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Author for correspondence:

Luz Boyero e-mail: luz.boyero@ehu.eus

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Biotic and abiotic variables influencing plant litter breakdown in streams: a global study

Luz Boyero^{1,2,3}, Richard G. Pearson³, Cang Hui^{4,5}, Mark O. Gessner^{6,7}, Javier Pérez¹, Markos A. Alexandrou⁸, Manuel A. S. Graça⁹, Bradley J. Cardinale¹⁰, Ricardo J. Albariño¹¹, Muthukumarasamy Arunachalam¹², Leon A. Barmuta¹³, Andrew J. Boulton¹⁴, Andreas Bruder^{15,16}, Marcos Callisto¹⁷, Eric Chauvet^{18,19}, Russell G. Death²⁰, David Dudgeon²¹, Andrea C. Encalada^{9,22}, Verónica Ferreira⁹, Ricardo Figueroa²³, Alexander S. Flecker²⁴, José F. Gonçalves Jr²⁵, Julie Helson²⁶, Tomoya Iwata²⁷, Tajang Jinggut²⁸, Jude Mathooko²⁹, Catherine Mathuriau³⁰, Charles M'Erimba²⁹, Marcelo S. Moretti³¹, Catherine M. Pringle³², Alonso Ramírez³³, Lavenia Ratnarajah^{13,34}, José Rincon^{35,36} and Catherine M. Yule²⁸

¹Faculty of Science and Technology, University of the Basque Country (UPV/EHU), 48940 Leioa, Spain ²IKERBASQUE, Basque Foundation for Science, 48013 Bilbao, Spain ³College of Marine and Environmental Sciences and TropWater, James Cook University, Townsville, Queensland 4811, Australia ⁴Centre for Invasion Biology, Department of Mathematical Sciences, University of Stellenbosch, Private Bag X1, Matieland 7602, South Africa ⁵African Institute for Mathematical Sciences, Muizenburg 7945, South Africa ⁶Department of Experimental Limnology, Leibniz Institute of Freshwater Ecology and Inland Fisheries (IGB), 16775 Stechlin, Germany ⁷Department of Ecology, Berlin Institute of Technology (TU Berlin), 10587 Berlin, Germany ⁸Ecology, Evolution, and Marine Biology, University of California, Santa Barbara, CA 93106, USA ⁹MARE-Marine and Environmental Sciences Centre, Department of Life Sciences, University of Coimbra, 3001-456 Coimbra, Portugal ¹⁰School of Natural Resources and Environment, University of Michigan, Ann Arbor, MI 48109, USA ¹¹Laboratorio de Fotobiologia, INIBIOMA, CONICET, Universidad Nacional del Comahue, Quintral 1250, 8400 Bariloche, Argentina ¹²Sri Paramakalyani Centre for Environmental Sciences, Manonmaniam Sundaranar University, Alwarkurichi, 627412 Tamil Nadu, India ¹³School of Biological Sciences, University of Tasmania, Private Bag 55, Hobart, Tasmania 7001, Australia ¹⁴Ecosystem Management, School of Environmental and Rural Science, University of New England, Armidale, NSW 2351, Australia ¹⁵Department of Aquatic Ecology, Eawag: Swiss Federal Institute of Aquatic Science and Technology, Überlandstrasse 133, 8600 Dübendorf, Switzerland ¹⁶Institute of Integrative Biology (IBZ), ETH Zurich, 8092 Zurich, Switzerland ¹⁷Laboratório de Ecologia de Bentos, Departamento de Biologia Geral, ICB, Universidade Federal de Minas Gerais, 30161-970 Belo Horizonte, MG, Brazil ¹⁸UPS, INPT; EcoLab, Université de Toulouse, 118 Route de Narbonne, 31062 Toulouse, France ¹⁹EcoLab, CNRS, 31062 Toulouse, France ²⁰Institute of Agriculture and Environment-Ecology, Massey University, 4442 Palmerston North, New Zealand ²¹School of Biological Sciences, The University of Hong Kong, Pokfulam, Hong Kong SAR, People's Republic of China ²²Laboratorio de Ecología Acuatica, Colegio de Ciencias Biologicas y Ambientales, Universidad de San Francisco de Quito, Campus Cumbayá, PO Box 17, 1200841 Quito, Ecuador ²³Faculty of Environmental Science and Water Research Center for Agriculture and Mining, University of Concepción, Box 160-C, Concepción, Chile ²⁴Department of Ecology and Evolutionary Biology, Cornell University, Ithaca, NY 14853, USA ²⁵Laboratório de Limnologia/AquaRiparia, Departamento de Ecologia, ECL/IB, Universidade de Brasilia, 70910-900 Brasilia, Distrito Federal, Brazil ²⁶Surface and Groundwater Ecology Research Group, Department of Biological Sciences, University of Toronto at

