## TITLE:

Variation in physiological function across source populations of a New Zealand

freshwater snail

# **AUTHORS:**

Emma S. Greimann<sup>1</sup>, Samuel Fahrner<sup>2</sup>, James D. Woodell<sup>1</sup>, Samantha K. Hennessey<sup>1</sup>, Michael R. Kline<sup>1</sup>, Jorge A. Moreno<sup>1</sup>, Madeline R. Peters<sup>1</sup>, Jennifer L. Cruise<sup>3</sup>, Maurine Neiman<sup>1</sup>, Joel Sharbrough<sup>1,4</sup>

<sup>1</sup> University of Iowa, Iowa City, IA, USA

<sup>2</sup> University of Minnesota, Saint Paul, MN, USA

<sup>3</sup> University of St. Thomas, Saint Paul, MN, USA

<sup>4</sup> Colorado State University, Fort Collins, CO, USA

# **CORRESPONDING AUTHOR:**

Joel Sharbrough

jsharbro@rams.colostate.edu

Colorado State University

Fort Collins, CO 80523

## **KEYWORDS**:

asexual reproduction, JC-1, mitochondria, oxygen consumption, *Potamopyrgus antipodarum*, sexual reproduction

## 1 ABSTRACT

2 The mitochondrial and nuclear-encoded genes responsible for cellular respiration are 3 expected to experience relatively intense purifying selection, meaning that variation in 4 these genes will often decrease fitness. Still, extensive variation for mitochondrial 5 haplotype and function persists in natural populations. We integrated physiological, 6 cellular, and behavioral approaches to quantify phenotypes relevant to mitochondrial 7 function across a diverse sample of Potamopyrgus antipodarum, a New Zealand snail 8 characterized by frequent coexistence between otherwise similar sexual and asexual 9 individuals. We found extensive across-lake variation in organismal oxygen 10 consumption and behavioral response to heat stress coupled with elevated 11 mitochondrial membrane potential in males vs. females. These data set the stage for 12 applying this important model system for sex, host-parasite interactions, invasion 13 biology, and ecotoxicology to novel tests of the relationships between mitochondrial 14 variation and performance in natural populations. 15

#### 16 **INTRODUCTION**

17 Mitochondrial function is a critical component of eukaryotic function and fitness [1]. 18 Despite its importance, genes underlying oxidative phosphorylation are often 19 polymorphic within species [1], which has important consequences for phenomena from 20 mitonuclear incompatibilities [2] to DNA barcoding [3]. Indeed, extensive variation for 21 mitochondrial function has been observed in a diverse array of species [4]. Although 22 some variation in metabolic and mitochondrial traits has been linked to specific 23 environmental correlates (e.g., altitude - [5], temperature - [6], energy source - [7]), we 24 lack a systematic understanding of the distribution of this variation across biogeographic 25 space. 26 Maternal transmission of mitochondrial genomes is expected to influence the 27 distribution of phenotypic variation because only female-derived mutations are 28 transmitted. This phenomenon has two primary consequences: 1) ~50% reduction in  $N_{\rm e}$ 29 relative to biparentally inherited genomes, and 2) sexually antagonistic mutations only 30 experience effective natural selection in females [8]. This latter phenomenon, the socalled "Mother's Curse", is predicted to result in the accumulation of mutations that are 31 32 neutral or beneficial in females, but deleterious in males [9]. The lack of widespread 33 evidence for Mother's Curse (but see [10]) may point to mechanisms that prevent the 34 spread of male-specific deleterious mutations in mitochondrial genomes [11, 12]. The 35 extent to which Mother's Curse shapes patterns of variation in mitochondrial function 36 therefore represents yet another important unanswered question in evolutionary biology. 37 Because mitochondrial function requires compatibility between nuclear and

38 mitochondrial gene products [13], reproductive mode can also dramatically impact the

