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Journal

Journal of Ecology, 110(6)

ISSN

0022-0477

Authors

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Publication Date 2022-06-01

DOI 10.1111/1365-2745.13878

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Nitrogen increases early-stage and slows late-stage decomposition across diverse grasslands

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- 42
- 43 Key words: grasslands, litter decomposition, nitrogen, nitrogen deposition,
- 44 Nutrient
- 45 Network (NutNet), phosphorus

46 Abstract

- 47 1. To evaluate how increased anthropogenic nutrient inputs alter carbon
- 48 cycling in grasslands, we conducted a litter decomposition study across 20
- 49 temperate grasslands on three continents within the Nutrient Network, a
- 50 globally distributed nutrient enrichment experiment
- 51 2. We determined the effects of experimental nitrogen (N), phosphorus (P),
- 52 and potassium plus micronutrient (K_{μ}) additions on decomposition of a
- 53 common tree leaf litter in a long-term study (maximum of 7 years; exact
- 54 deployment period varied across sites). The use of higher-order
- 55 decomposition models allowed us to distinguish between effects of nutrients
- 56 on early- versus late-stage decomposition.
- 57 3. Across continents, addition of nitrogen (but not other nutrients)
- 58 accelerated early-stage decomposition and slowed late-stage decomposition,
- 59 increasing the slowly decomposing fraction by 28% and the overall litter
- 60 mean residence time by 58%.

61 4. Synthesis. Using a novel, long-term cross-site experiment, we found 62 widespread evidence that nitrogen enhances the early stages of 63 aboveground plant litter decomposition across diverse and widespread temperate grassland sites, but slows late-stage decomposition. These 64 findings were corroborated by fitting the data to multiple decomposition 65 66 models and have implications for nitrogen effects on soil organic matter 67 formation. For example, following nitrogen enrichment, increased microbial 68 processing of litter substrates early in decomposition could promote production and transfer of low molecular weight compounds to soils, and 69 70 potentially enhance stabilization of mineral-associated organic matter. By 71 contrast, by slowing late-stage decomposition, nitrogen enrichment could 72 promote particulate organic matter (POM) accumulation. Such hypotheses 73 deserve further testing.

- 74 **Keywords:** grasslands, litter decomposition, nitrogen, nitrogen deposition,
- 75 Nutrient Network (NutNet), phosphorus

76 Introduction

77 Human activities are altering element fluxes at a global scale. For 78 example, agricultural and industrial activities have greatly increased global 79 cycling of reactive nitrogen (N) (Ackerman et al. 2019), increasing atmospheric N inputs relative to phosphorus (P). Yet, we lack a general 80 mechanistic understanding of how variation in nutrient inputs alters carbon 81 82 (C) cycling, hampering the development of Earth System Models to 83 accurately forecast global C dynamics in a changing world (Peñuelas et al. 84 2013). Such understanding if especially lacking for grasslands, which 85 represent ca. 30% of non-agricultural global land area and terrestrial net primary productivity (NPP) (Chapin et al. 2011) and store 20% of the world's 86 soil C (FAOSTAT 2009). Effects of N deposition on grassland C cycling will 87 88 likely depend on supplies of other nutrients, since NPP in grasslands is often 89 co-limited by multiple nutrients, with N limitation more prevalent at higher 90 latitudes (Fay et al. 2015, Cleland et al. 2019, Du et al. 2020). On the other 91 hand, surface soil C stocks in grasslands can be relatively insensitive to 92 short-term (ca. 3 years) N and P enrichment, but can increase with short-93 term addition of K with micronutrients (Crowther et al. 2019). This 94 inconsistency between nutrient effects on plant biomass production (NPP). 95 and thus C inputs to soils, and soil C accumulation indicates a gap in understanding nutrient effects on grassland litter and soil organic matter 96 97 (SOM) decomposition.

98 Past studies of nutrient effects on decomposition in grasslands have 99 been limited in geographic scope or duration. In studies at individual or 100 relatively few sites, N had varied effects (Hunt et al. 1988, Aerts et al. 2003, Liu et al. 2006, Hobbie 2008, Liu et al. 2010, Hou et al. 2021) and P had 101 102 neutral effects (Aerts et al. 2003) on decomposition. In a more widespread 103 but short-term study of tea leaf decomposition (90 days across 21 104 grasslands), N and P both weakly increased tea leaf decomposition rate and 105 the tissue stabilization factor, a modeled representation of the biochemically 106 transformed residues that persist at late stages of decomposition (Ochoa-

107 Hueso et al. 2020). Meta-analyses combining experiments done at single or 108 few sites to generate synthetic insights into nutrient effects on 109 decomposition have uncovered important among-study patterns (Knorr et al. 110 2005, Zhang et al. 2018, Gill et al. 2021). However, strong inference from 111 meta-analysis is limited because it combines data from studies using 112 different methods (e.g., substrate types, nutrient treatment levels). Most 113 available studies are of short duration and often lack information on key 114 covariates (climate, soil chemistry, plant productivity). For example, the 115 median study duration in a recent meta-analysis of 334 paired 116 decomposition sequences (from control and associated N-fertilized plots) was 117 2 years (Gill et al. 2021). In addition, past meta-analyses focused solely on N (with supply rates varying widely among studies); yet nutrients other than N 118 119 might be expected to influence decomposition or interact with N treatments. 120 While individual studies and meta-analysis provide important insights, 121 generating a general understanding of nutrient effects on grassland litter 122 decomposition requires long-term, geographically distributed experiments 123 that use standard methods and thus overcome the limitations to inference 124 imposed by the diverse methodologies (e.g., substrate, duration, nutrient 125 enrichment rates) inherent in meta-analyses (Borer et al. 2014). 126 Long-term studies are needed specifically to determine if nutrients

have different effects early versus later in the decay process. In forests, N 127 128 often reduces litter decomposition in its later stages (Fog 1988, Berg and 129 Matzner 1997, Carreiro et al. 2000, Hobbie et al. 2012, Gill et al. 2021). Such 130 N inhibition has been attributed to inhibition of oxidative enzyme activity 131 (Carreiro et al. 2000, Waldrop et al. 2004, Hobbie et al. 2012, Jian et al. 132 2016, Chen et al. 2018), which may be less strong in grasslands than in forests, because of their distinct fungal communities with reduced oxidative 133 134 enzyme capabilities (Sinsabaugh 2010). On the other hand, if N inhibition of 135 late-stage decomposition results from other mechanisms, such as decreased 136 pH and microbial biomass (Treseder 2008), or increased decomposer

137 necromass accumulation (Gill et al. 2021), such N inhibition might be138 expected in grasslands as well as in forests.