²⁷Department of Environmental Sciences, University of Yamanashi, Kofu, Yamanashi 400-8510, Japan

Scarborough, 1265 Military Trail, Toronto, Ontario, Canada M1C 1A4

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 28 School of Science, Monash University, Jalan Lagoon Selatan, Bandar Sunway, Selangor 47500, Malaysia

²⁹Department of Biological Sciences, Egerton University, PO Box 536, Egerton, Kenya ³⁰Instituto de Investigaciones en Ecosistemas y Sustentabilidad, Universidad Nacional Autónoma de México, Ciudad de México 04510, DF, México

³¹Laboratory of Aquatic Insect Ecology, University of Vila Velha, Vila Velha 29 102-920, Brazil
³²Odum School of Ecology, University of Georgia, Athens, GA 30602, USA

³³Department of Environmental Science, University of Puerto Rico, Rio Piedras, San Juan 00919, Puerto Rico

³⁴Institute for Marine and Antarctic Studies and Antarctic Climate and Ecosystems Cooperative Research Centre, University of Tasmania, Hobart, Tasmania 7001, Australia ³⁵Laboratorio de Contaminación Acuática y Contaminación Fluvial, Departamento de Biología, Facultad de Ciencias, Universidad del Zulia, Apartado Postal 526, Maracaibo, Venezuela ³⁶Programa Prometeo, Senescyt, Escuela de Biología, Ecología y Gestión, Universidad del Azuay, Apartado 981, Cuenca, Ecuador

(D) LB, 0000-0001-7366-9299

Plant litter breakdown is a key ecological process in terrestrial and freshwater ecosystems. Streams and rivers, in particular, contribute substantially to global carbon fluxes. However, there is little information available on the relative roles of different drivers of plant litter breakdown in fresh waters, particularly at large scales. We present a global-scale study of litter breakdown in streams to compare the roles of biotic, climatic and other environmental factors on breakdown rates. We conducted an experiment in 24 streams encompassing latitudes from 47.8° N to 42.8° S, using litter mixtures of local species differing in quality and phylogenetic diversity (PD), and alder (Alnus glutinosa) to control for variation in litter traits. Our models revealed that breakdown of alder was driven by climate, with some influence of pH, whereas variation in breakdown of litter mixtures was explained mainly by litter quality and PD. Effects of litter quality and PD and stream pH were more positive at higher temperatures, indicating that different mechanisms may operate at different latitudes. These results reflect global variability caused by multiple factors, but unexplained variance points to the need for expanded global-scale comparisons.

1. Introduction

Plant litter breakdown is a key process in organic matter recycling and supports food webs in terrestrial and aquatic ecosystems. Terrestrial plants produce *ca* 120 billion tonnes of organic carbon each year [1], 90% of which escapes herbivory and eventually enters the dead organic matter pool [2]. After death, the plant litter either can be decomposed, with its components recycled back into their inorganic forms, or the recalcitrant portion can be stored over long periods [3]. The rate at which plant litter is transformed to other forms of organic and inorganic carbon determines the rate of recycling of biologically essential nutrients, the capacity for carbon storage in ecosystems and the rate at which greenhouse gases such as carbon dioxide (CO₂) are outgassed—all processes that influence regulation of the global climate [4].

Knowing which biotic and abiotic factors drive plant litter breakdown is crucial for understanding how ecosystems function and how vulnerable they are to environmental perturbations such as climate warming, biodiversity loss and biological invasions [5,6]. In terrestrial ecosystems, the relative roles of climate and plant litter quality vary among biomes [7,8]. A recent synthesis showed that climate had a greater role in cold or dry ecosystems and deciduous forests, whereas litter quality traits (i.e. the carbon:nitrogen (C:N) ratio and specific leaf area (SLA)) were more important in humid grasslands and agro-ecosystems, both factors being equally important in tropical wet forests [9]. Another study also highlighted the main role of climate on breakdown in cold areas of the Northern Hemisphere [10], whereas several terrestrial studies at the global scale found that effects of litter quality on breakdown were greater than effects of climate [11–13]. Detritivores can also be important contributors to breakdown rates in terrestrial ecosystems [9], particularly in temperate and wet tropical climates where biological activity is not constrained by temperature or moisture [14].

