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1 Mutation independently affects reproductive traits and dauer
2 larvae development in mutation accumulation lines of
3 *Caenorhabditis elegans*

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6 Hills A. J., Green J. W. M. and Harvey S. C.

7

8 Biomolecular Research Group, Canterbury Christ Church University, Canterbury, CT1 1QU, UK.

9

10 Author for correspondence:

11 Simon C. Harvey

12 e-mail: simon.harvey@canterbury.ac.uk

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21

22 **Abstract**

23 Developmental decisions are important in organismal fitness. For the nematode *Caenorhabditis*
24 *elegans*, which is naturally found in the ephemeral food patches formed by rotting plant material,
25 correctly committing to dauer or non-dauer larval development is key to genotype survival. To
26 investigate the link between reproductive traits, which will determine how populations grow, and
27 dauer larvae formation, we have analysed these traits in mutation accumulation (MA) lines of *C.*
28 *elegans*. We find that reproductive traits of individual worms – the total number of progeny and
29 the timing of progeny production – are highly correlated with the population size observed in
30 growing populations. In contrast, we find no relationship between reproduction traits and the
31 number of dauer larvae observed in growing populations. We also do not observe a mutational
32 bias in dauer larvae formation. These results indicate that the control of dauer larvae formation
33 is distinct from the control of reproduction and that differences in dauer larvae formation can
34 evolve rapidly.

35

36 **Introduction**

37 In the wild, animals will encounter a wide variety of environmental conditions. This means that
38 no single phenotype can be optimal at all times. This can provide a selective advantage to
39 genotypes that are capable of modifying their phenotype in response to the environment
40 experienced (Pigliucci 2005). Given the importance of phenotypic plasticity, it is critical to
41 understand how such response are controlled and how they vary between genotypes. A model
42 important in understanding the genetic control of phenotypic plasticity is the developmental
43 switch between dauer and non-dauer larval development in nematodes. Dauer larvae are
44 specialised alternate third larval stage animals that are found in the Rhabditae and are related
45 to the infective larvae of many parasitic nematodes (Sudhaus 2010).

46

47 Within *Caenorhabditis elegans*, a species that is a coloniser of ephemeral bacterial blooms in
48 rotting vegetation (Frézal and Félix 2015), appropriate dauer larvae development is critical to
49 successful migration between ephemeral food resources. Dauer larvae form under conditions
50 that are not suitable for reproduction, specifically low food and high population density (Hu
51 2007). Individual worms assess population density based on the relative amounts of a complex
52 mix of ascarosides and other related molecules (von Reuss et al 2012). Most work on dauer
53 larvae development relies on exposing synchronised cohorts of larvae to defined amount of
54 exogenous pheromone. However, under natural conditions dauer larvae will form within growing
55 populations and hence the population dynamics will be important. Further, the production of
56 ascarosides – in terms of both amounts and types of molecule produced – varies depending on
57 the conditions experienced by the worms (von Reuss et al 2012). In a growing population of
58 worms, the dynamic production of ascarosides will therefore potentially allow individuals to
59 perceive, and hence respond to, a very nuanced picture of their environment.

60

61 The decision to develop as a dauer larvae is variable between isolates of *C. elegans* (Viney et
62 al. 2003; Green et al. 2013; Diaz and Viney 2015). This variation between isolates is a
63 consequence of differences between isolates in both the composition of the pheromone they
64 produce and their response to pheromone within the environment (Choe et al. 2012; Diaz et al.
65 2014). The importance of pheromone perception in this is supported by comparisons of dauer
66 larvae formation within growing populations and dauer larvae formation of synchronised cohorts
67 of larvae exposed to defined amount of exogenous pheromone (Green et al. 2015). In this
68 comparison, most quantitative trait loci (QTLs) affecting dauer larvae formation are detected in
69 both conditions, suggesting that the same variants are affecting the decision (Green et al.
70 2015).

