1   **Biodiversity modulates the size-abundance relationship in changing**

2   **environments**

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ABSTRACT

Organismal abundance tends to decline with increasing body size. Metabolic theory links this size

structure with energy use and productivity across levels of biological organization, predicting a size-

abundance slope of -0.75 that is invariant across environments. We tested whether the size-abundance

relationship is robust to a gradient of protist species richness (1 to 6 species), temperature (15 to 25°

C), and time. Our results support the expected slope of -0.75 for the size-abundance relationship, but

we found interactive effects indicating that the size-abundance relationship is not invariant. In high-

richness communities, temperature increased the abundance of small protists more than that of the

large protists, leading to a steeper size abundance slope. Temperature and size-dependent species

interactions were identified as potential drivers. Understanding variation in size-abundance

relationship hence provides novel insight into the underlying mechanisms shaping the size structure of

ecological communities under environmental change.

KEYWORDS: body size, protists, species richness, size-abundance relationship, temperature

INTRODUCTION

Smaller species are usually more abundant than larger species, that is, they have larger populations. The body size-abundance relationship, a well-established empirical pattern, describes this inverse relationship between the average body size of a species and its abundance (Damuth 1991; White *et al.* 2007). The slope of the size-abundance relationship is typically -0.75 (Brown *et al.* 2004; Nee *et al.* 1991). Body size-abundance relationships have become increasingly important in community ecology as they provide insights into the links between the size structure and abundance of communities (Arim *et al.* 2011; Enquist *et al.* 1998; Giometto *et al.* 2013; Gjoni *et al.* 2023; Jacquet *et al.* 2020; Li 2002; Long & Morin 2005; Meehan *et al.* 2004; Morán *et al.* 2010; White *et al.* 2004, 2007). The size-abundance relationship can also connect community-level ecological processes with the effect of various environmental factors and hence may be used to predict how communities will be reshaped by environmental change (Gjoni *et al.* 2023; Gjoni & Glazier 2020).

Theory postulates that size-abundance relationships link community size structure with ecosystem functioning through "energy equivalence" (Brown *et al.* 2004; Nee *et al.* 1991) when body size and abundance scale inversely to metabolic rate. This contrasting pattern was first recognized by Damuth (1981) who showed that abundance scaled with body size according to mass$^{-0.75}$, while metabolic rate is assumed to scale with mass with a power of 0.75. Energetic equivalence implies that the energy flux per unit area—calculated as the product of metabolic rate and abundance—remains constant and independent of body size. In other words, energy flux per unit area is independent of body size, i.e., mass^{0}, implying that population energy use is independent of body size. Energy equivalence is grounded in the notion that the abundance of a population directly depends on the availability of their resources. An increased resource input should thus lead to a corresponding increase in population abundance.

Despite these theoretical predictions, it remains unclear whether the expected size-abundance relationship is robust to environmental change and alterations in biodiversity. Body size and abundance are responsive to biotic and abiotic factors, which may lead to changes in the size-abundance scaling when either the abundance of small or large organisms is particularly affected (Gjoni *et al.* 2023). Indeed, deviations from the expected -0.75 slope for the size-abundance have been observed (Arim *et al.* 2011; Gjoni *et al.* 2017, 2023; Gjoni & Basset 2018; Gjoni & Glazier 2020; Long & Morin 2005; Meehan *et al.* 2004), suggesting that energy used may not be the same across all organism sizes (Nee *et al.* 1991) and that the size-abundance scaling is not as invariant as assumed. However, the abiotic and biotic drivers of size-abundance scaling are rarely systematically investigated.

Warming often leads to higher abundance of smaller organisms relative to larger ones, leading to a change in size-abundance scaling (Gjoni *et al.* 2023; Gjoni & Glazier 2020; Morán *et al.* 2010;