In contrast to terrestrial ecosystems, information on the relative roles of different drivers of plant litter breakdown in fresh waters is scarce. In freshwater ecosystems in general [15], and in streams and rivers in particular [16], large amounts of organic carbon are processed, contributing significantly to global carbon fluxes. For example, recent estimates show that mean CO_2 evasion from fresh waters is 2.1 Pg C yr⁻¹ (cf. \approx 9 Pg C yr⁻¹ from anthropogenic sources [15]), 86% of which comes from streams and rivers [17]. Breakdown of terrestrially derived plant litter is a pivotal component of stream ecosystem functioning but, despite numerous commonalities, the process is likely to differ in several respects between aquatic and terrestrial ecosystems, in part because of the contrast in water availability [18,19]. Specifically, water is not a limiting factor to the breakdown process within perennial streams, so climate is expected to affect breakdown rates mostly through the influence of temperature, and less so through precipitation (which can, however, cause floods that remove litter from stream reaches [20,21]).

Several local-scale stream studies (encompassing one or several streams within a single region) have demonstrated that litter quality traits (e.g. lignin or nutrient concentration) show tight relationships with litter breakdown rates [22-26]. The few comparable continental-scale and globalscale studies have shown that climate and detritivores can be key determinants of breakdown rates [27-30]. However, there have been few assessments of the relative roles of climate, litter quality and detritivores on litter breakdown in streams at continental or global scales, limiting our capacity to predict the effects of global change on the breakdown process. Furthermore, with one exception [28], global-scale breakdown studies have not considered the potential role of plant diversity on litter breakdown, although species loss is a major global concern [31] and can alter ecosystem processes [3]. The link between plant diversity and litter breakdown rate is still unclear and apparently weak [18,32,33], but virtually all studies examining plant diversity effects on litter breakdown have focused on species richness, neglecting other diversity components such as phylogenetic diversity (PD) [34].

Here, we present a global-scale study of litter breakdown in streams to assess the relative roles of biotic, climatic and other environmental factors on breakdown rates. We conducted an experiment in 24 streams on five continents encompassing a latitudinal range of 90° (47.8° N–42.8° S), using local litter mixtures differing in quality and PD (but not species richness), and an additional common litter type (black alder, *Alnus glutinosa* (L.) Gaertn.) to control to some

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extent for variation in litter quality and diversity across sites. Alder leaves were chosen because the genus is widely distributed across the North Temperate Zone, and because they are highly palatable to temperate and tropical stream detritivores [35]. We manipulated detritivore presence using coarse-mesh and fine-mesh bags, which allowed colonization of litter-consuming detritivores or excluded them, respectively.

We expected that the relative importance of litter quality and PD, detritivore presence, climate and other environmental factors would vary globally, based on the existing evidence from terrestrial ecosystems [9,11,13,14]. We hypothesized that (i) microbial breakdown would increase with temperature (through its effect on metabolic rate [36,37]), and hence decrease with latitude; (ii) detritivore-mediated breakdown would be greater at higher latitudes, where litter-consuming detritivores are more abundant and diverse [38]; (iii) litter of high quality would break down rapidly [24,26]; (iv) phylogenetically diverse plant litter would break down faster because of the potential presence of a wider range of species trait values, which increases the chance for niche partitioning [39]; (v) more basic water (higher pH) would enhance breakdown, as is typically the case [40]; (vi) breakdown would be faster in narrower streams where litter retentiveness is generally higher [41]; and (vii) climate would modulate the effects of other environmental factors and litter on breakdown rates; for example, higher temperatures could enhance any effects of litter quality or water chemistry [42].

2. Methods

(a) Field experiment

We conducted a litter breakdown experiment at 24 stream sites around the world with absolute latitudes ranging from 0.37° to 47.80° (electronic supplementary material, table S1 and figure S1); 14 sites were located within the tropics (less than or equal to 23.5° latitude) and 10 in temperate regions (more than 23.5°), comprising a large range of climatic patterns (electronic supplementary material, figure S2a,b). At each site, we chose a single stream reach draining a forested catchment that experienced little human influence. The experiment was run at a time of high litter inputs and low flood susceptibility at each site. We chose three native riparian tree species locally common in the riparian vegetation and well represented in stream leaf litter (electronic supplementary material, table S2). In total, we collected freshly abscised leaves of 70 species, as two species were shared between two sites each. Alder leaves were collected to serve as an approximate control for variation in litter quality across sites; however, the leaves were locally collected near the study sites or, when not locally available (tropical sites, Southern Hemisphere and some Northern Hemisphere temperate sites), shipped from either Portugal or Spain or collected in botanical gardens to avoid problems with import regulations. Although some intraspecific variation in alder litter quality across locations was likely to occur [25], we expected such variation to be much smaller than that among the 70 other species used in the experiment.

