

Modelling vegetation dynamics in managed grasslands: Responses to drivers depend on species richness

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Modelling vegetation dynamics in managed 1

grasslands: responses to drivers depend on 2

species richness 3

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- Thibault Moulin^a, Antoine Perasso^a, François Gillet^{a,b,*} 5
- 6 ^a Université Bourgogne Franche-Comté, UMR CNRS 6249 Chrono-environnement, Besançon,
- 7 France
- 8 ^b Ecole Polytechnique Fédérale de Lausanne, Ecological Systems Laboratory, Lausanne,
- 9 **Switzerland**
- 10 * Corresponding author
- E-mail addresses: thibault.moulin@univ-fcomte.fr (T. Moulin), antoine.perasso@univ-11
- fcomte.fr (A. Perasso), francois.gillet@univ-fcomte.fr (F. Gillet). 12

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Abstract

- 16 In mathematical grassland models, plant communities may be represented by a various
- number of state variables, describing biomass compartments of some dominant species or 17
- 18 plant functional types. The size of the initial species pool could have consequences on the
- 19 outcome of the simulated ecosystem dynamics in terms of grassland productivity, diversity,
- 20 and stability. This choice could also influence the model sensitivity to forcing parameters.
- 21 To address these issues, we developed a dynamic grassland model, DYNAGRAM, designed to
- 22 simulate seasonal changes in both aboveground biomass production and species composition
- 23 of managed permanent grasslands under various soil, climate and management conditions. We
- 24 compared simulation results from alternative instances of DynaGraM that only differ by the
- 25 identity and number of state variables describing the green biomass, here plant species. We
- 26 assessed the sensitivity of each instance of the model to key forcing parameters for climate,
- 27 soil fertility, and defoliation disturbances, using univariate and multivariate regression trees
- 28 and dynamic trees.
- 29 Results of 10-year simulations under various climate, fertility and defoliation conditions
- 30 showed that the final total biomass was tending to increase with the size of the species pool,
- 31 while species evenness and the proportion of surviving species was tending to decrease. We
- 32 found a positive correlation between the species survival ratio and the defoliation intensity,
- 33 and this correlation increased with the initial species richness. The sensitivity to forcing
- 34 parameters of community structure and species evenness differed markedly among alternative
- 35 models, showing a progressive shift from high importance of soil fertility (fertilisation level,
- 36 mineralization rate) to high importance of defoliation (mowing frequency, grazing intensity)
- 37 as the size of the species pool increased. By contrast, the key drivers of total biomass
- 38 production were independent of species richness and only linked to resource supply (nitrogen
- 39 and water).
- 40 These results highlight the need to take into account the role of species diversity to explain the
- 41 behaviour of grassland models.
- 42 **Keywords**: dynamic trees; grassland model; multivariate regression trees; plant succession;
- 43 sensitivity analysis; species pool.

1. Introduction

- 45 Semi-natural permanent grasslands present important ecological, cultural and economic
- values, and support many ecosystem services due to their biodiversity (Gaujour et al., 2012;
- 47 Mauchamp et al., 2012). Some temperate European grasslands, managed for a long time with
- 48 low-intensity grazing or mowing, are the plant communities with the world records for
- 49 species richness at fine scale (Wilson et al., 2012).
- Various diachronic studies have revealed recent changes in the structure and composition of
- grassland plant communities in Europe (Gillet et al., 2016; Homburger and Hofer, 2012;
- Mitchell et al., 2017; Wesche et al., 2012). The causes of these changes are related to climatic
- (warming, extreme events) and anthropogenic (technical and socio-economic changes in
- 54 management) constraints and disturbances. Land-use intensification or abandonment may lead
- 55 to a severe reduction of species diversity with adverse impacts on ecosystem services.
- Fertilisation and defoliation regimes have been identified as the main drivers of plant species
- 57 richness in productive grasslands (Gaujour et al., 2012; Mauchamp et al., 2016, 2014).
- According to field observations, local species richness of temperate natural and semi-natural
- 59 grasslands shows unimodal relationship with productivity, with a maximum biodiversity for
- 60 intermediate productivity (Dengler et al., 2014). This questions how grassland ecosystem
- stability and productivity are influenced by vascular plant species richness. Field observations
- and experiments on semi-natural grasslands suggest that the most productive communities are
- 63 the least diverse, but also the least resilient. By contrast, field experiments based on artificial
- species mixtures showed a positive correlation between species richness and productivity
- 65 (Hector et al., 1999; Reich et al., 2012). A recent meta-analysis of experimental studies on
- such artificial species assemblages revealed that species richness generally improves
- 67 resistance of grassland productivity to climatic extreme events, but not its recovery rate (Isbell
- et al., 2015). Increasing species richness in such artificial communities stabilizes whole
- 69 community biomass but destabilizes the dynamics of constituent populations (de Mazancourt
- 70 et al., 2013; Gross et al., 2013).
- 71 Since experimental studies have not been able to provide a general explanation to the
- 72 relationships observed between grassland diversity and their productivity and stability,
- dynamical models are powerful tools for testing predictions of ecological theories. A lot of
- 74 mathematical models of grassland dynamics based on explicit ecological processes have been
- developed. Some non-exhaustive reviews of grassland models are available (Snow et al.,
- 76 2014; Taubert et al., 2012; Wiegand et al., 2008). Grassland models published so far have
- targeted monospecific swards (Duru et al., 2009; McCall and Bishop-Hurley, 2003;
- 78 Schapendonk et al., 1998), mixtures of a grass and a legume species (Lazzarotto et al., 2009;
- 79 Thornley, 1998), or a multispecies community constant in time (Jouven et al., 2006; Riedo et
- al., 1998). In a review of thirteen grassland models (Taubert et al., 2012), only two considered
- 81 more than three different species or plant functional types: GRAS, a succession compartment
- model (Siehoff et al., 2011) and STEPPE, a gap model of semiarid grasslands (Coffin and
- Lauenroth, 1990); however, none of them simulate biomass. So far, very few grassland
- 84 models have been designed to simulate both composition and biomass changes in
- 85 multispecies semi-natural grasslands used as pastures or/and hayfields, a requirement for
- 86 understanding diversity-productivity-stability relationships. Indeed, most of current grassland
- 87 models focus on the impact of agricultural practices on forage production with many
- 88 ecophysiological and biophysical details inherited from crop models but do not consider
- 89 temporal changes in taxonomic and functional composition. An attempt (Lauenroth et al.,
- 90 1993) to couple individual-based gap models describing changes in species composition
- 91 (STEPPE) and ecosystem models describing nutrient cycling and biomass production
- 92 (CENTURY) or water balance (SOILWAT) led to the development of the ECOTONE model,

