Warming decouples eco-evolutionary feedback between body size and abundance in long

term natural ecosystems

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Abstract

Eco-evolutionary feedback, in which abundance influences trait evolution and evolving

traits, in turn, affect abundance dynamics, play a critical role in maintaining community

structure, function, and stability. A key foundation of this feedback is the bidirectional

relationship between body size and abundance. However, the integrity of this relationship is not

fixed but varies with climate warming and biotic factors. Using long-term monitoring data from

101 ecological systems spanning over 10-47 years and encompassing multiple generations of

plankton, fish, and terrestrial beetles, we provide the first synthetic evidence that warming

decouples body size–abundance feedback and this decoupling is stronger in smaller-bodied

species. Our findings suggest that warming disrupts trait–demography interactions and indicate

that the eco-evolutionary feedback may be vulnerable to anthropogenic pressures.

Keywords: Body size-abundance feedback, Eco-evolutionary feedback, Climate Warming,

Species traits.

Introduction

Eco-evolutionary feedback—the reciprocal interactions between ecological and

evolutionary processes—are increasingly recognized as a key driver of biodiversity patterns

population stability, species interactions, and ecosystem functioning (Fronhofer *et al.* 2023;

Govaert *et al.* 2019; Loreau *et al.* 2023). The feedback arises when traits under selection alter

ecological dynamics, which conversely modify selection pressures, creating feedback loops that

shape long-term species adaptation to changing environments (De Meester *et al.* 2019;

Zamorano *et al.* 2023). This feedback bridges the traditional divide between ecology and

evolution, showing that evolutionary change can occur rapidly enough to influence contemporary

ecological dynamics (Thrall & Blanc 2023).

At the heart of eco-evolutionary feedback lie functional traits that mediate both

ecological performance and evolutionary responses to environmental change (De Meester *et al.*

2019; Pantel & Becks 2023). These traits influence key population processes—such as

demographic rates and resource use—and thereby shape ecological dynamics (De Meester *et al.*

2019). The traits often evolve in response to environmental pressures, generating feedback that

further modify ecological processes (De Meester *et al.* 2019). Among them, body size is

particularly influential due to its pervasive effects across levels of biological organization, and

eco-evolutionary feedback often scale with body size, because of shorter generation times

facilitating faster evolutionary responses (De Meester *et al.* 2019). Evolutionary shifts in body

size affect survival, reproduction, and competitive ability, ultimately shaping population

abundance (Pelletier *et al.* 2009). Reciprocally, changes in abundance alter selection on body

size: high abundance intensifies resource competition, favoring smaller, more efficient

individuals, whereas low abundance can favor larger body sizes due to advantages in

competition, mate acquisition, or predator avoidance (Pelletier *et al.* 2009). The bidirectional

relationship between body size and abundance thus forms a key foundation of eco-evolutionary

feedback operating within and across ecological communities, shaping community structure,

buffering ecological systems against environmental fluctuations (Ahti *et al.* 2020; Carlson *et al.*

2011; Edeline & Loeuille 2021; Malerba *et al.* 2018).

Importantly, the strength of body size–abundance feedback is not fixed (Govaert &

Klauschies 2024; Han *et al.* 2023). It varies with environmental changes, biotic interactions, and

evolutionary dynamics (Fouqueau & Polechová 2024; Kaitala 2021). In stable external

environments, tightly coupled body size–abundance relationships can reinforce predictable eco-

evolutionary dynamics, as trait shifts propagate through demographic processes and ecological

feedback loops (Fouqueau & Polechová 2024). However, in the context of global change, the

robustness of these relationships is increasingly uncertain.