139 Cross-site experiments are needed to determine whether effects of 140 nutrients on litter decomposition are consistent across geographically 141 diverse sites, or depend on site characteristics such as climate and edaphic 142 factors. Regional analyses of litter decomposition in grasslands found no 143 relationships with precipitation or temperature (Bontti et al. 2009, O'Halloran 144 et al. 2013), although studies at larger spatial scales found evidence for both 145 (e.g., Tuomi et al. 2009). However, nutrient effects on decomposition might 146 be more pronounced in high rainfall areas where water does not constrain 147 microbial activity, as has been shown for net primary production (Burke et al. 148 1997, Harpole et al. 2007, La Pierre et al. 2016, Ren et al. 2017), or may 149 depend on ambient soil nutrient supply (Hobbie 2005).

150 Here we decomposed a standard oak leaf litter (to avoid any home-151 field advantage effects) for up to seven years in a replicated nutrient 152 addition experiment in grasslands across multiple continents that ranged 153 widely in climate and edaphic conditions, to (1) determine the effects of N 154 and P enrichment on different stages of litter decomposition, and (2) address 155 whether and how nutrient limitation of leaf litter decomposition varies 156 geographically. The long duration of the experiment allowed us to determine 157 which empirical decomposition model structures (single, double, and 158 asymptotic exponential or Weibull models) best described litter 159 decomposition dynamics and specifically whether effects of nutrient addition 160 differed between the early and late stages of decomposition as shown 161 previously for N, primarily in forests (Gill et al. 2021). Unlike in forests, we 162 did not expect N to inhibit the later stages of decomposition in these 163 grasslands because of low oxidative enzyme activity. In addition, we 164 expected that nutrient effects on decomposition would depend on soil 165 resource supply, such that N limitation of decomposition would be common 166 across temperate and alpine regions with young, P-rich soils, but not in areas 167 with high ambient N deposition or N-rich soils. On the other hand, we

- 168 expected that P limitation would be restricted to sites with highly weathered
- 169 soils that are low in available P (e.g., Australia) or to areas with high N
- 170 deposition. Furthermore, we hypothesized that nutrient limitation of
- 171 decomposition would be strongest in regions with high precipitation or where
- 172 moisture supply is relatively constant across the year.



this study. Site were located in North America, Europe, and Australia. (a) Map of site locations; (b) relationship between site MAT and MAP; (c) relationship between site MAT and precipitation distribution; (d) relationship between site MAP and precipitation distribution. Abbreviations as follows: MAT, mean annual temperature (°C); MAP, mean annual precipitation (mm); precipitation distribution (precipitation in wettest month/MAP).

173

174 Materials and Methods

175 Decomposition Experiment

176 In the Nutrient Network (NutNet) study, N is added as slow-release

177 urea (10 g N m 2 y 1), P as triple superphosphate (10 g P m 2 y 1), and K as

178 potassium sulfate (10 g K m⁻² y⁻¹) with micronutrients (Ca, Mg, S, B, Cu, Fe, 179 Mn. Mo. and Zn) (K_u) in a full-factorial manner annually to experimental plots. 180 Twenty sites in the United States, Canada, Australia, and Europe participated 181 in the present decomposition study (Fig. 1, Table S1), which began <1-2182 years following the initiation of fertilization treatments. Sites ranged 17-fold in atmospheric N deposition (1.1 – 18.9 kg N ha⁻¹ yr⁻¹), 7-fold in mean annual 183 184 precipitation (331 – 1898 mm vr^{-1}), and 60-fold in mean annual temperature 185 (0.3 – 18.4 °C). Precipitation and temperature were unrelated across sites, 186 allowing us to explore their independent effects. Treatments were applied to 187 5 m x 5 m plots in a randomized block design; for this study, three blocks 188 were used at all sites except Sierra Foothill, where five blocks were used. 189 We selected leaf litter of *Quercus ellipsoidalis* (Table S2) as a common 190 substrate for a long-term decomposition study because its N concentration (0.927 %) fell within the range of N concentrations of grassland leaf litter 191 192 (Wedin and Pastor 1993), it was readily abundant and, since it does not occur in any of the plots, should not exhibit home field advantage effects 193 194 that could confound among-site environmental effects. Freshly fallen leaf 195 litter of Q. ellipsoidalis was collected from several adjacent individual trees in 196 October 2008 at the Cedar Creek Ecosystem Science Reserve, Minnesota, 197 USA. Litter bags (20 cm x 20 cm, 1-mm mesh fiberglass window screen) were 198 filled with ca. 10 grams (dry weight) of sterilized leaf litter (autoclaved at 199 121°C for 15 minutes). Sufficient bags were constructed to allow annual 200 harvest over seven years and sent to individual site researchers in 201 September 2009. Bags were strung together in groups of seven and were 202 pinned to the ground (1 string per plot) using landscape staples. Subsamples 203 of initial litter were analyzed for (1) C fractions using an ANKOM Fiber 204 Analyzer (Ankom Technology, Macedon, New York, USA) (cell solubles, 205 hemicellulose+bound protein, cellulose, and lignin+other recalcitrants) 206 (n=5), (2) initial C and N concentrations on a Costech ECS4010 element 207 analyzer (Costech Analytical, Valencia, California, USA) (n=5), and (3) initial 208 concentrations of phosphorus (P), magnesium (Mg), calcium (Ca),

209 manganese (Mn), and potassium (K) by Inductively Coupled Argon Plasma

210 Emissions Spectrometry (ICP, Applied Research Laboratory 3560) following

211 digestion in 10% HCl (Munter and Grande 1981) at the University of

212 Minnesota's Research Analytical Laboratory (n=1) (Table S2).

213 Sites varied in accessibility and the frequency and timing of research 214 visits, thus the date of initial deployment ranged from December, 2009 to 215 October, 2010, Litter bags were sequentially harvested at approximately 216 annual intervals, and litter was cleaned of any material other than colonizing 217 microbes, dried (65°C to constant mass), weighed, and sent to the University 218 of Minnesota for further processing. At the University of Minnesota, 219 harvested litter was ground, milled, and analyzed for total carbon (C) and N 220 as above, and a subsample was ashed (600°C for 6 hours) to determine ash-221 free dry mass (AFDM). Proportion initial mass remaining was converted to 222 proportion initial C remaining for further analyses to account for soil 223 contamination, as soil has low C concentration relative to litter. This method 224 is similar to correcting for soil contamination using ash-free dry mass, since 225 they are highly correlated, as confirmed here (across all samples in this 226 study: %C = 0.75 + 0.50*%AFDM, $R^2 = 0.77$, P<0.0001). We handled missing 227 %C data and outlier proportion C remaining values as described in the 228 Supporting Information.

