

ELECTRONIC APPENDIX

This is the Electronic Appendix to the article

Strategic growth decisions in helper cichlids

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Electronic Appendix A

Strategic growth decisions in helper cichlids

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We created 32 artificial breeding groups by introducing a focal large helper male (standard length SL = 40-45 mm) and a small helper male (SL = 30-35 mm) in each compartment. Two days later, we added a breeding female (SL = 55-65 mm) and either a large or a small breeder male (each in half of the cases, SL = 70-75 mm or 46-51 mm respectively). Large breeder males were 30.0 mm (range 25.3-33.5 mm) and small breeder males were 6.2 mm (range 3.5-9.3 mm) larger than the focal large helper males. Non-accepted group-members were replaced, until a stable group with a size-dependent dominance rank formed (*i.e.* individuals showed the submissive behaviours tail-quivering and zig-zag swimming towards larger group members). We measured body mass (1 mg accuracy) and SL (0.5 mm accuracy, measured by all three authors and averaged), and repeated the measurements after 30 days to determine growth (sequence 1). After 30 days, all breeder males were replaced: half of the groups with a large breeder male now received a new large breeder male of similar size (27.4 mm larger than the focal helper on day 30, range 24.5-30.2 mm), the other half received a new small breeder male (6.8 mm larger than the focal helper on day 30, range 5.3-8.7 mm). Similarly, half of the groups with a small breeder male now received a new large breeder male (27.1 mm larger than the focal helper on day 30, range 25.8-30.7 mm),

the other half received a new small breeder male (6.3 mm larger than the focal helper on day 30, range 5.5-9.5 mm). Such breeder male replacements occur naturally (Taborsky & Limberger 1981; Balshine *et al.* 1998) and helpers were all accepted. Again, growth was measured after 30 days (sequence 2).

The four treatments were abbreviated as follows: SS, SL, LL, LS, where: S. or .S = small breeder male in sequence 1 or 2, L. or .L = large breeder male in sequence 1 or 2 (figure 1). Additionally, we created 8 single breeding pairs (SP, focal male and female breeder of SL = 40-45 mm, similar in size to ensure stable rank, pair building and breeding) and growth was determined on day 30 and 60 (sequence 1 and 2) to compare with the focal helpers and test for status-dependent growth. Each section included groups of all five treatments (figure 1), and no significant effects of section on the results were detected.

All statistical analyses were done with General Linear Models (GLM) in SPSS 11.0 (Lead Technologies Inc.). To assess the status-dependent growth hypothesis, male breeders from the SP treatment were compared with the focal male helpers in all other treatments (SS, SL, LL, LS) using Repeated Measures GLM. Two helper males were discarded from the analyses, since one helper died after the first sequence was completed and one helper changed status during the second sequence. As expected, male breeders grew faster than similar sized large helper males, in line with status-dependent growth (table A1).

To assess the strategic growth hypothesis (using $\ln[\text{helper growth}]$ as dependent variable, to account for exponential diminishing growth in fish), focal helper males breeding with differently sized breeding males were compared, controlling for random individual, fixed sequence and the \ln -transformed helper size at the start of each sequence effects in a Mixed GLM. Since the focal growth rate was highly sensitive to both the initial focal helper size at each sequence ($\ln[\text{helper}$

SL or mass]) and the exact difference in $\ln[\text{SL or mass}]$ between the focal helper and his breeder male, exact values or differences in values were fitted, instead of fixed treatment effects. All 32 helpers were measured at the start (initial size) and the end of the first and the second sequence. The only two exceptions were one male helper who died after the first sequence was completed and was replaced with a size-matched helper, and one helper who changed status during the second sequence; the results of this sequence were discarded in all analyses (hence sample size reduced from 64 to 63). The results are shown in table A2. The same analyses were performed on the small helper males (table A3). All 32 small helpers were measured at the start (initial size) and the end of the first and the second sequence, with one group discarded from the analyses as described above ($n = 63$). Since small helpers might adjust their growth rate not to the difference in initial size with the large helper or the breeding male, but rather to the growth rate of the large helper or the breeder male, we also constructed a model incorporating these two factors. However, growth of the large helper and breeder male did not affect growth of the small helper male ($\ln[\text{growth in SL}]$ of large helper: $P=0.60$, of breeder male: $P=0.31$; $\ln[\text{growth in mass}]$ of large helper male: $P=0.84$, of breeder male: $P=0.66$).