Climate warming, one of the most pervasive anthropogenic stressors, fundamentally

alters ecological and evolutionary processes, with body size frequently implicated as a trait

sensitive to warming (Gardner *et al.* 2011; Govaert & Klauschies 2024). Numerous studies

document shifts toward smaller body sizes in response to rising temperatures—a pattern

consistent with metabolic scaling theory, resource limitation, and life-history trade-offs (Gardner

et al. 2011; Sheridan & Bickford 2011). At the same time, warming can affect population

abundance via changes in reproduction, survival, and resource dynamics (Visser 2008; Walsh *et*

al. 2019). Despite extensive research on how warming affects body size and abundance

independently, whether climate change disrupts the coupling between these two processes—and

thus potentially erodes the eco-evolutionary feedback—remains unclear. Furthermore, species

diversity and traits can influence these two processes, driving shifts in both body size and

abundance. However, whether they disrupt the coupling between these two processes remains

unknown.

In this study, we present the first systematic evaluation of how climate warming

influences the feedback between body size and population abundance, potential effect on eco-

evolutionary feedback. We synthesize a continental-scale dataset encompassing 101 long-term

monitoring sites across aquatic and terrestrial systems—including plankton, fish, and ground

beetle species—spanning 10 to 47 years and encompassing multiple generations for each species.

Using this dataset, we evaluated the presence of bidirectional links between body size and

abundance through Convergent Cross Mapping (CCM) (Sugihara *et al.* 2012), a method for

detecting causal links (Fig. 1b). Last, we assessed how the proportion of bidirectional links

responds to warming and biotic drivers. Taken together, this study provides insight into whether

climate warming and biotic factors disrupt body size–abundance feedback, with implications for

body size–mediated eco-evolutionary feedback.

Methods

We compiled a long-term dataset encompassing 101 ecosystems: 51 aquatic planktonic

systems monitored for 10–40 years, 30 fish systems spanning 31–47 years, and 20 terrestrial

systems (beetles) spanning 9–11 years. The duration of these time series captures multiple

generations for each species. The inclusion of multiple generations enables the detection of

evolutionary changes in body size, beyond plastic or pure ecological responses. Such temporal

resolution captures concurrent shifts in ecological dynamics (e.g., abundance) and evolutionary

trait change (e.g., body size), fulfilling the criteria necessary to assess eco-evolutionary feedback.

Aquatic plankton Data. Body size and abundance data were compiled from 51 long-

term monitoring sites across aquatic planktonic ecosystems, comprising 20 lakes, 8 rivers, and 23

marine systems (Fig 2a). These datasets were accessed by North Temperate Lakes Long-Term

Ecological Research (Magnuson *et al.* 2023a, 2024a, 2022, 2023b), Pangaea (Felden *et al.* 2023),

USGS (Nejad *et al.* 2017), Greenland Ecosystem Monitoring (Christensen *et al.* 2017) and Data

Observation Network for Earth (Michener *et al.* 2011). The 51 ecosystems have been

consistently sampled for 10–40 years. The resident species include phytoplankton and

zooplankton, with maximum generation times typically less than a week and two months,

respectively (Elbrächter 1976; Gillooly 2000; Litchman 2000). Therefore, the 10–40 years of

sampling encompass numerous generations of these planktonic species. For each of the 51

aquatic planktonic ecosystems, there were three types of variables: (1) species abundance; (2)

species size (i.e. body length, biovolume, or biomass); (3) water temperature. For species size

(2), a single metric is reported for each species and subsequently used for further analysis.

Next, for consistency, each species' time series were standardized to seasonal sampling

frequencies (if multiple samplings were conducted per season, e.g., monthly and bimonthly

sampling data sets, those were averaged). Here, the seasonal resolution (trimonthly) is the shared

consensus that can be applied to all 51 data sets, and the seasonal average here is also the most

representative measure across all data sets, because the equal sampling interval is necessary for

Convergent Cross Mapping (CCM) analysis. After the seasonal average, there were 0.05%

missing points across the whole planktonic dataset (41 out of 85161 data points). Those missing

data points were linearly interpolated using *na.approx* function in the package of zoo (Karakoç *et*

al. 2020; Zhao *et al.* 2023). Then, the species abundance across all data sets was scaled to the

same unit (individuals per litre). Species length, biovlume and biomass were standardized to the

same unit (μm), (μm^3) and (mg/m^3) respectively.

