabolic rates [10].

47

48 Metabolic rates have been studied for over a century [11], yet the adaptive potential of this 49 variation in metabolism remains unclear [12]. Mixed evidence shows that metabolism is 50 sometimes under selection (e.g., [13–15]) and is somewhat heritable [16–18] and repeatable 51 [19,20], suggesting that the fitness consequences of slow and fast metabolic rates are 52 context-dependent [21,22]. It is unresolved whether metabolism has evolved as a driver or 53 simply a by-product of the pace of life. However, metabolic rates (often measured as oxygen 54 consumption or carbon dioxide production) reflect the energy use of an organism, so that 55 measures of metabolic rate are meaningful in linking the physiology of an individual with its 56 life history. Metabolic rates are not fixed across ontogeny however, and within-generation 57 acclimation can act to down-regulate metabolism under low food availability [23]. While this 58 metabolic suppression may slow the pace of life, it can also facilitate survival under stressful 59 conditions [24]. If there is a causal relationship between metabolism and the pace of life, 60 then context-dependent selection may drive a correlated suite of responses [8]. Elucidating the links between metabolic rate, the pace-of-life, and its fitness consequences, is critical for 61 62 understanding the capacity for organisms to respond to changing environments [25]. 63

The environment a parent experiences can shape the phenotype of their offspring, known
generally as parental effects [26]. This form of epigenetic inheritance across two or more
generations (termed inter- and trans-generational plasticity respectively) [27] can be

67 adaptive or maladaptive – acting as either a buffer or conduit to the effects of environmental

- 68 stress [28]. Adaptive Adaptive parental effects arise when parents anticipate and respond to
- 69 environmental cues, to produce shifts in their offspring's phenotype that maximise their
- fitness in the offspring environment [26].[26]. For example, when exposed to cool
- temperatures, mothers tend to produce larger offspring [29], leading to enhanced offspring
- survival in that same environment [30,31]. Alternatively, under a bet hedging strategy,
- 73 parents in stressful or unpredictable environments increase variance in their offspring
- 74 phenotypes, with variable consequences for offspring fitness, but overall enhancing parental
- 75 fitness [32]. If intergenerationall fintergenerational plasticity is adaptive such that it confers
- 76 fitness benefits for offspring, then shifts in parental provisioning should be in line with
- selection on offspring traits. Conversely, increased variance in parental investment that does
- 78 not enhance offspring fitness consistently may be indicative of a bet-hedging strategy to
- 79 maximise parental fitness. Overall trends across studies show that
- 80 intergenerational intergenerational plasticity on offspring phenotype are generally weak
- compared with the direct effects of the offspring environment [33,34], and caution needs to
- be exercised when inferring the adaptive value of trait plasticity [35,36]. Nonetheless, the
- transmission of the parental environment across generations is an important source of
- 84 phenotypic variation. In particular, when environmental conditions are correlated between
- generations, maternal effects can account for up to half of the phenotypic variation within
- 86 populations as additive genetic effects [37,38].
- 87

88 Positive anticipatory effects are thought to evolve in changing but predictable environments 89 to enhance offspring fitness [39], however formal selection analyses are lacking. Selection is 90 the phenotypic covariance between fitness and a trait [40], yet most transgenerational 91 studies have reported the effect of parental environment on an aspect of offspring 92 performance that may trade off with actual fitness [41]. Selection analysis uses multiple 93 regression of individual relative fitness on traits of interest to estimate standardised linear 94 and nonlinear selection coefficients [42]. Used in combination with experimental manipulation of environmental predictability across generations, selection analysis can 95 96 reveal the relative scope for evolutionary change on an offspring trait. If parents can 97 anticipate the environment their offspring will experience, and provision accordingly, then 98 selection on offspring metabolic rates should align with shifts in offspring investment. 99 However, in cases where the offspring environment differs unpredictably from the parental 100 environment, the direction, form, and strength of selection may not align with the mean and 101 variance of offspring phenotypes that parents produce. Selection analysis cannot clarify 102 whether shifts in offspring phenotype in response to parental environment have evolved in 103 response to selection (i.e., whether they are due to genetic or epigenetic causes), however it does provide a meaningful first step to understanding whether transgenerational plasticity islikely to be adaptive in a given environment.

106

107 Food availability and environmental temperature experienced by the parental generation are 108 known to alter parental investment with performance consequences for subsequent 109 generations [30,31]. Poor parental condition may elicit an adaptive response in offspring via 110 transgenerational plasticity, and offspring from parents exposed to low food may suppress 111 their metabolic rate, or alter energy allocation towards maintenance or growth, to 112 compensate for lower energy provisioning from the mother. Alternatively, investment in 113 offspring can be the direct result of parental condition transfer effects, which can be 114 adaptive, but are not contingent on environmental predictability across generations [43]. 115 Regardless of the source of offspring trait variation, the implications of intergenerational 116 plasticity are likely to be context dependent. For example, warmer temperatures increase the 117 metabolic rates of ectotherms and may thereby exacerbate the fitness consequences of 118 variation in energy acquisition and allocation in low resource environments [44]. Food 119 availability in the parental generation is likely to alter maternal energy allocation (e.g., 120 offspring size and composition and/or number) towards offspring as well as mediate the 121 physiology of the offspring; the same is true for environmental temperatures in the case of 122 ectotherms. However, it remains so far unclear as to the direction of these responses, 123 whether they are under selection, and whether they constitute an adaptive parental strategy