71

72 Here we use mutation accumulation (MA) lines to further investigate the relationship between
73 reproductive traits, which will determine rates of population growth, and dauer larvae formation
74 in growing populations. MA lines are maintained at very low population sizes – for *C. elegans*,
75 worms are maintained at a population size of one worm per generation – such that the effects of
76 natural selection are minimised. Here we report the variation between MA lines produced from
77 the N2 isolate in lifetime reproductive success (LRS), the intrinsic rate of increase (r), and the
78 population size and the number of dauer larvae at food exhaustion in growing populations.

79

80 **Methods**

81 *Worms*

82 N2 was obtained from the *Caenorhabditis* Genetics Centre. The 19 MA lines assayed here are a
83 subset of lines derived from N2 and obtained from Charles Baer (University of Florida). At
84 assay, lines had undergone 331 generations of MA – 250 prior to their arrival at Canterbury and
85 a further 81 generations after arrival – using methods as described by Baer et al (2005).
86 Ancestral “pseudolines” (see Vassilieva and Lynch 1999) were not created or analysed here.

87

88 *Assays*

89 All assays were performed at 20°C and were initiated with fourth larval stage worms (L4s)
90 grown from synchronised, arrested, L1s. Assays for LRS were undertaken on NGM plates with
91 *Escherichia coli* OP50 strain as food (Stiernagle 2006), with individuals moved to new plates
92 daily throughout the reproductive period and progeny counted after 2-3 days further incubation
93 at 20°C. LRS data was used to estimate r for each line, by iteration from $\sum e^{-rx} l_x m_x = 1$, where l_x
94 represents the age specific survivorship to day x and m_x represents the fecundity on day x
95 (Vassilieva and Lynch 1999). Assays for the population size and the dauer larvae number at
96 food exhaustion were performed as described by Green and Harvey (2012) and Green et al
97 (2013), with populations initiated on sloppy agar plates with a 4 g l⁻¹ agar concentration using

98 100µL of a 10% w/v concentration of OP50 from overnight culture. At food exhaustion, identified
99 by the worms dispersing from the exhausted area, population size and dauer larvae number
100 were determined. Within each experiment, plates were blind- coded and treatments (lines) were
101 randomised. Plates that became contaminated or on which the population had failed to grow
102 were discarded.

103

104 *Statistics*

105 Data from the MA lines were analysed using a random-effects GLM, with line fitted as a random
106 factor, and the among-line component of variance used as the relevant measure of genetic
107 variation to calculate the mutational variance V_m (determined as among-line variance/[2 x
108 number of generations]). To compare between traits, we determined mutational heritability,
109 determined as V_M/V_E , and mutational variability, determined as $V_m/[\text{trait mean}]^2$ (Houle et al.
110 1996), for the number of dauer larvae, the population size and LRS. Differences between MA
111 lines and N2 were determined by weighted pairwise Student t-tests for each trait. The
112 relationship between traits was also investigated by Pearson correlation. Analyses were
113 undertaken in Minitab 17 (Minitab Inc.).

114

115 **Results and Discussion**

116 As expected, given previous work on MA lines, significant variation between the 19 MA lines
117 were observed for all traits (Figure 1) (LRS: $F_{18,237} = 23.35$, $p < 0.001$; r : $F_{18,231} = 38.23$, $p <$
118 0.001 ; population size: $F_{18,67} = 6.07$, $p < 0.001$; number of dauer larvae: $F_{18,67} = 3.51$, $p < 0.001$).
119 Estimates of mutational variability (CV^2) were similar for the population size and for LRS
120 (0.00011 and 0.00014, respectively), and are in line with previous estimates for fitness-related
121 traits. Again, similar estimates of mutational heritability (V_M/V_E) were seen for these traits
122 (0.0017 and 0.0025, respectively). In contrast, the mutational variability for the number of dauer
123 larvae was very high (0.00114). These estimates are however predicated on the assumption

124 that there is no residual among-line variance in the ancestor. Given that no ancestral
125 pseudolines were analysed here, these values may therefore be overestimates. The estimates
126 of mutational heritability and mutational variability for the number of dauer larvae also need to
127 be treated with caution as the residuals are not normal. Traits closely associated with fitness are
128 likely to exhibit strong mutational biases, as they are typically expected to be under strong
129 directional selection (Keightley and Eyre-Walker 1999; Lynch et al. 1999) and thus the majority
130 of mutations will move the trait directionally away from the original value. Here, LRS, r and
131 population size at food exhaustion all decrease in the MA lines in comparison to N2 (for all MA
132 lines these traits are all either significantly lower than N2 or do not differ from N2). This matches
133 what has been seen in previous studies of fecundity in *C. elegans* MA lines (e.g. Vassilieva and
134 Lynch 1999; Baer et al. 2005) and shows that this is also found when the size of growing
135 populations is considered.