Last, we removed the long term linear and seasonal trend from each time series.

Specifically, we first removed the long-term linear trend from each time series by using the

residuals from a linear regression against time (Chang *et al.* 2020, 2022). Next, we removed

seasonality by scaling against the mean and standard deviation of values occurring in the same

year, $D_{season}(t) = (O(t) - \mu_{season})/\sigma_{season}$, where $D_{season}(t)$ is the deseasoned time series,

$O(t)$ is the original time series, μ_{season} is the year mean, and the σ_{season} is the year standard

deviation (Chang *et al.* 2020, 2022). Finally, each time series (both species abundance and size)

was scaled to zero mean and unit variance.

Fish Data. Fish size and abundance data were compiled from 30 monitoring sites,

including six rivers, five lakes and 19 marines. These datasets were accessed by North Temperate

Lakes Long-Term Ecological Research (Magnuson *et al.* 2024c, b), International Council for the

Exploration of the Sea (ICES) (Fish trawl surveys 2025), and Reef Life Survey (Edgar & Stuart-

Smith 2014). These ecosystems have been consistently sampled for 31–47 years. Among all fish

species, Atlantic cod exhibit the longest generation time (≤8 years), medium-sized sportfish

range from approximately 5–7 years, and the remaining small, fast-reproducing species have

generation times of less than 2 years. Thus, the 31–47 years of sampling encompass numerous

generations across all species. Again, for each Fish ecosystem, there were three types of

variables: (1) species abundance; (2) species size (i.e. body length); (3) water temperature.

For consistency each species time series of abundance and size were standardized to the

yearly sampling frequency (if multiple samplings were conducted per year, e.g., quarterly

sampling data sets, those were averaged). Again, the yearly resolution is shared across all data

sets. After the yearly average, there were no missing points across the fish dataset. Then, the fish

abundance across all data sets was standardized to the same unit (caught per effort). Species

length was standardized to the same unit (mm). As all fish data were standardized to yearly

intervals, there was no need to remove long-term linear or seasonal trends from each time series.

Therefore, the original data were utilized for further analysis. Finally, each time series (both

species abundance and size) was scaled to zero mean and unit variance.

Terrestrial species. Terrestrial species size and abundance data were compiled from 20

monitoring sites, all focused on ground beetles. All data were retrieved from BioTIME (Dornelas

et al. 2018). These ground beetles were consistently sampled for 10–11 years. Given a maximum

generation time of 1–3 years, the sampling period encompasses numerous generations across all

species. Again, for each site, there were three types of variables: (1) species abundance; (2)

species size (i.e. biomass, mg/m^3); (3) surface temperature. Each site reports both abundance

and size. The daily temperature and precipitation for each site were retrieved from NOAA's

National Centers for Environmental Information (GHCN-Daily, Version 3 2012).

Next, to ensure consistency, each species' time series were standardized to seasonal

sampling frequencies (if multiple samplings were conducted per season, those were averaged).

After calculating seasonal averages, 0.52% of data points were missing in the dataset. These

missing values were subsequently linearly interpolated using the *na.approx* function from the

zoo package (Karakoç *et al.* 2020; Zhao *et al.* 2023). Then, long-term linear and seasonal trends

were removed from each time series using the same methodology applied to the plankton data.

Finally, each time series, including both species abundance and size, was standardized to have a

mean of zero and a unit variance.