Pomeranz *et al.* 2022). The metabolic theory of ecology (MTE) predicts that increased temperatures accelerate biochemical reactions, speeding up metabolism exponentially and hence requiring more energy to maintain basic physiological functions (Brown *et al.* 2004; Gillooly *et al.* 2001). As a consequence, warmer conditions decrease the carrying capacity (i.e. biomass at equilibrium) because the same energy supply can only support a smaller number of individuals (Allen *et al.* 2002). If temperature affects all organisms similarly regardless of size, as assumed by the MTE, then temperature should not alter the size-abundance relationship (Fig. 1a). The temperature-size rule describes that organisms often exhibit smaller body sizes in warmer conditions (Adams *et al.* 2013; Atkinson 1995; Daufresne *et al.* 2009; Finkel *et al.* 2010; Gardner *et al.* 2011; Sheridan & Bickford 2011). This reduction in size leads to lower absolute energy requirements, hence more individuals can be supported by the same amount of energy. However, if all organisms shrink similarly due to temperature, as implied by the temperature-size rule, we do not expect any change in the size scaling. However, empirical evidence shows that the expected -0.75 slope in the size-abundance relationship often varies (Gjoni *et al.* 2023; Morán *et al.* 2010; Perkins 2022; Pomeranz *et al.* 2022; Saito *et al.* 2021). Furthermore, it is known that species interactions are sensitive to the environmental context (reviewed by Tylianakis *et al.* 2008), in particular, to temperature (Burnside *et al.* 2014) and size (Freckleton & Watkinson 2001).

Few studies on size-abundance relationships have so far explicitly considered the biotic context of, for example, community composition and species richness (but see Long *et al.* 2006; Long & Morin 2005). Biodiversity can play a critical role in shaping the size-abundance relationship due to species interactions that determine community resource use and structure. Research on biodiversity-ecosystem functioning has, for example, revealed that increased biodiversity, particularly species richness, consistently leads to higher community abundance (Hooper *et al.* 2005; Tilman *et al.* 2001). Higher community abundance can be driven by the selection effect where highly productive species are more likely to occur in species-rich communities, and the complementarity effect, where resource partitioning among species can reduce competition and lead to higher community biomass (Blackburn & Gaston 1996; Tilman *et al.* 2001). How community size structure is related to biodiversity effects is less well understood. Size variation of, for example, competitive communities is often small (Tilman *et al.* 2004) and there is little evidence that size structure is linked to complementarity or selection effects (but see Norkko *et al.* 2013) suggesting that size-abundance scaling should not be affected by species richness per se (Figure 1a). However, larger organisms are often competitively superior, as increased richness intensifies asymmetric resource competition, but the ecological context can mediate whether larger or smaller species are favored (Lawton & Hassell 1981; Gómez & González-Megías 2002). Therefore, high richness may lead to either a shallower or steeper size-abundance slope depending on whether larger or smaller species gain a competitive edge in more species-rich communities (Fig. 1c).

Temperature and diversity can also interactively shape the size-abundance relationship, if they
disproportionally affect the interactions between larger and smaller organisms (Fig. 1b). Higher
temperatures tend to reduce the number of larger individuals, due to their higher metabolic demands,
creating a competitive advantage for smaller individuals due to their lower resource requirements
(Petchey *et al.* 1999). Temperature may also counteract some of the positive biodiversity effects
observed at higher richness due to direct negative physiological effects on biomass, leading to
shallower or even negative biodiversity-abundance relationships (Parain *et al.* 2019; Pennekamp *et al.*
2018). Since interactive effects of temperature and diversity on the size-relationship are difficult to
disentangle in observational data, dedicated experimental approaches are required to address this
question.

Effects of biotic and abiotic factors on the size-abundance scaling may further vary through
time, since ecological communities are complex, adaptive systems which exhibit temporal dynamics
in response to both biotic and abiotic factors (McGrady-Steed *et al.* 1997). Over time, changes in
temperature and biodiversity may alter the relative proportions of organisms of different sizes within a
community. Due to the equilibrium assumptions of the energy equivalence (Brown *et al.* 2004; Nee *et*
al. 1991), we would expect the slope to converge to -0.75 in the long run. Indeed, experimental
studies have shown that closed communities with controlled species richness (e.g., low vs. high
richness) converge to a stable size-abundance relationship (Fig. 1d; Long & Morin 2005). In contrast,
natural communities can exhibit fluctuating size-abundance relationships over time, driven by
dynamic species interactions and environmental variability (Fig. 1e; White *et al.* 2004) that deviate
from the expected slope.   Whether temporal changes in the size-abundance scaling are further
modulated by environmental change is currently unknown.