- to maximise offspring fitness.
- 125

126 Despite evidence that metabolic rates are under selection, it is yet to be established whether 127 parents can modify the metabolic rates of their offspring in adaptive ways. Recent work on 128 ectotherms has shown evidence for both the presence [45,46] and absence [47] of 129 transgenerational responses of metabolic rates to temperature. However, offspring fitness in 130 these studies was measured indirectly as growth [46,47] or aerobic scope [45], which may 131 trade off with actual fitness, and in [45] some treatments showed extensive mortality, hence 132 results may be due to selective mortality. Formal tests of whether transgenerational plasticity 133 aligns with selection on offspring metabolic rates, via measures of offspring metabolism and 134 fitness under different environments, are currently lacking. Further, it is often unclear how 135 selection on metabolic rates may be mediated by its correlation with traits that set the pace 136 of life, such as developmental and growth rates. Here we manipulate parental food 137 availability and temperature in zebrafish to determine whether context-dependent selection 138 on offspring metabolic rates is in line with intergenerational plasticity on metabolic rates and 139 traits that set the pace of life. Here we manipulate parental food availability and temperature 140 in zebrafish to determine whether context-dependent selection on offspring metabolic rates

- is in line with intergenerational plasticity on metabolic rates and traits that set the pace of life.
- 142 We hypothesise that warm offspring temperatures will select for smaller embryo size and
- 143 lower metabolic rates, while selection at the cool (benign) offspring temperature will be
- 144 relaxed. Shifts in parental investment should mirror selection on offspring phenotype when
- their environments match thus parents in the warm environment should produce smaller
- 146 offspring with lower metabolic rates compared to parents from the cool temperature (Figure
- 147 1A). Further, we predict that parental effects should be exaggerated when parental food
- 148 availability is low, under which conditions parents should produce offspring with lower
- 149 metabolic rates than parents from the high food availability environment.

150 Materials and Methods

151 *Parent maintenance and treatments*

152 All procedures were approved by the University of Sydney Animal Ethics Committee 153 (protocol number: 2021/1932). Adult zebrafish were obtained from a commercial supplier 154 (Livefish, Childers, QLD, Australia), and housed in a controlled temperature room (22 °C 155 with 12L:12D). The supplier maintained zebrafish at 22-24°C in large communal ponds, and 156 so the parental fish were unlikely to have been closely related. The experiment was run in 157 two replicate blocks, one month apart. Within each block, fish were first allocated randomly 158 across four 35 I tanks (35-38 fish per tank) for two weeks to acclimate. Fish were then sexed as per [48], and 60 females and 60 males were allocated evenly across 12 experimental (11 159 160 I) tanks; each tank was filled with aged water and contained a sponge filter and a plastic 161 plant. We conducted four parental treatments (with three replicate tanks each) in a fully factorial design (Figure 1B). Parents were held at either 30 °C or 24 °C temperature, referred 162 163 to hereon as 'high' and 'low' parent temperature, respectively, and either a high feeding frequency (three times per day, five days per week) or low feeding frequency (once per day, 164 165 four days per week). Previous studies have shown that 30 °C is higher than optimal, and that 166 the low food regime was sufficient to allow growth but at a submaximal level [49,50]. The 24 167 °C treatment represents a relatively low but benign temperature previously shown to

168 facilitate normal growth [51].

169

170 To validate the feeding treatments used, measures of parent body mass and length taken at the end of the experiment were used to assess condition [52]. Parents were weighed (to the 171 172 nearest 0.001 g) and total body length (to the nearest 0.1 cm) measured, and the exponent 173 for the slope of In(length) and In(mass) calculated as 2.79. Measures of body condition were then calculated as mass/length^{2.79}. To maintain fish in stable temperature treatments, tanks 174 175 were held within water baths, containing three submersible heaters (Aqua One 200 W; 176 Techden, Sydney, Australia) and a powerhead water pump (Aqua One maxi). Temperature 177 loggers recording every 15 min were placed into two tanks per temperature treatment. Tanks were maintained within ±1.5 °C of their target temperature for the duration of the treatment. 178 179 Fish were fed flake food (5mg per fish; Supervit Fish Flakes, Tropical, Chorzów, Poland) [49] 180 at each feeding event according to the regime described above and at randomised times 181 between 8am – 8pm each feeding day. A 50% water change was conducted twice per week. 182 The adult food and temperature treatments were applied for eight weeks, after which adult 183 fish were bred.

184

185 The evening before breeding, all fish from each replicate tank were transferred into 10L 186 plastic breeding tanks containing a coarse mesh base, through which fertilised eggs could 187 pass to avoid being eaten by adults. Maintaining males and females in the same tank 188 promotes the release of pheromones that stimulate ovulation and oviposition in females and 189 spawning by males [53]. The next morningbreeding tanks were inspected within 1h post-190 fertilisation, and eggs were filtered through a sieve onto a petri dish containing buffered E3 191 medium as per standard procedure for embryo rearing [54]. Unfertilised eggs or dead 192 embryos were immediately removed.

193

194 Embryo and yolk size measurements, treatments, and rearing

195 Within one hour of collection from parental tanks, individual fertilised embryos were sampled 196 by sifting gently through a sieve, and then photographed under a dissecting microscope (x30 197 magnification; Leica S9D stereomicroscope with FLEXACAM C3 camera). Developmental 198 stages of *D. rerio* are easily identifiable due to the transparency of embryos. The sphere 199 stage shows a flat border between the blastodisc and yolk, and total embryo area and yolk 200 area were measured to the nearest μm^2 . The ratio of yolk area to total embryo area was 201 consistent among treatments (Figure S2). Hence, assuming density of embryo tissue did not 202 change with embryo size, we calculated embryo mass (μg) from embryo diameter at the 203 sphere stage using a relationship previously determined for *D. rerio* [50]. Embryos were then 204 placed individually into wells of 24-well culture plates containing E3 medium. For each of the 205 four parental treatment combination, 72 embryos were randomly allocated to each of two 206 offspring temperature treatments (24°C and 30°C), resulting in a total of 576 embryos 207 equally divided across 8 treatment groups: two parental temperatures (24 °C versus 30 °C) x 208 two parental conditions (low versus high) x two offspring temperatures (24 °C versus 30 °C) 209 [Figure 1B]. Offspring were maintained in incubators (Eurotherm Micro Digital Control Model 210 i-80, Steridium, Australia) on a 12L:12D light cycle for the remaining duration of the 211 experiment. Since offspring were removed from the parental treatment, photographed, and 212 placed into their treatment temperatures within three hours of fertilisation (approximately 2-213 4% of their total development time), we were able to separate the parental and offspring 214 environment.