Detecting the bidirectional link among body size and abundance. To ensure the

applicability of empirical dynamic modeling (EDM), we included only species with sufficiently

long and continuous time series data (i.e. length $n \geq 30$) (Sugihara *et al.* 2012). For each species

(plankton, fish or terrestrial species), we identified the bidirectional causal link among species

size and its abundance (i.e. size →abundance, abundance →size) using convergent cross-

mapping (CCM) (Fig 1). CCM is based on Takens's theorem for dynamic ecosystems, which

infers the causal relationship among variables from their empirical time series (Deyle & Sugihara

2011; Sugihara *et al.* 2012) (see three brief videos for a summary introduction:

http://tinyurl.com/EDM-intro). If the causal variable X (e.g. species abundance) and affected

variable Y (e.g. species size) belongs to the same dynamical system, information on the causal

variable X is encoded in the affected variable Y. Thus, one can predict the states of causal

variable X using the affected variable Y. CCM infers causality by measuring the extent to which

the causal variable X has left signatures in the time series of the affect variable Y; a procedure

known as cross mapping (cross-prediction) (Sugihara *et al.* 2012). In this study, the appropriate

embedding dimensions for cross-mapping E were determined by the optimized the hindcast

cross-mapping in which X(t) projected one-step backward to Y(t-1), examining values of E from

2 to square root of n, where n is the length of the time series (Karakoç *et al.* 2020). Thus, we

computed n as the geometric mean time series length across all data sets. E was finally examined

from 2 to 7 across all data sets. We used simplex projection to select the best E that gave the

highest prediction skill (Hsieh *et al.* 2005).

To accommodate the fact that the causal variables (e.g., species abundance) can exhibit

time-delayed effects on the affected variables (e.g., species size), we carried out 0 to 1 year (i.e.

0~4 time point for plankton and terrestrial species but 1 time point for fish) lagged CCM

analyses, in which we retained the CCM with the time lag resulting in the highest cross mapping

skill ρ (Ye *et al.* 2019). The criterion for testing causality is the convergence toward higher

cross-mapping skill with increasing library length (i.e., the number of points used for state space

reconstruction) (Sugihara *et al.* 2012). An increase in library length enhances the density of

points in the reconstructed attractor, facilitating more precise identification of nearest

neighboring points for predictions. This, in turn, leads to improved predictive accuracy (Sugihara

& May 1990). Convergence can be assessed by evaluating whether there is a significant

monotonic increasing trend in cross-mapping skill ρ as library length increases using Kendall's τ

test (Chang *et al.* 2020), and whether the ρ at the largest library length is significantly greater

than that the ρ at the smallest library length using Fisher's Z test (Chang *et al.* 2020). In this

study, library length was set from the minimum (E) to the maximum length (i.e., the length of the

entire time series). Throughout this study, an interaction link (e.g., X→Y) was considered causal

if both Kendall's τ test and Fisher's Z test were statistically significant ($P < 0.05$) for testing

convergence (Chang *et al.* 2020).

Finally, the warming rate, representing the long-term warming trend of water temperature

or land surface temperature in each ecosystem, was measured using the Theil–Sen median-based

trend estimator, implemented via 'sens.slope' from 'trend' package (Chang *et al.* 2020; Mohsin

& Gough 2010). This method was chosen due to its widespread use in quantifying warming rates

(Pinsky *et al.* 2025; Tong *et al.* 2023), its robustness to episodic extreme events, and its ability to

provide stable estimates of long-term trends in the presence of outliers (Mohsin & Gough 2010).

Species mean body mass was computed as average across whole time series, estimated from

biovolume, assuming that organisms have the same density as water, if unavailable, was obtained

from published sources (Cohen & Lough 1981; Huntley 1992).

Bidirectional links in responses to climate change and biotic drivers. We employed a

generalized linear mixed model (using the glmmTMB function from the glmmTMB package

(Brooks *et al.* 2017)) with a binomial error distribution to test the main effects of warming, mean

temperature, species richness, mean species body mass, and their pairwise interactions on the

proportion of observed links—either from body size to abundance (size →abundance) or from

abundance to body size (abundance → size). To ensure normality, mean temperature, species

richness, and mean species body mass were log-transformed using base 10. In the model, the

proportion of each link was used as the response variable, consisting of binary values: 1

indicating the presence of a causal link and 0 indicating the absence of a link. Last, we set

species groups (e.g. plankton) nested within sample locations (e.g. Trout Lake) as random

intercepts. Details of statistics see Table S1.