Leaves were air-dried, weighed, enclosed in coarse-mesh (10 mm) and fine-mesh (0.5 mm) bags (ca 1 g per species per bag in three-species mixtures, or ca 3 g for alder leaves) and secured in streams. Three coarse-mesh and three fine-mesh bags were retrieved on each of four dates: day 0 (to determine any mass loss due to handling), and approximately days 14, 28

and 56. Bags were collected with a net (0.5 mm mesh) and taken to the laboratory where leaves were cleaned, oven-dried and weighed. Ash-free dry mass was not estimated for logistical reasons; however, it is unlikely that the litter breakdown data were biased by mineral particles associated with the leaves as there was no indication of calcite precipitation in our streams (most of them being soft-water systems) or association of mineral particles with leaves (most streams having coarse substratum). In addition, all leaves were thoroughly cleaned under water before drying. At each site, we recorded pH as a measure of water chemistry, and wetted stream width as an estimate of stream size.

(b) Climatic and leaf quality data

Comprehensive water temperature data were not available for all sites, so mean annual air temperature data were extracted from the WorldClim database v. 1.3 [43] at the highest resolution (2.5 min of arc) using DIVA-GIS software, 7.5.0.0. (http://www.diva-gis.org). We assessed leaf quality by using the mean SLA, which is the ratio of leaf area (cm²) to leaf dry mass (g). SLA is a key ecological and physiological plant trait [44] that often correlates with leaf toughness, nutrient concentration and breakdown rate [45,46]. To measure SLA, we scanned 20 leaves of each plant species, estimated their areas with IMAGEJ 10.2, dried them to constant mass and weighed them to the nearest 0.1 mg. We then calculated the mean SLA at each site.

(c) Plant phylogenetic diversity

We used PD, defined as the total phylogenetic distance among species, as a measure of plant litter diversity [47]. To estimate PD among the three species selected at each site, we constructed a molecular phylogeny of the 70 species and 8935 bps of DNA (including five outgroups) using partial 18S ribosomal DNA, rbcl, matK, atpB, trnl, rpl16, rpoB and rpoC1 sequences available in GenBank (electronic supplementary material, table S3). We used these markers as they provided the most comprehensive sets of data for the target species. Data for several species were not available in GenBank, so we chose closely related congeners as replacements. We constructed alignments independently for each gene using MUSCLE v. 3.8.31 [48]. Nucleotide substitution models were selected for each gene using the Akaike information criterion (AIC) as implemented in jMODELTEST v. 0.1.1 [49]. We searched for the maximum-likelihood phylogeny using RAxML v. 7.2.8 [50], partitioning the dataset by gene. Random starting trees were used for each independent tree search, and topological robustness was investigated using 100 bootstrap replicates. We used a rate-smoothed Bayesian phylogeny, estimated using BEAST v. 1.6.2 [51], assuming a relaxed uncorrelated lognormal clock with all other parameters set as default. The relaxedclock analysis was used to estimate relative divergence times, thereby converting branch-length values from the substitutions per site to an estimate of time since divergence from a common ancestor. The Bayesian Markov chain Monte Carlo ran for 10 million generations sampled every 1000 generations, whereas stationarity and effective sample sizes (ESS > 200) were examined using TRACER v. 1.6 [52], discarding all trees under the asymptote as burn-in. Finally, we constructed a consensus tree (electronic supplementary material, figure S3) with mean node heights from the posterior distribution using TREEANNOTATOR v. 1.6.2 [51]. To calculate PD per community, we used the comdist function in the R package 'picante' [53].

(d) Data analysis

We used linear regression to detect latitudinal trends in litter variables, mean annual temperature and other environmental variables. Breakdown rates of local litter mixtures and alder were estimated for each site using the breakdown coefficient (k)resulting from the exponential decay model $m_f/m_i = e^{-kt}$, where $m_{\rm f}$ and $m_{\rm i}$ are the final and initial litter mass (g), respectively, *t* is time in days (d) and *k* is the breakdown rate coefficient. We estimated the relative contributions of different factors to litter breakdown rates using general linear models, with key predictors selected based on the AIC following a stepwise search in backward and forward directions, using the stepAIC function in the R package 'MASS' [54]. The initial models for litter mixtures included litter variables (mean SLA and PD), mean annual temperature (hereafter temperature), other environmental variables (pH and stream width) and the interactions between temperature and the other variables. We included these interactions to explore the directional effects of climate on other variables. The models for alder included temperature and other environmental variables (and their interactions) but no litter traits, because SLA was expected to vary little within species (compared with variation in SLA across the total of 70 species) and PD = 1. We examined multicollinearity of the variables to be included in the models with the variance inflation factor (VIF) using the VIF function in the R package 'fmsb' [55].