215

216 Offspring metabolic rate measures

The rate of oxygen consumption ($\dot{V}O_2$) was measured as a common proxy for metabolic rate

218 (MR) of the offspring at three developmental stages: 1) 25% through embryonic

- 219 development (14 and 30 hours post-fertilisation (hpf) for embryos incubated at 30 °C and 24
- 220 °C, respectively, 2) 1-4 hours post-hatching (hph) and 3) one week post-hatching (wph),

221 hereon referred to as MR_{embryo}, MR_{hatch}, and MR_{larva}, respectively. Individual offspring of 222 known identification were photographed to measure diameter (MR_{embryo}) or length (MR_{hatch}, 223 and MR_{larva}) to the nearest μ m, then placed into individual 80 μ l (MR_{embrvo} and MR_{hatch}) or 500 224 μ l (MR_{larva}) glass vials containing Milli-Q water and a nonconsumptive O₂ sensor spot. We 225 used two 24-channel PreSens sensor dish readers (SDR2, PreSens, Germany), each with 226 24-chamber glass microplates (Loligo Systems Aps, Tjele, Denmark) to measure $\dot{V}O_2$ in 40 227 offspring and four blank vials simultaneously over a 2-h interval at their respective treatment 228 temperature (24 °C or 30 °C). For a detailed description of methods, see [50]. To calculate 229 the most linear rates of decrease in oxygen concentration within each timeseries dataset 230 (adjusted for background oxygen extraction), we used the RespR package, designed for processing closed chamber aquatic respirometry data in R [55]. Slopes were then converted 231 232 into rate of oxygen consumption, accounting for oxygen solubility of 5.91 ml O₂ at 24 °C and 5.29 ml O₂ at 30 °C (0 ppt salinity, STDP) [56]. 233

234

235 Offspring hatching time and survival measures

236 Eggs were held in their individual wells of the culture plates to allow recording of embryo 237 development time (time in hours from fertilisation until hatching; hpf) and survival; their water was changed daily using a solution of Milli-Q water with 0.5 g l⁻¹ of red sea salt at the 238 239 treatment temperature. Based on hatching time pilot data, we monitored embryos every two 240 hours from 30 hpf at 30 °C and 90 hpf at 24 °C, until all embryos were recorded as either 241 hatched or deceased. Within two hours of hatching, larvae were photographed for measures 242 of larval length (0 hph) and moved into larger 6-well culture plates filled with fresh water and 243 placed back into incubators at their respective treatment temperature. At four days post-244 hatching (dph), once feeding structures were fully formed, offspring were fed paramecium (4-245 5 dph), egg yolk (5-14 dph), flake food (5-14 dph), and Artemia sp. (from 15 dph) ad libitum. 246 Larvae were measured again at one week post hatchling (1wph) to obtain measures of 247 growth rate (mm day⁻¹ = (length at 1wph / length at 0hph) / 7). Larvae were monitored for 248 survival daily until two weeks post hatching. Sample sizes for all measures are provided in 249 Table S1.

250

251 Analysis of parent and offspring treatment effects on parent and offspring phenotypes

All analyses were conducted in R v4.2.3 [57]. Linear mixed effects models using the "Imer"

function within the *Ime4* package [58] were used to analyse the effect of parental

environment (feeding frequency and temperature) on parent body condition. The effect of

255 feeding frequency (low/high), temperature (24 °C/30 °C), and their interaction on body

256 condition was tested, as well as the random effect of "Tank ID" within block (three per

257 treatment). We also used linear mixed effects models to analyse the effect of parental 258 condition (low/high feeding frequency), parental temperature (24 °C/30 °C), offspring temperature (24 °C/30 °C), and all interactions, on offspring phenotypes. The significance of 259 260 parent "Tank ID" as a random effect was tested for all responses. We focussed on four key 261 offspring traits: 1) embryo mass (parental investment), 2) metabolic rates (MR_{embryo}, MR_{hatch}, 262 and MR_{larva}), 3) development time (time from fertilisation until hatching), and 4) growth rate 263 (length at two weeks post hatching divided by length at hatching). All candidate models for 264 offspring responses are provided in Table S2S2. Embryo mass (µg) was included as a covariate in metabolic rate and development time models (m2 and m3; Table S2). We used 265 266 embryo mass since we only had estimates of length and area for larvae at hatching and one 267 week post-hatching and have previously shown this to be an important indicator for hatch 268 and larval mass [59]. We used Akaike Information Criteria (AIC) for model ranking and 269 averaged models with Δ conditional AIC (AICc) <22 using the R package MuMin [60–62] 270 (Table S3), and the estimated marginal means from the best-fitting model were used for all 271 post-hoc comparisons using the emmeans package [63]] 272

273 Correlations between developmental and growth rates with metabolic rates

274 To explore within-individual associations among measures of developmental and growth 275 rates with metabolic rates, we ran repeated measures correlations using the package *rmcorr* 276 [64]. Using a repeated measures framework accounts for the non-independence of 277 observations measured on the same individuals. To explore within-individual associations 278 among measures of developmental and growth rates with metabolic rates, we ran repeated 279 measures correlations using the package rmcorr [64]. Using a repeated measures 280 framework accounts for the non-independence of observations measured on the same 281 individuals.