Here, we investigated the size-abundance relationship across temperature and biodiversity
(i.e., initial species richness) in controlled experimental microcosms over a six-week period
(Pennekamp *et al.* 2017, 2018). All protist communities were initially started from the same set of
species, thus changes in size and abundance were integrating both phenotypic change in size and
abundance. Based on a null model informed by the MTE, we hypothesize the following (1) size-
abundance relationship follows a slope of -0.75, (2) this slope remains unchanged regardless of
temperature or species richness, with only the intercepts varying, and (3) the slope converges to -0.75
over time. To test these predictions, we tracked the cross-community size relationship across
communities varying in richness (from 1 to 6 species) and temperature (from 15 to 25°C). We
examined cross-community size relationships that aggregate size and abundance at the community
level, by relating the average individual size of a community with its total abundance across different
communities (White *et al.* 2007). We assessed the role of species interactions by comparing
experimentally assembled multi-species communities to simulated "communities without species
interactions," where single-species data were pooled to match the richness and composition of the

experimental communities. Unlike previous observational studies, this controlled approach allowed us
to directly test how temperature and diversity jointly shape size-abundance scaling. Our results
confirmed that the initial slope of the size-abundance relationship approximated -0.75, but
temperature and diversity interactively altered it, challenging its assumed invariance. Notably, the
slope was stable at first but became shallower over time, suggesting that species interactions mediate
the effects of temperature, diversity, and temporal dynamics on size-abundance scaling.

Figure 1 Conceptual overview of size-abundance relationships across temperature and diversity. a) Temperature
and diversity may affect abundance and size regardless of the size of individuals, which may lead to changes in
the intercept, but not the scaling. In contrast, smaller or larger individuals may be favored at b) higher (T1) or
lower temperature (T2), or c) lower or higher species richness, leading to changes in the size-abundance scaling.
Over time, the size-abundance relationship may be stable or converge to the expected slope of -0.75 (d), or show
more variable trajectories with endpoints that differ from the expected slope.

MATERIALS & METHODS

We used microcosms to test the effect of temperature and species richness on size-abundance scaling.
To this end, we factorially manipulated temperature (15, 17, 19, 21, 23, and 25 °C) and species
richness (1 to 6 species) in communities of six bacterivorous ciliate species: *Colpidium striatum,*
Dexiostoma campylum, Loxocephalus sp*., Paramecium caudatum, Spirostomum teres,* and
Tetrahymena thermophila. These species span a cell length gradient ranging from ~30 μm to 400 μm.

The species were cultured in organic protist pellet medium on which the freshwater bacterium
Serratia fonticola was inoculated as the sole protist food source. Due to the non-sterile handling of
cultures, additional bacteria may have colonized the microcosms throughout the experiment. The
organic medium consisted of protist pellets (Carolina Biological Supplies, Burlington, NC) at a
concentration of 0.55 g per L of Chalkley's medium, and two wheat seeds for slow nutrient release
(for details on cultivation and experimental procedures, see also Altermatt *et al.* (2015)). Stocks of
protist cultures (monocultures) were established before the start of the experiment to inoculate the
microcosms. Single species cultures (richness = 1) were started at a density of three individuals per
mL in 100 mL medium kept in 250 mL Duran bottles. Multispecies communities with species
richness ranging from 2 to 6 species were initiated as a fixed fraction of the species-specific carrying
capacity. Therefore, communities were assembled at fixed initial richness. Since communities were
closed to dispersal, richness can only decrease, and species may be lost due to extinctions. We
replicated each level of species richness and composition at least twice for all levels including an
additional replicate for the lowest and additional three replicates for the highest levels of species
richness resulting in 115 experimental units per temperature. Cultures communities were randomly
assigned to their temperature level and placed in climate-controlled incubators (Pol-Eko Aparatura,
Wodzislaw, Poland) set to 15, 17, 19, 21, 23, or 25 °C. We used two independent incubators for each
temperature level.

To measure abundance and size of individual cells, we sampled each experimental unit every
day for the first 7 days, then 3 times per week for the following 50 days, and a final sampling 7 days
later. Each sample was analyzed with automated video species classification analysis (Pennekamp *et*
al. 2015, 2017) where we measured the abundance and size of every population using five-second
videos (at 25 frames per second). Using the video microscopy data we calculated the average size
(i.e., volume) and abundance of each species (Supplementary material section S1). We calculated
individual cell volume based on cell length and width measurements using the following equation:
cell volume = 4/3 * pi * width /2) ^ 2 * length/2 (Laakso *et al.* 2003). Volume is reported in
microlitre. Abundance was calculated by counting the number of individuals of a given species on the
50th frame of a video, thus avoiding double counting of individuals, and then upscaled to density per
mL.