We tested separate models for overall breakdown in coarsemesh litter bags (k_c), microbial breakdown in fine-mesh bags (k_f) and detritivore-mediated breakdown (k_d), which was calculated from the difference in the proportion of litter mass remaining between coarse-mesh and fine-mesh bags at each sampling date [29]. Normality of residuals was examined with Shapiro–Wilk's test; three variables (k_c for alder and k_f for litter mixtures and alder) showed lognormal distributions, which became normal after loge-transformation. Main and interaction effects were visually explored using the *visreg* and *visreg2d* functions in the R package '*visreg*' [56]. We further examined the role of detritivores on breakdown of local litter mixtures and alder by comparing k_c and k_f for the global dataset and separately for tropical (n = 14) and temperate sites (n = 10), using Wilcoxon signed-rank tests.

3. Results

Mean annual temperature decreased with latitude (r = -0.73, p < 0.001), whereas the other environmental or litter variables showed no latitudinal trend (r < 0.26, p > 0.22 in all cases). VIFs were all less than 2, indicating the absence of multicollinearity (electronic supplementary material, table S4). Our final models included different factors and interactions (table 1). The overall model for total breakdown of the local litter mixtures was significant (p = 0.0499) and showed significant effects of litter SLA and pH and a significant interaction between temperature and SLA (figure 1*a*,*b*). The overall model for microbial breakdown of litter mixtures was significant (p = 0.042) and showed a significant effect of pH (figure 1c). For detritivore-mediated breakdown of litter mixtures, the overall model was not significant (p = 0.107) but it showed significant effects of litter PD and its interaction with temperature (figure 1*d*). For total alder breakdown, the overall model was significant (p = 0.046) and showed the effects of temperature and its interaction with pH (figure 1f). The overall model for microbial breakdown of alder was significant (p = 0.029) and showed an effect of temperature (figure 1e). Finally, detritivore-mediated breakdown of alder was not significantly affected by any factor (p = 0.90).

Litter breakdown rates were higher in coarse-mesh than in fine-mesh bags for litter mixtures and alder across all sites (litter mixtures: Z = -3.19, p = 0.0014; alder: Z = -3.87, p < 0.001) and at temperate sites (litter mixtures: Z = -2.76,

p = 0.0058; alder: Z = -3.36, p < 0.001). In the tropics, breakdown was faster in coarse-mesh bags for alder, but the difference was weaker than in temperate sites (Z = -2.21, p = 0.027) and there was no difference for litter mixtures (Z = -1.81, p = 0.070; figure 1*g*,*h*).

4. Discussion

(a) Climatic effects on global patterns of litter breakdown in streams

The analysis of our global dataset revealed influences of several abiotic and biotic factors on litter breakdown rate in streams, paralleling findings for terrestrial ecosystems [9,11,14]. The only factor having no influence on breakdown was stream width, counter to our expectations [41], probably because we sampled only during stable base-flow periods. Mean annual temperature was a key influence on litter breakdown of alder, having a main effect on total and microbial breakdown. As expected, microbial breakdown rate increased with temperature and hence was higher towards lower latitudes. A companion study found that this rate increased with contemporaneous water temperature, and the relationship conformed to the metabolic theory of ecology [27]. Here, we found the relationship to hold with long-term air temperature data.

Although we did not detect any variation with temperature for detritivore-mediated breakdown when it was examined separately, total breakdown of alder increased towards cooler streams, and this pattern was most likely related to the higher abundance and diversity of litter-consuming detritivores at higher latitudes [38,57]. Accordingly, our comparison of coarse-mesh and fine-mesh bags revealed a greater role of detritivores on litter breakdown in temperate than in tropical streams, although this difference was most obvious for litter mixtures.

(b) Major influence of litter quality on breakdown of litter mixtures

While breakdown of a single substrate type (alder leaves) across latitudes was mostly influenced by environmental factors and especially temperature, litter variables (SLA and PD) were of major importance to explain variation in breakdown of litter mixtures. This agrees with the results of other global-scale studies [12] and a comprehensive meta-analysis [11] based on data from terrestrial ecosystems. Nevertheless, for litter mixtures, temperature was important as well in that it modulated litter effects on breakdown.