All analyses were done with R (ver. 4.1.2). The CCM analysis was examined using the

rEDM package (version 1.2.3).

Results

We used a binary response variable to indicate the presence (1) or absence (0) of a link

between body size and abundance, with links identified using Convergent Cross Mapping.

Across 101 ecological systems monitored over 10 to 47 years—encompassing multiple

generations for each of 5858 species—our analysis revealed that warming decouples body size–

abundance feedback, as evidenced by a negative association between warming and the

proportion of both link types (abundance → size and size → abundance) (Fig 3a-d). This pattern

suggests that warming can reduce the potential for body size–abundance feedback, potentially

weakening body size based eco-evolutionary feedback.

Notably, the effect of warming body size–abundance feedback varied with species body

size and species richness. Warming more strongly decoupled body size‑abundance feedback in

smaller-bodied species, as evidenced by steeper negative slopes for both abundance → size and

size → abundance links (Fig 3a-b). In contrast, this decoupling effect was weaker and in some

cases reversed—among larger-bodied species (Fig 3a-b), indicating a potential shift toward

feedback coupling. These findings suggest that warming may more strongly weaken body size–

mediated eco-evolutionary feedback in smaller-bodied species, while potentially reinforcing it in

larger-bodied species. However, species richness showed no consistent effects on body size‑

abundance feedback. Warming more strongly decoupled the size → abundance link at higher

species richness (Fig 3c), while its decoupling effect on the abundance → size link was stronger

under lower species richness (Fig 3d). Last, we found mean temperature had no effect on either

link of abundance → size or size → abundance (Table S1).

Discussion

Our study provides the first systematic evaluation of body size–abundance feedback—a key foundation of eco-evolutionary dynamics at population and community levels—in response to environmental change, and demonstrates the potential for warming to disrupt these feedbacks. By analyzing long-term field monitoring data from 101 ecosystems—each encompassing multiple generations per species—we captured sufficient temporal resolution to detect potential evolutionary responses. Our results show that warming decouples body size–abundance feedback, suggesting that warming temperatures can decouple body size–mediated eco-evolutionary dynamics. Notably, this decoupling effect can be stronger in smaller-bodied species, indicating that trait dependent eco-evolutionary feedback to climate warming.

Body size evolve rapidly to environmental changes (Merckx *et al.* 2018) and can exert a significant influence on population abundance (Martins *et al.* 2023). Previous experimental studies have documented that the body size and abundance feedback are tightly linked in single-species populations (Gibert *et al.* 2022, 2023), a two-species competitive interactions (Govaert & Klauschies 2024), and a microbial food web (Han *et al.* 2023). Our study further showed that warming tended to decouple body size and abundance feedback (Fig 3). Previous evidence has documented weaker body size–abundance feedback at higher temperatures in an experimental protist food web. (Han *et al.* 2023). This may be attributed to the differing response rates of body size and abundance to warming (Genner *et al.* 2010). Higher temperatures can lead to a significant reduction in species body size (Audzijonyte *et al.* 2020; Terry *et al.* 2022), while it can have slight effect on population abundance (Sebastian *et al.* 2012). Similarly, higher temperatures have shown to accelerate growth, leading to an rapid increase in population abundance (Brandenburg *et al.* 2019; Han *et al.* 2023), while simultaneously either leaving body size unaffected or slightly altered body size (Crozier *et al.* 2010; Genner *et al.* 2010). These different responses rate among size and abundance may weaken the link between them (Han *et al.* 2023).