39 evolution of variation in mitochondrial function because only sexual reproduction allows 40 for the regular movement of mitochondrial genomes across diverse nuclear genomic 41 backgrounds. Sexual reproduction between distantly related parents can give rise to 42 hybrid offspring harboring mitonuclear incompatibilities [14]. On the other hand, 43 inbreeding and asexual reproduction can reduce the efficacy of selection on 44 mitochondrial genomes [15, 16]. While reduced mitochondrial function and organismal 45 fitness in hybrid lineages is well documented [2], there are few empirical tests of 46 whether and how inbreeding or asexuality affects mitochondrial function, leaving an 47 important gap in our understanding of the evolutionary consequences of changes in 48 reproductive mode. Surveys of mitochondrial genomes of asexual lineages [17, 18] 49 have revealed elevated rates of accumulation of nonsynonymous mutations in 50 mitochondrial genomes compared to sexual lineages. Determining whether these 51 mutations actually result in reduced function will have profound implications for our 52 understanding of the maintenance of sex. 53 Potamopyrgus antipodarum, a New Zealand freshwater snail [19], is ideally

54 suited to answer these outstanding questions regarding mitochondrial function. There 55 is both extensive mtDNA population structure in their native range [20] and evidence for 56 local adaptation of snails to their source lakes [21, 22]. Temperature in particular 57 appears to be a primary determinant of the geographical distribution of *P. antipodarum* 58 within New Zealand [23]. Because asexuality has arisen multiple times within P. 59 antipodarum [20, 24], and because sexual and asexual lineages frequently coexist in 60 nature [24], asexual lineages can be treated as repeated "natural experiments" into the 61 evolutionary consequences of asexuality.

Here, we tested whether lake of origin, reproductive mode, or sex affect
mitochondrial and behavioral function under laboratory conditions in field-collected *P. antipodarum*.

65

### 66 MATERIALS AND METHODS

67 Because definitive determination of reproductive mode requires snail sacrifice, we 68 sampled field-collected snails from lakes known to harbor sexual and asexual 69 individuals, with populations at least ~10% male [24]. Upon arrival at the University of 70 Iowa, snails were housed at 16°C on a 18hr light/6 hr dark schedule, and fed Spirulina 71 algae 3x per week, as described in [25]. We arbitrarily selected adult snails from each 72 lake collection and isolated each snail in a 0.5 L glass container with 300ml carbon-73 filtered H<sub>2</sub>O. Water was changed weekly. Assay sampling details (sample size, etc.) are 74 summarized in Table 1. Reproductive mode was determined after assay completion, 75 following the flow cytometry protocol outlined in [24]. We measured oxygen consumption as described in [25] for 57 wild-caught snails 76 77 from each of six lakes at three different water temperatures: 16°C (not stressful, and 78 similar to New Zealand lake temperatures), 22°C (moderately stressful), and 30°C 79 (stressful) [25]. At each temperature, we assayed snails in a randomly determined order, 80 and only snails that completed all temperature trials were included in analyses. Mean 81 wet mass for each individual was calculated from the three separate temperature trials. 82 Righting behavior [25] and time to emergence following a startling stimulus (M

Neiman pers. obs.) increase with temperature in *P. antipodarum*, indicating that both
assays are effective proxies for heat stress. We quantified righting and emergence

times under each of the same three temperature treatments as for oxygen consumption
in 46 wild-caught *P. antipodarum*.

JC-1 is a small, positively charged molecule that diffuses down the
electrochemical gradient of the inner mitochondrial membrane and fluoresces green
when dispersed and red when aggregated inside the mitochondrial matrix [26].
Therefore, the ratio of red: green fluorescence in freshly isolated mitochondria can
serve as a proxy for mitochondrial membrane potential. We measured red: green ratios
in JC-1-treated mitochondrial extracts from 46 wild-caught *P. antipodarum* as described
in [25] using a Becton Dickenson LSR II flow cytometer.

