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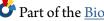
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Experimental Manipulation of the Cost of Thermal Acclimation in Drosophila melanogaster

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Experimental manipulation of the cost of thermal acclimation in *Drosophila melanogaster*

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Acclimation to environmental change can impose costs on organisms. One potential cost is the energy and nutrients consumed by a physiological response, e.g. the resources required for expression of heat-shock proteins (Hsps). We examined the significance of this cost by genetic manipulation. We isolated four isofemale lines from a Drosophila melanogaster population previously transformed with a hsp70-lac2 fusion. Lines were similar in Hsp70 expression but differed in β -galactosidase expression upon heat shock, and replicates of each line were reared on a high quantity and low quantity medium. Multiple heat shock reduced survival in all lines, but did not increase developmental time. Variation in expression of β -galactosidase among lines, which differed more than 4-fold in response to heat treatment, was unrelated to the decreased survival. Thus the predicted effects of β -galactosidase expression on components of fitness were not evident. The superimposition of costs upon those normal for acclimation had no effect on mortality or developmental time, even when resources were especially limiting.

ADDITIONAL KEY WORDS:—genetics - heat shock - metabolism - stress - trade-offs.

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INTRODUCTION

Individuals of many species can change their phenotype when they encounter a novel thermal environment. Such change, termed thermal acclimation, is sometimes adaptive in that it can improve organismal performance in the novel thermal environment. A growing number of studies, however, report that thermal acclimation is neutral or deleterious with respect to fitness in some circumstances, i.e. acclimated individuals perform no better or even worse than non-acclimated individuals (Bennett & Lenski, 1997). One of several possible mechanistic explanations for such nonadaptive outcomes of thermal acclimation is that the expression of a new thermal phenotype (Feder, 1996) can consume too much energy or nutrients, monopolize the expression apparatus and related cellular pathways, and thereby jeopardize other cellular functions. For example, in expression of heat-shock proteins (Hsps), a near-universal response to high temperatures, Hsps can be virtually absent from the cell before heat stress and suddenly undergo 10 000-fold increases in expression (Lindquist, 1993). These proteins may accumulate to account for 10–15% of soluble protein in the cell (Loomis & Wheeler, 1982; Palter et al., 1986), displacing routine protein synthesis. The negative consequences of this intense protein expression for performance and fitness can be profound: *Drosophila* cells that express Hsps at benign temperatures grow more slowly than normal cells (Feder et al., 1992), a yeast strain that cannot express Hsp104 grows faster than its wild-type counterpart on some media (Sanchez et al., 1992), fecundity declines in adult Drosophila melanogaster (Meigen) treated to induce the heat-shock response (Krebs & Loeschcke, 1994), and overexpression of Hsp70 reduces larva-to-adult survival of D. melanogaster (Krebs & Feder, 1997a). Heckathorn et al. (1996a) found that when nitrogen availability is limiting, other proteins are catabolized to provide amino acids for stress protein production. Consequently, loci such as the hsp genes face antagonistic selection, as costs of stress tolerance potentially cancel out benefits of acclimation (Calow, 1991; Coleman, Heckathorn & Hallberg, 1995; Hoffmann, 1995; Krebs & Loeschcke, 1996; Parsons, 1996).

These negative outcomes of thermal acclimation, while consistent with an excessive energetic cost of Hsp expression, are also consistent with other candidate mechanisms such as toxicity of Hsps at high concentration (Krebs & Feder, 1997a). Therefore, to examine the energetic cost explanation explicitly, we exploited Drosophila that had been genetically engineered with a hsp70-lacZ fusion. Even without this transgene, Drosophila larvae express Hsps massively in response to a mild heat shock, such as those that may occur in nature (Junge-Berberovic, 1996; Feder, 1997; Feder, Blair & Figueras, 1997). With the transgene, larvae also express \(\beta\)-galactosidase in response to mild heat shock. β-galactosidase expression typically does not benefit Drosophila, but its production consumes energy and amino acids. From a mass population of this strain, we founded four isofemale lines that express variable amounts of this innocuous protein as a response to heat. Thus these four lines are expected to differ primarily in their cost of thermal acclimation. Replicates of each line were reared with and without repeated induction of β-galactosidase expression, and at high and low resource levels (manipulated by diluting available food at equal volumes). If the resource cost of β-galactosidase expression is non-negligible to fitness, replicates reared with repeated induction of \beta-galactosidase expression should perform less well than lines without β-galactosidase expression, and this difference should be more pronounced in replicates in the low resource treatment than in the high resource treatment.