Counter to our prediction, litter SLA had a negative overall influence on total breakdown of litter mixtures. This result contrasts with those of previous research showing that breakdown is greater in leaves with lower lignin content [19,22], which tends to be correlated with SLA [45]. However, the positive interaction of SLA with temperature indicates that small SLAs enhanced breakdown only at low temperatures, whereas the opposite was true in streams of warmer regions, where breakdown of litter mixtures with a higher mean SLA was faster (figure 1*b*). This result is not easily explained but might reflect a relatively larger influence of detritivores at lower latitudes that use tough leaves for purposes other than nutrition, such as case construction by caddisflies [58]. Table 1. Results of general linear models (selected based on the Akaike information criterion, AIC, following a stepwise search in backward and forward directions) examining the influence of key predictors (mean annual temperature, T; mean specific leaf area, SLA, and litter phylogenetic diversity, PD, as litter variables; and pH and wetted width, SW, as stream environmental variables) on total litter breakdown rates of litter mixtures and alder in coarse-mesh bags (k_c) , microbial breakdown rates in fine-mesh bags (k_f) and detritivore-mediated breakdown rates (k_d) . The AIC, adjusted r^2 , *F*-test and *p*-value are shown for each model, and the mean estimate and standard error, *t*-statistic and *p*-value for each factor (italic entries indicate significant results at the p < 0.05 level).

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	$F_{7,12} = 2.92$ $p = 0.0499$ crobial $k_f \sim T + SLA + pH + T \times SLA$ $AIC = -18$ adjusted $r^2 = 0.32$	SLA PD pH T \times SLA T \times PD T \times PH	- 0.000321 - 0.135900 0.023480 0.000019 0.0009196 - 0.000975	0.000110 0.088550 0.010510 0.00006 0.000559 0.000559	-2.92 -1.54 2.24 2.95 -1.75 -4.07	0.013 0.151 0.045 0.012 0.167 0.106 0.106	
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	crobial $k_{ m f} \sim { m T} + { m SLA} + { m pH} + { m T} imes { m SLA}$ ${ m Alc} = -18$ adjusted $r^2 = 0.32$	<i>рН</i> 7 × SLA 1 × PD T × рН	0.023480 0.000019 0.0009196 	0.010510 0.000006 0.006253 0.000559	2.24 2.95 1.47 -1.75 -4.07	0.045 0.012 0.167 0.106 0.106	
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It $1 \times \beta H$ 0.0097 0.0059 -1.75 0.105 fcola $k \sim T + SIA + \beta H + T \times SIA$ interept -6.650175 1.632409 -4.07 0.005 dusted $r^2 = 0.32$ $1 = 0.028414$ 0.00557 -1.18 0.005 -1.18 0.005 dusted $r^2 = 0.32$ $1 = 0.02$ $1 = 0.023$ 0.00557 0.00557 -1.18 0.025 $r_{a,5} = 3.25$ 0.0052 0.00529 0.00557 0.00557 0.0055 0.00557 0.0055 $r_{a,5} = 0.042$ $A \sim T + P0 + PH + SW + T$ $T \times SIA$ 0.00529 0.00529 0.00539 0.0055 0.00552 0.0055 0.0056	crobial $k_{\rm f} \sim {\rm I} + {\rm SLA} + {\rm PH} + {\rm I} \times {\rm SLA}$ AIC = -18 adjusted $r^2 = 0.32$	T × pH	-0.000975	0.000559	-1.75 -4.07	0.106	
cobial $k_{\gamma} - T + SIA + pH + T \times SIA$ adiated $k_{\gamma} - T + SIA + pH + T \times SIA$ intercept -6.650175 1.62409 -4.07 000 $A(C = -18)$ $A(C = -18)$ intercept -6.650175 1.632490 -1.18 0.025 $A(C = -18)$ $A(C = -18)$ $T_{415} = 3.25$ $S1A$ -0.00188 0.00575 -1.18 0.025 $F_{415} = 3.25$ $S1A$ 0.00539 0.00575 -1.78 0.025 $F_{415} = 3.25$ D_{41} D_{42} $D = 0.012$ <	crobial $k_f \sim T + SLA + pH + T \times SLA$ AIC = -18 adjusted $r^2 = 0.32$				-4.07	0.001	
	AIC = -18 adjusted $r^2 = 0.32$					0.001	
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		Т	— 0.058414	0.049330	- 1.18	0.255	
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	$F_{4,15} = 3.25$	SLA	-0.010108	0.005675	-1.78	0.095	
T × SIA 0.000629 0.000325 1.94 0.01 intivore-mediated $k_a \sim T + PD + PH + SW + T$ 0.00032 0.00032 1.94 0.003 intivore-mediated $k_a \sim T + PD + PH + SW + T$ 0.0003 0.00032 0.040 0.003 0.043 0.053 0.043 0.063 0.044 0.063 0.044 0.063 0.044 0.063 0.044 0.063 0.044 0.063 0.044 0.063 0.044 0.063 0.044 0.063 0.044 0.063 0.044 0.063 0.044 0.063 0.044 0.063 0.044 0.064 0.064 0.064 0.064 0.064 0.064 0.044 0.064 0.044	p = 0.042	Ηq	0.422662	0.166398	2.54	0.023	
Introne-mediated $k_{\rm d} \sim T + D + \mu H + SW + T$ Intervent		T imes SLA	0.000629	0.000325	1.94	0.072	
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	tritivore-mediated $k_{\rm d} \sim {\rm T} + {\rm PD} + {\rm pH} + {\rm SW} + {\rm T}$						
$\begin{tabular}{lllllllllllllllllllllllllllllllllll$	imes PD + T $ imes$ PD + T $ imes$ SW						
adjusted $r^2 = 0.25$ T0.0005620.0012660.440.663 $F_{7,16} = 2.08$ $F_{7,16} = 2.08$ PD -0.077460 0.028080 -2.76 0.014 $p = 0.107$ $p = 0.107$ PD -0.07460 0.028080 -2.76 0.014 $p = 0.107$ $p = 0.107$ PD 0.004855 0.003296 1.47 0.160 $r > P$ $r > PD$ 0.00021 0.00021 0.00011 1.88 0.078 $r > P$ $r > PD$ 0.005619 0.00119 1.88 0.009 $r > P$ $r > PD$ 0.005619 0.00173 2.97 0.009 $r > P$ $r > PD$ 0.00305 0.00173 -1.76 0.009 $r > P$ $r > SW$ -0.00001 0.000173 -1.60 0.009	AIC = -208	intercept	— 0.009987	0.024860	— 0.40	0.693	
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	adjusted $r^2 = 0.25$	Т	0.000562	0.001266	0.44	0.663	
$p = 0.107$ $p = 0.002$ $p = 0.002296$ 1.47 0.160 SW 0.00021 0.00210 1.88 0.078 $T \times PD$ 0.005619 0.001892 2.97 0.009 $T \times PH$ -0.000305 0.00173 -1.76 0.008 $T \times PH$ -0.000305 0.000173 -1.76 0.009 $T \times SW$ -0.00001 0.00001 -1.60 0.0001 -1.60 0.0001	$F_{7,16} = 2.08$	D	— 0.077460	0.028080	- 2.76	0.014	
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	p = 0.107	Нd	0.004855	0.003296	1.47	0.160	
$\begin{array}{cccccccccccccccccccccccccccccccccccc$		SW	0.000021	0.000011	1.88	0.078	
$\label{eq:relation} \begin{split} \Gamma \times pH & -0.00305 & 0.00173 & -1.76 & 0.098 \\ \Gamma \times SW & -0.00001 & 0.00001 & -1.60 & 0.129 \end{split}$		T imes PD	0.005619	0.001892	2.97	0.009	
$T \times SW$ -0.00001 0.00001 -1.60 0.129		T imes pH		0.000173	-1.76	0.098	
		T imes SW	- 0.000001	0.000001	- 1.60	0.129	