94 We used a mixed-effects model framework to quantify the relationships between 95 oxygen consumption and behavioral metrics with categorical variables for temperature 96  $(16^{\circ} \text{ C}, 22^{\circ} \text{ C}, 30^{\circ} \text{ C})$ , lake of origin (n = 3-6 depending on the analysis), reproductive 97 mode (asexual, sexual), sex (male, female; only fit in models pertaining to behavior 98 assays), and a continuous variable for mass (g; only fit in model pertaining to oxygen 99 consumption). We modeled a term for snail identity as a random intercept to account for 100 repeated measures on individuals across temperatures. Finally, we modeled 101 mitochondrial membrane potential, measured as the ratio of red: green fluorescence, as 102 a function of lake, reproductive mode, and sex using analysis of variance (ANOVA). 103 We developed final models using backwards selection until only predictors with 104 p-values less than 0.05 remained. To test assumptions of normality and heteroscedacity 105 of errors, we graphically inspected residuals and log- or square-root-transformed 106 response variables when necessary. We performed all statistical analyses in R [27], 107 fitting fixed-effect models with the Im function, fitting mixed-effects models using the

108 Ime4 package [28], and estimating degrees of freedom for mixed-effect models using
109 Satterthwaite's approximation via the ImerTest package [29].

110

### 111 **RESULTS**

All model-fitting results are detailed in Table 2. We found that temperature (p < 0.0001),

mass (p = 0.00154), and lake of origin (p = 0.0072), but not reproductive mode, were

- significantly associated with the rate of oxygen consumption (Figure 1a). Temperature
- 115 was a significant predictor of both righting ability (p < 0.0001, Figure 1b) and emergence
- time (*p* < 0.0001, Figure 1c). Lake of origin was a significant predictor of righting ability
- 117 (p = 0.0155), but not of emergence time. Neither reproductive mode nor sex were
- significantly associated with behavioral responses to heat stress. Sex, but not lake of
- 119 origin or reproductive mode, was a significant predictor of mitochondrial membrane
- potential (p = 0.0070, Figure 2), with higher mitochondrial membrane potential in males
- 121 vs. females.

122

#### 123 DISCUSSION

Here, we report the first evidence of population-structured variation for mitochondrial and behavioral function in *P. antipodarum*. Combined with population structure in mitochondrial genes [18, 20], this result suggests that mitochondrial function could be locally tuned in *P. antipodarum*. We also find that males have higher mitochondrial membrane potential than females. This result suggests that male *P. antipodarum* do not suffer from Mother's Curse, at least with respect to their ability to generate a proton motive force.

- 131 These data set the stage for future studies addressing multiple important
- 132 evolutionary questions, including invasiveness, response to parasite infection,
- 133 ecotoxicology, and the evolutionary maintenance of sex. Chief among these questions
- 134 is whether and how asexuality might influence mitonuclear coevolution. While asexuality
- 135 should reduce the efficacy of natural selection in both nuclear [30] and mitochondrial
- 136 genomes [16, 31], stable transmission of mitonuclear genotypes may also facilitate
- 137 rapid mitonuclear coadaptation and thereby local adaptation [32]. Importantly, the strong
- 138 lake effect implies that extensive intrapopulation sampling is necessary for evaluating
- 139 mitochondrial function in *P. antipodarum*.
- 140

## 141 ACKNOWLEDGEMENTS

- 142 Flow cytometry was performed at UI's Flow Cytometry Facility. We thank Laura Bankers,
- 143 Kaitlin Hatcher, Katelyn Larkin, and Kyle McElroy for snail collections.
- 144

## 145 **DATA Availability**

- 146 Data to be archived on Dryad.
- 147

#### 148 AUTHOR CONTRIBUTIONS

- 149 ESG, JTS contributed to all aspects of study, SF to statistical analyses and manuscript
- 150 drafting, JLC, MN to concept, statistical design, and manuscript drafting, and JDW, SKH,
- 151 MRK, JAM, and MRP to data collection and manuscript editing.

152

153 FUNDING

- 154 Work was supported by National Science Foundation (NSF: MCB 1122176; DEB -
- 155 1310825), Iowa Academy of Sciences (ISF #13-10), and Dewey Stuit Fund for
- 156 Undergraduate Research.
- 157
- 158 COMPETING INTERESTS
- 159 We have no competing interests.
- 160
- 161 ETHICAL STATEMENT
- 162 Not applicable to mollusks.