282

283 Selection analysis

284 We used a classic multiple regression approach derived from evolutionary theory to 285 characterise temperature-dependent selection acting on embryo metabolic rates, within each 286 parental environment [42]. This framework allows for standardised and comparable 287 estimates of both linear (β) and nonlinear (γ) selection coefficients. For each form, we 288 estimated the direction (sign of coefficients), and strength (magnitude of coefficients) of 289 selection acting on offspring mass and mass-independent metabolic rate (MIMR),), across 290 incubation temperatures, as per [42]. These measures have been used previously to provide 291 a more complete picture of the adaptive landscape for offspring metabolic rates [9]. 292

293 Fitness was measured as survival from fertilisation to two weeks post-hatching. This period 294 of life typically shows greatest mortality rates in egg-laying fish and is considered a 295 bottleneck to reproduction, and therefore fitness [65]. Survival was treated as binary data -296 offspring that survived to two weeks post hatching were assigned "1", whereas offspring that 297 died before two weeks were assigned "0". First, autocorrelation between traits was checked 298 to determine which traits should be included in the analysis. Metabolic rates at each 299 ontogenetic stage were significantly correlated (when embryo mass was included as a covariate; $F_{3,1724} = 3434$, p < 0.0001), particularly between MR_{embryo}, and MR_{hatch} ($r^2 = 0.71$). 300 We decided to use mass-independent metabolic rate (MIMR) since recent work has shown 301 302 that including both mass and metabolic rate in selection analyses can overestimate the 303 strength of selection on metabolic rates (Cameron and Marshall, In Review). Correlations 304 between embryo mass, MIMR_{embryo}, and MIMR_{larva} were relatively weak and variance inflation 305 factors were less than 5, hence both MIMR_{embryo}, and MIMR_{larva} were included, but MIMR_{hatch} 306 was excluded from the analysis. To prepare data for selection analysis, we followed the 307 method of [42]: first, within each combination of parent and offspring treatment, we 308 converted predictor variables of embryo mass, MIMR_{embryo}, and MIMR_{larva} into units of 309 standard deviation (mean of 0, standard deviation of 1), and divided each measure of 310 absolute fitness by mean absolute fitness to mean-centre survival.

311

312 Survival data were fitted using logistic regression in a generalised linear model using the 313 "glm" function. We ran a series of nested models to test for differences in linear and 314 nonlinear forms of selection. We first tested whether there were significant differences in selection among parental and offspring environments, via a sequential model fitting method 315 [66,67]. We then tested for significant interactions between selection (linear and nonlinear) 316 317 and environment (parental condition, parental temperature, and offspring temperature). 318 Since we only found significant interactions between selection and offspring temperature. 319 fitness data were mean-centred (see details above) within offspring temperature, and 320 selection coefficients were estimated for offspring incubated at 24 °C and 30 °C separately. 321 Selection coefficients from the logistic regression were transformed into linear estimates as 322 per [68]. Following [69], we doubled guadratic regression coefficients and their standard 323 errors before reporting selection gradients.

324

325 Results

326 Effects of parental environment on parent body condition and offspring size

- 327 Parents in the low feed treatment showed significantly lower body condition than individuals
- 328 within the high feed treatment (t = -6.44, df = 7.34, p < 0.001), however, there were no
- 329 differences in condition between high and low temperature treatments (Figure S1). Despite
- low body condition, parents held under the low feed frequency regime produced embryos
- that were heavier than those from high-condition parents (Table 1). Although there appeared
- to be a trend for heavier offspring from cool-reared parents (Figure 2A), there was no
- significant effect of parent temperature on embryo mass (Table 1). Table 1).
- 334

335 Effects of parental and offspring environments on offspring metabolic rates

336 Offspring reared at the high temperature treatment showed significantly higher metabolic

- rates than those reared at the low temperature treatment (Table 1, Figure 2B). We also
- found significant parental environment effects on offspring metabolic rates (Table 1, Figure
- 2B). Parents exposed to the high temperature or low food treatments produced offspring with
- lower metabolic rates at embryo, hatch, and larval stages. We also found a significant
- 341 interaction between embryo mass and parent feed frequency, where the slope between
- embryo metabolic rate and mass was steeper in offspring from low-feed frequency parents(Table 1, Figure 3).
- 344

345 Effects of parental and offspring environments on offspring developmental and growth rates 346 Offspring incubated at the low temperature took almost twice as long to develop than those 347 incubated at the warm temperature (Table 1, Figure 2C). More interestingly, development 348 time at a given offspring temperature was affected by the parental temperature as well as 349 the parental feed frequency, being extended in offspring from low-feed frequency or high-350 temperature parents; thus hatching was delayed by 9h on average when offspring reared at 351 the cool temperature came from low feed compared with high feed parents (Table S1, Figure 352 2C). We also found significant interactive effects between offspring temperature and parent 353 feed treatment, and between offspring temperature and parent temperature on development 354 time. High food treatment parents produced offspring that developed faster when reared at 355 the low offspring temperature, but not high offspring temperature treatment (Table 1, Figure 356 2C, t = -2.34, p = 0.088). Embryos developing at the cool treatment developed faster when 357 their parents were also from the cool temperature, relative to parents from the warm 358 temperature (Table 1, Figure 2C, t = -0.90, p = 0.807).

359

Larval growth rates during the second week post-hatching were faster in offspring from highfeed parents (Table 1, Figure 2D). We also found a significant three-way interaction between

- 362 parental temperature, parent condition, and offspring temperature for larval growth rate:
- 363 growth was slowest in offspring from the low offspring temperature treatment and when
- 364 parents were from both low feeding frequency and from the low temperature (Table 1).
- 365

366 Correlations between offspring traits

367 We found significant positive correlations between all metabolic rates (embryo, hatch, larval),

368 and between larval growth rates and these three metabolic rates (Figure 4). In contrast,

369 embryo development time was significantly negatively correlated with metabolic rates. We

- 370 found no significant correlation between embryo development time and larval growth rate
- 371 overall, however they were significantly negatively correlated within offspring temperature
- 372 treatments (Figure S3).
- 373

374 Effects of parental environment on offspring survival

375 Overall, we found that survival was lowest in offspring from parents in the low feed frequency 376 regime, but offspring and parent temperatures showed no effect on offspring survival to two 377 weeks post hatching (Table 1). Although parents in the low feed treatment produced larger 378 offspring, embryo mass did not itself predict survival.