We found that warming more strongly decoupled body size–abundance feedback in smaller-bodied species, whereas this effect was weaker—and in some cases reversed—among larger-bodied species. This pattern suggests that warming may more strongly weakens body size–based eco-evolutionary feedback in smaller-bodied species, while potentially reinforcing it in larger-bodied species. To date, no empirical evidence has been documented to validate this

trend. Given their shorter generation times and higher sensitivity to environmental fluctuations

(Gillooly 2000; Zeuss *et al.* 2017), these species may exhibit more rapid but less stable trait–

abundance dynamics under warming, increasing the likelihood of feedback decoupling.

In contrast, at higher levels of species richness, warming more strongly decoupled the

link of size → abundance, while exhibiting a weaker decoupling effect on link of abundance →

size. Previous experimental studies have demonstrated that increased species richness reduces

the proportion of links of size→ abundance (Govaert & Klauschies 2024). Here, enhanced

species richness can subject organisms to a broader spectrum of ecological interactions, such as

intensified interspecific competition and predation, which can disrupt the link of size→

abundance within species (Govaert & Klauschies 2024). However, there is currently no empirical

evidence documenting the interactive effects between warming and species richness, nor

established mechanisms to explain their interactions on both links.

Our analysis further revealed that, unlike indoor experiments where body size and

abundance are nearly always linked (Gibert *et al.* 2022, 2023), this feedback is less tightly

coupled in natural ecosystems (Fig. S1). Across 101 ecosystems, 25.1% of species exhibited a

causal link from size to abundance, while 20.3% showed a link from abundance to size. The

weak coupling between body size and abundance suggests that body size based eco-evolutionary

dynamics may be loosely linked at natural ecosystems. Moreover, the bidirectional feedback

between body size and abundance appears asymmetrical, with a higher proportion of size →

abundance links compared to the reverse (25.1% vs. 20.3%, Fig S1). Taken together, these results

suggest that body size and abundance may exhibit independent, rather than simultaneous,

responses to environmental change in natural systems. Indeed, previous evidence has

documented independent shifts in body size and abundance in response to urbanization (Merckx

et al. 2018) and habitat fragmentation (Warzecha *et al.* 2016; Zastavniouk *et al.* 2017). Species'

size can even provide greater accuracy and serve as an earlier warning signals for predicting

population collapses than abundance, under environmental deterioration (Clements *et al.* 2017,

2019; Clements & Ozgul 2016).

In summary, long-term field data spanning over 10 years and covering multiple

generations provides sufficient temporal resolution to detect evolutionary responses in natural

systems. While controlled experiments are well-suited for isolating causal eco-evolutionary

feedback, the long-term field studies used in this analysis are essential for capturing these

dynamics in natural ecosystems under sustained environmental change. By revealing that climate warming decouples body size and abundance relationships, this study emphasizes the vulnerability of size-based eco-evolutionary feedback to anthropogenic pressures.

Competing interests

The authors declare no competing interests.

Author Contributions

Q.H.Z., F.D.L. and Y.X.G.W designed the research. Q.H.Z. and Y.X.G.W performed the study and analyzed the data. Q.H.Z and G.S performed the CCM analysis. Q.H.Z., F.D.L. and Y.X.G.W led the writing, with contributions from C.X. and G.S.

Data and code availability

The data reproducing all the results in this study are available on GitHub (https://github.com/QZhao16/link.abundance-size).

Acknowledgements

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Figures

Figure 1. **An overview of the methods and sample locations**. **a**, The sample location for
plankton, fish and terrestrial beetles. **b**, Conceptual illustration of the bidirectional feedback
between body size and abundance. Evolutionary shifts in body size influences key fitness
components—survival, reproduction, and competitive ability—that shape population abundance.
In turn, changes in abundance modify selection pressures on body size through effects on
resource competition, mate acquisition, and predator avoidance. This reciprocal interaction
among body size and abundance forms a core mechanism underpinning eco-evolutionary
feedback. In this study, we investigate body size-abundance feedback at the individual focal
species (e.g. *Gambusia affinis*) rather than across species.

