


REVIEW

Population cycles in voles and lemmings: state of the science and future directions

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ABSTRACT

1. Despite nearly a century of research, the causes of population cycles in Arvicoline rodents (voles and lemmings) in northern latitudes are not yet fully understood. Theory tells us that delayed density-dependent feedback mechanisms are essential for rodent population cycles, suggesting vegetation–rodent, rodent–parasite or rodent–predator interactions as the most likely drivers of population cycles.
2. However, food provisioning, carried out either indirectly through fertilisation treatments of the habitat or directly through food supplementation, has failed to alter population cycles substantially, suggesting that variation in food supply by itself is not necessary or sufficient to cause cyclic fluctuations in abundance.
3. Predator exclusion experiments conducted in Fennoscandia have succeeded in slowing population crashes and increasing autumn densities, implicating predation as the most likely cause of rodent cycles in this region. However, experimental removal of specialist predators in northern England had no discernible effect on a cyclic vole population, casting doubt on the notion that predation is a necessary explanation of rodent population cycles.
4. Population cycle research has contributed substantially to our current understanding of the dynamics, regulation and persistence of biological populations, but we do not yet know with certainty what factors or processes cause multiannual population fluctuations or if population cycles are driven by the same mechanisms everywhere. Recent theoretical and empirical studies suggest that extrinsic factors (primarily food supply and predator abundance) may interact with population intrinsic processes (e.g. dispersal, social behaviour, stress response) to cause multiannual population fluctuations and to explain biological attributes of rodent population cycles.
5. Solving the enigma of population cycles may necessitate identifying factors and processes that cause phase-specific demographic changes, and performing conclusive experiments to ascertain the mechanisms that generate multiannual density fluctuations.

3 INTRODUCTION

Although periodic outbreaks in rodent numbers have been observed throughout recorded history, Elton (1924) was the first to scrutinise the phenomenon of multiannual population fluctuations scientifically. In this exceptionally insightful paper, Elton (1924) not only synthesised information on cyclic fluctuations in abundance of many animal species, but also extensively explored causes and consequences of population cycles with a particular focus on

lemmings and voles. He suggested that the causes of population cycles in lemmings ‘lie either with the lemmings themselves or with their environment’, but that the cause of periodicity must ‘lie with the environment’, in light of the spatial synchrony in lemming cycles. A careful reading of his paper reveals that Elton envisioned many of the modern ideas in animal ecology, including the concept of population regulation, prey switching, trophic interactions and cascades, and climatic forcing. In essence, Elton’s (1924, 1942) work on population cycles laid the foundation of

	
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1 the 'Eltonian' tradition of animal ecology, and influenced
2 generations of ecologists. Nearly a century has elapsed since
3 the publication of Elton's seminal work, but population
4 cycles remain enigmatic even today.

5 The often contentious history of population cycle re-
6 search has been thoroughly discussed in the literature,
7 and the scientific progress has been reviewed periodically
8 (e.g. Krebs et al. 1973, Tamarin 1978, Lidicker 1988, Batzli
9 1992, Krebs 1996, Stenseth 1999, Berryman 2002, Hanski
10 & Henttonen 2002, Lambin et al. 2002, Turchin 2003,
11 Kelt et al. 2018). Here, I provide a thorough review of
12 published research on rodent population cycles, highlight
13 interesting features of cyclic rodent populations, review
14 progress, discuss the possible reasons for our failure to
15 ascertain causes of population cycles, and identify knowl-
16 edge gaps that must be filled to identify necessary and
17 sufficient causes of population cycles.

19 CHARACTERISING RODENT POPULATION 20 CYCLES

21 The most prominent feature of cyclic populations are multi-
22 annual fluctuations in abundance with cyclic phases (increase,
23 peak, decrease and low phases) occurring every 3–5 years
24 (Krebs 2013). Various statistical approaches have been used
25 to characterise population cycles, including the s-index,
26 wavelet analysis, and time-series analysis (see Appendix S1
27 for details). Field evidence shows that most, if not all, cyclic
28 rodent populations are characterised by phase-related changes
29 in body mass, social behaviours, age structure, age at sexual
30 maturation, survival and reproductive rates (Chitty 1952,
31 1960, Krebs et al. 1973, Krebs & Myers 1974, Boonstra &
32 Krebs 1979, Boonstra 1994, Prevot-Julliard et al. 1999,
33 Norrdahl & Korpimäki 2002); together, they constitute the
34 'biological definition' of population cycles (Krebs 1996). At
35 high densities, sexual maturation is delayed, the length of
36 breeding season shortens, and juvenile survival rates and
37 reproductive rates are reduced, the proportion of young
38 animals in the population declines, and the mean age of
39 reproductive females increases; these changes precede and/
40 or accompany population crashes (Krebs et al. 1973, Boonstra
41 1994, Krebs 1996, Prevot-Julliard et al. 1999, Ergon et al.
42 2001a, b, Norrdahl & Korpimäki 2002). At low densities,
43 these patterns are reversed. Another feature of cyclic popu-
44 lations is the phase-related changes in body mass: individuals
45 are substantially heavier (up to 30% heavier) during the
46 high-density phase, and this is often referred to as the
47 'Chitty effect' (Chitty & Chitty 1962, Oli 1999). Also, social
48 behaviour may be affected (animals in high-density phases
49 are much more aggressive than those in low-density phases;
50 Krebs 1970, Cockburn 1988, Lidicker & Ostfeld 1991). These
51 syndromes of population cycles are sometimes ignored or
52 dismissed as unimportant by some theoretical ecologists