Monocultures and polycultures had different starting densities; therefore, we aligned the time
series to make the dynamics directly comparable. To do so, we only consider monoculture time series
after reaching 20% of their carrying capacity, in correspondence with the polyculture time series as
done previously (Pennekamp *et al.* 2018). The alignment did not influence the qualitative patterns
observed over the full experimental period. Time series were cut to have a maximum length of 42
days. Alignment of mono- and polyculture time series led to differences in the time points where
abundance and size were measured. We used cubic hermite splines to interpolate the mean abundance
and mean volume of species in each experimental microcosm. We manually checked that

interpolations closely agreed with the size and abundance dynamics observed. In rare cases, when
either negative abundances or volumes resulted from the interpolation, we discarded the observation.
For all subsequent analyses (size-abundance scaling and its temporal dynamics), we only considered
every second time point to minimize temporal autocorrelation and to not artificially increase the
sample size due to the interpolation. We then calculated the cross-community size relationship
relating total community abundance (sum of the abundance of all species in the community) and the
weighted mean size of an individual in the community (White *et al.* 2007).

We used a regression approach to quantify the relationship between abundance (response) and
size, richness and temperature in our controlled microcosm communities. We scaled size, temperature
and richness treatments (by subtracting the respective mean from all values) to facilitate the
interpretation of model coefficients (Schielzeth 2010). The random structure for all models consisted
of a random intercept for community composition and a random intercept for incubator. We then
perform four analyses: 1) we estimated the overall cross-community scaling relationship across protist
communities, regressing total abundance on mean individual size (White *et al*. 2007). 2) We regressed
total abundance on mean individual size, temperature, and richness as well as all interactions between
main effects averaged over time. We included all possible interactions of the 3-way interactions
models (cell volume x temperature x richness). 3) We fitted the previous model (3 main effects and all
possible interactions, same random effects) to each time point to understand how estimates change
through time. 4) To understand the role of species interactions on the size-abundance scaling
relationship, we simulated polycultures without interactions by pooling monoculture data with the
same richness and composition as the real polycultures that we studied experimentally. We then used
the same model as 3) to compare the effects of size, richness, and temperature, as well as their
interactions between the simulated and real communities. Deviations in coefficient estimates would be
putatively driven by species' interactions. All analyses were performed in the statistical computing
environment R (version 3.6.1).

RESULTS

Interacting temperature and richness effects on size-abundance relationship
Averaging over time, we found a strong negative relationship between total abundance and average
cell volume (b = -0.71, CI = -0.777 to -0.644) across community richness and temperature treatments,
undistinguishable from the expected value of b = -0.75 (Fig. 2A). Although deviations from the linear
fit are visible, they disappear when data was stratified by richness (Fig. 2B).
Including temperature and richness in the model (Table 1), the size-abundance slope (b = -
0.749 CI = -0.813 to -0.685) perfectly matched the expected value. However, temperature and
richness jointly changed the effect of size on abundance (volume:temperature coefficient = -0.015, CI
= -0.024 to -0.005 and volume:richness coefficient = -0.055, CI = -0.100 to -0.010). The effect of size
on abundance was furthermore mediated by a three-way interaction (volume:temperature:richness

coefficient = -0.012, CI = -0.018 to -0.007). In other words, at low richness, the abundance of small

and large protists was not affected by temperature (similar slope), whereas in more species rich

communities, temperature increased the abundance of smaller protists relative to larger ones, resulting

in a steeper size-abundance scaling (Fig. 3, Table 1). For a complementary view, stratifying by

richness within temperature, refer to section S2 of the supplementary material.

Figure 2 The relationship between centered average log10 cell volume ($\mu m\ \mu L^{-1}$) and average $\log_{10}$ total

abundance of protist communities approximated with a simple linear fit (A). Deviations are visible at the lowest

(< -0.5 log cell volume) and largest average cell sizes (> 0.5 log cell volume) but disappear when we stratify by

richness (B).

Table 1 Temperature and species richness interact to affect the size-abundance relationship across protist

communities. All predictors were mean centered for the analysis. Confidence intervals (CI) not overlapping with

zero are significant. All terms including the effect of volume on abundance are shown in bold.