Not all participating sites were able to sample litter bags for the full duration of the study due to investigator turnover, natural disasters (floods, herbivory, fires), and unnatural disasters (e.g., loss of sites to development). In some cases, investigators were unable to visit sites annually. Therefore, the total number of sites included in the final dataset varied from year to year (Table S1). Herein we present data from 20 sites that contributed data from at least three, and up to seven, harvests (Fig. S1).

236 Temperature, precipitation, solar radiation, and soil physical and chemical237 properties

238 To understand variation in decomposition across and within sites as 239 well as interactions between nutrient addition treatments and potential 240 covariates, we considered modeled site-level predictors, including mean 241 annual temperature (MAT), mean annual precipitation (MAP), potential 242 evapotranspiration (PET), aridity index (AI, MAP/PET) (Hijmans et al. 2005), 243 and precipitation distribution (Precip. Dist., precipitation in wettest 244 month/MAP; unitless), atmospheric N deposition (Ackerman et al. 2019), and 245 annual top of atmosphere insolation (NASA GISS 246 https://data.giss.nasa.gov/modelE/ar5plots/srlocat.html; Hansen et al. 2005). 247 We also considered the following plot-level data: pretreatment soil %C and 248 %N (determined as above, Crowther et al. 2019), Mehlich 3-extractable P, Mn, and base cations (K, Ca, Mg, Na) (Waypoint Analytical, Memphis, TN, 249 250 USA, Mehlich 1984), and pH (1:1 soil:water suspension, Waypoint Analytical, 251 Memphis, TN, USA). Aboveground plant biomass was measured by clipping, 252 drying, and weighing all live and dead plant biomass from two 0.1 x 1 m

253 strips/plot at peak biomass annually (2010-2016).

254 Overview of Analytical Approach

255 In our analytical approach, we (1) assessed fit of proportion C mass 256 remaining to four different decomposition models (single, double, and 257 asymptotic exponential; Weibull) that differ in how they represent 258 decomposing litter (e.g., as a single pool decomposing at one rate versus as 259 multiple pools decomposing at different rates); (2) fit litter N dynamics to a 260 guadratic function to determine the extent and timing of litter N 261 immobilization during decomposition; (3) analyzed effects of factorial N \times P \times 262 K_{μ} treatments on parameters obtained from the model-fitting described in (1) 263 and (2), using treatment-only mixed-effects models, which included N, P, and 264 K_u treatments as fixed effects and site and experimental block as nested 265 random effects; (4) analyzed for potential interactive effects of N, P, and K_{μ} 266 fertilization and edaphic and environmental site characteristics on 267 decomposition model parameters, using treatment-covariate models, which

268 included single covariates in mixed-effects models; and (5) used model

269 selection to identify edaphic and environmental variables that most strongly

270 explained cross-site variation in decomposition model parameters.

271 Decomposition model fitting

272 Having up to seven harvests over seven years allowed us to fit litter 273 proportion C remaining over time to four alternative decomposition models 274 that vary in mathematical structure, and thus in their representation of the 275 biophysical process of litter decomposition and chemical composition of the 276 litter (Olson 1963, Wieder and Lang 1982, Cornwell and Weedon 2014). This 277 allowed us to explore whether nutrients had different effects in the early 278 versus later stages of decomposition. The single exponential decay model 279 describes litter as a single pool with a constant decay rate, k_s , and assumes a 280 constant fraction of litter mass is decomposed per unit time, t. The 281 proportion of initial litter C remaining, X, is a function of k_s and t:

- $X = e^{-k_s t}$ Eqn.
- 283

282

1

 $X = i^{-k_1 t} + (1 - L) e^{-k_2 t}$

The *double* and *asymptotic exponential models* divide litter into two pools that decompose at different rates. In the double exponential model, a labile fraction, *L*, decomposes at rate k_1 , while the remaining slow fraction (*1-L*) decomposes at rate k_2 :

288

Eqn. 2

289 In the asymptotic model, *A* describes the slow fraction, which decomposes at 290 rate zero, while the remaining labile fraction (1-A) decomposes at rate k_a :

291

 $X = A + (1 - A) e^{-k_a t}$ Eqn. 3

Although the decomposition rate of A would not realistically be zero, A
approximates a fraction of litter for which the decomposition rate is
negligible (Berg 2000).

These first three models represent litter as having one or two discrete (pools", each with a constant decomposition rate. A fourth model, the *Weibull model*, characterizes litter decay as a continuous distribution of

298 residence times, as might be expected given the diverse chemistry of

299 constituents found in litter. This distribution is described by a scale

300 parameter, β , and a shape parameter, α (Fréchet 1927, Weibull 1951,

301 Cornwell and Weedon 2014):

4

 $X = e^{-\left(\frac{t}{\beta}\right)^{a}}$

302

303

304 The shape parameter, α , indicates degree of divergence of decomposition 305 rates between the early versus later stages: $\alpha > 1$ indicates that 306 decomposition is slow early on and then accelerates in the later stages, as 307 might occur if there is a lag period before decomposition begins, whereas α 308 < 1 indicates more rapid early-stage decomposition relative to the later 309 stages, as might occur if more labile constituents decompose first, followed 310 by more complex constituents. When α equals 1, the Weibull model collapses 311 to a single-exponential decay model where the instantaneous decay rate is

Eqn.

Eqn.

312 constant and
$$k_s = \frac{t}{\beta}$$
.

313 As the rate of decomposition described by the Weibull model changes 314 continuously with time, we cannot compare individual k values associated 315 with discrete litter pools. We therefore used the Weibull framework to 316 calculate the time to 10%, 25%, and 50% mass loss ($t_{1/10}$, $t_{1/4}$, $t_{1/2}$, respectively) 317 and the litter mean residence time (MRT) (Cornwell and Weedon 2014) as: 318 Eqn. 5 11

- 319 and
- 320
- $MRT = \beta \Gamma \left(1 + \frac{1}{\alpha} \right)$
- 321

where p is the proportion of litter mass remaining at time t and Γ is the 322 323 gamma function.

324 To determine the most appropriate decomposition models to describe 325 litter decay dynamics, we initially fit decomposition models to pooled data 326 from each site x treatment combination (3-5 plots per pool) and compared

327 model AIC_c values to assess the capacity of individual decomposition model 328 structures to describe decomposition dynamics across treatments (Burnham 329 and Anderson 2002). Differences of >3 between model AIC_c values (Δ AIC_c) 330 were used to identify significant differences between model fits. Exponential 331 models were fit using the *mle2* function in the *bblme* package (Bolker 2020) 332 and Weibull models were fit using the *nls* function. To evaluate whether 333 nutrient addition changed which model best fit the data, we compared which 334 model was the best fit to the data for all plots that did vs. did not receive N, 335 P, or K_{μ} , in three separate contingency analyses.