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Figure 2. **The body size and abundance feedback in response to warming**. Feedback

identified by CCM were coded as 1, and the absence of a link as 0. The solid lines and shaded

error bands represent the significant best-fit trendlines and corresponding 95% confidence

intervals from two-sided logistic regressions across 101 ecosystems. Black points represent the

binary link values (0 or 1) between abundance and body size for each species across 101

ecosystems (total points per panel: n = 5858). Points are vertically jittered to reduce overlapping.

A generalized linear mixed model with a binomial error distribution was used to test

the effect of warming on the proportion of each link (size → abundance or abundance

→ size). The response variable was binary, with 1 indicating the presence of a link and

0 indicating its absence. Species groups (e.g., fish) nested within sampling locations

(e.g., Trout Lake) were included as random intercepts. The effect of warming is visualized

by using the ggpredict function from the ggeffects package. ***P< 0.001, **P<0.01, *P<0.05.

Figure 3. **The body size and abundance feedback in response to biotic drivers. a-d,** The
feedback in response to the interactive effects of warming and mean species body mass (**a–b**)
and interactive effects of warming and species richness (**c–d**). Feedback identified by CCM were
coded as 1, and the absence of a link as 0. The solid lines and shaded error bands represent the
significant best-fit trendlines and corresponding 95% confidence intervals from two-sided
logistic regressions across 101 ecosystems. Points represent the binary link values (0 or 1)
between abundance and body size for each species across 101 ecosystems (total points per panel:
n = 5858). Points are vertically jittered to reduce overlapping. The interaction among predictors
(**a-d**) is visualized using the ggpredict function from the ggeffects package. For statistical results
see table S1.

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Supplementary Figures

Figure S1. **Proportion of body size–abundance feedback across 101 ecosystems. a,** At the

species level, mean proportions of causal links are shown for aquatic plankton, fish, and

terrestrial species. **b**–**c**, At the ecosystem level, frequencies and proportions of causal links from

abundance → size (**b**) and size → abundance (**c**) across 101 ecosystems. For detailed

proportions of causal links within each ecosystem, see Figures S2–S3.

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Figure S2. **The bidirectional link between species size and abundance in each of aquatic**

ecosystems. Red points indicate causal links identified by CCM, while blue points indicate the

absence of a link. Letters on the x-axis represent sampled ecosystems.

Figure S3. **The bidirectional link between species size and abundance in each of terrestrial**

ecosystems. Red points indicate causal links identified by CCM, while blue points indicate the

absence of a link. Letters on the x-axis represent sampled ecosystems.

Table S1. Generalized Linear Mixed Model (glmmTMB function in glmmTMB package) with a binomial error distribution was employed to test the main effects of warming, mean temperature, species richness, species mean body mass and pair wise interaction among warming, species richness and species mean body mass on proportion of either link (size →abundance or abundance →size). In the model, the proportion of each link was treated as the response variable, consisting of binary values: 1 indicating the presence of a causal link and 0 indicating the absence of a link. This model set species groups (e.g. fish) nested within sample locations (e.g. Trout Lake) as random intercepts. Mean temperature, species richness, and mean species body mass were log-transformed using base 10 to ensure normality. ***$P<$ 0.001, **$P<0.01$, *$P<0.05$.

	proportion of size → abundance			proportion of abundance → size		
	coefficient	Z	P value	coefficient	Z	P value
(Intercept)	-1.25	-6.95	**<0.001*****	-1.81	-10.90	**<0.001*****
Species richness	-0.01	-0.09	0.93	0.31	2.19	**0.03***
Warming	-0.88	-4.90	**<0.001*****	-1.71	-8.06	**<0.001*****
Mean body mass	0.18	1.25	0.21	0.35	2.32	**0.02***
Mean Temperature	0.03	0.23	0.82	-0.07	-0.56	0.57
Species richness* Warming	-0.35	-2.25	**0.02***	0.79	3.24	**0.001****
Warming* Mean body mass	0.73	4.15	**<0.001*****	1.41	5.79	**<0.001*****

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