379

380 Selection on offspring metabolic rates

381 Offspring from low-food parents showed greater survival when they had relatively low 382 embryo metabolic rates, as shown by significant negative directional selection (Table 2, 383 Figure 5E-H). Across all offspring high-temperature treatments, we found evidence for 384 negative directional selection on embryo metabolic rates (Table 2, Figure 5B,D,F,H). We 385 also found positive directional selection on offspring embryo mass when they were reared at 386 the low temperature from high-feed parents (P24HO24 and P30HO24; Figure 5A and 5C) or 387 they were reared at the high temperature but from high-feed and low temperature parents (P24HO30; Figure 5B). Conversely, we found negative directional selection on embryo mass 388 389 when offspring originated from parents reared at the high temperature and low food 390 treatments (P30LO24 and P30LO30; Figure 5G-H). There was also evidence for stabilising 391 selection on embryo metabolic rate in P30LO24 (Figure 5G), as shown by a significant 392 negative guadratic coefficient (Table 2). We found no significant directional selection on 393 larval metabolic rates, however there was significant positive correlational selection for 394 embryo and larva metabolic rates in P24LO30 (Table 2, Figure 5F), suggesting that 395 consistently lower metabolic rates were favoured in this environment...

396 Discussion

397 Intergenerational effects can be an important source of offspring phenotypic variation - here we provide evidence of adaptive intergenerational plasticity for offspring metabolic rates. We 398 399 found that low parental food availability negatively impacted offspring survival, but also 400 altered offspring metabolic phenotypes in a direction that aligned with selection on offspring 401 traits. The low feeding frequency treatment in our study produced low condition parents that 402 invested in larger offspring, compared with parents from the high feeding frequency 403 treatment. We also found that when parents were reared under either the warm (30 °C) 404 temperature, low feeding frequency treatment, or both, they produced offspring with lower 405 metabolic rates. Warm developmental temperatures generally increase the metabolic rates 406 of offspring; however, we show that at these temperatures selection acts to decrease 407 offspring metabolism, and that parents modify their offspring accordingly. 408

409 Parental condition and offspring temperature increased selection on offspring metabolism

Overall, we found that low parental food levels increased the presence and strength of
negative selection acting on embryo metabolic rate (MR_{embryo}), such that offspring with lower

412 MR_{embrvo} were more likely to survive a critical period of early development (compare Fig. 4 E-

- 413 H with A-D). Previous work has clearly demonstrated the direct effects that environmental
- 414 temperature and food availability produce on metabolic rates [21,24,70,71]. Acute effects of
- 415 warming generally increase metabolic rates in ectotherms, yet acclimatisation or adaptation
- 416 can act to suppress energy expenditure [72,73]. Similarly, low food availability often selects
- 417 for reduced metabolic rates [74], presumably to conserve energy reserves. Further,

temperature and food availability can interact to affect metabolism in complex ways, with

- 419 evidence for temperature mediating both an increase and decrease in metabolism with
- 420 increases in food availability [75–77].

421

422 Intergenerational plasticity is adaptive when environments are consistent across generations 423 We found similar patterns between intergenerational plasticity and selection on offspring 424 metabolic rates when parent and offspring temperatures matched. Parents reared under the 425 warm temperature treatment produced offspring with lower metabolic rates, which were 426 more likely to survive than warm-reared offspring from cool-reared parents. Consequently, 427 offspring with slower metabolic rates showed greater survival in warm developmental 428 temperatures, particularly when they originated from parents from the low food treatment. 429 The downregulation of offspring metabolism is likely to be particularly crucial when food 430 availability is low, where offspring are more likely to be reliant on internal energy reserves to 431 fuel early life growth, maintenance, and development. The alignment of intergenerational 432 plasticity and selection provides evidence that shifts in offspring metabolic phenotypes can

433 be adaptive when the environment in the parent generation matches that of the offspring 434 generation. This has often been assumed in studies measuring performance metrics, such 435 as growth or aerobic capacity, which may trade off with actual fitness [31,39,78]. Through 436 use of a selection analysis, our study provides standardised, comparable estimates of 437 selection, showing that parents can program their offspring with metabolic phenotypes that 438 enhance early life survival. Our findings, however, have worrying implications for 439 environmental mismatches between generations. We acknowledge that ourour study was 440 conducted on zebrafish reared under stable lab conditions, and thatthat wild-caught fish or 441 other taxa may respond differently [79]. However, under However, under increasingly warmer 442 and more variable climates, parents may not be able to keep pace with provisioning their 443 offspring to enhance survival during a vulnerable life stage, and there may be increasing 444 reliance on thermal acclimation to buffer populations to environmental change.