Table 1. Status-dependent growth in the cichlid *Neolamprologus pulcher*. Results of SPSS Repeated Measures GLMs on 30 day growth in SL and mass as dependent variables, showing breeder males ($n = 8$) that were initially the same size as large helper males ($n = 30$) grow faster. All interactions were non-significant.

| | Mean Square | d.f. | F | P | Coefficient \pm s.e. (Sequence 1 / 2) |
|--|-------------|------|-------|----------|--|
| <i>Dependent Variable: growth SL</i> | | | | | |
| Tests of Between-Subjects Effects | | | | | |
| Intercept | 882.501 | 1 | 797.7 | < 0.0001 | Seq. 1: 3.317 \pm 0.212 Seq. 2: 4.150 \pm 0.187 |
| Status | 10.039 | 1 | 9.08 | 0.005 | Seq. 1: 1.143 \pm 0.462* Seq. 2: 0.640 \pm 0.407* |
| Error | 1.106 | 36 | | | |
| Tests of Within-Subjects Effects | | | | | |
| Sequence | 4.271 | 1 | 4.27 | 0.077 | |
| Sequence * Status | 0.799 | 1 | 0.62 | 0.436 | |
| Error(Sequence) | 1.288 | 36 | | | |
| <i>Dependent Variable: growth mass</i> | | | | | |
| Tests of Between-Subjects Effects | | | | | |
| Intercept | 44626304.1 | 1 | 884.8 | < 0.0001 | Seq. 1: 737.99 \pm 36.94 Seq. 2: 874.10 \pm 42.07 |
| Status | 904002.2 | 1 | 17.92 | <0.001 | Seq. 1: 306.64 \pm 80.50* Seq. 2: 228.40 \pm 91.68* |
| Error | 50.437.1 | 36 | | | |
| Tests of Within-Subjects Effects | | | | | |
| Sequence | 118837.764 | 1 | 2.73 | 0.107 | |
| Sequence * Status | 19330.839 | 1 | 0.44 | 0.51 | |
| Error(Sequence) | 43575.056 | 36 | | | |

* Large helper males are the reference category: coefficient is set to zero.

Box's M Test of Equality of Covariance Matrices, Growth SL: $M = 3.561$, d.f.1 = 3, d.f.2 = 2256.262, $F = 1.057$, $P = 0.366$. Growth mass: $M = 4.368$, d.f.1 = 3, d.f.2 = 2256.262, $F = 1.297$, $P = 0.274$.

Levene's Test of Equality of Error Variances, Growth SL: Sequence 1, $F_{1,36} = 1.261$, $P = 0.269$; Sequence 2, $F_{1,36} = 2.101$, $P = 0.156$. Growth mass: Sequence 1, $F_{1,36} = 0.005$, $P = 0.945$; Sequence 2, $F_{1,36} = 2.586$, $P = 0.117$.

Table A2. Strategic reduced growth in large male helpers of the cichlid *Neolamprologus pulcher*. Results of SPSS Mixed GLMs on 30 day growth in ln-transformed SL and mass as dependent variables, showing that large helper males grew more slowly when helping small breeder males ($n = 63$). ‘Sequence’ was fitted as a fixed factor, ‘Individual’ was fitted as a random factor, whereas the ln-transformed initial helper size at sequence 1 or 2 (abbreviated ln[helper size]) and the ln-transformed difference in size between the breeder male and helper male (abbreviated ln[difference size]) were both fitted as continuous factors. All interactions were non-significant.

| | Mean Square | d.f. | F | P | Coefficient \pm s.e. (Sequence 1 / 2) |
|--|--------------------|-------------|----------|----------|--|
| <i>Dependent Variable: ln[growth SL]</i> | | | | | |
| Intercept | 2.771 | 1 | 82.79 | < 0.0001 | 56.914 \pm 6.227 |
| Sequence | 3.487 | 1 | 104.2 | < 0.0001 | -1.442 \pm 0.141* |
| Individual | 0.191 | 31 | 5.7 | < 0.001 | |
| Ln[helper SL] | 2.664 | 1 | 79.6 | < 0.001 | -14.702 \pm 1.648 |
| Ln[difference SL] | 0.271 | 1 | 8.11 | 0.008 | 0.125 \pm 0.044 |
| Error | 0.033 | 28 | | | |
| <i>Dependent Variable: ln[growth mass]</i> | | | | | |
| Intercept | 1.701 | 1 | 32.95 | < 0.001 | 30.746 \pm 5.342 |
| Sequence | 1.3 | 1 | 25.18 | < 0.001 | -1.203 \pm 0.240* |
| Individual | 0.138 | 31 | 2.68 | 0.005 | |
| Ln[helper mass]** | 1.026 | 1 | 19.87 | < 0.001 | -3.093 \pm 0.694 |
| Ln[difference mass]** | 0.011 | 1 | 0.22 | 0.643 | 0.019 \pm 0.042 |
| Error | 0.052 | 28 | | | |

* Sequence 2 is the reference category: coefficient is set to zero.