for convenience (e.g. Stenseth 1999, Hanski et al. 2001,
Turchin 2003), but most field studies show that these bio-
logical changes are nearly ubiquitous and indispensable
components of rodent populations that exhibit cyclic or
otherwise large-scale population fluctuations. Other features
of cyclic populations include latitudinal gradients in the
degree of cyclicity (e.g. populations in northern Europe
tend to be more cyclic than southern ones) and a broad
range of spatial synchrony within and among species
(Stenseth 1999, Krebs 2013). Spatial patterns in cyclic dy-
namics range from travelling waves (e.g. Lambin et al. 1998)
to highly synchronous fluctuations over large geographic
areas (e.g. Angerbjörn et al. 2001). Thus, any hypothesis
attempting to explain population cycles must explain phase-
specific changes in population characteristics as well as
broader spatial and temporal patterns in abundance that
define population cycles.

THEORETICAL FOUNDATIONS

The theoretical impetus for population cycle research has
been provided by two complementary ecological theories:
the theory of population regulation and the theory of preda-
tor–prey (or, more broadly, consumer–resource) dynamics.
The former embodies the 'density paradigm', and the latter
the 'mechanistic paradigm' of population ecology proposed
by Krebs (2002). The theory of population regulation pro-
vides a conceptual foundation for our understanding of
how biological populations are regulated, and attempts to
explain why populations fluctuate as they do. Factors or
processes that prevent unlimited population growth have
long been debated in the literature (Nicholson 1933,
Andrewartha & Birch 1954, White 2001, Turchin 2003),
but population regulation theory suggests that density-
dependent feedback mechanisms that permit populations
to grow at faster rates at low densities but reduce popula-
tion growth rates at high density are necessary and often
sufficient for preventing unlimited population growth
(Royama 1992, Turchin 1999, 2003). Although the impor-
tance of abiotic (density-independent) factors in limiting
populations is widely recognised, Royama (1992) argued
that population regulation necessarily implies density-
dependence, and that unregulated populations cannot persist.
The structure of density-dependence determines the pattern
of population fluctuations (Royama 1992). Direct (or first-
order) density-dependence typically leads to stable equilib-
rium whereas delayed (or second-order) density-dependence
can generate a variety of dynamical patterns, including cyclic
fluctuations, depending on the strength of direct and delayed
density-dependence. The primary analytical tools used within
this framework are time-series models such as autoregressive
or autoregressive-moving average models (Royama 1992,
Turchin 1999, 2003; Appendix S1).

The theory of predator-prey dynamics was motivated by the Lotka–Volterra predation model, which showed that interaction between prey and their specialist predators can cause cyclic fluctuations in prey abundance (Lotka 1924). Motivated by the cyclic dynamics generated by predator–prey models (Lotka 1924, May 1973), the theory of predator–prey dynamics assumes that interactions between predators and their prey underlie rodent population cycles, and focuses on understanding coupled dynamics of predators and the rodent prey. Because models of trophic–interactions or consumer–resource dynamics (predator–prey, herbivore–consumer and host–parasite) often exhibit cyclic dynamics, models of predator–prey, rodent–vegetation and rodent–parasite dynamics have been studied more extensively than any other (Hanski et al. 2001, Turchin 2003). Modelling frameworks adopted in mechanistic models of rodent population cycles include differential equation models (Hanski et al. 2001, Turchin & Batzli 2001, Reynolds et al. 2013), difference equation models (Inchausti & Ginzburg 1998, 2009), age-structured or stage-structured matrix population models (Klemola et al. 2003), individual-based population models (Radchuck et al. 2016), and integral projection models (van Benthem et al. 2017).

Capture–mark–recapture modelling approaches, especially those developed by Pradel (1996), are particularly useful for the study of cyclic populations (Pradel 1996, Nichols 2016) for several reasons. First, they permit direct estimation of realised population growth rate and its components: recruitment and apparent survival. Second, recruitment, apparent survival and realised population growth rates can be directly modelled as functions of temporal covariates (e.g. rodent and/or predator abundance, season, phase of a cycle, food addition) and site covariates (e.g. habitat quality). Third, this approach provides a framework for representing competing hypotheses as statistical models, and for using information-theoretic approaches for assessing empirical support for each of the competing hypotheses (Burnham & Anderson 2002). Fourth, questions about the relationship between sets of demographic parameters (e.g. abundance and survival) can be addressed in a manner that deals adequately with sampling variances and covariances associated with the different parameters. Unfortunately, these models have seldom been applied to the cyclic rodent populations (but see Yoccoz et al. 1998, Ozgul et al. 2004, Goswami et al. 2011). Long-term capture–mark–recapture studies of fluctuating populations are challenging, and it is particularly difficult to obtain adequate sample size during the low phase.