93 able to simulate transitions between biomes (Peters, 2002). Several biophysical pasture models can be used to simulate growth of composite pasture types of several species under 94 95 various management strategies, such as CLASS-PGM, a generic pasture growth model (Vaze 96 et al., 2009), or DAIRYMOD and the SGS Pasture Model (Johnson, 2016), but these models are 97 rather complex, limited to five "species" and not optimized for mowing management. GEMINI 98 (Soussana et al., 2012) is another very detailed biophysical model of grassland ecosystems 99 that explicitly simulates competition among any number of clonal populations for two key 100 resources (light and nitrogen) along vertical canopy and soil profiles; plant functional traits 101 are used to explain the response of aboveground net primary production and of plant 102 community structure to various soil, climate and management conditions, including grazing, 103 cutting and fertilization. Some recent mathematical models are based on stochastic differential 104 equations adapted from the Lotka-Volterra equations to simulate changes in aboveground 105 biomass of plant species mixtures and to explain community dynamics by environmental and 106 demographic stochasticity (de Mazancourt et al., 2013; Lohier et al., 2016). 107 Taking into account the whole species diversity of grasslands in a mathematical model is 108 quite impossible, given the large number of vascular plant species that may potentially co-109 occur in such ecosystems. For example, the regional species pool in permanent grasslands of 110 the French Jura Mountains exceeds 1000 and the average number of species per local 111 grassland is about 30, ranging from 3 to 92 (Mauchamp et al., 2012). To reduce the number of 112 state variables describing the plant community in the model, a common solution is to consider 113 plant functional types (PFTs) instead of individual species. PFTs can be viewed as theoretical "average species" combining parameters from a set of more or less functionally related 114 115 species, such as legumes, perennial grasses or erect forbs. Typically, a PFT is a group of 116 species with similar functional traits explaining their response to constraints and disturbances, 117 such as management practices, and their effect on ecosystem function, such as forage production (Cruz et al., 2010, 2002; Duru et al., 2009; Graux et al., 2016). The GRAS model 118 119 (Siehoff et al., 2011) simulates the succession of both dominant species and PFTs described 120 by their cover dynamics, contrary to most agronomic grassland models, such as MODVEGE 121 (Calanca et al., 2016; Jouven et al., 2006), which simulate seasonal changes in one or several 122 biomass compartments, assuming a constant assemblage of species or PFTs. Individual-based, 123 spatially explicit community grassland models such as IBC-GRASS (Weiss et al., 2014) are 124 able to simulate metacommunities of many PFTs over hundred years, but at the price of a 125 simplified representation of edaphic and climatic drivers. By contrast, (Confalonieri, 2014) 126 suggested a model of plant community dynamics based on a single instance of a generic crop 127 simulator, state variables being only simulated for the whole community. EcoHyD (Lohmann 128 et al., 2017) is an ecohydrological model of semi-arid grasslands and savannas, able to 129 simulate cover dynamics of a large number of PFTs, but focuses on hydrological processes and does not implement nutrient stress. 130 131 In this paper, we address a general issue that could be raised in any modelling study: the 132 appropriate choice of the level of detail of the model, i.e. the number of state variables and 133

In this paper, we address a general issue that could be raised in any modelling study: the appropriate choice of the level of detail of the model, i.e. the number of state variables and associated parameters to be taken into account for describing the dynamical system, here the plant community. Although this choice is usually motivated by model objectives (e.g., with a focus on forage production or on biodiversity conservation) and by technical constraints (e.g., data availability, model performance), what are its consequences on the simulated dynamics of the virtual community in terms of grassland productivity, diversity, and stability? This question is especially relevant for permanent grasslands used for grazing or mowing and subject to seasonal changes in species composition and forage production, because of the high number of species coexisting in such ecosystems.

141 Thus, the first aim of the present study is to examine the consequences of increasing the

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number of state variables (i.e. the size of the local species pool) describing plant community

composition on model behaviour, regarding grassland productivity, diversity and stability. Specifically, we formulated four hypotheses: (H1) the inclusion of more species in the model increases its *productivity*, hence the final total biomass after ten-year simulations with various values of control parameters; (H2) the inclusion of more species in the model decreases its final species evenness, the component of community *diversity* measuring equitability in biomass and resource partitioning among species; (H3) the inclusion of more species in the model decreases its *stability*, hence the final proportion of surviving species (species survival ratio); (H4) species survival ratio increases with intensity and frequency of disturbances, and this positive correlation increases with the size of the species pool.

The second purpose is to assess the consequences of increasing the size of the species pool on the model sensitivity to key control parameters regarding climate, soil fertility and defoliation. We hypothesized that (H5) the hierarchy of influential parameters depends on species identity but not on the size of the species pool.

For these purposes, we need a simple mechanistic model in which the main ecological processes are implemented (seasonal mechanisms of plant growth, soil resource competition, response to climatic constraints and agricultural management) and in which we can vary the number and the identity of species in the initial plant community. As no such model was available, we developed DYNAGRAM, a simple and flexible dynamical model based on ordinary differential equations (ODEs), able to simulate various combinations of climate, soil and management conditions. The aim of this paper is not to present this model in detail along with results of calibration and validation with experimental data, which will be the topic of a next paper, but to use it as a tool to address two general relevant questions in ecological modelling. We simply ensured we properly modelled competition processes among species, under various forms of land use, by comparing simulated results to expert knowledge of grassland ecosystems (Cruz et al., 2010).

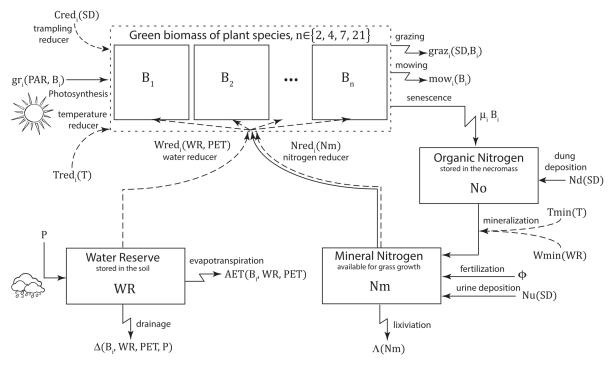


Fig. 1 Structure of the DYNAGRAM model. Solid lines denote flows of matter or energy and dashed lines denote reducers (mineral nitrogen acts both as a flow and as a reducer). State variables and forcing climatic variables are listed in Table 1 and a description of all parameters can be found in Tables A.1 and A.2 (Appendix A).