Term	Estimate [95% CI]
Intercept	4.564 [4.524, 4.603]
volume	-0.749 [-0.813, -0.685]
temperature	-0.019 [-0.023, -0.014]
richness	0.080 [0.051, 0.109]

volume:temperature	-0.015 [-0.024, -0.005]
volume:richness	-0.055 [-0.100, -0.010]
richness:temperature	-0.003 [-0.004, -0.001]
volume:temperature:richness	-0.012 [-0.018, -0.007]

Figure 3 Temperature and richness interact to alter size-abundance relationships of protist communities. The dashed line indicates the expected size-abundance scaling of -0.75. The figure shows the fitted size-abundance scaling based on the model presented in Table 1.

The size-abundance relationship through time

At the beginning of the experiment, the size abundance slope was steeper (b = -0.9) than expected but then stabilized around the expected slope of b=0.75 until day 20. Afterwards the size-abundance slope became considerably shallower than expected (b = -0.35). Similar temporal patterns were observed when we used community evenness accounting for species extinctions and changes in relative abundance (see Section S3). In other words, initially, communities with small average individual size tended to have higher abundance, then, average community size led to expected abundance, while at the end, communities with higher average size had higher abundance than expected (Fig. 4A).

Figure 4: Temporal dynamics of effect sizes (and 95% confidence intervals) of a linear mixed-effects model where abundance is modelled as a function size (V = volume), temperature (= T) and richness (= R) and their interactions for the real (observed) and simulated communities (in blue and red, respectively). The x-axis shows the day. A) Effect of cell size (V) on abundance, B–C) effect of two-way interactions between size and temperature (V x T), size and richness (V x R) on abundance. D) Three-way interaction between size, temperature and richness (V x T x R) on abundance. The dashed lines indicate a slope of -0.75 in panel A and zero in panels B–D (a deviation from zero would indicate a slope modification).

Temporal analysis of the size-abundance slope revealed that temperature (V x T) initially
promoted larger organism sizes (Figure S3). However, after day 15, higher temperatures favored
smaller organisms. By the end of the experiment, the temperature effect was no longer significant. In
contrast, species richness (V x R) initially reduced the effect of cell size. After day 20, the effect
reversed, with richness promoting larger organisms. Finally, the interactive effect of temperature and
species richness (V x T x R) on size structure was negative after day 10, became positive from days
14 to 20 and 25 to 35, and was non-significant in between. These changes resulted in changes in the
predicted slopes through time, with high richness communities being more strongly affected by
temperature than low richness communities (see section S4). Finally, we compared the temporal
dynamics of the real, observed polycultures in which species engage in positive or negative
interactions, with simulated polycultures without such species interactions (Fig. 4). At the beginning
of the experiment, simulated communities had a steeper size-abundance slope than expected,
indicating a higher relative abundance of smaller organisms. After day 14, the slope of simulated
communities became shallower, approaching -0.40 by day 20, in contrast to the real communities that
maintained a slope of -0.75 until day 20 (Fig. 4A). By day 30, the slope of simulated communities
returned to the expected value, while real communities became increasingly shallow. The combined
effect of temperature and richness (V x R x T) showed no significant influence throughout the
experiment, indicating that the impact of temperature on the size-abundance slope was not dependent
on community richness, unlike the more dynamic responses seen in real protist communities (Fig.
4D).

DISCUSSION

Our study revealed that the size-abundance relationship of protist communities is mediated by
temperature, biodiversity (i.e., initial species richness), and time. The shift in the size-abundance slope
with temperature and diversity challenges the widely held assumption that size-abundance
relationships remain stable within a trophic level across time and space (Damuth 1981; Enquist &
Niklas 2001; Niklas & Enquist 2001). As expected, we found a negative relationship (b ∼ -0.75)
between abundance and body size across species richness, and temperature, in line with the expected
slope of the energy equivalence (Damuth 1991; Nee *et al.* 1991). This is consistent with other studies
that have used cross-community size scaling to test energetic equivalence and found the same slope
value across phytoplankton (Li 2002), macroinvertebrate, amphibian, and fish (Arim *et al.* 2011)
communities, as well as across rodent (White *et al.* 2004) and tree communities (Enquist *et al.* 1998).
The size-abundance relationship is typically mirrored by the metabolic rate, which follows a 0.75
scaling exponent with mass (Brown *et al.* 2004), implying a slope of -0.75 for the size-abundance
relationship. This alignment is seen in natural and experimental phytoplankton communities, where
the scaling slope is close to -0.75 (Malerba *et al.* 2017; Perkins *et al.* 2019). In contrast, an

experimental study using protist communities initially divided into low and high species richness
showed no change over time   with a cross-community size scaling of -1, which converged over time
(Long & Morin 2005).