HYPOTHESES AND EMPIRICAL EVIDENCE

Krebs (2013) placed hypotheses of rodent cycles into five broad categories based on the causal factors invoked:

food, predation, disease, self-regulation and multifactorial hypotheses. Hypotheses of rodent cycles have also been organised under the banners of various historically significant ‘schools of thought’ (not mutually exclusive; Batzli 1992, Krebs 1996, 2013): 1) intrinsic school, which proposed that factors internal to the population (e.g. genetics, dispersal, social behaviour, stress response) are necessary and sufficient to cause population cycles (e.g. Chitty 1960); 2) extrinsic school, which proposed that extrinsic factors such as food, predators and parasites are the cause of population cycles (e.g. Hanski et al. 2001, Berryman 2002); 3) single factor hypotheses, which suggested that a single intrinsic or extrinsic factor is the primary cause of population cycle with additional factors playing only secondary roles (Courchamp et al. 2000, Hanski et al. 2001); and 4) multi-factorial school, which argued that rodent population cycles are too complex to be explained by single factor and that multiple intrinsic or extrinsic factors interact to cause population cycles (e.g. Lidicker 1988). Here, I briefly review relevant hypotheses and models (when appropriate), and summarise the empirical evidence, if any.

Predator–prey dynamics and the specialist predator hypothesis

Early models of predator–prey dynamics suggested that interaction between specialist predators and their prey can lead to cyclic population dynamics (Lotka 1924, May 1973). These results motivated the specialist predator hypothesis, which posits that a delayed numerical response of resident specialist predators to changes in prey abundance leads to multiannual population cycles. Models of predator–prey dynamics are the most intensively studied mechanistic models of population cycles (Hanski et al. 2001, Turchin 2003). Predator–prey models of cyclic populations are typically formulated as two-dimensional systems of differential equations. Various versions of predator–prey models have been developed and analysed; they differ mostly in the way functional response is modelled and in whether or not the effects of generalist predators are included (Turchin & Hanski 1997, 2001, Hanski et al. 2001, Turchin 2003). A version of the predator–prey model (Appendix S1), referred to as the Hanski model, can generate prey population dynamics that are strikingly similar to cyclic dynamics exhibited by voles in Fennoscandia. The Hanski model has been extended to include seasonality and environmental stochasticity; these extensions generate more realistic dynamics (Turchin & Hanski 1997, Hanski et al. 2001).

The predation hypothesis assigns a prominent role to predators and postulates that non-migratory, specialist predators are necessary, and a combination of predation

1 by specialist and generalist predators are both necessary
2 and sufficient, for causing rodent population cycles and
3 the latitudinal gradient in cyclic amplitudes observed in
4 Fennoscandia (Korpimäki & Norrdahl 1998, Hanski et al.
5 2001, Korpimäki et al. 2002, Lambin 2017). Specialist
6 predators are thought to be the primary drivers of popu-
7 lation cycles (hence, this idea is often referred to as the
8 specialist predator hypothesis), because it takes time for
9 specialist predators to respond numerically (via reproduc-
10 tion) to changes in prey abundance, which introduces the
11 time-lag required to generate second-order dynamics and,
12 consequently, population cycles. The diversity and abun-
13 dance of generalist predators are thought to determine
14 the cycle length and amplitude. Because generalist pred-
15 ators can affect vole populations instantaneously by increas-
16 ing predation rate (and thus reducing survival) as the
17 vole density increases (and the voles' own survival and
18 reproduction does not depend on the abundance of a
19 particular prey species), these generalist predators generate
20 direct density-dependence in vole population dynamics.
21 Together, specialist and generalist predators create a pat-
22 tern of autocorrelation in prey population dynamics that
23 is thought to be characteristic of 3–5 year rodent cycles
24 (Bjørnstad et al. 1995, Hanski et al. 2001, Turchin 2003).
25 The predation hypothesis has also been suggested to ex-
26 plain the latitudinal gradient in rodent population cycles
27 in Fennoscandia, because the degree of cyclicity increases
28 in northern latitudes, which generally corresponds with a
29 decrease in the diversity and abundance of generalist pred-
30 ators of cyclic rodents (e.g. Hanski et al. 1991, Bjørnstad
31 et al. 1995, Stenseth 1999).

32 The predation hypothesis has been tested by using ob-
33 servational and modelling studies, as well as field experi-
34 ments (reviewed by Hanski et al. 1991, 2001). Perhaps
35 the strongest support for this idea comes from the studies
36 conducted by Erkki Korpimäki's research group in Finland.
37 In a series of experiments, Korpimäki's research group
38 found (Norrdahl & Korpimäki 1995a, b, Korpimäki &
39 Norrdahl 1998, Sundell et al. 2000, Korpimäki et al. 2002):
40 1) nomadic avian predators did not substantially affect
41 vole abundance, and these predators tracked vole abun-
42 dance without time lag, leading to the conclusion that
43 these avian predators could not cause rodent cycles al-
44 though they might cause geographic synchrony in popula-
45 tion cycles; 2) experimental reduction in densities of all
46 main predators slowed vole population declines, whereas
47 populations continued to decline in control areas and at
48 sites where the abundance of only the specialist predator
49 (weasel *Mustela nivalis*) was reduced; and 3) reduction in
50 the abundance of all main predators during summer and
51 autumn increased the autumn vole density fourfold during
52 the low phase and twofold during the peak phase, and
53 retarded the initiation of the decline phase (Korpimäki

et al. 2002). These results were taken as evidence 'for the
hypothesis that specialist predators drive summer decline
of rodent populations in northern Europe' (Korpimäki &
Norrdahl 1998). Also, these findings and results of time
series analyses led Korpimäki et al. (2002) to conclude
that cyclic rodent populations may be transformed to
non-cyclic by manipulating predator abundance, and that
predators may generate the rodent population cycles ob-
served in Fennoscandia.