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2. Materials and methods

2.1. Model structure and main assumptions

- 177 DYNAGRAM is a process-based succession model that simulates the aboveground vegetation
- dynamics of a temperate grassland, both in term of composition and of forage production.
- Using a common structure (Fig. 1), we developed four alternative versions of the model,
- which correspond to different levels of diversity.
- We built DYNAGRAM under some key assumptions, in the way to capture only essential
- processes, and keeping as reduced as possible the dimension of the parameter space.
- Grassland vegetation is described by n state variables B_i representing the green biomass
- (leaves) for each plant species i. We did not consider the belowground biomass (roots) and the
- reproductive parts of the plants. Hence, we assumed no seed production and no recruitment
- from the seedbank. Two other state variables describe available resources in the soil: mineral
- nitrogen Nm and water reserve WR (Table 1).
- Basically, DYNAGRAM is a mechanistic model of resource competition (Tilman, 1980)
- operating at a daily time step. We mainly based the regulation of green biomass and plant
- 190 competition from the standpoint of resource dynamics. A single nutrient, mineral nitrogen
- 191 (nitrate or ammonium), is consumed by plants for growth and is supplied by a pool of soil
- organic nitrogen No, itself supplied by the senescence of standing biomass and litter
- 193 decomposition.

We considered four forcing climatic variables defined as simple sinusoidal functions of time

- and acting on different processes (Table 1). This simplified formulation of the annual cycle
- allows the manipulation of a reduced number of climatic parameters. Average daily
- temperature T acts as a limiting factor for plant growth and litter decomposition, daily rainfall
- 198 P fills the soil water reserve, photosynthetic active radiation PAR is a driver of biomass
- production by photosynthesis, and potential evapotranspiration *PET* determines the loss term
- in the soil water budget equation that may cause drought stress.

Description

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State

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Table 1 State variables and forcing climatic variables implemented in DYNAGRAM.

variables	•	•
Green biomass	B_i	Green biomass of plant species <i>i</i>
Organic nitrogen	No	Amount of organic nitrogen in the soil
Mineral nitrogen	Nm	Amount of mineral nitrogen in the soil available for plant growth
Water reserve	WR	Amount of water in the soil

Symbol

Forcing climatic variables	Symbol	Description
Temperature	T	Daily average air temperature acting on plant growth and mineralization rates
Precipitation	Р	Daily rainfall, the supply to the water reserve
Photosynthetically active radiation	PAR	Portion of the solar radiation spectrum driving photosynthesis and determining potential growth
Potential evapotranspiration	PET	Function that determines the loss term in the water budget

Green biomass dynamics for each species is described by the following ODE (1). For clarity reason, we only wrote the dependence on state variables:

$$\frac{dB_i}{dt} = gr_i(B_i) Nred_i(Nm) Tred_i Wred_i(WR) Cred_i - \mu_i B_i - mow_i(B_i) - graz_i(B_i)$$
(1)

- where μ_i is a constant leaf senescence rate, as assumed in the GRAS model (Siehoff et al.,
- 208 2011). A comprehensive description of DYNAGRAM will be the subject of a next paper.
- Below we provide details on the three functions gr_i , mow_i and $graz_i$, useful to better
- 210 understand the sensitivity analysis we performed. A complete description of all parameters is
- 211 given in Tables A.1 and A.2 (Appendix A).
- We assumed plant growth $gr_i(PAR, B_i)$ to be driven by an Ivlev function describing the
- 213 photosynthesis process, as defined in MODVEGE (Jouven et al., 2006) and in LINGRA
- 214 (Schapendonk et al., 1998). This function defines the maximal growth obtained in optimum
- 215 conditions regarding resources (nutrients, water) and temperature, and in absence of
- 216 disturbances (trampling). It describes the efficiency of use of solar radiation for
- 217 photosynthesis and is modelled as the product of the Photosynthetically Active Radiation
- 218 (PAR), the maximal Radiation Use Efficiency (RUE), which accounts for a decrease in
- efficiency at light intensity higher than 5 MJ m⁻², and an Ivlev function of the Surface Leaf
- Area (SLA_i) and of the plant biomass (B_i) that accounts for the proportion of light intercepted
- by the sward. This optimum plant growth function is given by equation (2).

$$gr_i(PAR(t), B_i(t)) = PAR(t)RUE(t)\left(1 - e^{-\alpha SLA_i \frac{B_i(t)}{10}LAM}\right)10,$$

$$RUE(t) = RUE_{max} \min(1, 1 - \gamma_1(PAR(t) - \gamma_2))$$
(2)

- Four reducers decrease biomass productivity. The first reducer, $Nred_i(Nm)$, is modelled by
- an Holling function depending on the biomass. The second reducer, $Wred_i(WR, PET)$, is a
- function of the water reserve and of the potential evapotranspiration. The third growth
- reducer, $Tred_i(T)$, depends on temperature. The fourth reducer, $Cred_i(SD)$, represents a
- decrease in plant growth due to trampling and is function of the livestock density, SD.
- Defoliation disturbances, i.e. mowing $mow_i(B_i)$ and grazing $graz_i(SD, B_i)$, act as a loss in
- green biomass.
- The first defoliation disturbance consists on moving events with a periodical cut of the
- canopy, following a pre-specified mowing schedule. All B_i variables related to green biomass
- are pulse-wise reduced and a proportion λ_i of all species biomass is removed. We assume this
- proportion to be independent of the biomass value and only depends on the species-specific
- canopy height. For a given mowing event, we get:

$$mow_i(B_i(t)) = \lambda_i B_i(t) \tag{3}$$

- 234 The second defoliation disturbance is due to grazing by livestock, following a predefined
- schedule. At every grazing day, a fixed amount κSD of standing biomass is removed by
- grazing. This loss of biomass is distributed among all species according to their appetence and
- their biomass, following a Holling type III function. During a given grazing period, we get:

$$graz_{i}(SD(t), B_{i}(t)) = \kappa SD(t) \frac{\rho_{i} B_{i}^{ng}(t)}{1 + \sum_{j=1}^{n} \rho_{j} B_{j}^{ng}(t)}$$

$$\tag{4}$$

- Soil processes are summarised by the following ODEs (5) to (7). Again, we only wrote the
- 239 dependence on state variables.