Our experiment revealed a size-abundance scaling close to -0.75, suggesting that the scaling
of metabolic rate with body size may be a possible explanation. However, when we explicitly
considered the abiotic and biotic context of communities, we found that the size-abundance
relationship of protist communities is mediated by temperature, biodiversity and their interaction.
Interacting effects between size-abundance scaling and temperature as well as diversity, are
inconsistent with energetic equivalence. In our high-richness communities, small organisms were
favored (more abundant than expected) with high temperature, whereas large organisms were favored
(more abundant than expected) when the temperature was low. Smaller organisms have lower
absolute resource requirements (Gillooly *et al.* 2001; Reuman *et al.* 2014), allowing for a higher
number of smaller organisms to be sustained (Li *et al.* 2006; Morán *et al.* 2010). Temperature may
lead to smaller organisms on average according to the temperature-size rule (Atkinson *et al.* 2003),
however, this physiological effect will affect organisms regardless of their size. Therefore, this cannot
explain the observed changes in size-abundance scaling, requiring additional explanations. Species
richness increased abundance, which is in line with experimental evidence that increasing the number
of species (i.e., in this case, competing for the same resources, such as bacteria) increases the total
acquisition of resources due to complementarity (Gómez & González-Megías 2002). However,
richness did not interact with size, suggesting that complementarity and selection effects are not size-
dependent in our communities. So how can we explain changes in size-abundance scaling across
temperature and species richness gradients?

Species interactions are major determinants of community structure and have been found to
be strongly temperature-dependent (Burnside *et al.* 2014; Jiang & Morin 2004). The observed
interactions may hence be driven by temperature-dependent size asymmetries in resource competition
(Cotgreave 1993; Nee *et al.* 1991; Russo *et al.* 2003), since high richness communities with stronger
interactions were particularly affected. Asymmetric competition results from the differential ability of
species to occupy higher positions in a competitive hierarchy (Freckleton & Watkinson 2001).
Competitive asymmetries between small and large species further depend on various biotic and
abiotic ecological factors (Gjoni & Glazier 2020; Morán *et al.* 2010; Pomeranz *et al.* 2022). Resource
acquisition is fueled by metabolism, and the metabolism increases with higher temperatures by
accelerating the rates of biochemical reactions (Brown *et al.* 2004). Temperature hence may intensify
competition and also affect which species is the dominant competitor for resources (Carson & Root
2000; Clements *et al.* 2014; Gillooly *et al.* 2001; Jiang & Morin 2004; Long & Morin 2005; Reuman
et al. 2014; Schmitz 2003). Consequently, these processes may drive the observed dominance of
smaller organisms in more species rich communities at high temperature, while larger organisms

dominate in communities at low temperature, where they remain competitively superior due to larger
access to resources.

Previous analyses on protist communities have indeed shown that species interactions become
more asymmetric when temperatures rise. Tabi et al. (2020) have shown that this leads to lower
coexistence of protist communities and a decoupling between abundance in isolation and abundance
in communities. We provide indirect evidence of this process by comparing the temporal dynamics of
size-abundance scaling in simulated communities and observed polycultures. In real polycultures, the
initial slope aligned with the expected slope of -0.75, and over time, there was an interactive effect of
size, temperature and richness. In contrast, this pattern did not occur in simulated polycultures, as they
lack the influence of species interactions. Maynard et al. (2020) estimated coefficients that describe
temperature effects on species interactions, showing mostly negative effects of temperature and
decreasing coefficients across the temperature gradient. Jointly, this suggests that the observed
steepening of the size-abundance scaling in high richness communities at high temperature is driven
by species interactions that are both temperature-dependent and size asymmetric.