Empirical evidence for the role of specialist predators
(or predators generally) outside Fennoscandia are mixed.
Fauteux et al. (2016) conducted a fencing experiment in
Bylot Island, Canada, to test the idea that predators limit
cyclic brown lemming *Lemmus trimucronatus* populations.
Their results revealed that summer density and survival
of lemmings, as well as nest density during the winter,
were higher in experimental grids (i.e. with reduced pre-
dation) than in control grids, providing evidence that
predators may limit the lemming populations. However,
these results do not show that predators regulate lemming
populations in a delayed density-dependent fashion so as
to cause population cycles. A similar experiment at Pearce
Point, Northwest Territory, Canada (Reid et al. 1995)
revealed that, although the collared lemming population
protected from predators via fencing experienced higher
survival, it did not grow faster than the unprotected popu-
lation. These authors suggested their failure to exclude all
predators and the disruption of dispersal events as expla-
nations for these results.

The strongest evidence contradicting the specialist preda-
tor hypothesis comes from a long-term experiment con-
ducted in the Kielder Forest, northern England (Graham
& Lambin 2002). In a replicated field experiment, Graham
and Lambin (2002) experimentally reduced the density of
weasels, the specialist predator of the field vole *Microtus
agrestis* in the Kielder Forest, by live-trapping and remov-
ing them; they then monitored the abundance of both
voles and weasels. Results revealed that 1) weasel removal
did not prevent the crash of the population during the
decline phase of the cycle; 2) vole populations fluctuated
similarly in control and manipulated sites, and 3) there
was no evidence to suggest that the numerical response
of weasels to temporal changes in vole abundance was
delayed density-dependent, a condition necessary for pre-
dation to cause cyclic fluctuations in abundance. These
results led Graham and Lambin (2002) to conclude that
'changes in weasel predation rate were not responsible for
driving the population cycles of field voles observed in
Kielder Forest'. However, it is important to note that
Graham and Lambin's (2002) experiment reduced the
abundance of the specialist predator, but not of generalist
predators that prey on voles to varying degrees. Also,
Kielder Forest vole cycles are thought to be somewhat

1 different from those in Fennoscandia (Korpimäki et al.
2 2005). Thus, Graham and Lambin's (2002) results only
3 contradict the specialist predator hypothesis as a universal
4 explanation of population cycles; they do not necessarily
5 rule out the possibility that specialist predators interact
6 with other intrinsic and extrinsic factors to generate popu-
7 lation cycles in Kielder Forest or elsewhere.

8 9 10 **Vegetation–rodent dynamics and the food 11 hypothesis**

12 The idea that bottom-up processes could regulate rodent
13 numbers, either directly through changes in quantity or
14 quality of food, or indirectly via soil nutrients, has been
15 around for a long time. Periodic overexploitation of food
16 resources by rodents can potentially cause populations to
17 crash due to starvation; time required for vegetation to
18 regrow or soil nutrients to be replenished could introduce
19 the delayed effect needed to cause population cycles.
20 Consequently, rodent–vegetation interactions have long
21 been considered as an important factor causing, or con-
22 tributing to, population cycles (Summerhayes 1941, Pitelka
23 1957, 1964).

24 Turchin and Batzli (2001) developed and analysed a
25 suite of rodent–vegetation models (represented as two- or
26 three-dimensional differential equations) with the goal of
27 understanding quantitative characteristics of rodents and
28 their food plants that can cause herbivore cycles. They
29 assumed that recovery of vegetation following overgrazing
30 can take time, which can introduce time-lags that are
31 necessary for a consumer resource system to generate cyclic
32 dynamics. They considered a variety of scenarios, because
33 many aspects of rodent–vegetation dynamics are not well
34 understood (e.g. the recovery or growth pattern of rodent
35 food sources following overgrazing, the functional response
36 of rodents to changes in vegetation abundance). Although
37 some of these models generated cyclic population dynam-
38 ics, none yielded periodicity of 3–5 years. Because vegeta-
39 tion–herbivore dynamics can be affected by rodent–predator
40 dynamics, they also considered multitrophic models (veg-
41 etation–rodents–specialist predator dynamics), which gen-
42 erated cyclic rodent dynamics with realistic periodicity
43 (Turchin & Batzli 2001). However, they advised caution
44 in interpreting these results.

45 The first rigorous experiment to test the food hypothesis
46 was conducted by Summerhayes (1941) under the supervi-
47 sion of Charles Elton. Some experimental plots were ex-
48 cluded from rodent herbivory using fences; the vegetation
49 within and outside the enclosures was compared to test
50 the idea that overgrazing precedes population crash. The
51 results revealed that vegetation within and outside of ex-
52 closures did not differ substantially, and that there was
53 no evidence of overgrazing or of starvation prior to the

population crash (Chitty 1996, Krebs 2013). Exclusion of
collared lemmings *Dicrostonyx groenlandicus* and brown
lemmings *Lemmus trimucronatus* and other herbivores by
fencing actually reduced diversity and biomass of vascular
plants near Barrow, Alaska, USA (Johnson et al. 2011).
A study using experimental enclosures found that winter
grazing by collared lemmings and brown lemmings had
no discernible effect on the biomass of vascular plants or
moss in the Canadian Arctic (Bilodeau et al. 2014). There
has been no consistent evidence that overgrazing affects
food quality, that overgrazing adversely influences rodent
reproduction, or that overgrazing necessarily reduces rodent
abundance. Contrary to expectations, Klemola and col-
leagues (Klemola et al. 2000a, b) found that voles in
previously overgrazed enclosures reproduced just as well
as those inhabiting control sites (i.e. areas with no previ-
ous overgrazing), and that vole abundance was higher in
previously overgrazed areas than in control sites.