$$\frac{dNo}{dt} = \sum_{i=1}^{n} \delta_i \, \mu_i \, B_i + Nd - \, \theta \, No \, Tmin \, Wmin(WR) \tag{5}$$

$$\begin{split} \frac{dNm}{dt} &= \theta \ \textit{No} \ \textit{Tmin} \ \textit{Wmin}(\textit{WR}) + \textit{Nu} + \Phi - \Lambda(\textit{Nm}) \\ &- \sum_{i=1}^{n} \delta_{i} \ \textit{gr}_{i}(\textit{B}_{i}) \ \textit{Nred}_{i}(\textit{Nm}) \ \textit{Tred}_{i} \ \textit{Wred}_{i}(\textit{WR}) \ \textit{Cred}_{i} \end{split}$$
(6) $\frac{dWR}{dt} = P - AET(B_i, WR) - \Delta(B_i, WR)$ (7)We introduced feedback loops for the two resources, available nitrogen and water reserve in the soil. The loss of green biomass due to senescence constitutes the litter, which fills a pool of organic nitrogen No stored in the soil organic matter. Depending on surface temperature Tmin(T) and soil moisture Wmin(WR), this organic nitrogen is mineralized in mineral nitrogen Nm (nitrate and ammonium) that becomes available for plant uptake. These two reducer functions affect the mineralization process and should not be confused with *Tred* and Wred that reduce plant growth. For each unit of biomass produced, we assumed that a fixed proportion δ_i of mineral nitrogen is consumed (Lazzarotto et al., 2009). Lixiviation $\Lambda(Nm)$ limits the amount of mineral nitrogen in the soil solution. Presence of grazing animals in the pasture brings organic Nd(SD) and mineral Nu(SD) nitrogen restitution proportionally to the stock density SD. The mineral nitrogen pool is also filled by mineral fertilization events, Φ . The water reserve WR is filled by precipitations P and is emptied by the actual evapotranspiration $AET(B_i, WR, PET)$ and by drainage $\Delta(B_i, WR, PET, P)$. AET corresponds to the amount of water lost daily by evapotranspiration, which depends on green biomass, water reserve WR and potential evapotranspiration PET. Drainage corresponds to the loss of water by infiltration once the water reserve in the soil exceeds the water holding capacity WHC. To summarize, if n is the size of the species pool, DYNAGRAM is defined by a system of 3 + n ordinary differential equations associated with 3 + n state variables, with $9 \times n$ speciesspecific parameters, 31 global parameters, and 13 parameters for climatic forcing functions. The three primary C-S-R plant strategies defined by Grime (Grime, 1977; Pierce et al., 2017) are implicitly included in DYNAGRAM. Competitive ability (C strategy) occurs without any stress or disturbance and mainly depends on the potential growth rate of each species and their efficiency to achieve photosynthesis (parameters SLA_i , μ_i). Stress tolerance (S strategy) is represented by several species-specific parameters included in the reducers of this potential vegetative growth regarding three limiting factors (temperature, mineral nitrogen and water) and a trampling effect (parameters $T_{l,i}$, k_i , η_i , σ_i). Resistance to disturbance (R strategy) is represented by species-specific parameters concerning resistance to defoliation (mowing and grazing, which correspond to an explicit loss in green biomass; parameters ρ_i , λ_i). To show the wide diversity of life strategies and plant functional types among the 21 species considered in DYNAGRAM, we plotted their location within the CSR triangle in Fig. 2. The DYNAGRAM model complies with five out of six criteria introduced in a recent review of grassland models (Taubert et al., 2012), by taking into account: (1) species richness, (2)

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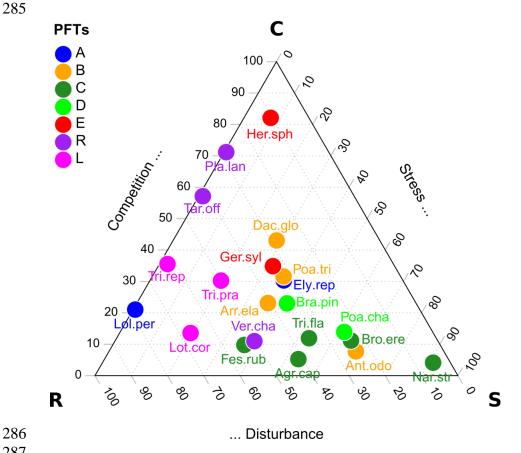
considered in DYNAGRAM, we plotted their location within the CSR triangle in Fig. 2.

The DYNAGRAM model complies with five out of six criteria introduced in a recent review of grassland models (Taubert et al., 2012), by taking into account: (1) species richness, (2) resource limitation (light, nutrients, water), (3) management activities (mowing, grazing, fertilizing), (4) model simplicity, and (5) interspecific competition for resources. To keep the model simple, we did not comply with the sixth criterion concerning the representation of belowground competition processes.

We implemented DYNAGRAM in R language (R Core Team, 2017). Numerical integration of the ODE system was computed with the function 'ode()' of the R package 'deSolve' (Soetaert et al., 2010), using the Euler method and an integration time step of one day to correspond to the length of discrete management events included in the model. This choice of discrete time and difference equations was also made to reduce the simulation time in batch runs, after

having verified that the model outputs were similar to the ones obtained with a smaller fixed or variable time step and a solver considering continuous time, such as 'lsoda', taking into account events (sudden changes in a state variable, e.g. mowing or fertilization).

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Fig. 2 Ternary plot of the CSR triangle indicating the strategy of the 21 species included in the full model (Pierce et al., 2017) and grouped into seven plant functional types (PFTs). Abbreviated names of species and PFTs refer to Table A.2.