Changes in performance were assayed by effects on survival and developmental time. Survival, we predicted, would decline where costs are extreme (Gebhardt & Stearns, 1988), while developmental time, which varies in response to minor changes in nutrient resources (Robertson, 1960; Bakker, 1961), would be a more sensitive assay of energetic costs.

MATERIAL AND METHODS

Origin of hsp70-lacZ lines

Simon & Lis (1987) constructed *Drosophila* lines that express bacterial β -galactosidase as a response to heat. In summary, the transformation vector, inserted within a P-element, contained wild-type hsp70 sequence from -194 to the lacZ fusion point and has the hsp70 poly(A) + signal region downstream of lacZ. The fusion protein produced has only the first 7 amino acids of Hsp70 and a functional β -galactosidase with improved expression relative to original trials (Simon *et al.*, 1985).

Although Simon & Lis (1987) originally screened lines to eliminate those with atypical expression, new variation developed over the 10+ years of maintenance. After subculturing singly mated females from the original line, preliminary assays identified several isofemale lines with relatively high and low β -galactosidase expression, of which we chose four for further study.

Chemical and fitness assays

From the same group of parental flies from each line, we collected larvae to assay β-galactosidase activity, Hsp70 expression, survival to adult and developmental time. Flies oviposited on petri dishes containing a yeast-cornmeal-molasses-agar medium, from which larvae could easily be collected from the surface. The assays of β galactosidase activity and Hsp70 expression each required 10–15 2nd-instar larvae per replicate. These larvae were transferred to microtubes with 10 µl phosphate buffered saline and treated either for 1 h at 36°C and 2 h at 25°C before assaying β-galactosidase activity, or for 1 h at 36°C and 1 h at 25°C for Hsp70 concentration. These treatments induce near-maximal levels of expression of each protein (Krebs & Feder, 1997b, unpublished data). Larvae were frozen in liquid nitrogen and stored at -80°C for subsequent biochemical analysis. To assay survival and developmental time, we transferred groups of 40 larvae to a series of vials containing either a high quantity resource (8 ml of undiluted yeast-cornmeal-molasses-agar medium) or a low quantity resource (8 ml of the same batch of medium diluted to 25% with 1% agar). Larvae in half the vials within each resource treatment group developed in a constant 25°C rearing environment (controls), and half received three acclimation induction treatments, 1 h at 36°C, in a thermostatted water bath, at 2, 24 and 48 h after collection. These treatment times correspond to the larval developmental stages, 1st-instar, early 2nd-instar and early 3rd-instar. When not undergoing heat treatment, all larvae developed within a large covered container at 25°C with moistened toweling to ensure high humidity.

To assay β-galactosidase activity, we first sonicated each sample of 10–15 larvae in reaction buffer (Simon & Lis, 1987) on ice and then centrifuged the lysate at 4°C for 30 min at 13 000 rpm. β-galactosidase activity of the supernatant was determined according to Simon & Lis (1987); reactions contained 6.125 μl supernatant and 1 mM chlorophenol red-β-D-galactopyranosidase in 200 μl volume reaction buffer, and were run in triplicate at 37°C. Absorbance of the coloured reaction product was measured at 405 nm with a microplate reader. The protein content of the supernatant was determined with the BCA assay (Pierce Biochemical), and results expressed as mOD₅₆₂·min⁻¹·μg soluble protein⁻¹. Hsp70 expression was determined for 6 or 7 groups of 10–15 larvae per line by enzyme linked immunosorbent assay (ELISA), which has been described elsewhere (Welte et al., 1993; Feder et al., 1996; Krebs & Feder, 1997c). Results were expressed as a percent of a standard, expression in S2 Drosophila cells treated at 36.5°C for 1 h and 1 h at 25°C.

For each line, we determined the proportion of larvae that emerged as adults from each vial; data underwent arcsine square-root transformation before statistical analysis. We estimated the mean developmental time per vial by averaging that for males and females, each determined as \log^{-1} of the log day-of-emergence per individual.