A variant of the food hypothesis is the nutrient–recovery
hypothesis, originally proposed to explain population cycles
of brown lemmings in Barrow, Alaska (Pitelka 1964). The
nutrient–recovery hypothesis states that high rodent den-
sities decimate the vegetation, leading to starvation and a
population crash. Lemming populations do not experience
population growth until soil nutrients are accessible to
plants, plant production and accumulation of litter have
increased and nutrient levels in food plants return to
normal levels (Pitelka 1964, Pitelka & Batzli 2007). This
idea was tested using a fertilisation experiment in Alaska,
which showed that lemming populations did not respond
positively to the experimental fertilisation (Pitelka & Batzli
2007). A similar fertilisation experiment failed to produce
expected results in another cyclic herbivore, the snowshoe
hare *Lepus americanus* in the Kluane region of Canada
(Krebs et al. 1995).

A more direct approach to test the food hypothesis is
to provide supplemental food to experimental populations,
with the expectations that food supplementation should
stop or drastically alter population cycles (Krebs 2013).
Several food supplementation experiments have been con-
ducted to test this idea (e.g. Cole & Batzli 1978, Taitt &
Krebs 1981, 1983, Getz et al. 2006; for reviews, see Boutin
1990, Prevedello et al. 2013). Taken together, results of
food supplementation studies show that food supply af-
fects many aspects of rodent ecology (e.g. density, move-
ment and spacing behaviours, and in some cases, re-
production) but food supplementation does not stop or
substantially alter population cycles, suggesting that changes
in food supply by itself are not necessary or sufficient
for population cycles to occur. However, Johnsen et al.
(2017) reported that bank voles *Myodes glareolus* went
extinct in all study plots that did not receive supplemental
food during the second winter of the experiment, whereas

1 populations persisted in plots receiving supplemental food.
2 This result and evidence from multifactor experimental
3 studies suggests that food supply can interact with other
4 factors, such as predation, to affect population cycles (see
5 below).

6 Many plants produce secondary compounds (e.g. silica,
7 phenolic compounds) as a deterrent to or defence against
8 herbivory, and these defence mechanisms may affect her-
9 bivore survival and reproduction (Huitu et al. 2014,
10 DeAngelis et al. 2015, Hartley & DeGabriel 2016). Massey
11 et al. (2008) found that grazing by field voles at high
12 vole densities led to increased silica levels in a winter
13 food plant, and eating food with high silica levels led to
14 reduced vole body mass. Based on these findings, Massey
15 et al. (2008) suggested that inducible, silica-based defences
16 in grasses may drive field vole population cycles. Grazing
17 by field voles (Reynolds et al. 2012) and root voles *Microtus*
18 *oeconomus* (Wieczorek et al. 2015) has been shown to
19 increase silica levels in their food plants; the silica levels
20 are lagged and a nonlinear function of the grazing inten-
21 sity. Using coupled grass-vole differential equation models,
22 Reynolds et al. (2012, 2013) showed that delayed, silica-
23 based plant defensive responses to grazing intensity can,
24 in theory, generate population cycles. However, there exists
25 no empirical evidence that grazing-induced changes in
26 silica (or other measures of food quality) are necessary
27 for, or substantially affect, rodent population cycles (Ruffino
28 et al. 2018).

30 Host–parasite dynamics and the disease 31 hypothesis

32 Rodents, like many animal species, are often infected by
33 parasites (or communities of parasites) that can potentially
34 affect survival, reproduction or both, which can in turn
35 affect population dynamics. This possibility did not escape
36 the notice of population cycle pioneers; indeed, recurring
37 epidemics were one of the first possible explanations of
38 population cycles in rodents (Chitty 1996, Krebs 2013).
39 Although early studies based on primitive detection meth-
40 ods indicated that parasites such as *Toxoplasma* could be
41 important in regulating vole populations in the UK (Elton
42 et al. 1935), they were rejected as plausible explanations
43 of population cycles due to lack of evidence (Elton et al.
44 1935, Chitty 1954). Diseases and parasites have been studied
45 in several cyclic rodents (e.g. *Myodes* spp. and *Microtus*
46 spp. and *Arvicola* spp.), with some studies showing that
47 diseases can affect demographic parameters and others
48 showing the lack of evidence for such effects (e.g. Telfer
49 et al. 2005, Cerqueira et al. 2007, Deter et al. 2007, 2008).