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2.2. Choice of the biomass state variables

As our aim was to capture the dynamics of the taxonomic and functional community composition and to assess the importance of community diversity, we considered various combinations of species to represent biomass state variables. In this way, we wrote four alternative versions of DynaGRAM that only differ by the number $n \in \{2, 4, 7, 21\}$ of these state variables. We designed several assemblages containing a number n of species selected as follows among a list of 21 representative herbaceous species (Table A.2 in Appendix A). A first model version (DYNAGRAM2) considers a mixture of two species, namely perennial ryegrass (Lolium perenne), a grass, and white clover (Trifolium repens), a legume. These two species are frequent in intensively managed grasslands and are included in several agronomic grassland models, such as PROGRASS (Lazzarotto et al., 2009), which derives from the pasture simulation model PASIM (Calanca et al., 2007; Graux et al., 2011; Riedo et al., 1998). We selected a set of frequent and potentially dominant individual species, differing by their functional traits and ecological preferences, representative of the species pool in mid-

mountain European pastures and hayfields to be included in a second version of the model

- 307 (DYNAGRAM21) with 21 plant species. As shown by Fig. 2, these species belong to various
- 308 PFTs and present a wide range of CSR strategies.
- Finally, for DYNAGRAM4 and DYNAGRAM7, we kept the two species present in
- 310 DYNAGRAM2 (*Lolium perenne* and *Trifolium repens*), to which we added ten combinations of
- 311 two or five other species, respectively, randomly selected among the 19 remaining species
- included in DYNAGRAM21. Thus, by keeping in all 22 alternative instances of the model (the
- basic DynaGraM2, the full DynaGraM21, and ten replicates of DynaGraM4 and
- 314 DYNAGRAM7) a same grass and a same legume species we ensured a minimum common
- 315 heterogeneity in the functional composition of each instance, and by our random selection of
- 316 other species we avoided any subjective bias.

2.3. Selection of the key control parameters

- When performing a sensitivity analysis, we look at the sensitivity of a model output $\mathbf{Y} = f(\mathbf{X})$
- 319 to variations of the control parameters in a matrix **X**. Each line of the matrix **X** defines a set of
- parameters for which we solve the ODE system. In a first identification analysis, we assessed
- the sensitivity of model outputs (total biomass and species evenness) to a deviation of $\pm 20\%$
- of all parameters, except species-specific parameters (Online Appendix B, Tables B.1, B.2
- 323 and B.3).

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- From the results of this first analysis, we selected the six most influential control parameters
- 325 to be used for a more detailed sensitivity analysis, grouped in three categories: (i) *climate*
- parameters include mean monthly temperature Tm and mean monthly sum of precipitation
- 327 Pm; (ii) soil fertility parameters include the mineralization rate of organic nitrogen θ and the
- 328 amount of mineral nitrogen dropped off each year on the soil as fertilizers Φ; (iii) *defoliation*
- disturbance parameters include the livestock density SD during each grazing period and the
- number of mowing events per year *ncut*. The **X** matrix consisted of all 1728 combinations of
- three or four values for the six control parameters (Table 2).
- We considered a management scenario that is identically repeated each year. Cattle is
- continuously present in the grassland during one long grazing season, from the 1st of July to
- 334 the 31st of August. The first mowing event appears on the 1st of June, and a second mowing
- event on the 2^{nd} of October. The amount of fertilizer, $\Phi = 90 \text{ kg N ha}^{-1}$, is split in six fractions
- spread on the grassland (May 1st, May 15th, June 10th, June 20th, September 1st and September
- 337 15th).

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Table 2 Set of values of the selected control parameters in the matrix **X** of the sensitivity analysis. Columns of **X** correspond to the six key parameters. Lines of **X** are built from all combinations of those values, leading to $3^3 \times 4^3 = 1728$ lines. LU: livestock unit (1 LU is equivalent to a dairy cow).

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Parameter	Category	Values in matrix X Unit				
Tm	Climate	7	9	11		°C
Pm	Climate	50	80	110	140	mm month ⁻¹
θ	Soil fertility	4 10-4	7 10-4	1 10-3		d^{-1}
Φ	Soil fertility	0	20	50	100	kg N ha ⁻¹ a ⁻¹
SD	Defoliation	0	0.5	1	1.5	LU ha ⁻¹
ncut	Defoliation	0	1	2		a^{-1}

2.4. Selection of the response variables

- 345 The next step was thus to extract the response variables from the simulation outputs after
- running each alternative version of the model. Those response variables are either a matrix Y
- 347 (if built from several output variables per simulation) or a vector y (if built from a single
- 348 output variable).

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- Each line of the input matrix **X** corresponds to a set of parameters, for which we run the
- model and generated one response matrix Y and three response vectors y extracted from the
- model output at time $t_f = 3559$, that is the 1st of October at year 10 of the simulation. This day
- is selected just before the second yearly cut and after all other management practices having
- occurred in the year.
- Y is a *n*-columns matrix of biomass values of all species at day t_f . \mathbf{y}_1 is a vector of the total
- biomass values B_{tot} at day t_f , simply computed as the sum of the green biomass values of all
- 356 present species:

$$B_{tot}(t_f) = \sum_{i=1}^{n} B_i(t_f)$$
(8)

- 357 \mathbf{y}_2 is a vector of Simpson evenness values B_{eve} of the plant community at day t_f . Simpson
- evenness is a taxonomic diversity index ranging between 0 and 1 and measuring the
- proportion of dominant species in the community according to differences in biomass (Hill,
- 360 1973; Jost, 2006). A value of 1 means that all species initially introduced are equally
- distributed in biomass in the grassland, whereas a value close to 0 means the persistence of
- only one single species that outcompeted all others. We computed Simpson evenness by
- dividing the inverse Simpson diversity index by n, the number of species initially present in
- 364 the model:

$$B_{eve}(t_f) = \frac{1}{n \sum_{i=1}^{n} \left(\frac{B_i(t_f)}{B_{tot}(t_f)}\right)^2}$$

$$\tag{9}$$

- 365 \mathbf{y}_3 is a vector of the ratio S of the number of surviving species at day t_f to the initial species
- pool n, considering as virtually absent a species with a biomass less than 1 g DM ha⁻¹:

$$S(t_f) = \frac{\#\{B_i(t_f) > 10^{-3}\}}{n} \tag{10}$$

- 367 S represents the final proportion of surviving species, also called 'species survival ratio'
- 368 thereafter.

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369 2.5. Assessment of the model's response to variation in species richness

- 370 To relate the variation of the computed indexes B_{tot} , B_{eve} and S to the size of the initial species
- pool, we performed Kruskal-Wallis tests with post-hoc multiple comparisons on the
- 372 simulation outputs over all combinations of values for the six climate, fertility and defoliation
- parameters. Differences in total final biomass B_{tot} would allow to know if grassland
- productivity is impacted by increasing species richness in the model, whereas differences in
- evenness B_{eve} and survival species S would provide information about the impact on
- 376 community diversity and resilience.