# 163

#### Table 1. Summary of source populations of Potamopyrgus antipodarum.

Oxygen consumption assay							
Lake	Latitude, Longitude	Sexual	Asexual	Male	Female		
Alexandrina	-43.900476, 170.453978	14	2	-	16		
Clearwater	-43.602131, 171.043917	-	4	-	4		
Kaniere	-42.832886, 171.14759	16	-	-	16		
Paringa	-43.713068, 169.411348	5	-	-	5		
Rotoroa	-41.855414, 172.637882	-	17	-	17		
Selfe	-43.237765, 171.520449	-	3	-	3		

#### Behavior and mitochondrial membrane potential assays<sup>1</sup>

Lake	Latitude, Longitude	Sexual	Asexual	Male	Female			
Alexandrina	-43.900476, 170.453978	3	-	3	-			
Ellery	-44.046898, 168.654261	2	3	-	5			
Kaniere	-42.832886, 171.14759	5	1	4	2			
Mapourika	-43.315212, 170.204061	8	2	6	4			
Rotoroa	-41.855414, 172.637882	4	1	-	5			
Selfe	-43.237765, 171.520449	9	8	9	8			

<sup>1</sup> – Same individual snails were used in behavioral and mitochondrial membrane

potential assays

164

Table 2. Linear and mixed-effects models of select predictors on oxygen consumption, righting

xygen consumption	n <sup>1</sup>			
Factor	X <sup>2</sup>	df	р	Non-significant predictors
Intercept	2.700	1	0.1004	
Temperature	39.038	2	< 0.0001	Depreductive Mede
Mass	11.061	1	0.0009	Reproductive Mode
Lake of Origin	15.280	5	0.0092	
Factor <sup>§</sup>	X <sup>2</sup>	df	р	Non-significant predictors
Intercept	2.164	1	0.141	
Temperature	27.336	2	< 0.0001	Reproductive Mode
Mass	4.365	1	0.0367	
ghting time <sup>2</sup>				

Factor	X	ar	р	Non-significant predictors <sup>*</sup>
Intercept	59.205	1	< 0.0001	
Temperature	73.661	2	< 0.0001	Reproductive Mode, Sex
Lake of Origin	14.020	5	0.0155	
Factor <sup>§</sup>	X <sup>2</sup>	df	p	Non-significant predictors*
Intercept	94.767	1	< 0.0001	
Temperature	69.655	2	< 0.0001	Reproductive Mode, Sex
Lake of Origin	13.429	4	0.0094	
Factor <sup>†</sup>	X <sup>2</sup>	df	p	Non-significant predictors*
Intercept	364.550	1	< 0.0001	Reproductive Mode, Sex,
Temperature	52.370	2	< 0.0001	Lake of Origin

#### **Emergence Time<sup>3</sup>**

Factor	X <sup>2</sup>	df	p	Non-significant predictors*
Intercept	448.891	1	< 0.0001	Reproductive Mode,
Temperature	46.646	2	< 0.0001	Sex, Lake of Origin
Factor <sup>§</sup>	X <sup>2</sup>	df	р	Non-significant predictors*
Intercept	363.860	1	< 0.0001	Sex, Reproductive Mode,
Temperature	34.608	2	< 0.0001	Lake of Origin
<b>Factor</b> <sup>†</sup>	<b>X</b> <sup>2</sup>	df	р	Non-significant predictors*
Intercept	59.205	1	< 0.0001	
Temperature	73.661	2	< 0.0001	Reproductive Mode, Sex
Lake of Origin	14.020	5	0.0155	

## Mitochondrial membrane potential<sup>4</sup>

Factor	Sum of Squares	df	F	р	Non-significant predictors*	
Intercept	55.876	1	305.1038	< 0.0001	Deproductive Mode	
Sex	1.466	1	8.0035	0.0070	Reproductive Mode, Lake of Origin	
Residuals	8.058	44			Lake of Oligin	

_	Factor <sup>§</sup>	Sum of Squares	df	F	р	Non-significant predictors*
	Intercept	55.876	1	329.344	< 0.0001	Deproductive Mode
	Sex	1.026	1	6.046	0.0183	Reproductive Mode,
	Residuals	6.956	41			Lake of Origin
	Factor <sup>†</sup>	Sum of Squares	df	F	p	Non-significant predictors*
	Intercept	96.564	1	482.37	< 0.0001	Reproductive Mode, Lake of
	Residuals	6.406	32			Origin, Sex

<sup>1</sup> – Type III Repeated-Measures Analysis of Deviance  $\chi^2$  Test of O<sub>2</sub> consumption per hour