Three of the four decomposition models (single-exponential, 336 337 asymptotic, Weibull) best described litter decomposition across sites and 338 treatments (see Results), so we then fit each of the three models to the 339 proportion C remaining from all litter bags harvested over time from an 340 individual plot. To identify outliers within the individual plot models and 341 obtain more precise parameter estimates for decomposition models, we 342 calculated the sum of the residual distances between the proportion C 343 remaining at each litter bag harvest and the predicted value in the Weibull 344 model, adjusted for the number of predicted values (n_{Pred}) in each model 345 according to the following relationship:

346

 $Model Deviance = \frac{\Sigma [X_{Pred} - X_{Obs}]^2}{n_{Pred}}$ Eqn.

347

348 Individual plot models with the highest 2.5% of all deviance scores 349 were evaluated individually for outliers. In these cases, we iteratively 350 removed individual data points and re-fit Weibull models to identify predictor 351 sets that resulted in the lowest model deviance. Individual harvest points 352 were removed from 22 of 494 decomposition curves and their removal did 353 not affect analyses presented below. We eliminated entire plots from the 354 analysis when outlier removal reduced the number of harvest points included 355 in the model prediction below three (2% of models fit to three points).

356 Model fitting to estimate changes in litter N dynamics

Nitrogen is often immobilized initially before being released from
decomposing litter (Staaf and Berg 1981). To describe changes in the litter N
pool throughout the decomposition period, we fit plot-level changes in litter
N pool size through time to quadratic functions with the form:

361

362

$$Z = (\beta_1 t + \beta_2 t^2) + \beta_0$$

363 where Z is the litter N pool (grams), t is the time since the start of the study (years), β_1 and β_2 are coefficients on the *t* and t^2 terms, respectively, and β_0 is 364 365 the initial litter N pool (grams) in each plot. Models were fit to plot-level 366 decomposition sequences. We used the guadratic function to calculate the maximum litter N pool (N_{max}) and the time to N_{max} (t_{Nmax}) during the litter 367 368 decomposition period. We excluded data from individual litter bag harvests 369 in which the litter C:N ratio fell outside the 95% confidence interval of the 370 mean for each harvest time point. We also excluded plots with three or fewer 371 data points describing litter N pool size.

372 Nutrient effects on litter decomposition

373 We evaluated effects of N, P, and K_{μ} addition on parameters describing early (Weibull $t_{1/10}$, $t_{1/4}$, and $t_{1/2}$; asymptotic k_a), late (asymptotic A), and 374 375 overall (single exponential k_s ; Weibull α , MRT) litter decomposition using 376 general mixed models (nlme R package) (Pinhiero et al. 2019) in which site 377 and block were included as nested random effects. These *treatment-only* 378 statistical models were specified as a full-factorial block design in which N, P, and K_{μ} treatments were included as separate fixed factors. Treatment-only 379 380 models were also used to analyze treatment effects on litter N dynamics, 381 N_{max} and t_{Nmax} .

Using *treatment-covariate* statistical models, we considered whether inclusion of single covariates – soil pH, soil C (%), atmospheric N deposition rate (kg N ha⁻¹ y⁻¹), soil P (ppm), soil Mn (ppm), MAT (°C), MAP (mm),

385 precipitation distribution, annual insolation (W m⁻²) or aboveground biomass 386 $(q m^{-2})$ – interacted with nutrient addition treatments or altered the 387 significance of treatment effects. We fit individual three-way ANCOVA models 388 that included N, P, and K_u fertilization treatments in factorial combination 389 with each covariate (covariates were considered individually because of the 390 high number of covariates relative to the number of sites; site and block 391 were included as nested random effects; *Ime* function; *nIme* R package; 392 Pinhiero et al. 2019). All predictor and response variables were log or squareroot transformed to approximate normality, as appropriate. 393

394 Cross-site variation in litter decomposition

395 We used *model selection* to identify predictor variables that best 396 described cross-site variation in decomposition model parameters (Weibull α_{i} 397 MRT, $t_{1/10}$, $t_{1/4}$, and $t_{1/2}$; asymptotic k_a , A; single-exponential k_s). We considered 398 predictors describing climate and edaphic conditions, including MAT, MAP, AI, 399 PET, and precipitation distribution; soil C, N, P, Mn, and base cation content; 400 soil pH; and modeled N deposition rate. After screening potential predictors 401 for collinearity (Pearson R > |0.70|, Table S3; Hmisc R package; Harrell 402 2020), we excluded AI, which was highly correlated with MAT; PET, which 403 was highly correlated with MAT and AI; soil N, which was highly correlated 404 with soil C; the sum of the extractable base cations, which was highly 405 correlated with pH; and annual insolation, which was highly correlated with 406 MAT (Pearson's R > 0.70). The final set of predictors is presented in Table 407 S3.

We fit individual global mixed-effects models with all potential
predictors (MAT, MAP, Precip. Dist., soil C, soil P, soil Mn, soil pH, annual N
deposition rate, and aboveground plant biomass; Table S3) for each
response variable using the *lme* function in the *nlme* R package (Pinhiero et
al. 2019). We then generated a full submodel set from the global model
using the 'dredge' function and considered all two-way interactions (*MuMIn* R
package) (Nakagawa and Schielzeth 2013). McLaughlin Natural Reserve and

Hopland Research and Extension Center in California, USA, did not report
initial soil chemistry, and thus sites were excluded from the model selection
analysis, while Val Mustair in Europe was excluded as modeled N deposition
data were unavailable. Experimental block nested within site was included as
a random effect in each model, but fertilization treatments were excluded
from models.

We present the top models with \triangle AIC < 3 and conditional model average associated with each parameter. For individual candidate models with \triangle AIC < 3, we calculated the variance explained by the fixed effects in isolation (marginal R²) and that explained by both the fixed and random effects in combination (conditional R²; *MuMIn* R package) (Nakagawa and Schielzeth 2013). All analyses were conducted using R version 3.5.3 (R Foundation for Statistical Computing, Vienna, Austria, 2018).