50 **4** Smith (2008) modified the classic ‘susceptible, infected
51 and recovered’ model of host–pathogen dynamics to in-
52 clude two types of recovered host: hosts that have recovered
53

from the disease and gained immunity but cannot yet
reproduce; and hosts that have recovered and can repro-
duce, but at a rate lower than the healthy individuals
(Appendix S1). Their host–parasite model was a system
of four differential equations, and assumed that the disease
influenced both survival and reproduction. They concluded
that diseases with brief infection periods but slow recovery
of reproductive function once hosts recover from the dis-
ease can generate high-amplitude, multiannual population
fluctuations. Based on these modelling results and the field
evidence that some diseases of field voles (or infection of
voles by multiple potentially interacting parasites) show
density-dependent cyclic fluctuation, the authors offered
vole–parasite interactions as a potential explanation of
population cycles in northern England (Smith et al. 2008,
2009).

Perhaps the most extensive study of the potential effect
of diseases on the dynamics of cyclic populations was
conducted by the research groups of Xavier Lambin, Sandra
Tefler and Michael Begon in the UK. Studies of field
voles in northern England showed that voles are infected
by an interacting community of parasites, that some of
these parasites show density-dependent (and sometimes
cyclic) prevalence patterns, and that some of these infec-
tions can adversely affect voles’ population parameters
(Cavanagh et al. 2004, Telfer et al. 2005, 2011). Using
models of host–parasite dynamics parameterised with field
data, Smith et al. (2008, 2009) showed that diseases can
potentially cause cyclic fluctuation in vole abundance. These
studies also revealed complex interactions among com-
munities of pathogens mediated by competition for re-
sources and host immune response (Begon et al. 2009).
However, most studies of the effects of disease on cyclic
rodent population dynamics have been observational, and
there has been no convincing experimental evidence that
diseases or parasites cause rodent population cycles.
Experimentally testing disease effects on cyclic population
would necessarily require manipulating a community of
parasites or vaccinating voles; such experiments would be
logistically challenging and have not been attempted (but
see Smith et al. 2008, Turner et al. 2014).

Maternal effects model and hypothesis

Inchausti and Ginzburg (1998, 2009) suggested that ma-
ternal effects, defined as non-genetic transmission of quality
from mother to offspring, can cause population cycles,
and developed a model to formalise this idea. The Inchausti-
Ginzburg model assumes that the ‘quality’ of mothers
(vaguely defined) changes in a density-dependent fashion,
and, in turn, influences population dynamics. The model
is formulated as two systems of difference equations (each
system consisting of two difference equations), one to

1 model the coupled dynamics of rodent population and
2 the quality of mothers in spring, and another to model
3 the same dynamics in autumn. The 'quality' of mothers
4 is assumed to be transmitted from mothers to offspring,
5 the phenotypic transmission of quality from others to
6 offspring. However, the periodicity of cycles generated by
7 this model is generally inconsistent with those observed
8 in nature for realistic parameter values. Turchin (2003)
9 notes that realistic parameter values lead to 1–2 year cycles,
10 and that unrealistically high survival rates (>0.95 per
11 month) would be needed to generate cycles with periodic-
12 ity of 3–5 years.

13 Of several intrinsic (or self-regulation) hypotheses pro-
14 posed to explain rodent population cycles, one (the Chitty
15 hypothesis) has been rejected (Boonstra & Boag 1987),
16 and evidence for others have been mixed or inconclusive
17 (Appendix S2). Except for the maternal effect hypothesis,
18 mathematical or theoretical models representing intrinsic
19 hypotheses do not currently exist. Based on the findings
20 of a reciprocal transplant experiment showing that past
21 environments had no substantial effect on field vole life
22 history variables, Ergon et al. (2001a) rejected intrinsic
23 mechanisms as an explanation of population cycles. Because
24 intrinsic factors are difficult to manipulate, experimental
25 tests of intrinsic hypotheses have been rare (Krebs 2013).

26 27 **Models invoking multiple factors and the** 28 **multifactorial hypotheses** 29

30 Rodent population dynamics are inherently complex, and
31 highly variable over time and space. Populations of the
32 same species exhibit multiannual cycles at some times and
33 in some places, but seasonal or erratic fluctuations at other
34 times or in other places. These universally accepted ob-
35 servations, combined with the fact that experiments in
36 which one or two factors were manipulated have failed
37 to explain population cycles satisfactorily, led Lidicker
38 (1978, 1988, 2000) to conclude that rodent population
39 cycles are too complex to be explained by one or two
40 factors and that multiple factors interact to cause the
41 observed population cycles. He rejected arguments regard-
42 ing whether intrinsic or extrinsic factors are the primary
43 cause of population cycles as unhelpful, proposed that
44 four intrinsic and four extrinsic factors are involved in
45 driving the demographic machinery underlying cyclic popu-
46 lation dynamics in California voles *Microtus californicus*,
47 and argued that the relative roles of these factors vary
48 across density phases and over time and space. The mul-
49 tifactorial perspective is conceptually appealing, especially
50 in light of the failure of hypotheses invoking one or two
51 factors to explain population cycles satisfactorily. In ad-
52 dition, this perspective can explain annual or multiannual
53 population fluctuations, spatio-temporal variations, and

biological attributes of cycles. Experimentally manipulating
many factors simultaneously can be logistically challenging;
consequently, there have been few experimental tests of
the multifactor hypothesis in its original form (Krebs 2013).
Nonetheless, the potential influence of two or three factors
thought to be the key drivers of population cycles have
been tested using both mathematical models and
experiments.