<sup>2</sup> – Type III Repeated-Measures Analysis of Deviance  $\chi^2$  Test of log-transformed righting times

 $^{3}$  – Type III Repeated-Measures Analysis of Deviance  $\chi 2$  Test of square-root-transformed emergence times

<sup>4</sup> – Type III Analysis of Variance F Test of log-transformed ratios of red: green in mitochondrial extracts

\* - Non-significant predictors listed in order of elimination from the model

 $\ensuremath{\S}$  – Model fit only included lakes from which both sexual and as exual snails were assayed

† - Model fit only included lakes from which both male and female snails were assayed

165

# 166 FIGURE LEGENDS

# 167 Figure 1. Physiological responses to heat stress for *P. antipodarum* across

- 168 **source lakes**. a) Oxygen consumption/hour/gram. b) Righting time. c) Emergence time
- 169
- 170 Figure 2. Mitochondrial membrane potential in field-collected *P. antipodarum*.
- 171 Ratios of red: green fluorescence of JC-1-treated mitochondrial extracts for a) snails
- 172 from all six New Zealand lakes, b) male vs. female snails from all lakes, and c) male vs.
- 173 female snails from three New Zealand lakes with replication for sex.

174

#### 175 **REFERENCES**

- 176 [1] Dowling, D.K., Friberg, U. & Lindell, J. 2008 Evolutionary implications of non-neutral
- 177 mitochondrial genetic variation. *Trends Ecol Evol* **23**, 546-554.
- 178 [2] Sloan, D.B., Havird, J.C. & Sharbrough, J. 2017 The on-again, off-again relationship
- between mitochondrial genomes and species boundaries. *Mol Ecol* **26**, 2212-2236.
- 180 [3] Hebert, P.D., Ratnasingham, S. & deWaard, J.R. 2003 Barcoding animal life:
- 181 cytochrome c oxidase subunit 1 divergences among closely related species. *Proc Roy*
- 182 Soc Lond B **270 Suppl 1**, S96-99.
- 183 [4] Dobler, R., Rogell, B., Budar, F. & Dowling, D.K. 2014 A meta-analysis of the
- strength and nature of cytoplasmic genetic effects. *J Evol Biol* **27**, 2021-2034.
- 185 [5] Simonson, T.S., Yang, Y.Z., Huff, C.D., Yun, H.X., Qin, G., Witherspoon, D.J., Bai,
- 186 Z.Z., Lorenzo, F.R., Xing, J.C., Jorde, L.B., et al. 2010 Genetic evidence for high-
- 187 altitude adaptation in Tibet. Science **329**, 72-75.
- 188 [6] Clarke, A. & Johnston, N.M. 1999 Scaling of metabolic rate with body mass and
- 189 temperature in teleost fish. *J Anim Ecol* **68**, 893-905.
- 190 [7] Montooth, K.L., Marden, J.H. & Clark, A.G. 2003 Mapping determinants of variation
- in energy metabolism, respiration and flight in *Drosophila*. *Genetics* **165**, 623-635.
- 192 [8] Frank, S.A. & Hurst, L.D. 1996 Mitochondria and male disease. *Nature* **383**, 224-224.
- 193 [9] Camus, M.F., Clancy, D.J. & Dowling, D.K. 2012 Mitochondria, maternal inheritance,
- 194 and male aging. *Curr Biol* **22**, 1717-1721.