RESULTS

Lines varied in β -galactosidase activity after treatment at 36°C (Fig. 1A, $F_{3,16}$ = 22.4, P<0.001), and each line differed from all others (Tukey's multiple comparisons test, P<0.05). By contrast, Hsp70 expression varied little (Fig. 1B, $F_{3,22}$ =2.0, NS). These lines also varied in larva-to-adult survival (Fig. 2A, $F_{3,112}$ =3.2, P<0.05) and in developmental time (Fig. 2B, $F_{3,104}$ =3.2, P<0.05). The low quantity medium reduced survival ($F_{1,112}$ =63.2, P<0.001) and lengthened larval development ($F_{1,104}$ =235, P<0.001).

The predicted effects of β -galactosidase expression on components of fitness were not evident. Repeated exposure to heat shock, with consequent induction of both Hsps and β-galactosidase, did not affect developmental time in a combined analysis of all lines ($F_{1.104} = 1.0$, NS). Only in line 2 did repeated induction of β -galactosidase expression affect development time. This effect was opposite in direction (decrease) to that expected if resources were limiting, occurred in both high and low quantity medium, and was in a line with intermediate β-galactosidase expression. Although repeated heat shock reduced larva-to-adult survival ($F_{1.112} = 13.8$, P < 0.001), other findings were inconsistent with predictions: First, effects of repeated heat shock were greater in the high quantity medium than in the low quantity medium $(F_{1,112}=6.5,$ P<0.05), in which resource limitation should have been more problematic. Second, while lines varied in response to medium quantity $(F_{3,112} = 4.3, P < 0.01)$, the depression in larva-to-adult survivorship when grown on the low-quantity medium bore no consistent relationship to the β -galactosidase expression in the four lines. In fact, the line with by far the greatest β -galactosidase expression was intermediate to the other lines in this respect. Thus, despite producing very different levels of βgalactosidase, all lines responded to heat similarly in both traits (line \times heat-treatment interactions; survival, $F_{3,112} = 0.1$, NS; developmental time, $F_{3,104} = 1.2$, NS) in the high and low quantity environments (line x heat treatment x medium quantity, survival, $F_{3,112} = 0.7$, NS; developmental time, $F_{3,104} = 0.4$, NS).

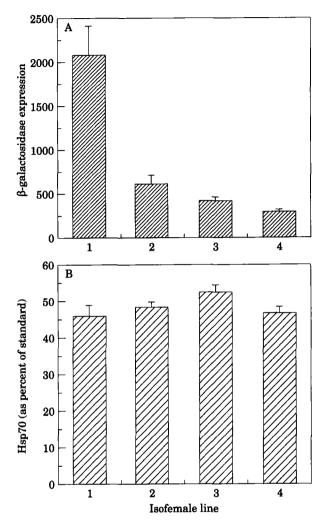


Figure 1. Mean concentration (\pm SE) of (A) β -galactosidase after 1 h at 36°C and 2 h at 25°C and (B) Hsp70 after 1 h at 36°C and 1 h at 25°C for 2nd-instar larvae of four isofemale lines. Protein concentrations are relative to mOD per μ g soluble protein for β -galactosidase, or as a percent of the concentration of Hsp70 found for S2 *Drosophila* cells treated 1 h at 36.5°C and 1 h at 25°C.

DISCUSSION

Multiple heat treatment reduced survival in lines transformed with $hsp70-lac\mathcal{Z}$, but differential expression of β -galactosidase cannot explain fitness variation among lines. β -galactosidase varied more than a factor of four among lines, but neither mortality nor developmental time differed substantially between the line with the highest expression and others. Although thermal stress causes many physiological changes (Huey & Bennett, 1990; Feder, 1996), short heat treatments did not lengthen developmental time of any line and, while mortality increased following heat treatment, the increase did not covary with β -galactosidase expression. Had conditions become limiting, developmental time should have varied (Bakker, 1969).