** Fitting ln[helper SL] and ln[difference SL] instead of mass gave essentially the same results.

Table A3. No evidence for adjustments in growth were detected in the small male helpers of the cichlid *Neolamprologus pulcher*. Results of SPSS Mixed GLMs on 30 day growth in ln-transformed SL and mass as dependent variables ($n = 63$). ‘Sequence’ was fitted as a fixed factor, ‘Individual’ was fitted as a random factor, whereas the ln-transformed initial helper size at sequence 1 or 2 (abbreviated ln[helper size]) and the ln-transformed difference in size between the breeder male and small helper male (abbreviated ln[difference size bs]) or the difference in size between the large helper male and small helper male (abbreviated ln[difference size ls]) were all three fitted as continuous factors. All interactions were non-significant.

| | Mean Square | d.f. | F | P | Coefficient ± s.e. (Sequence 1 / 2) |
|--|--------------------|-------------|----------|----------|---|
| <i>Dependent Variable: ln[growth SL]</i> | | | | | |
| Intercept | 1.351 | 1 | 16.21 | < 0.001 | 42.060 ± 10.406 |
| Sequence | 1.326 | 1 | 15.9 | < 0.001 | -1.328 ± 0.333* |
| Individual | 0.112 | 31 | 1.34 | 0.22 | |
| Ln[helper SL] | 1.278 | 1 | 15.33 | 0.001 | -11.483 ± 2.933 |
| Ln[difference SL bs] | 0 | 1 | 0.02 | 0.897 | 0.016 ± 0.121 |
| Error | 0.083 | 28 | | | |
| <i>Dependent Variable: ln[growth SL]</i> | | | | | |
| Intercept | 0.595 | 1 | 7.26 | 0.012 | 36.143 ± 13.344 |
| Sequence | 0.733 | 1 | 8.94 | 0.006 | -1.177 ± 0.394* |
| Individual | 0.105 | 31 | 1.29 | 0.25 | |
| Ln[helper SL] | 0.642 | 1 | 7.84 | 0.009 | -10.019 ± 3.579 |
| Ln[difference SL ls] | 0.041 | 1 | 0.5 | 0.484 | 0.347 ± 0.489 |
| Error | 0.082 | 28 | | | |
| <i>Dependent Variable: ln[growth mass]</i> | | | | | |
| Intercept | 0.712 | 1 | 10.47 | 0.003 | 19.223 ± 5.907 |
| Sequence | 0.424 | 1 | 6.23 | 0.019 | -0.907 ± 0.363* |
| Individual | 0.11 | 31 | 1.61 | 0.102 | |
| Ln[helper mass]** | 0.304 | 1 | 4.46 | 0.044 | -1.755 ± 0.831 |
| Ln[difference mass bs]** | 0.084 | 1 | 1.23 | 0.276 | -0.075 ± 0.068 |

| | | | | | |
|--|-------|----|------|-------|-----------------|
| Error | 0.068 | 28 | | | |
| <hr/> | | | | | |
| <i>Dependent Variable: ln[growth mass]</i> | | | | | |
| Intercept | 0.375 | 1 | 5.29 | 0.029 | 18.819 ± 8.152 |
| Sequence | 0.271 | 1 | 3.83 | 0.061 | -0.902 ± 0.461* |
| Individual | 0.108 | 31 | 1.52 | 0.134 | |
| Ln[helper mass]** | 0.231 | 1 | 3.25 | 0.082 | -1.731 ± 0.960 |
| Ln[difference mass ls]** | 0 | 1 | 0.03 | 0.867 | -0.057 ± 0.335 |
| Error | 0.071 | 28 | | | |

* Sequence 2 is the reference category: coefficient is set to zero.

** Fitting ln[helper SL] and ln[difference SL] instead of mass gave essentially the same results.