2.6. Tree-based sensitivity analysis

- Regression tree is an iterative process of dividing the data into two partitions, and then
- 379 splitting it up further on each of the branches using discriminant explanatory variables (De'ath
- and Fabricius, 2000). Each split corresponds to a node. A regression tree can be interpreted as

- a decision tree, in which terminal leaves group records with homogenous response variables
- that can be explained by a series of explanatory variables (here model parameters). The
- 383 homogeneity of each node is characterized by its impurity, a measure that takes the value zero
- 384 for completely homogeneous nodes and increases as homogeneity decreases. Then
- maximizing homogeneity in a given group is equivalent to minimizing their impurity. The
- predicted value of the output variable for a particular node is the sample average of the output
- variable consisting of records included in that node. The splitting process continues until each
- 388 node reaches a user-specified minimum node size and becomes a terminal node (Makler-Pick
- 389 et al., 2011).
- 390 Dynamic trees were introduced to provide Bayesian inference for regression trees that change
- with the arrival of new observations (Taddy et al., 2011). Dynamic trees provide an efficient
- measure of a covariate's first-order effect, i.e. variance reduction directly attributed to splits
- on that variable. However, these indices are not appropriate for all applications of sensitivity
- analysis, as non-constant leaf prediction models, such as for linear trees. This issue has been
- 395 corrected by a technique of estimation of sensitivity indices (Gramacy et al., 2013), which is
- model-free and can be constrained to subsets of the input space. Thus, DT computes the Sobol
- index associated to the sensitivity analysis (Ravalico et al., 2005), and graphically estimates
- 398 how each parameter from the set of the matrix **X** impacts the output variable in vector **y**.
- 399 To investigate the influence of the number of state variables describing the plant community
- 400 (aboveground biomass of individual species) on the model sensitivity to key parameters
- 401 (climate, soil fertility, defoliation), we performed a tree-based sensitivity analysis for each
- level of complexity of DYNAGRAM, with 2, 4, 7 and 21 species, based on the matrix **X** and **Y**
- 403 (or vector **y**) described previously.
- 404 Our sensitivity analysis was based on three kinds of trees: (i) Univariate Regression Tree
- 405 (URT); (ii) Multivariate Regression Tree (MRT) and (iii) Dynamic Tree (DT). URT and DT
- 406 consider a vector \mathbf{y} containing one single synthetic variable (B_{tot} , B_{eve} , or S) contrary to MRT
- 407 (De'ath, 2002) that considers a matrix Y including several response variables (biomass of
- 408 each species). URT has been used to rank a set of influential control parameters, here {Tm,
- 409 Pm, θ , Φ , SD, ncut}, from top to bottom, according to their impact on a single output variable
- among the set $\{B_{tot}, B_{eve}, S\}$ for performing global sensitivity analysis (Eynaud et al., 2012;
- 411 Makler-Pick et al., 2011).
- We applied URT and DT to assess the sensitivity of the three response variables (total
- biomass B_{tot} , Simpson evenness B_{eve} and survival rate S at time t_f) to the variation of the six
- 414 control parameters in matrix **X**.
- To assess the sensitivity of species composition to the variation of the same parameters we
- 416 applied MRT to the multivariate response matrix **Y** of all species biomasses, constrained by
- 417 the input parameter matrix **X**. To our knowledge, it is the first application of MRT to model
- 418 sensitivity analysis.
- We pruned each tree to plot the result with six terminal nodes. We determined the most
- 420 influential parameters selected by univariate and multivariate regression trees by retaining the
- three discriminant parameters up to the second splits of each URT or MRT, respectively. We
- 422 compared the average number of discriminant parameters of each category (climate, soil
- fertility or defoliation) among the four species richness levels (2, 4, 7 and 21).
- 424 All analyses were performed in the R environment (R Core Team, 2017) with the packages
- 425 'mvpart' (univariate and multivariate regression trees) and 'dynaTree' (dynamic trees).

3. Results

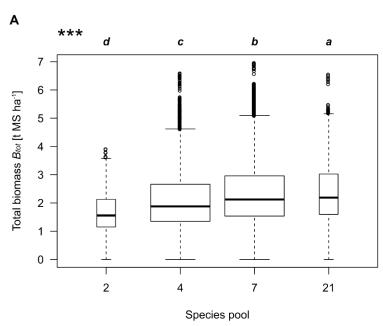
3.1. Model response to variation in initial species richness

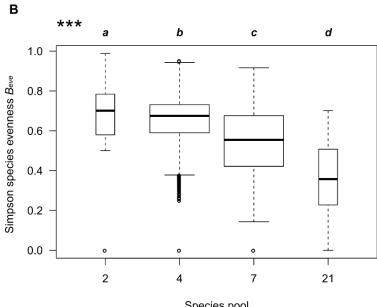
Under various climate, fertility and defoliation conditions given by all combinations of the six control parameters, the final total biomass after ten years of simulation B_{tot} globally increased with the number of species in the grassland (Fig. 3A).

By contrast, Simpson evenness B_{eve} tended to decrease with initial species richness, showing that the more the species pool was large, the less the final green biomass was equally distributed among the species (Fig. 3B).

The proportion of surviving species *S* tended also to decrease with the size of the species pool (Fig. 3C). However, for any initial species richness, we observed that some combinations of parameter values led to cases with null values for each response variable, i.e. with the extinction of all species due to excessive stress or disturbance.

For 21 species in the initial species pool, the survival rate increased with the disturbance intensity, which is the product of defoliation parameters *SD* and *ncut* (Fig. 4A). Moreover, the positive Kendall rank correlation between the disturbance intensity and the species survival rate increased with the size of the species pool (Fig. 4B).







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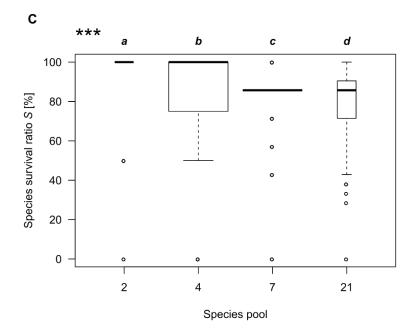


Fig. 3 Boxplots of final total biomass $B_{tot}(\mathbf{A})$, Simpson species evenness $B_{eve}(\mathbf{B})$ and proportion of surviving species $S(\mathbf{C})$ at the end of simulations from alternative DYNAGRAM models with all combinations of control parameters given in Table 2. All results of Kruskal-Wallis tests were highly significant (***: P < 0.001). Different letters indicate significant differences among groups of simulations, based on 1728 simulations for 2 and 21 species, and 10 series of 1728 simulations for different assemblages of 4 and 7 species.