Klemola et al. (2003) developed and analysed demo-
graphically based, density-dependent, stage-structured ma-
trix population models that incorporated the potential
influence of vegetation, specialist predators and generalist
predators on population dynamics of voles and lemmings.
They showed that tri-trophic interactions can produce
cyclic changes in abundance. However, the modelled dy-
namics did not adequately capture the shape and amplitude
of cycles observed in cyclic rodent populations; the authors
suggested that assumptions about phase-dependence in
trophic interactions or some population-intrinsic factors
would be needed to generate realistic population cycles
(Klemola et al. 2003). Therefore, trophic interactions may
be necessary but not sufficient to cause population cycles
similar to those observed in nature. This latter observation
is consistent with the idea that multiple factors may act
synergistically to generate population cycles (Lidicker 1978,
1988, Andreassen et al. 2013).

Using an individual-based modelling approach (Railsback
& Grimm 2011), Radchuck et al. (2016) tested models **5**
representing combinations of two intrinsic factors (sociality
and dispersal) and one extrinsic factor (predation) for
their potential to generate population cycles comparable
to those observed in the field. They found that only the
full model – including sociality, dispersal and predation
– yielded population cycles with periodicity, amplitude
and autumn densities comparable to those observed in
North Fennoscandia. This study is the only study that
used an individual-based modelling approach with empiri-
cally estimated parameters and simultaneously considered
both intrinsic and extrinsic factors.

The interactive effects of two or three factors on rodent
population dynamics have been tested experimentally. The
most common of these experiments manipulated the two
obvious potential drivers of rodent cycles: food and preda-
tors. Results of these experiments have revealed consistently
that the population-level effect of food supplementation
and predator exclusion (or predator removal) was almost
always greater than the singular effect of either treatment
(Klemola et al. 2000a, Huitu et al. 2003, Krebs 2013,
Prevedello et al. 2013). It is conceivable that primary
productivity or food supply modulates the effect of preda-
tors on prey populations (e.g. Oksanen & Oksanen 2000).
However, none of these experiments succeeded in stopping
or substantially altering cyclic population dynamics. Forbes

et al. (2015) experimentally tested for the effect of food and a pathogen (*Bordetella bronchiseptica*) on field vole population growth rate. Contrary to expectations, they found that the vole population that received supplemental food was more severely affected by the experimentally introduced pathogen than the population that did not. Few studies have tested for effects of three or more factors on cyclic rodent populations (but see Taitt & Krebs 1983, Krebs et al. 1995, Batzli et al. 2007).

DEMOGRAPHIC MECHANISMS: THE MISSING LINK

Changes in population size are caused by changes in survival, reproduction and immigration rates, which, in turn, are functions of extrinsic and intrinsic factors (Caswell 2001, Krebs 2002, Oli & Armitage 2004). Thus, cyclic changes in abundance are caused by cyclic changes in population growth rate, which are, in turn, a consequence of cyclic changes in vital demographic rates (Caswell 2001, Krebs 2002). Indeed, it is widely accepted that population cycles are characterised by phase-related changes in several life-history parameters. Arguably, the identification of demographic mechanisms underlying population cycles can facilitate ascertainment of factors and processes that cause multi-annual fluctuations in abundance (Oli & Dobson 1999, 2001, Krebs 2013). As an example, consider a population where phase-related changes in survival are the primary determinants of population growth (e.g. Korpimäki & Norrdahl 1998, Korpimäki et al. 2002). This knowledge is sufficient to exclude factors such as social behaviour or stress response as possible determinants of population cycles, because these factors are unlikely to cause direct mortality, and to conclude that natural enemies (e.g. predators or pathogens) are the likely cause of the phase-related changes in survival, and thus, of population cycles. Consider, on the other hand, a population where phase-related changes in age at first reproduction are the primary demographic drivers of population growth (Oli & Dobson 1999, 2001, Norrdahl & Korpimäki 2002, Krebs 2013). This information is sufficient to exclude predation as the cause of population cycles, because direct predation is unlikely to influence age at sexual maturation. Indeed, phase-related changes in reproductive parameters, rather than survival rates, are thought to be the demographic drivers of vole population cycles in Kielder Forest, England and in Chize, France (e.g. Smith et al. 2006, 2008, Ergon et al. 2011, X. Lambin, personal communication). This knowledge and logical reasoning leads to the conclusion that social factors (e.g. social suppression of reproduction, kinship, spacing behaviour), stress response or food resources are the likely drivers

of population cycles. Finally, consider a situation where population crashes are characterised by sharp increases in predator abundance, but rapid population growth is accompanied by early sexual maturation and a preponderance of young animals in the population (e.g. Boonstra 1994). This knowledge allows one to isolate the mostly likely environmental causes of population cycles and devise experiments to test them.

Unfortunately, phase-specific demographic patterns are yet to be rigorously quantified in most cyclic rodents; we do not yet know what demographic changes underlie transitions from one phase to another. Data are particularly scarce, and our knowledge is limited, for decline and low phases (Krebs 2013). A research programme that integrates the identification of demographic mechanisms of population cycles with environmental causes of demographic changes could help devise focused experiments that may resolve the enigma of rodent population cycles (Oli & Dobson 1999, 2001).

WHAT HAVE WE LEARNED AND WHERE DO WE GO FROM HERE?

Despite the recognition of rodent population cycles as significant ecological phenomena, we have failed to determine the underlying causative factors thus far. This failure is not for the lack of trying; theoretical and empirical research has produced many hypotheses and 1000s of publications (Batzli 1992, Krebs 2013). What have we learned from nearly a century of research on population cycles, and what do we need to do to solve the enigma of population cycles? A brief summary follows.