Our results revealed that size-abundance scaling is dynamically changing through time.
Contrary to our expectations, the slope of the size-abundance relationship was remarkably stable
during the first half of the experiment but deviated from the value of -0.75 by the end of the
experiment, hence indicating a higher proportion of larger individuals in the community. These results
are inconsistent with the MTE which predicts that given a sufficient amount of time, communities
should converge on a line of constant yield and energy use should be independent of body size
(Brown *et al.* 2004; Brown & Gillooly 2003). Our findings showing variation in the size-abundance
relationship over time implies that larger individuals acquire more energy than smaller individuals
after sufficient time for community development. These findings align with previous research, which
proposed that larger individuals may gain greater access to resources compared to the smaller ones
when resources become scarce in protist communities (Tabi *et al.* 2019). Furthermore, in experiments
with phytoplankton conducted over time, larger individuals tend to show a higher proportion relative
to the smaller ones (Malerba & Marshall 2019). The dominance of larger protists relative to smaller
ones may be explained by their ability to efficiently store resources (e.g., food intake), enabling them
to better endure periods of resource scarcity (Fenchel 1980; Thurman *et al.* 2010). The hypothesis
(Millar & Hickling 1990) that larger species may have more endurance under fasting conditions and
their ability to accumulate more resources, provides a compelling explanation for the deviation from
the -0.75 slope observed towards the end of the experiment. This hypothesis is plausible because
experimental studies have shown that larger cells, relative to smaller ones, promote the use of storing
resources in microbial organisms (Malerba *et al.* 2018). Thus, they maintain resource use efficiency
even as nutrient level change over time (Marshall *et al.* 2022). Similar findings were reported for a
field study, where changes in the average size of individual species and changes in species
composition influenced the evolving size-abundance relationship over time (White *et al.* 2004).

Our study system differs in certain ways from natural systems, which should be considered
when interpreting the results. First, we do not assess community turnover as an important mechanism
affecting community size structure under warming (Khaliq *et al.* 2024), since we did not allow for
colonization by warm-adapted species/genotypes. Second, microcosms are closed systems whose
supply of energy, despite replenishment, is not constant through time. While all microcosms have
initially received the same amount of resources, protist feed on bacteria, which could influence the
composition of the ciliate communities (Saleem *et al.* 2013) and hence potentially their dynamics. The
temporal effects observed therefore needed to be judged cautiously; in contrast to changing
temperature and biodiversity, they hence do not provide strong evidence against energetic
equivalence. Third, we do not know whether the observed changes in size-abundance scaling are only
driven by short-term responses (i.e., phenotypic plasticity), or also include evolutionary responses that
may have occurred over the timescale of the experiment. Despite these differences, our approach
provides important complementary insights to analyses of size-abundance scaling in natural
communities.

CONCLUSION

Our experiments suggest that temperature and species richness play an important role in shaping the
size structure of ecological communities through time. We hence caution against assuming that
energy flux is size-independent in dynamic ecosystems, as changes in the biotic and abiotic
environment may change community size structure and hence the flow of energy. Testing these
assumptions with experimental communities provided new insights into the mechanisms driving
community size responses to environmental change, improving our ability to predict how biodiversity
may influence the size structure of communities in the face of global environmental change.

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 SUPPLEMENTARY MATERIAL for

Biodiversity modulates the size-abundance relationship in changing

environments

Materials & methods

S1: Video microscopy and processing to calculate abundance and individual size

The videos were taken using a stereomicroscope (Leica M205 C) with a 16-times magnification

mounted with a digital CMOS camera (Hamamatsu Orca C11440, Hamamatsu Photonics, Japan). We

used the BEMOVI package (Pennekamp *et al.* 2015, version 1.0.2) to process the 18,720 videos

collected during the experiment and extract the raw trajectories. Global segmentation and tracking

parameters were defined for the automated processing of videos (Pennekamp *et al.* 2017). The

difference lag was defined to be 2 seconds, particle size was restricted to 5 to 2000 pixels, and the

grey threshold was set to an intensity value of 10. For particle linking, we specified a link range of 3

frames and a displacement of 20 pixels. These settings were optimized using a subset of videos

(spanning sampling dates of all single, two, and six species combinations at 15 °C, 21 °C, and 25 °C).

Video settings were optimized to rather include false positives rather than exclude true positives at

this step, and exclude false positives later in the processing pipeline. For further details regarding

video processing, please refer to (Pennekamp *et al.* 2017). After tracking, trajectories were filtered to

remove artifacts such as spurious trajectories (e.g., floating debris). Trajectories for analysis were

required to show a minimum net 7 displacement of at least 50 μm, a duration greater than 0.2 seconds,

and a detection rate of 80 percent (for a trajectory with a duration of 10 frames the individual has to be

detected on at least 8 frames), a median step length greater than 2 μm and a minimum mean speed of

50 pixels per second.

We applied machine learning (random forest classification) on quantitative properties such as

cell size and shape to predict to which species an individual belongs. We thus identified individuals

from all species automatically from the videos and quantified their cell size. Details of the procedure

can be found in the supplement (Pennekamp *et al.* 2017) for a detailed description of the experimental

design, setup of micro-ecosystems, and sampling regime.