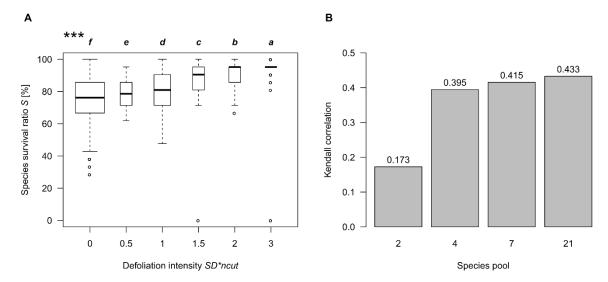
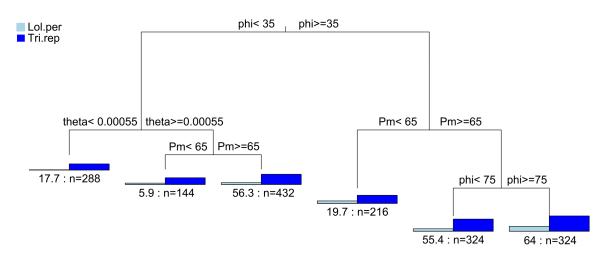


Fig. 4 The relationship between the defoliation intensity and the species survival ratio. **A**: Boxplots of the proportion of surviving species S at the end of simulations from DYNAGRAM21 with all combinations of control parameters, in function of the level of defoliation intensity, with results of Kruskal-Wallis and post-hoc tests (***: P < 0.001); different letters indicate significant differences among levels, based on 1728 simulations for 21 species. **B**: Bar plot of the Kendall rank correlations between the size of the species pool n and the species survival ratio S, based on 1728 simulations for 2 and 21 species, and 10 series of 1728 simulations for different assemblages of 4 and 7 species.

3.2. Sensitivity analysis of DYNAGRAM for 2 and 21 species

To illustrate some details of the sensitivity analysis we performed, let us focus at first on the results of simulations with the extreme values of the species pool, i.e. 2 and 21 species.

Α



Error: 0.374 CV Error: 0.383 SE: 0.0109

B

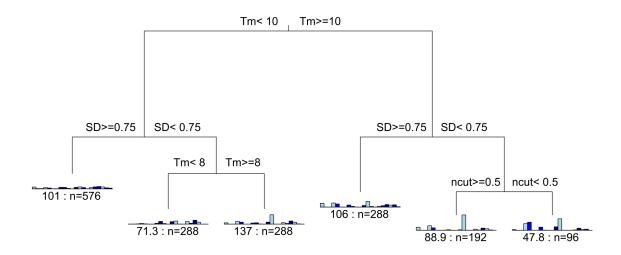


Fig. 5 Multivariate regression trees of a response matrix **Y** containing final green biomass of 2 (**A**) or 21 (**B**) species explained by control parameters out of matrix **X**. Barplots show biomass means for all species, with deviance and number of cases at each terminal node.

Error: 0.557 CV Error: 0.573 SE: 0.0129

3.2.1. Multivariate regression trees

The hierarchy of the most influential parameters that explained differences in final species composition differed completely between DYNAGRAM2 and DYNAGRAM21 (Fig. 5). Variations in quantitative composition of the two-species modelled grassland (Fig. 5A) were mainly driven by fertility parameters (fertilizer input Φ and mineralization rate θ converting organic N to mineral N) and by water supply Pm. By contrast, defoliation parameters (especially stock density SD) and temperature Tm played the major role in explaining the final composition of the species-rich grassland with 21 species (Fig. 5B). These results show that processes driving the assemblage may be strongly related to species richness.

3.2.2. Univariate regression trees

Like species composition, species evenness and survival rate depended mainly on soil fertility $(\Phi \text{ and } \theta)$ for DynaGraM2, but on defoliation (SD) for DynaGraM21 (Table 3).

By contrast, the hierarchy of parameters influencing final total green biomass was independent of the size of the species pool: the first discriminant parameter was in each case fertilisation intensity Φ , followed by mineralization rate θ and mean precipitation Pm. The defoliation regime played no important role in explaining the final standing biomass of a simulated grassland made of 2 or 21 species.

Table 3 Hierarchy of key control parameters resulting from a sensitivity analysis of DYNAGRAM based on MRT and URT on the final biomass of 2 and 21 species. Response variables were individual species biomasses B_i , total biomass (B_{tot}), Simpson evenness (B_{eve}), and survival rate ratio (S) at the end of 10-year 1728 simulations for each species pool. Numbers in brackets are thresholds applied to split the tree at the first (first line, in bold) or the second level (second line). Climate parameters: Tm (temperature), Pm (precipitation); soil fertility parameters: Φ (fertiliser input), θ (mineralization rate); defoliation parameters: SD (livestock density), ncut (mowing frequency).

2 species	21 species
Φ {35}	Tm {10}
$\theta \{5.5 \ 10^{-4}\} $ $Pm \{65\}$	$SD\{0.75\}$ $SD\{0.75\}$
Φ {35}	Φ {35}
$\theta \{5.5 \ 10^{-4}\} $ $Pm \{65\}$	$\theta \{5.5 \ 10^{-4}\} $ $Pm \{65\}$
Φ {75}	SD {0.75}
$\theta \ \{5.5 \ 10^{-4}\} \qquad \theta \ \{5.5 \ 10^{-4}\}$	$SD\{0.25\}$ $Tm\{10\}$
θ {5.5 10-4}	SD {0.75}
Φ {35}	Φ {35} Tm {10}
	$Φ$ {35} $θ$ {5.5 10^{-4} } Pm {65} $Φ$ {35} $θ$ {5.5 10^{-4} } Pm {65} $Φ$ {75} $θ$ {5.5 10^{-4} } $θ$ {5.5 10^{-4} }

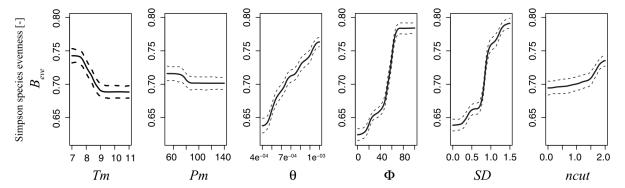
3.2.3. Dynamic trees of species evenness

As the first results of the sensitivity analysis based on composition (MRT) and on Simpson evenness (URT) were very similar, we will focus on this last relevant index of community structure B_{eve} to investigate in more detail the main effects of the six parameters using dynamic trees (Fig. 6). Results for total biomass and survival rate are provided as Supplementary Material (Online Appendix C, Fig. C.1 and C.2).