428 Results

429 Across all sites and treatments, the median percentage C remaining at 430 the end of the experiment was 36%. Sites with the fastest decomposition 431 were Cowichan, British Columbia; Spindletop Farm, Kentucky; University of 432 North Carolina; and Hall's Prairie, Kentucky (with 4, 7, 12, and 14% initial C 433 remaining on average across treatments, respectively). Sites with the slowest decomposition were Sheep Experiment Station, Idaho; Kinypanial, 434 435 Victoria: and Sagehen Creek Field Station, California (with 55, 59, and 62%) 436 initial C remaining, respectively). Formal comparisons of mass loss among 437 treatments and sites were not possible because of different duration of litter 438 deployment across sites. Hereafter, we focus on analyses of parameters 439 obtained from fitting decomposition models.

440 Decomposition Model Fits

Using AIC_c was to compare model fits, within the exponential decay
model family, single and asymptotic decomposition models provided the best
fits to the data, whereas the double exponential model was a poor fit to the
data (Table S4.A). Considering the Weibull decomposition model along with

the exponential decay models, the Weibull model fit the data as well as the
best exponential model fit in 61% of all *site*treatment* combinations (Table
S4.B). Because AIC_c was consistently higher for the double exponential
model than the other models, we therefore fit the single, asymptotic, and
Weibull models to all individual plots.

450 While neither N, P, nor K_u changed which decomposition model best fit 451 the data (Table S5.A), by contrast, the best model to describe litter 452 decomposition differed across sites (Table S5.B). The single exponential 453 model best described the majority of the treatments at 13 sites, the 454 asymptotic model at three sites, and the Weibull model at one site. At two 455 sites, the single exponential and Weibull models were the best fit for an 456 equal number of treatments. The double exponential was never the best 457 model for any treatments at any sites.

458 Effects of experimental nutrient addition on early stage litter decay ($t_{1/10}$, $t_{1/4}$, 459 $t_{1/2}$, k_a)

460 In treatment-only statistical models, N addition (but not P or K_{μ}) greatly 461 accelerated early-stage decomposition, decreasing the Weibull time to 10% 462 and 25% mass loss ($t_{1/10}$ and $t_{1/4}$) and increasing the initial decay rate (asymptotic k_a), by 16%, 12%, and 23% respectively (Fig. 2, Table S6). 463 464 Similarly, in treatment-covariate models, N fertilization (but neither P nor K_{μ}) 465 reduced $t_{1/10}$ and $t_{1/4}$ and increased asymptotic k_a (Table S7). However, in 466 contrast to our expectations, N-induced increases in early-stage litter 467 decomposition were not stronger in sites with high P or base cation 468 availability. There were some weak interactive effects of fertilization and 469 other site factors on early-stage decomposition (Table S7), with N more 470 strongly increasing early-stage decomposition (k_a) in sites with low soil Mn 471 content (Fig. S1) and pH (Fig. S2). K_{μ} increased $t_{1/10}$ and $t_{1/4}$ in sites with more 472 evenly distributed precipitation (not shown), while P and K_{μ} increased $t_{1/2}$ and 473 decreased $t_{1/4}$, respectively, in sites with high pH (Fig. S2).



475 Effects of nutrient addition on late stage and overall litter decay (α , MRT, k_s ,

476 *A*)

474

477 In contrast to the early stages of litter decomposition, in treatment-

478 only models, there were no significant main effects of any nutrients on the

- 479 single exponential decomposition rate (k_s) (Fig. 3; Table S6). However, both N
- 480 and P, but not K_{μ} , increased the Weibull MRT, and N decreased Weibull α and
- 481 increased the fraction of slowly decomposing C (asymptotic A). Similarly, in
- 482 treatment-covariate models, N increased Weibull MRT and A (Table S7),
- 483 indicating that N caused rates to slow as decomposition proceeded.
- 484 However, N effects on early and late-stage decomposition were largely

- 485 offsetting, as N had only a small effect on the single-exponential decay
- 486 model (k_s), increasing it by 6% on average (Fig. 3, Fig. 4, Table S7). The
- 487 decrease in Weibull α with added N, from values > 1 to values closer to 1,
- 488 indicated that N accelerated early-stage decomposition (Fig. 4). Phosphorus
- 489 also generally increased Weibull *MRT* and sometimes increased the fraction
- 490 of slowly decomposing C (A), indicating it also tended to slow the later stages
- 491 of decomposition. Phosphorus also affected litter half-life $(t_{1/2})$ in models that
- 492 included covariates, but complex N x P x K_{μ} interactions made those effects
- 493 difficult to interpret (Table S7).



Figure 3. Effects of nutrient addition on decomposition parameters describing the entire time course of decomposition, (a) single exponential k_s , (b) Weibull *MRT*, and (c) Weibull α ; or later stages of decomposition, (d) asymptotic *A*. For visualization purposes, means (S.E. bars) of untransformed data are shown for each treatment (pooled across sites). Mixed-effects treatment-only model results are shown in Table S6 and treatment-covariate models are shown in Table S7. Shaded region shows mean and standard error in control treatment to make visual comparison among treatments easier

- 494
- 495 Fertilization interacted with several site factors in its effects on late-496 stage and overall decomposition dynamics (Table S7). N increased single 497 exponential k_s more at sites with low than high N deposition (albeit weakly, 498 Fig. S3). Nitrogen also increased k_s in sites with low Mn (Fig. S1), but reduced 499 k_s and Weibull α , and increased Weibull $t_{1/2}$ and *MRT*, and asymptotic *A* more 500 where precipitation was more evenly distributed across the year (low Precip.

- 501 Dist; Fig. S4). Finally, N reduced Weibull α and k_s and increased Weibull *MRT*
- 502 more at sites with higher aboveground biomass (Fig. S5).



⁵⁰³

Nitrogen was typically immobilized in leaf litter in the initial stages of decomposition. Plots with faster early-stage decomposition (higher asymptotic k_{a} , lower Weibull $t_{1/10}$) exhibited higher maximum litter N pools (N_{max} , Table S8). Higher N_{max} was also associated with increased litter *MRT* and asymptotic *A*, as well as reduced Weibull α . In treatment-only models, N fertilization increased litter N_{max} (N: p < 0.0001), as well as the time to the maximum litter N content (p = 0.031, Table S9, Fig. S6).