136

137 There is however no mutational bias seen in the number of dauer larvae produced at food
138 exhaustion in the MA lines (Figure 1). Here, fewer dauer larvae are observed in some MA lines
139 than are seen in N2, whilst other MA lines have a greater number of dauer larvae. Retesting of
140 two MA lines and N2 controls replicates this result (Figure 2) and indicates that the difference in
141 dauer larvae number is not a consequence of an increased population size – line 15 having the
142 smallest number of worms and the largest number of dauer larvae at food exhaustion (Figure 2).
143 This suggests that dauer larvae development may be being selected for some optimal and
144 intermediate value.

145

146 Comparison between traits in the MA lines indicates that the various reproductive traits are
147 highly correlated (Table 1). That is, lines that produce a lower number of progeny are also likely
148 to do so more slowly and hence to also have a low r . This comparison also indicates that the
149 MA lines with a low LRS and a low r are likely to exhaust food at a lower population size than

150 lines with a high LRS and high r , suggesting that they are less efficient at utilizing the available
151 food. In contrast to the observed links between the various reproductive traits, the number of
152 dauer larvae observed at food exhaustion was not correlated with any of the other traits (Table
153 1). A positive correlation between the numbers of worms and of dauer larvae might have implied
154 that a response to the size of the population was driving dauer larvae numbers, *i.e.* that
155 differences in the number of dauer larvae were a trivial result of changes in the population size
156 (see also Figure 2). That this is not seen suggests that the differences between MA lines in
157 dauer larvae numbers are the result of either changes in how they are perceiving their
158 environment or in how such information is assessed in the dauer/non-dauer decision. This
159 supports the view that the majority of the differences between *C. elegans* lines in dauer larvae
160 formation within growing populations will be a consequence of changes in dauer-specific traits
161 (Green et al. 2015).

162

163 The development of dauer larvae under natural ecological conditions is not well understood, but
164 there is significant variation in the response of strains to pheromone under laboratory conditions
165 (Viney et al. 2003; Diaz et al 2014). It has also been shown that the number of dauer larvae
166 produced in growing populations varies between isolates (Green et al. 2013). The results from
167 our comparisons of the MA lines indicates that dauer larvae formation can change quickly and
168 that this is largely independent of changes in other traits. Given the limited number of mutations
169 that will have been sampled here, this implies that it is likely that *de novo* mutation alone would
170 be sufficient to allow the rapid evolution of both increased and decreased dauer larvae
171 formation. Dauer larvae formation might therefore represent a good trait in which to look for
172 local adaptation in *C. elegans*.

173

174 Conflict of Interest: The authors declare that they have no conflict of interest.

175

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219 modular library of small-molecule signals in *C. elegans*. *Journal of the American Chemical*
220 *Society*, 134(3), pp.1817-1824.

221 Table 1. Relationship between lifetime reproductive success (LRS), the intrinsic rate of increase
 222 (r), and the number of worms and of dauer larvae at food exhaustion in growing populations for
 223 mutation accumulation lines of *C. elegans*. Shown are the Pearson correlation and the
 224 associated p values.

225

	Number of dauer larvae	LRS	R
LRS	-0.199 0.401		
r	-0.081 0.735	0.93 <0.001	
Number of worms	-0.12 0.61	0.88 <0.001	0.90 <0.001

226

227

228 Figure legends

229

230 Figure 1: Dauer larvae development and reproductive traits vary between MA lines. Shown are
231 box plots of lifetime reproductive success (LRS), the intrinsic rate of increase, and the
232 population size and number of dauer larvae at food exhaustion in growing populations for N2
233 (shaded box, with median also shown by the vertical dotted line) and 19 MA lines. Lines are
234 ordered by LRS.