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litter breakdown	selected model	Tactor	estimate	s.e.	L	р
alder						
total	$k_{ m c} \sim { m T} + { m pH} + { m SW} + { m I} imes { m pH}$					
	AIC = -22	intercept	3.578300	3.945612	0.91	0.376
	adjusted $r^2 = 0.27$	Т	— 0.475873	0.209678	- 2.27	0.036
	$F_{4,18} = 3.00$	pH	— 1.029913	0.541762	-1.90	0.073
	p = 0.046	SW	0.000984	0.000525	1.87	0.077
		T × pH	0.068805	0.028986	2.37	0.029
microbial	$K_f \sim T$					
	AIC = -16	intercept		0.372650	- 13.32	0.000
	adjusted $r^2 = 0.17$	Г	0.046910	0.019960	2.35	0.029
	$F_{1,21} = 5.52$					
	<i>p</i> = 0.029					
detritivore-mediated	$k_{ m d} \sim { m T} + { m pH} + { m SW} + { m T} imes { m pH} + { m T} imes { m SW}$					
	AIC = -223	intercept	0.029220	0.051040	0.572	0.574
	adjusted $r^2 = -0.19$	F	— 0.001610	0.002725	— 0.591	0.562
	$F_{5,17} = 0.31$	рН	- 0.003204	0.006897	— 0.465	0.648
	<i>p</i> = 0.899	SW	0.000015	0.000022	0.665	0.515
		T imes pH	0.000247	0.000369	0.667	0.514
		$\mathrm{T} imes \mathrm{SW}$	- 0.000001	0.000001	— 0.663	0.516

Table 1. (Continued.)