511 Variables predicting litter decay dynamics across sites

512 In the model selection analysis, climatic and edaphic factors were 513 unable to explain substantial variation in decomposition model parameters 514 describing the early stages of decay (Weibull $t_{1/10}$, $t_{1/4}$; asymptotic k_a ; Table 515 S10). Less than 10% of the cross-site variation in these parameters was 516 captured by fixed effects, indicating that most of the variation among sites in 517 the initial decomposition rates was caused by unmeasured site factors (Table 518 S10). 519 By contrast, MAT, background N deposition, and precipitation variation

520 were modest to strong predictors of Weibull half-life ($t_{1/2}$), parameters

- 521 describing later-stage decomposition (asymptotic A), or parameters
- 522 describing the entire decomposition curve (Weibull α , MRT; single
- 523 exponential k_s) (Table S10). Generally, warmer sites had faster
- 524 decomposition (lower $t_{1/2}$, lower Weibull *MRT*), sites with more variable
- 525 seasonal precipitation (higher Precip. Dist.) had slower decomposition
- 526 (longer Weibull *MRT*, lower k_s), and sites with higher levels of atmospheric N
- 527 deposition had faster decomposition (lower Weibull MRT, lower asymptotic A,
- 528 Tables S7, S10).

529 Discussion

530 Contrasting effects of N on early versus late-stage decomposition

In a long-term (seven-year) experiment replicated across grasslands 531 532 spanning diverse biotic, edaphic, and climatic conditions, N consistently 533 accelerated early-stage decomposition. This points to N limitation of 534 microbes decomposing the N-poor Quercus litter, whereby added N 535 increased decomposer C demand. Neither P nor K_µ, alone or in combination 536 with N, limited decomposition in its early stages, in contrast to nutrient co-537 limitation of NPP in grasslands (Fay et al. 2015, Cleland et al. 2019, Du et al. 538 2020). Decomposer N demand likely was met via fungal translocation of soil 539 N in control plots (Frey et al. 2000, Chigineva et al. 2011, Xiong et al. 2014), 540 but via fertilizer-supplied N in N-treated plots. Accordingly, litter in the N-541 fertilized plots exhibited less initial delay of decomposition and greater N 542 immobilization than litter in control plots. Our findings corroborate 543 observations that across diverse litter types, low-N litter types exhibited 544 lagged decomposition dynamics relative to high-N litter (Cornwell and 545 Weedon 2014). In addition to alleviating N limitation, N may have affected 546 early-stage decomposition via other mechanisms. A study in semi-arid 547 grassland attributed positive effects of N on decomposition to acidification 548 that increased Mn availability and altered the microbial community (Hou et 549 al. 2021). However, while N fertilization also reduced soil pH (at 10/20 sites) 550 in the present study, the magnitude of pH reduction was unrelated to the

effects of N on decomposition model parameters (data and analyses notshown).

553 Our findings corroborate past studies demonstrating widespread N 554 stimulation of early-stage decomposition, and are the first to show this using 555 a long-term experiment in diverse grasslands. A short-term study of 556 decomposing tea leaves across 21 Nutrient Network sites also found positive 557 N effects on early-stage decomposition (Ochoa-Hueso et al. 2020). 558 Phosphorus addition also positively affected early-stage decomposition in 559 that study perhaps because the relatively high N concentrations in tea 560 increased microbial P demand. In a meta-analysis of N fertilization effects on 561 decomposition in diverse ecosystems, N fertilization reduced $t_{1/10}$ and $t_{1/4}$ and 562 increased Weibull MRT and asymptotic k_a , as found here (Gill et al. 2021). 563 Our findings are also consistent with results from some local to regional cross-site studies, predominantly in forests, showing N stimulation of initial 564 565 decomposition (Berg and Matzner 1997, Hobbie et al. 2012).

566 In contrast to our hypothesis, N inhibited later-stage decomposition 567 (increasing asymptotic A and increasing Weibull *MRT*), as found in many 568 forest studies (Berg and Matzner 1997, Whittinghill et al. 2012, Gill et al. 569 2021), and in a short-term grassland study, where N increased the Tea Bag 570 Index stabilization factor (a modeled representation of the biochemically 571 transformed tea leaves that persist at late stages of decomposition) (Ochoa-572 Hueso et al. 2020). Inhibition of late-stage decomposition by added N has 573 been attributed to several interrelated mechanisms including suppression of 574 oxidative enzyme activity: formation of recalcitrant compounds from 575 accumulated microbial necromass or reactions of inorganic N with litter 576 metabolites; Mn limitation; and suppression of microbial biomass by acidification (Fog 1988, Berg and Matzner 1997, Carreiro et al. 2000. 577 578 Treseder 2008, Zak et al. 2008, Hobbie et al. 2012, Entwistle et al. 2018, 579 Morrison et al. 2018, Whalen et al. 2018, Gill et al. 2021). The concordance 580 of results presented here with those from forest and short-term grassland

581 studies points to the generality that elevated N slows late-stage582 decomposition in terrestrial ecosystems.

583 We hypothesized that N would not inhibit late-stage decomposition in 584 grasslands because of relatively low plant lignin content, and associated low 585 basidiomycete abundance and lignin-degrading oxidative enzyme activity 586 (Sinsabaugh 2010). Indeed, meta-analyses published well after our study 587 began show little evidence that N inhibits oxidative enzyme activity in 588 grasslands (Jian et al. 2016, Chen et al. 2018, Xiao et al. 2018). However, 589 depressive effects of N enrichment on grassland microbial biomass (Ramirez 590 et al. 2012) could have slowed decomposition.

591 It is also possible that alleviation of microbial N limitation early in 592 decomposition led to necromass accumulation on litter surfaces and 593 contributed to the apparent reduction in late-stage litter because of 594 necromass constituents that decompose more slowly than the plant 595 compounds consumed by those decomposers (Liang et al. 2019, Gill et al. 596 2021). Increased N immobilization in response to N fertilization, which was 597 associated with reduced rates of late-stage decomposition, provides indirect 598 support for this mechanism. Concurrent measures of enzyme activity and 599 decomposer necromass are needed to distinguish among these potential 600 mechanisms.

601 Contrasting N effects early and late in decomposition, as found here, 602 have implications for N effects on SOM. Early in decomposition, N could 603 promote litter processing and transfer of low molecular weight compounds to 604 soils, potentially enhancing stabilization of mineral-associated organic matter 605 (MAOM). By contrast, larger slowly decomposing litter fractions could 606 promote accumulation of particulate organic matter (POM) and total SOM 607 (Cotrufo et al. 2015). Interestingly, a study conducted in nine North American 608 NutNet sites found no effects of N on MAOM or POM (Keller et al. 609 unpublished), raising uncertainty about whether differential effects of N over 610 the course of litter decomposition have consistent consequences for SOM.