235

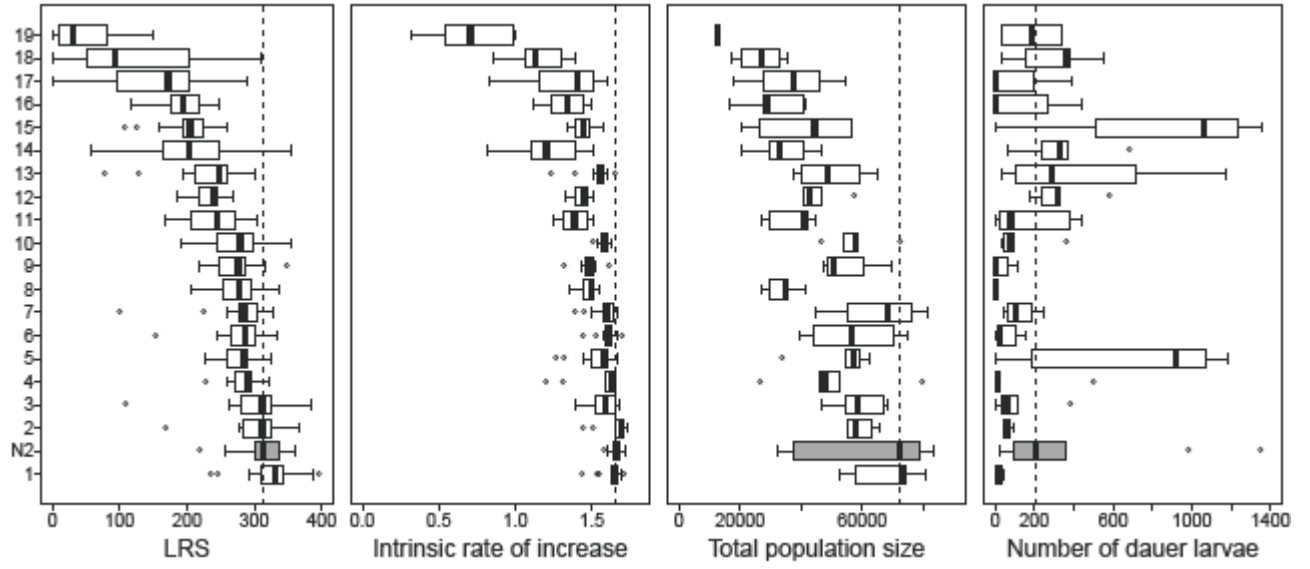
236 Figure 2: Dauer larvae development does not depend on population size in the MA lines. Shown
237 are the population size and number of dauer larvae at food exhaustion in growing populations
238 for N2 and 2 MA lines.

239

240

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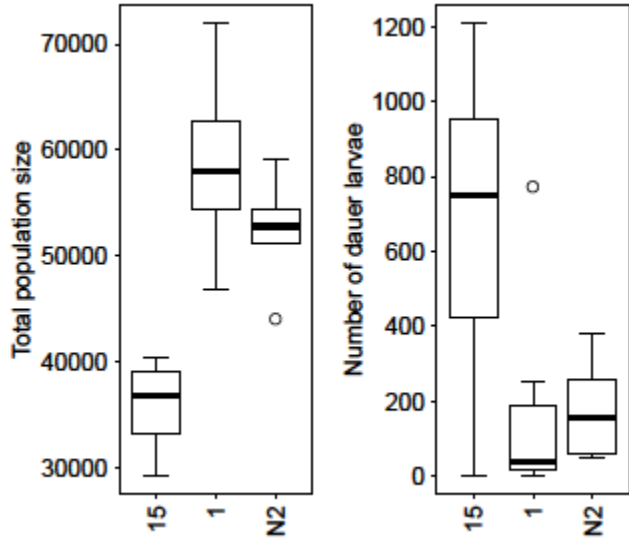
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