6

7



Figure 1. (a-f) Significant main effects (top panels) and interaction effects (middle panels) of mean annual temperature (T), mean specific leaf area (SLA), litter phylogenetic diversity (PD) and stream pH on total breakdown rates in coarse-mesh bags (k_c) , microbial breakdown rates in fine-mesh bags (k_f) and detritivore-mediated breakdown rates (k_d) of litter mixtures (a-d) and alder (e,f). When a variable had a main effect and an interaction in the same model, only the interaction is shown. (g,h) Box plots show litter breakdown rates (k) in coarse-mesh bags (grey boxes) and fine-mesh bags (white boxes) for the global dataset and for tropical and temperate sites separately. Horizontal lines show medians, boxes show 25th and 75th percentiles, error bars show 10th and 90th percentiles, and points show outliers. Brackets and asterisks denote significant differences in breakdown rates of litter between coarse-mesh bags (Wilcoxon signed-rank tests: *p < 0.05, **p < 0.001; n.s., not significant).

(c) Detritivore-mediated breakdown driven by phylogenetic diversity in litter mixtures

Detritivore-mediated breakdown was mostly driven by PD of litter mixtures, but apparently in the direction opposite to our prediction. The negative effect of PD on breakdown suggests that phylogenetic proximity of litter mixtures enhances detritivore consumption. This finding could be due to lower phylogenetic distance between litter species resulting in a higher concentration of high-quality resources, whereas larger phylogenetic distance caused the dilution of such resources [59]. Similarly, one of the few studies to explore PD effects on breakdown found that a lower phylogenetic distance in litter mixtures promoted microbial biomass and litter nutrient concentration [60].

However, this pattern was inconsistent across climates, as occurred for SLA. Breakdown of litter mixtures composed of species showing higher phylogenetic proximity was faster only in cooler streams, whereas the opposite was true at higher temperatures (figure 1*d*). This discrepancy might be related to a higher specialization of litter-consuming detritivores at temperate latitudes [61], which would benefit from the concentration of their preferred resources, whereas more generalist detritivores in the tropics [62] could be relatively more efficient when more varied resources are available.

(d) Higher breakdown in more alkaline waters

Stream pH, which typically reflects basic catchment lithology [63,64], also influenced litter breakdown rates, especially for litter mixtures. Breakdown was faster at higher pH, which agrees with the well-known effect of greater breakdown in more basic waters [65]. The effect of pH on alder breakdown was, however, modulated by climate: while breakdown was faster in more alkaline waters at higher temperatures, the opposite was true in cooler streams (figure 1*f*). This interaction may reflect the facts that (i) microbial decomposition is most important in the tropics and is boosted by higher calcium concentration [66,67], and (ii) major litter consumers such as stoneflies are more important in circumneutral and acidic streams at higher latitudes [68,69], while they are rare in the tropics [57]. Consistent with this explanation, caddiflies, which dominate the guild of litter-consuming detritivores in tropical streams [57], tend to be more sensitive to low pH [70]. However, targeted experiments are required to test these possibilities.

5. Conclusion

Our study is one of very few to assess litter breakdown in streams at the global scale [27,28,30,71]. The number of samples was relatively small for such broad scope (24 study sites), limiting statistical power of the analyses; nevertheless, we were able to show that multiple biotic and abiotic factors influence rates of litter breakdown in streams at the global scale, with (i) a large positive influence of temperature on microbial breakdown of alder leaves; (ii) a greater role of litter-consuming detritivores on breakdown towards high latitudes, where these detritivores are typically more abundant and diverse; (iii) a notable influence of litter quality and PD on breakdown of litter mixtures that varies across climates, possibly through different effects on microbial and detritivore assemblages at different latitudes; and (iv) generally faster breakdown in more basic waters with some influence of temperature, possibly owing to interactions with microbial activity, which is the most important mechanism for litter breakdown in warmer waters. Our models explained up to 41% of the total variance in the datasets, indicating that our explanatory variables could be important drivers of litter breakdown. However, substantial unexplained variation indicates that further comparative research is required to develop a

comprehensive picture of litter breakdown in streams at the global scale. Such understanding is vital to appropriate management of these ecosystems in the face of multiple anthropogenic stressors [72,73].

Data accessibility. Data are available in the electronic supplementary material and the Dryad Digital Repository: http://dx.doi.org/10. 5061/dryad.jg8r0.

Authors' contributions. L.B. and R.G.P. conceived the study; L.B. managed the overall project; L.B., C.H., J.P., M.A.A. and R.G.P. analysed the data; L.B. and R.G.P. wrote the manuscript with substantial contribution from M.O.G., M.A.S.G., J.P., B.J.C., C.H. and M.A.A.; and all other authors (listed alphabetically) locally coordinated or performed research and commented on the manuscript; all authors gave final approval for publication.

Competing interests. We have no competing interests.

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