Results

Section S2: Predicting size-abundance scaling across temperature and richness gradients

Figure S1 Temperature and richness interact to alter size-abundance relationships across protist

communities. This figure is another visualization (see Figure 2) of the mixed model predicting total

log10 abundance (number of individuals) with mean log10 cell volume (μm μL-1), temperature (15

°C, 17 °C, 19 °C, 21 °C, 23 °C, and 25 °C), and community richness (1, 2, 3, 4, 5, and 6 species), as

well as interactions between those predictors based on the model presented in table 1. The dashed line

indicates the expected slope of -0.75.

Section S3: Using evenness as a proxy of realized species richness

We experimentally manipulated initial species richness, but since community assembly and

coexistence are dynamic processes where abundances change through time and even extinctions may

occur, we also calculated species evenness as a proxy of realized species richness. Since some species

may not go extinct, but only reach very low densities, this measure is more appropriate than species

richness which turned out to be pretty stable over the duration of the experiment, although some

extinctions did occur. Figure S2 shows qualitatively similar trajectories when realized species richness

is used rather than initial species richness.

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Figure S2: Temporal dynamics of effect sizes (and 95% confidence intervals) of a linear mixed-effects
model where abundance is modelled as a function size (V = volume), temperature (= T) and realized
richness (= R) and their interactions for the observed communities. The x-axis shows the day. A) Effect of
cell size (V) on abundance, B-C) effect of two-way interactions between size and temperature (V x T), size
and realized richness (V x R) on abundance. D) Three-way interaction between size, temperature and
realized richness (V x T x R) on abundance. The dashed line indicates a slope of -0.75 in panel A and zero
in panels B–D.

Section S4: Interactions between size, temperature and richness through time
Temperature and richness influenced the size-abundance relationship in distinct, time-dependent ways
when comparing low- and high-richness communities. On day 11, the slope and intercept of the
relationship remained stable in both low- and high-richness communities, suggesting that temperature
had minimal influence on the relative abundance of small vs. large protists (i.e., no interaction, Fig.
S3A). By day 17, temperature caused a steeper slope, indicating a relative increase in smaller protist
abundance compared to larger ones, especially across high richness communities (i.e., negative
interaction, Fig. S3C). On day 23, size-abundance scaling became shallower with higher temperatures,
increasing the abundance of larger individuals compared to smaller individuals in low but not high
richness communities (Fig. S3B). On day 29, this pattern was apparent in both low and high richness
communities (i.e., positive interaction, Fig. S3D). These shifts highlight the dynamic influence of

temperature and richness on community structure, where the balance between small and large protist

abundance is regulated by time-dependent physiological and ecological processes (i.e., species

interactions).

Figure S3: Temperature and community richness alter size-abundance relationships of protist communities

across time. This figure is a visualization of the mixed model predicting total log10 abundance with mean log10

cell volume ($\mu m\ \mu L^{-1}$), temperature (15 °C, 17 °C, 19 °C, 21 °C, 23 °C, and 25 °C), and richness (low = 1 and

high = 6 species number), as well as interactions between those predictors across time (A. day = 11, B. day = 17,

C. day = 23 and D. day = 29; see Fig. 4D). All predictors were mean-centered for the analysis, but here the

actual richness and temperature values are shown). The dashed line indicates the expected slope of -0.75.

Interpretation:

It seems that in warm environments under intense competition, smaller individuals have higher

resource-use efficiency in the short-term, whereas, in the long-term, this is true for the larger ones,

potentially in combination with higher survival due to the ability to store resources. Previous work has

shown that intense competition selects for high resource efficiency and lower metabolism, leading

larger individuals to a competitive advantage (Bierbaum *et al.* 1989; Mueller & Ayala 1981; Pianka

1970). In contrast, our study showed that this is true only in cooler conditions and the long-term. This

implies that in the short-term and with increased temperature smaller individuals gain a competitive

advantage due to their lower resource demands. Smaller organisms require less energy for growth,

reproduction, and activity, allowing them to efficiently acquire resources and outcompete larger

individuals. In contrast, larger individuals experience higher resource demands that escalate with

temperature, making them less efficient. However, in the long term, larger individuals may survive

better compared to smaller ones, being able to exploit available resources. This suggests that, over

time, larger organisms are better able to balance resource use under elevated temperatures.

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