611 Utility of higher-order models in describing litter decomposition dynamics

612 Multi-parameter empirical models illuminated divergent N effects early 613 vs. late in decomposition. Thus model structure choice influences 614 interpretation of litter decomposition dynamics, with implications for 615 experimental design and hypothesis testing (Cornwell and Weedon 2014). Single-parameter models readily fit data from experiments with fewer 616 617 collection points, but may mischaracterize decomposition dynamics that shift 618 through time. Herein, the single exponential model often provided equivalent 619 AIC_c fit to two- or three-parameter models, suggesting that the single 620 exponential model sufficiently described the complexity of grassland litter 621 decomposition. However, the Weibull model structure fit the data equally 622 well or better than the single exponential model for 54% of site-treatment 623 combinations, and the asymptotic model fit the data equally well or better for 83%. As the fertilization responses observed here and in other recent 624 625 syntheses (Gill et al., 2021) emerged only when models allowed 626 decomposition rates to change through time, these higher-order model 627 structures deserve consideration in both experimental design and data 628 analysis.

629 Whether N addition would differently affect *Quercus* litter 630 decomposition relative to native litter at each site due to variation in litter 631 chemistry, lack of home-field advantage effects (Palozzi and Lindo 2018), or 632 sterilization prior to deployment, is unclear. In a study across eight locations 633 at one of our study sites (Cedar Creek), where the litter used here originated, 634 litter of *Ouercus ellipsoidalis* and *Schizachvrium scoparium*, a dominant C4 635 grasses in Cedar Creek grasslands and savannas, had similar initial acid-636 nonhydrolyzable fractions (lignin and other recalcitrants) and N concentrations, similar rates of decomposition, and responded 637 638 idiosyncratically to added N in grasslands in the early and late stages of 639 decomposition (Hobbie 2005, 2008). However, the *Quercus* litter used here 640 had higher concentrations of lignin (Table S2) than would be expected for 641 herbaceous species on average (e.g., Oliveira et al. 2020), which could have

accentuated negative effects of N on the later stages of decomposition if the
underlying mechanism was related to N inhibition of oxidative enzyme
activity.

645 Interactions between nutrient treatment effects and site factors

646 We found limited support for hypothesized interactions between site 647 factors and nutrient treatments on decomposition, suggesting that N effects 648 (and lack of P and K_{μ} effects) were general across diverse sites. We had expected that nutrients would limit decomposition more in warm, wet 649 650 regions, yet N limitation of early-stage decomposition was largely 651 independent of climate. However, negative effects of N on late-stage 652 decomposition were strongest in sites where precipitation was more evenly 653 distributed across the year. As the frequency of wetting events regulates 654 nutrient diffusion to litter bags, more consistent precipitation regimes may 655 support elevated N environments throughout the year, contributing to 656 elevated microbial biomass and higher necromass stabilization. Negative N 657 effects also increased with aboveground biomass, which may protect litter 658 from UV radiation and prevent decomposer desiccation, supporting increased 659 necromass stabilization.

Across 20 sites with widely varying geography, results did not support
our hypothesis of greater N limitation of decomposition in younger, less
weathered sites, and greater P limitation in older more weathered sites.
Instead, N increased early-stage decomposition regardless of soil chemistry
and climate, and the only consistent P effect was to increase Weibull *MRT*.

665 Variation in decomposition across sites

Mean annual temperature, precipitation variability, and atmospheric N
deposition all contributed to variation in decomposition rates across sites,
mostly later in decomposition. Faster decomposition in warmer sites,
observed previously in global decomposition syntheses (Gholz et al. 2000,
Zhang et al. 2008, Kwon et al. 2021), is expected because of greater
biological activity, longer growing seasons, and higher nutrient availability.

672 Greater photodegration in warmer sites could also have contributed to faster 673 decomposition in those sites, given that MAT was highly correlated with 674 annual insolation. Faster decomposition in sites where precipitation was 675 distributed more evenly relative to annual precipitation likely reflects the 676 importance of moisture in limiting decomposition in grasslands, especially at 677 times of year when temperatures are favorable (Meentemeyer 1978, Gholz 678 et al. 2000). Indeed, a manipulation of rainfall variability showed that lower 679 precipitation variability, resulting in higher average soil moisture, led to more 680 rapid decomposition (Walter et al. 2013).

681 Positive relationships between k_s and N deposition provide further 682 evidence that N limits decomposition across grasslands, and contrast results 683 from a cross-site study of tea leaf decomposition worldwide, which showed 684 negative relationships between atmospheric N deposition and mass loss over 685 3 and 12 months in temperate sites (Kwon et al. 2021). Effects of N from 686 deposition and fertilizer differed in key ways. First, whereas fertilizer N 687 reduced the initial lag period of slow decomposition, atmospheric N 688 deposition was associated with a longer lag period. This discrepancy could 689 have arisen if long-term effects of chronic N addition were not yet apparent 690 in fertilized plots, which began receiving fertilizer <1 to 2 years before litter 691 bags were deployed. For example, N deposition might have altered microbial 692 communities, as seen in long-term (> 8 y) fertilization studies (Frey et al. 693 2004, Hesse et al. 2015, Entwistle et al. 2018, Whalen et al. 2018, Hou et al. 694 2021), in ways that slowed colonization of litter by decomposer organisms 695 and reduced litter-soil hyphal connections. Other effects of chronic N 696 deposition associated with soil acidification (e.g., reduced base cation 697 supply) (Hou et al. 2021) might also have manifested slowly in N fertilization 698 treatments.

699 While sites with higher N deposition had smaller litter slow pools, N 700 fertilization increased the slow pool. As discussed previously, numerous long-701 term decomposition studies have demonstrated that N fertilization increases 702 the slow pool (asymptotic *A*). Fewer studies evaluated relationships of *A* with

703 ambient N deposition rates. Rates of N addition in fertilization studies,

704 including ours, are often higher than the highest rates of N deposition in

705 polluted regions. Herein, N was added at a rate of 100 kg N ha⁻¹ y⁻¹, whereas

706 the highest rate of N deposition across sites was ca. 20 kg N ha⁻¹ y⁻¹.

707 Although the specific mechanism by which added N increased the slow pool

is unknown, perhaps its effects only occur at relatively high rates of added N.