445

446 Potential proximal mechanisms underlying intergenerational effects on offspring metabolism 447 Metabolic suppression as a means to conserve energy has been well documented, yet 448 intergenerational mechanisms are less well explored [24,72,73]. Across generations, 449 epigenetic mechanisms such as changes in DNA methylation can facilitate developmental 450 thermal plasticity to buffer offspring from stressful temperatures [80-82]. One clear 451 mechanism by which parents may alter the transgenerational thermal sensitivity of offspring 452 metabolic and life-history traits is through changes in the density and efficiency of 453 mitochondria [46]. Fasting and warm temperature regimes can enhance mitochondrial 454 efficiency, such that a greater amount of ATP is produced per amount of oxygen consumed 455 [49]. For species that provision their offspring with finite energy reserves in eggs, energy-456 demanding warm temperatures may elicit an adaptive response in parents to produce 457 energy-efficient offspring. It may be that parents can program their offspring with more 458 efficient mitochondria to compensate for a predicted energetically costly environment as 459 reflected by lower metabolic rates [46]. We found that metabolism until two weeks post-460 hatching was unrelated to growth rates, supporting previous work that these two rates can 461 be decoupled and that low metabolic rates do not necessitate slow growth rates because it is 462 mitochondrial efficiency rather than metabolic rate per se that determines availability of ATP 463 for growth [83]. While fitness benefits of reduced metabolism were observed within this 464 study, trade-offs with such as oxidative stress may manifest later in life, affecting fitness-465 enhancing processes [84]. While our study did not detect any negative consequences of 466 metabolic suppression for early life survival in zebrafish, previous work has shown that slow 467 metabolic phenotypes possess lower competitive ability, compared with fast metabolic 468 phenotypes [9]. What is needed now is to go beyond measures of oxygen consumption to

investigate the capacity for parents to alter the efficiency of ATP production in their offspringand mediate fitness under warmer and more nutrient poor environments.

471

472 The presence and form of selection on metabolism varied across ontogeny

473 Despite clear evidence for selection on metabolic rate during embryonic development, we 474 found that, across all environments, directional selection on larval metabolic rate (MR_{larva}) 475 was absent. A recent meta-analysis showed limited evidence for selection on metabolic rate, 476 where the majority of selection coefficients were measured during the adult life stage [85]. 477 Variation or flexibility in metabolic rate may confer a fitness advantage, particularly under 478 selection regimes that change across time and space [24]. Metabolic rate is not a single trait. 479 hence metabolic rates expressed at particular life stages may also affect fitness [12,22]. 480 Metabolic rates may be repeatable, such that they are correlated across the life history, yet 481 we found differences in selection on metabolic rates measured one week apart. In our study, 482 we fed hatched larvae ad libitum, which may have relaxed selection on larval traits. 483 Alternatively, it may be that there are fitness consequences for a low or high larval metabolic 484 rate that were not measured in this study. Survival is a key component, but not an absolute 485 measure of fitness, and further measures are needed of both metabolism across ontogeny 486 and lifetime reproductive output. We did, however observe negative correlational selection 487 on embryo mass and MR_{larva}, in offspring reared at the cool (24 °C) temperature, from high 488 condition parents also reared at 24 °C. Offspring mortality was greatest in smaller embryos 489 with relatively high MR_{larva}, possibly because the reduced endogenous energy reserves often 490 attributed to smaller offspring were insufficient to sustain higher metabolism in the larval 491 stage. Variation in the strength, form, and direction of selection on combinations of early life 492 traits across environments reveals the diversity of adaptive landscapes that organisms may 493 enter, and the challenges that parents face when matching offspring phenotype to enhance 494 performance within a given environment.

495

496 Potential indirect selection on developmental and growth rates

497 We found that metabolic rates measured from the embryo stage through to one week post 498 hatching were consistently negatively correlated with development time and positively 499 correlated with growth rate, but that developmental and growth rates were onlyonly 500 correlated within offspring temperature treatments. Pace-of-life theory proposes that natural 501 selection should favour the integration of a suite of life-history and metabolic traits that 502 together enhance fitness [86]. In our study, warm and low condition parents produced 503 offspring with lowerPace-of-life theory proposes that natural selection should favour the 504 integration of a suite of life-history and metabolic traits that together enhance fitness [86]. In 505 our study, warm and low condition parents produced offspring with lower metabolic rates,

- 506 with evidence for a slower pace-of-life, including extended development time and reduced
- 507 growth rates. Potential mechanisms underlying this response from parents include
- 508 epigenetic modification such as DNA methylation in gametes or early developmental stages,
- or genetic constraints. Our finding that selection acts to reduce embryo metabolic rate in the
- 510 warm offspring treatment may inadvertently also act to reduce the pace of life if these traits
- are both phenotypically and genetically, correlated. There is evidence however, that pace-of-
- 512 life traits can be decoupled, whereby growth and developmental rates, for example, can
- 513 evolve independently [87]. Further measures of multivariate selection will help to disentangle
- the underlying drivers of correlated traits related to the pace of life [88]
- 515

516 Conclusions

517 Our study shows the importance of intergenerational plasticity as a source of variation in

- 518 metabolic rates during early life stages. When parent and offspring environments match,
- 519 parents can program offspring to express metabolic phenotypes that align with selection on
- 520 embryonic metabolic rate. Offspring with lower metabolic rates showed greater survival
- 521 when reared under warm temperatures, and this response was particularly evident when
- 522 offspring originated from low condition parents. Our findings support previous evidence that
- 523 the unpredictability of offspring environment may in part explain why anticipatory parental
- 524 effects are not always, or only weakly, observed. However, identifying the mechanistic basis
- 525 of parental effects on variation in metabolic rate is an important next step.

526	Table 1. Output from best fitting linear mixed effects models. Estimates provided for
527	fixed effects of Parent ('P') temperature (24 °C or 30 °C), parent feeding frequency (High; H
528	or Low; L) and offspring ('O') temperature (24 $^{\circ}$ C or 30 $^{\circ}$ C) on offspring phenotypes: 1)
529	Embryo mass, 2) Metabolic rates (a. MR _{embryo} , b. MR _{hatch} , c. MR _{larva}), 3) Development time, 4)
530	Growth rate, and 5) Survival to two weeks post hatching. For survival, logistic generalised
531	linear mixed effect regression was used and individuals were assigned either "1" for alive at
532	two weeks post hatching or "0" for dead. Parental Tank ID was included as a random effect
533	in all models. All candidate models are provided in Table S2 and ranked in Table S3). All
534	comparisons are made in relation to 'L' parent feed frequency and 30 $^\circ$ C parent and

535 offspring temperature. Significance level set at p < 0.05.