1. We do not yet know what factors and processes are necessary and sufficient to cause population cycles in rodents, because no experiment has succeeded in stopping population cycles anywhere, although some have prevented or delayed the population crash. Given the complexity of the ecological systems of which cyclic rodents are a part, experimentally stopping population cycles (although desirable) is perhaps unrealistic; however, the hypothesised factor(s) must substantially alter cyclic dynamics for it to be a necessary cause of population cycles.
2. Modelling and experimental studies in Fennoscandia have established predation by specialist and generalist predators as the most likely cause of population cycles in northern Europe. However, tests of the predator hypothesis elsewhere have yielded mixed results, suggesting that predation is not a universal explanation of population cycles or that population cycles do not have a universal explanation.

3. A large body of theoretical work has led to a plethora of mathematical models that can generate cyclic population fluctuations. These theories and models have provided many insights, and have highlighted the role of seasonality and the potentially interacting effects of predators, parasites and intrinsic factors.
4. Although it is widely recognised that social factors and predator-induced stress responses can strongly influence rodent population processes, there have been very few experimental tests of intrinsic hypotheses of population cycles. For example, the chronic stress hypothesis, as modified by Rudy Boonstra's research group (Boonstra et al. 1998, Sheriff et al. 2017, Peers et al. 2018), is amenable to experimental testing and should be tested in cyclic rodent populations (see also Appendix S2).
5. Results of a modelling study by Radchuck et al. (2016) suggested that both intrinsic and extrinsic factors may be necessary to explain rodent population cycles adequately, yet few attempts have been made to test the multifactorial hypothesis due to logistical difficulties.
6. It is widely accepted that population cycles are accompanied by several life-history and behavioural traits that vary across cyclic phases. Whether these changes are necessary for population cycles to occur, or are simply epiphenomena associated with fluctuating populations, remains unknown and deserves further research.
7. Loss of cyclicity and/or dampening of cyclic amplitudes has been reported in some cyclic rodents in Europe. It now appears that the collapse of population cycles or dampening of cyclic amplitudes were localised and temporary phenomena (Appendix S3, Brommer et al. 2010).
8. Earth's climate is changing; climate change will inevitably affect the distribution and abundance of many plant and animal species via changes in the abiotic environment, primary productivity and species interactions (Boonstra et al. 2018). There exists some evidence that climate change may dampen the dynamics of some cyclic rodent populations (e.g. Kausrud et al. 2008, Schmidt et al. 2012, Cornulier et al. 2013). However, the direction or magnitude of climate change effects on population cycles remains unclear.
9. Despite a long history of population cycle research, many data gaps remain. Few researchers have quantified the phase-specific demography of cyclic populations, and we know very little about the decline and low phases of the cycle. Filling these data gaps, and making complete demographic descriptions of population cycles (which are essentially population processes), is critical for further progress in population cycle research.

Some theoreticians have argued that theories and models are sufficient to understand and explain population cycles,

or that population cycles have been fully explained by predator-prey models (e.g. Turchin 2003). This view-point is rejected by empirical ecologists, who emphasise the importance of experiments and long-term field studies (e.g. Krebs 2013). Population cycle research would greatly benefit from a better integration of theoretical and empirical work: experiments guided by theories and testable theories that explain observed facts are needed. The application of statistically rigorous modelling approaches, such as capture-mark-recapture analyses, has increased in recent years. Capture-mark-recapture methodologies now permit direct estimation and modelling of population growth rate, as well as age- and state-specific survival and reproductive rates. These approaches should be helpful in discerning demographic mechanisms underlying population cycles. Other unresolved questions in population cycle research include (also see Krebs 2013): 1) is there a universal explanation of rodent cycles? 2) Are there key experiments that can solve the enigma of population cycles once and for all? What are they? 3) Should experiments attempt to stop population cycles? Or is it enough to alter population growth rates? And 4) what constitutes the conclusive test of a hypothesis proposed to explain population cycles?

As carefully articulated by Krebs (2013), the process of explaining population cycles must begin with rigorous and quantitative characterisation of phase-specific demography. This knowledge can help exclude implausible explanations from further consideration. Population cycle research over the last 30 years has been focused almost exclusively on numerical changes, with little attention for the biological attributes of cycles. A demographically based research agenda that integrates numerical dynamics with phase-related demographic changes, and experiments that manipulate both intrinsic and extrinsic factors, will be needed to solve the enigma of population cycles, an ecological problem that fascinated Charles Elton and generations of ecologists. Population cycle research has a contentious history (Tamarin 1978, Stenseth 1999, Krebs 2013), since most leading research groups in the past seemed to subscribe to a specific idea. Rejection of data or ideas that do not support a particular view-point seemed to occur more often in population cycle research than in most other ecological disciplines. I concur with Krebs' (2013) suggestion that a dose of humility would benefit us all, because 'no one knows where we will finish, and so much still remains to be done'.

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SUPPORTING INFORMATION

Additional supporting information may be found in the
online version of this article at the publisher's
web-site.

Appendix S1. Details of autoregressive time series models,
and models of predation and diseases.

Appendix S2. Intrinsic hypotheses of population cycles.

Appendix S3. Are population cycles dampening or collapsing?

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