709 Unexplained site variation

710 Initial decomposition rates varied widely among sites, but in ways not 711 explained by measured site climate or edaphic factors. Across six grassland 712 sites in the US Central Great Plains, decomposition of common aboveground 713 substrates was similarly unrelated to climate factors (Bontti et al. 2009) and 714 a cross-sites study of decomposing tea leaves similarly found a high 715 percentage of unexplained variation across grassland sites (Kwon et al. 716 2021). What other factors might have contributed to variation in initial 717 decomposition across sites? Variation in the decomposer community (Prober 718 et al. 2015) and how guickly it colonized the litter could have contributed to 719 site variation in initial decomposition because of variation in metabolic 720 capabilities of microbial decomposers (Strickland et al. 2009, Gessner et al. 721 2010, Bray et al. 2012, Glassman et al. 2018). Other factors, such as the 722 contributions of soil fauna to decomposition or vascular plant community 723 composition, might also have contributed to site-to-site variation in 724 decomposition (García-Palacios et al. 2013). For example, the site with the 725 highest total decomposition (Cowichan) is dominated by densely growing 726 rhizomatous grasses, whose roots completely encased the litter bags after 727 several years. Previous decomposition trials at this site also observed rapid 728 biomass decomposition (Ziter and MacDougall 2013). Photodegradation is 729 also important for aboveground decomposition in grasslands and could have 730 contributed to site variation in decomposition dynamics (Austin and Vivanco 731 2006, Adair et al. 2017), although greater aboveground biomass was 732 associated with faster k_s in control plots (but not in plots receiving N),

733 opposite what might be expected if photodegradation contributed 734 substantially to initial decomposition rates. The fiberglass mesh used in this 735 study transmits about 45-50% of solar radiation (Adair et al. 2017), so photodegradation could have contributed to mass loss. We were unable to 736 737 evaluate the effects of PAR at the ground surface on decomposition across 738 sites because of lack of data, so we cannot rule this out as a potential 739 mechanism. Nevertheless, variation in irradiance, soil contamination of the 740 litter bags (Brandt et al. 2010, Adair et al. 2017), and position of the bags 741 relative to the ground surface, standing, and fallen vegetation (Lin and King 742 2014) could have caused variation in photodegradation and biotic 743 decomposition.

744 Concluding Remarks

745 In a long-term (seven-year) study of decomposition of a common leaf 746 litter substrate in a factorial fertilization experiment replicated at 20 747 grassland sites spanning a wide range of biotic, edaphic, and climatic 748 conditions, N fertilization consistently increased early-stage and slowed late-749 stage decomposition, as often seen in forests. These patterns were 750 discernable because of the long duration of the study, the large number of 751 sites, and the use of decomposition models that could distinguish early- and 752 late-state decomposition dynamics. These findings corroborate those of Gill 753 et al. (2021) in showing that fitting decomposition data to single exponential 754 models can obscure effects of environmental factors on decomposition 755 dynamics. Implications of contrasting early- versus late-stage effects of N on 756 litter decomposition for SOM pools and dynamics deserve further empirical 757 study and exploration using ecosystem models.

758 Acknowledgments

759 This work was supported by grants from the National Science Foundation,

- 760 including the NSF Ecosystems Studies program (NSF DEB-1556529), the
- 761 Cedar Creek Long Term Ecological Research project (NSF DEB-0620652,
- 762 DEB-1234162), and the Research Coordination Network program (NSF DEB-

- 763 1042132). Additional support for coordination and data management came
- 764 from the University of Minnesota's Institute on the Environment (DG-0001-
- 765 13). We thank the Minnesota Supercomputer Institute for hosting project
- 766 data and the Institute on the Environment for hosting Network meetings. Gill
- 767 was supported by a University of Minnesota College of Biological Sciences
- 768 Grand Challenges Postdoctoral Fellowship. We are grateful to numerous
- 769 people who helped construct, deploy, collect, and process litter bags, and
- 770 prepare litter for and conduct laboratory analyses, including, but not limited
- to, Lori Biederman, Jennifer Firn, Paul Frater, Eric Lind, and Bonnie McGill.

772 Author Contribution

- 773 SEH and ALG contributed equally. ETB, EWS, WSH, SEH, and KSH designed
- the research. ETB, CRB, EEC, CMD, KFD, DSG, WSH, SEH, ASM, RLM, BAM,
- 775 JLM, JWM, ACR, MS, EWS, JW, and LHY performed the research. PBA, ETB,
- 776 CRB, EEC, CMD, KFD, ALG, DSG, WSH, SEH, ASM, RLM, BAM, JLM, JWM, ACR,
- 777 MS, EWS, JW, and LHY contributed to data analysis, collection or
- 778 interpretation. PBA, ETB, EEC, CMD, KFD, ALG, WSH, SEH, KSH, ASM, RLM,
- ACR, MS, EWS, JW, and LHY contributed to writing the manuscript.

780 Data Availability

- 781 Data are archived and available: <u>https://www.cedarcreek.umn.edu/research/</u>
 782 <u>data/dataset?aige247</u>,
- 783 doi:10.6073/pasta/2ebd7eaf2fac27e1c7eee9678baa7940. Code to replicate
- 784 analyses and figures are available:
- 785 <u>https://github.com/gill20a/NutNetLitterDecomp</u>.

786 Figure Legends

- 787 Figure 1. Distribution and meteorological conditions of 20 Nutrient Network
- 788 sites included in this study. Site were located in North America, Europe, and
- 789 Australia. (a) Map of site locations; (b) relationship between site MAT and
- 790 MAP; (c) relationship between site MAT and precipitation distribution; (d)
- 791 relationship between site MAP and precipitation distribution. Abbreviations as

- 792 follows: MAT, mean annual temperature (°C); MAP, mean annual precipitation
- 793 (mm); precipitation distribution (precipitation in wettest month/MAP).

794

795 Figure 2. Effects of nutrient addition on early stages of litter decomposition. 796 (a) Weibull time to 10% mass loss $(t_{1/10})$, (b) Weibull time to 25% mass loss 797 $(t_{1/4})$, and (c) initial litter decay rate (asymptotic k_a). For visualization 798 purposes, means (S.E. bars) of untransformed data are shown for each 799 treatment (pooled across sites). Mixed-effects treatment-only model results 800 are shown in Table S6 and treatment-covariate models are shown in Table 801 S7. Shaded region shows mean and standard error in control treatment to 802 make visual comparison among treatments easier.

803 Figure 3. Effects of nutrient addition on decomposition parameters describing 804 the entire time course of decomposition, (a) single exponential k_s , (b) Weibull 805 *MRT*, and (c) Weibull α ; or later stages of decomposition, (d) asymptotic A. 806 For visualization purposes, means (S.E. bars) of untransformed data are 807 shown for each treatment (pooled across sites). Mixed-effects treatment-only 808 model results are shown in Table S6 and treatment-covariate models are 809 shown in Table S7. Shaded region shows mean and standard error in control 810 treatment to make visual comparison among treatments easier. 811 Figure 4. Comparison of proportion C remaining over 10 years predicted by 812 (a) single exponential, (b) asymptotic exponential, and (c) Weibull models of

- 813 decomposition between treatments receiving (+N, green line) or not
- 814 receiving (-N, grey dashed line) N. Model parameters were obtained by
- 815 averaging across all replicates, sites, and appropriate treatments.
- 816

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