Predictors	Estimate	SE	df	t-value	p-value
	1. Embryo m	ass			
Intercept	56.70	1.49	15.03	38.12	<0.0001***
P feed (L)	8.45	1.06	298.95	7.98	<0.0001***
P temperature (30)	0.18	1.26	233.14	0.14	0.89
P feed (L) \times P temperature (30)	0.37	1.16	367.29	0.25	0.81
	2a. Log ₁₀ MRe	mbryo			
Intercept	-4.27	0.08	184.05	-56.04	<0.0001***
Log ₁₀ Embryo mass	0.69	0.04	174.21	16.02	<0.0001***
P feed (L)	-1.01	0.11	340.08	-9.30	<0.0001***
O temperature (30)	0.06	0.00	505.68	25.81	<0.0001***
P temperature (30)	-0.03	0.00	13.88	-6.02	<0.0001***
Log_{10} Embryo mass \times P feed (L)	0.51	0.06	333.03	8.27	<0.0001***
2b.	Log ₁₀ MR _{hatch}				
Intercept	-4.00	0.13	105.98	-30.44	<0.0001***
Log ₁₀ Embryo mass	0.72	0.07	103.55	9.79	<0.0001***
O temperature (30)	0.07	0.00	442.76	17.79	<0.0001***
P feed (L)	-0.78	0.20	181.16	-3.93	<0.001**
P temperature (30)	-0.01	0.01	12.80	-2.48	0.03
Log ₁₀ Embryo mass x P feed (L)	0.38	0.11	179.19	3.43	<0.001***
2c.	Log ₁₀ MR _{larva}				
Intercept	-3.19	0.14	198.41	-22.60	<0.0001***
Log ₁₀ Embryo mass	0.35	0.08	187.11	4.38	<0.0001***
O temperature (30)	0.03	0.01	370.10	6.09	<0.0001***
P feed (L)	-0.06	0.01	14.94	-8.31	<0.0001***
P temperature (30)	-0.02	0.01	9.29	-2.75	0.02*
3. Development time					
Intercept	110.66	0.96	24.89	115.63	< 0.0001***
O temperature (30)	-48.77	0.82	564.05	-59.61	< 0.0001***
P feed (L)	6.79	1.17	76.00	5.79	< 0.0001***
P temperature (30)	6.24	1.24	55.30	5.03	< 0.0001***
O temperature (30) × P feed (L)	-5.72	1.19	548.31	-4.80	< 0.0001***
O temperature (30) × P temperature (30)	-6.49	1.16	565.52	-5.59	<0.0001***
P feed (L) \times P temperature (30)	0.65	1.54	124.97	0.42	0.68
O temperature (30) \times P feed (L) \times P	2.40	1.81	486.64	1.32	0.19
temperature (30)					
	Growth rate	;			
Intercept	89.78	0.65	460	138.73	<0.0001***
O temperature (30)	-0.10	0.93	460	-0.10	0.92
P feed (L)	-3.04	1.01	460	-3.02	0.003**
P temperature (30)	0.57	0.95	460	0.60	0.55
O temperature (30) \times P feed (L)	2.32	1.41	460	1.64	0.10
O temperature $(30) \times P$ temperature (30)	-0.12	1.37	460	-0.09	0.93

P feed (L) \times P temperature (30)	1.37	1.43	460	0.96	0.34
O temperature (30) \times P feed (L) \times P temperature (30)	-4.50	2.03	460	-2.22	0.02*
	5. Survival				
Intercept	1.08	0.20		5.39	<0.0001***
O temperature (30)	-0.104	0.21		-0.49	0.63
P feed (L)	-0.72	0.21		-3.42	<0.001***
P temperature (30)	-0.08	0.22		-0.37	0.71
O temperature $(30) \times P$ temperature (30)	0.16	0.35		0.48	0.66
P feed (L) \times P temperature (30)	-0.14	0.36		-0.40	0.69
O temperature (30) \times P feed (L	0.11	0.36		0.32	0.75

Table 2. Selection coefficients (mean and standard error). Direction and strength of linear (β) and nonlinear (γ) selection on embryo mass and mass-independent metabolic rates across two life stages (MIMR_{embryo} and MIMR_{larva}; μ IO₂h⁻¹) in *Danio rerio*. Fitness was measured as survival to two weeks post hatching. Results shown for each combination of Parent ('P') temperature (24 °C or 30 °C), parent feeding frequency (High; H or Low; L) and offspring ('O') temperature (24 °C or 30 °C). Significant selection gradients (*p* < 0.05) shown in bold.