- 195 [10] Innocenti, P., Morrow, E.H. & Dowling, D.K. 2011 Experimental evidence supports
- a sex-specific selective sieve in mitochondrial genome evolution. *Science* **332**, 845-848.
- 197 [11] Kuijper, B., Lane, N. & Pomiankowski, A. 2015 Can paternal leakage maintain
- sexually antagonistic polymorphism in the cytoplasm? *J Evol Biol* **28**, 468-480.
- 199 [12] Wade, M.J. & Brandvain, Y. 2009 Reversing Mother's Curse: selection on male
- 200 mitochondrial fitness effects. *Evolution* **63**, 1084-1089.
- [13] Rand, D.M. 2001 The units of selection on mitochondrial DNA. *Annu Rev Ecol Syst*32, 415-448.
- 203 [14] Burton, R.S., Pereira, R.J. & Barreto, F.S. 2013 Cytonuclear genomic interactions
- and hybrid breakdown. *Annu Rev Ecol Evol Syst* **44**, 281-302.
- 205 [15] Wright, S.I., Ness, R.W., Foxe, J.P. & Barrett, S.C.H. 2008 Genomic consequences
- of outcrossing and selfing in plants. *Int J Plant Sci* **169**, 105-118.
- 207 [16] Neiman, M. & Taylor, D.R. 2009 The causes of mutation accumulation in
- 208 mitochondrial genomes. *Proc Roy Soc Lond B* **276**, 1201-1209.
- [17] Henry, L., Schwander, T. & Crespi, B.J. 2012 Deleterious mutation accumulation in
  asexual *Timema* stick insects. *Mol Biol Evol* 29, 401-408.
- [18] Neiman, M., Hehman, G., Miller, J.T., Logsdon, J.M., Jr. & Taylor, D.R. 2010
- 212 Accelerated mutation accumulation in asexual lineages of a freshwater snail. *Mol Biol*
- 213 Evol 27, 954-963.

- [19] Winterbourn, M. 1970 Population studies on the new zealand freshwater gastropod,
- 215 Potamopyrgus antipodarum (Gray). J Mollus Stud **39**, 139-149.
- 216 [20] Neiman, M. & Lively, C.M. 2004 Pleistocene glaciation is implicated in the
- 217 phylogeographical structure of *Potamopyrgus antipodarum*, a New Zealand snail. *Mol*
- 218 *Ecol* **13**, 3085-3098.
- [21] Krist, A.C., Bankers, L., Larkin, K., Larson, M.D., Greenwood, D.J., Dyck, M.A. &
- 220 Neiman, M. 2017 Phosphorus availability in the source population influences response
- to dietary phosphorus quantity in a New Zealand freshwater snail. *Oecologia*. **185**, 596605.
- [22] Lively, C.M. & Jokela, J. 1996 Clinal variation for local adaptation in a host-parasite
  interaction. *Proc Roy Soc Lond B* 263, 891-897.
- [23] Winterbourn, M. 1969 Water temperature as a factor limiting the distribution of
- 226 Potamopyrgus antipodum (Gastropoda Prosobranchia) in the New Zealand thermal
- 227 region. *NZ J Mar Fresh* **3**, 453-458.
- [24] Neiman, M., Paczesniak, D., Soper, D.M., Baldwin, A.T. & Hehman, G. 2011 Wide
- 229 variation in ploidy level and genome size in a New Zealand freshwater snail with
- coexisting sexual and asexual lineages. *Evolution* **65**, 3202-3216.
- [25] Sharbrough, J., Cruise, J.L., Beetch, M., Enright, N.M., & Neiman, M. 2017 Genetic
  variation for mitochondrial function in the New Zealand freshwater snail *Potamopyrgus*
- 233 antipodarum. J Hered **108**, 759-768.

- 234 [26] Garner, D.L. & Thomas, C.A. 1999 Organelle-specific probe JC-1 identifies
- 235 membrane potential differences in the mitochondrial function of bovine sperm. Mol
- 236 *Reprod Devel* **53**, 222-229.
- [27] Team, R.C. 2017 R: A language and environment for statistical computing [Internet].
- 238 Vienna, Austria; 2014.
- [28] Bates, D., Sarkar, D., Bates, M.D. & Matrix, L. 2007 The Ime4 package. *R package version* 2, 74.
- [29] Kuznetsova, A., Brockhoff, P.B. & Christensen, R.H.B. 2015 Package 'ImerTest'. *R package version* 2.
- [30] Charlesworth, B. 2012 The effects of deleterious mutations on evolution at linked
  sites. *Genetics* **190**, 5-22.
- [31] Gabriel, W., Lynch, M. & Burger, R. 1993 Muller's Ratchet and Mutational
- 246 Meltdowns. Evolution 47, 1744-1757.
- [32] Neiman, M. & Linksvayer, T.A. 2006 The conversion of variance and the
- evolutionary potential of restricted recombination. *Heredity* **96**, 111-121.
- 249