The fitness consequences of energy and nutrient reallocation depend on the

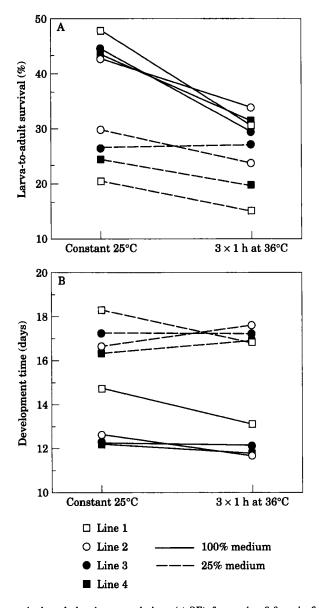


Figure 2. Mean survival and developmental time (\pm SE) for each of four isofemale lines either reared at constant 25°C, or at 25°C interspersed with three 1 h treatments at 36°C. Points are connected to indicate treatment responses for each line. Heat treatment induces both Hsp70 expression and expression of the inserted hsp70-lacZ transgene in these lines.

resource environment in which traits are measured. At times, *Drosophila* larvae live in an infinite pool of energy and nutrients. When orchard fruits are not maturing, *Drosophila* must migrate to natural refuges, which consequently may lead to virtually unlimited resources oscillating with severe larval competition on discrete and ephemeral breeding sites (Atkinson, 1979, 1985). Increasing competition or larval density decreases both the probability of any individual becoming an adult, lengthens developmental time and affects body size and other traits in those who are successful

(Barker, 1983). As the environment declines, genetic parameters responsible for variation in and covariation among these traits may also change (Gebhardt & Stearns, 1989). In our experiments, the low resource treatment effectively reduced survival and increased time of development, which is consistent with resource limitation, and high temperatures or other stresses should increase demands further (Koehn, 1991).

Koehn & Bayne (1989) and Hawkins (1991) suggest that any increase in energy demands, however small, can be a problem. With respect to heat shock, which induces rapid synthesis of Hsps, Coleman and colleagues view allocation of organismal resources as a conflict primarily between Hsp expression and other protein synthesis (Heckathorn et al., 1996b). Indeed, they find that some proteins decline after heat shock and that this loss increases proportionally under conditions of low nitrogen, which Heckathorn et al. (1996a) interpret as a potential reallocation of amino acids for Hsp production. Free amino acid pools therefore may be insufficient for rapid production of new Hsps during the heat-shock response, but whether fitness similarly declines after this reallocation of resources is unknown. In our experiments, superimposing additional costs upon those normal for acclimation had no effect on mortality or developmental time, even when resources were especially limiting. A possible explanation for this outcome is that neither Hsp production, which can have massive costs, nor the additional expression of β-galactosidase exhausted supplies of energy and nutrients. Even inducing these proteins three times failed to increase developmental time in any of the lines.

Energy and nutrient storage may buffer the acute resource demands of thermal acclimation even for larvae on a less concentrated medium. When less food is available, slower larval development may enable similar levels of nutrient storage, as also suggested by the lack of energy limitation in growth despite seasonal variation in resource uptake in mussels (Widdows & Hawkins, 1989; Kreeger et al., 1995). During heat shock in *Drosophila*, a coordinate depression in synthesis of proteins other than Hsps (Solomon et al., 1991) also may spare otherwise limiting stores of amino acids for Hsp expression. Both assembly of energy stores and the consumption of resources in excess of immediate needs can themselves be costly, but these costs may be paid over an entire lifetime rather than acutely. Lengthening the time over which energetic costs are paid may reduce effects on fitness. This explanation also may apply to the recovery in growth after heat shock by larvae that over-express Hsp70 relative to those that express normal levels (Krebs & Feder, 1997a).

The only 'positive' outcome of repeated pretreatment, reduced survival, is equally well consistent with toxicity of the changes associated with thermotolerance. β-galactosidase was one of many proteins produced during heat shock of the lines, and effects could be due either to all the aggregate changes or to one in particular. Hsp70, the protein most highly induced by heat, increased similarly in all four lines. High concentrations of this protein and other family members are sufficient to raise mortality (Feder et al., 1992; Krebs & Feder, 1997a), inhibit protein secretion (Dorner, Krane & Kaufman, 1988, 1992), and in vitro, promote protein aggregation (M. Borrelli and J. Lepock, pers. comm.). Nonetheless, raising Hsp70 levels can improve thermotolerance (Feder et al., 1996) although fitness benefits after stress may trade-off with costs in its absence (Krebs & Feder, 1997c). Our results add to the growing work on the evolutionary consequences of variation in Hsp70 and inducible thermotolerance by providing additional evidence that the heat-shock response has costs to fitness as well as benefits. Moreover, at least in the present

study, limited availability of either energy or nutrients are at most a minor contributor to these costs.

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