Parent environment	Offspring environment		β		γ	
				Embryo mass	MIMR _{Embryo}	MIMR _{Larva}
		Embryo mass	0.134 (0.041)	-0.053 (0.114)	0.010 (0.055)	-0.020 (0.043)
	O24	MIMREmbryo	-0.061 (0.041)		-0.054 (0.092)	0.082 (0.048)
D0411		MIMR _{Larva}	-0.013 (0.033)			0.011 (0.075)
PZ4H				Embryo mass	MR _{Embryo}	MR _{Larva}
	O30	Embryo mass MIMR _{Embryo} MIMR _{Larva}	0.039 (0.036) -0.157 (0.040) -0.055 (0.035)	-0.117 (0.105)	0.052 (0.057) 0.051 (0.113)	-0.030 (0.060) -0.084 (0.065) 0.181 (0.095)
			. ,	Embryo mass	MR _{Embryo}	MR _{Larva}
	O24	Embryo mass MIMR _{Embryo} MIMR _{Larva}	0.180 (0.056) 0.052 (0.033) -0.002 (0.041)	0.371 (0.224)	0.040 (0.125) 0.002 (0.068)	0.004 (0.129) 0.056 (0.079) -0.179 (0.119)
			, , , , , , , , , , , , , , , , , , ,	Embryo mass	MR _{Embryo}	MRLarva
P30H	O30	Embryo mass	-0.109 (0.089)	0.210 (0.215)	-0.165 (0.181)	0.181 (0.258)
		MIMREmbryo	-0.167 (0.074)		-0.091 (0.074)	0.320 (0.218)
		MIMR _{Larva}	-0.021 (0.045)			0.100 (0.073)
				Embryo mass	MR _{Embryo}	MR _{Larva}
	O24	Embryo mass	-0.092 (0.070)	-0.106 (0.166)	-0.083 (0.124)	-0.160 (0.095)
		MR _{Embryo} MR _{Larva}	-0.122 (0.047) 0.046 (0.059)		0.346 (0.281)	0.157 (0.138) 0.090 (0.159)
P24L				Embryo mass	MR _{Embryo}	MR _{Larva}
	000	Embryo mass	-0.200 (0.057)	-0.343 (0.270)	-0.329 (0.222)	0.043 (0.099)
	030	MR _{Embryo} MR _{Larva}	-0.070 (0.035) -0.050 (0.037)		0.329 (0.320)	0.297 (0.149) -0.060 (0.093)
			. ,	Embryo mass	MR _{Embryo}	MR _{Larva}
		Embryo mass	-0.122 (0.056)	-0.039 (0.113)	0.038 (0.095)	0.077 (0.086)
	O24	MR _{Embryo}	-0.161 (0.054)		-0.353 (0.138)	-0.038 (0.093)
		MR _{Larva}	-0.017 (0.038)			-0.051 (0.081)
P30L				Embryo mass	MR _{Embryo}	MR _{Larva}
		Embryo mass	-0.166 (0.056)	0.218 (0.233)	0.171 (0.111)	-0.173 (0.125)
	O30	MR _{Embryo}	-0.120 (0.042)		-0.168 (0.104)	-0.008 (0.065)
		MR _{Larva}	0.022 (0.030)			0.144 (0.120)



545 Figure 1. A. Conceptual diagram: predicted responses of temperature-dependent selection 546 on and intergenerational plasticity of embryo metabolic rates at cool and hot offspring 547 temperatures. We hypothesise that hot offspring temperatures will select for lower metabolic 548 rates, while selection at the cool offspring temperature will be relatively relaxed (positive but 549 not significant correlation between fitness and metabolic rate shown). If intergenerational 550 plasticity aligns with selection when environments across generations match, then similar 551 trends in the direction and strength of selection should be observed. We therefore predict 552 that parents in the warm environment (pink) will produce offspring with lower metabolic rates 553 (smaller curved arrows) compared to parents from the cool temperature (blue), and that this 554 will be correlated with development time and growth rates, with fitness benefits for offspring. 555 B. Experimental design: parents were held under one of four treatment combinations: 24 °C or 30 °C and low or high feeding frequency then bred to produce offspring that were 556 557 reared at either 24 °C or 30 °C. Embryo size (diameter, area, mass) and yolk area were measured at 1-4 hours post fertilisation (hpf), and metabolic rates (measured as rate of 558 oxygen consumption) measured at three stages: 25% of embryonic development (MR_{embryo}), 559 560 1-4 hours post hatching (hph; MR_{hatch}), and 1 week post-hatching (1wph; MR_{larva}). Offspring 561 were then monitored for survival up to two weeks post hatching.





D. Embryo growth rate (mm day⁻¹)

563

0.0015

24-H-24

24-H-30

562

564 Figure 2. Offspring phenotypes in response to parent and offspring treatments.

24-L-30

30-H-24

30-H-30

30-1-24

565 Responses of A. Embryo mass, B. Metabolic rate (MRembryo, MRhatch, MRlarva), C.

24-L-24

- 566 Development time, and D. Growth rate, measured across combinations of parent
- 567 temperature (24 °C or 30 °C), feeding frequency (Low; 'L' or High; 'H'), and offspring
- temperature (24 °C or 30 °C). First number in treatment description refers to parent 568
- 569 temperature and the second refers to offspring temperature. Note that metabolic rates
- 570 shown in panel B do not account for the significant effect of embryo mass.

30-L-30





576 Figure 3. The relationship between offspring metabolic rates (Log₁₀ MR_{embryo}, Log₁₀

577 MR_{hatch}, Log₁₀ MR_{larva}) and Log₁₀ Embryo mass in response to parent and offspring

578 treatments. Coloured data points and lines reflect parent temperature (24 °C or 30 °C),

579 feeding frequency (Low; 'L' or High; 'H'), and offspring temperature (24 °C or 30 °C). Grey

bars are standard error. First number in treatment description refers to parent temperature

and the second refers to offspring temperature.

582



584 **Figure 4. Correlation plots for offspring phenotypes**. Pairwise correlations between

585 offspring traits: metabolic rates (MR_{embryo}, MR_{hatch}, MR_{larva}), embryo development time, and

586 larval growth rate, across combinations of parent and offspring treatments. Coloured plots

587 represent significant correlations between traits.



Figure 5. Selection surface plots. Selection on embryo mass (μ g) and metabolic rate (MR_{embryo}; μ IO₂h⁻¹) across combinations of parent (P)

temperature (24 °C or 30 °C) and feeding frequency (high; H or low; L) and offspring (O) temperature (24 °C or 30 °C) environments.

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603 Competing interests

- 604 The authors have no competing interests to declare.
- 605

606 Data accessibility statement

- All data and code have been made publicly available for peer review on the Open Science
- 608 Framework: <u>https://osf.io/6357s/?view_only=9c6e1ac841fb4e6188fd297aeeaa2733</u>.

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