Dynamic trees confirmed the strong positive effect of livestock density SD on species evenness with 21 species (Fig. 6B), as compared to other parameters. With two species (Fig. 6A), the main positive effects were also due to the amount of fertilizer Φ and the maximal mineralization rate θ , which were much less influential with a large species pool. The number of mowing events *ncut* showed opposite effects on evenness: slightly positive for two species and negative for 21 species. In both cases, mean temperature Tm had a negative effect on evenness.

Despite common trends easily explained by their common structure, the sensitivity analysis revealed strong divergences between the two alternative versions of DYNAGRAM, depending on the size of the species pool, suggesting a shift in key parameters influencing community organization.

Α





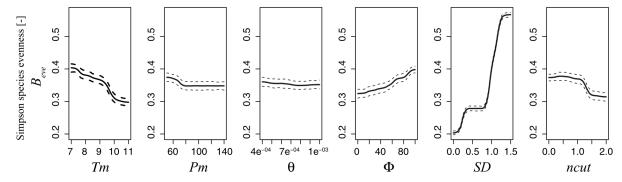


Fig. 6 Main effects of six key parameters on Simpson species evenness B_{eve} , from dynamic trees performed on DYNAGRAM2 (**A**) and DYNAGRAM21 (**B**) outputs. Climate parameters: Tm (temperature), Pm (precipitation); soil fertility parameters: Φ (fertiliser input), θ (mineralization rate); defoliation parameters: SD (livestock density), ncut (mowing frequency).

3.3. Impact of species number on the hierarchy of influential parameters

Tree-based sensitivity analysis of DYNAGRAM2 and DYNAGRAM21 suggested a shift in key parameters determining the grassland composition according to the size of the species pool. To assess the impact of species number, irrespective to species identity, on the hierarchy of the key parameters, we have to include in the comparison the results of regression trees for DYNAGRAM4 and DYNAGRAM7, each with ten random samples among the full set of 21 species. Results of this comparison are summarized by bar plots (Fig. 7).

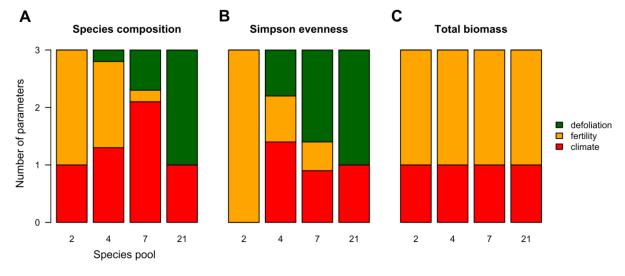


Fig. 7 Bar plots of the three most influential control parameter categories on species composition (**A**), Simpson evenness (**B**) and total green biomass (**C**) at the end of simulations with 2, 4, 7 and 21 species. Three key parameters were extracted from the first and second splits of each MRT (**A**) or URT (**B**, **C**). Results for 4 and 7 species are mean numbers from ten random species samples.

A progressive shift from high importance of soil fertility to high importance of defoliation as the number of species increased was evidenced for both species composition (Fig. 7A) and Simpson evenness B_{eve} (Fig. 7B). A small increase in species number from 2 to 4 induced a small decrease of fertility importance in favour of defoliation and climate. Increasing further species number from 4 to 7 enhanced this trend. Finally, by including all 21 species the

- influence of soil fertility parameters vanished in favour of defoliation parameters. The shift
- 547 we evidenced in relative parameter influence followed a gradient only driven by the size of
- 548 the species pool, whatever the species identity.
- By contrast, results for total biomass B_{tot} (Fig. 7C) confirmed the absence of any trend,
- already observed in Table 3: parameters influencing final total green biomass were totally
- independent of the size of the species pool. B_{tot} always mainly depended on parameters linked
- to resource supply: input of Nm by soil fertilization Φ , maximal mineralization rate θ of No to
- Nm, and water supply by rainfall Pm.

4. Discussion

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4.1. Final model outputs depend on initial species richness

- The positive relationship between the size of the species pool and the final green biomass
- after 10 growing seasons (Fig. 3A) suggests a better overall productivity of species-rich
- modelled ecosystems in favourable conditions, whatever species identity. This result is in
- accordance with our first hypothesis and with the findings of field experiments with artificial
- plant communities (Hector et al., 1999; Reich et al., 2012). An analysis of these biodiversity
- experiments (Gross et al., 2013) suggested that interspecific competition generates
- compensatory dynamics in grasslands that help stabilize community biomass. This stabilizing
- 563 effect of species richness on community biomass arises because increasing species richness
- increases mean biomass by a larger amount than it increases variance, thus increasing
- stability. The contrast with observations in (semi-)natural grasslands, which generally
- conclude to a negative correlation between species richness and productivity, can be
- explained by the same reasons that those given in the case of these biodiversity experiments:
- our simulation experiment considers artificial assemblages of a relatively small number of
- species (21 species is a relatively low species richness as compared to the one of semi-natural
- grasslands), which are not likely to coexist in extent plant communities resulting from abiotic
- and biotic species filtering (Lortie et al., 2004). This contrast between results of simulations or
- experiments and observations highlights the need to consider multi-species sward in grassland
- models, which was identified as one of the fifteen key challenges by a panel of European
- grassland modellers and experts (Kipling et al., 2016b), and a priority for further research.
- 575 The simulated effect of species diversity on forage production confirms that modelling multi-
- species swards, along with grassland quality and the impact of management changes, requires
- 577 further development (Kipling et al., 2016a; van Oijen et al., 2018).
- The negative relationship between initial species richness and final species evenness (Fig. 3B)
- 579 confirms our second hypothesis and the ability of our model to capture competition processes
- that could lead to dominance of some species. DYNAGRAM also simulates species extinction,
- and this diversity loss is an expression of competitive exclusion, with dominant species that
- outcompete some others, depending on environmental conditions.
- Furthermore, we observed a decrease in the proportion of surviving species as the size of the
- species pool increases (Fig. 3C), confirming our third hypothesis. This trend is also in
- accordance with theory and experiments that suggest that increasing species richness should
- destabilize the dynamics of individual populations (Gross et al., 2013; Tilman, 1999). This
- decrease in the species survival ratio means that for high diversity levels not all species
- initially present can survive. However, contrary to Tilman's tradeoff-based niche theory
- rooted on a simple resource-reduction competition model (Tilman, 1980), which states that
- two different species cannot coexist on a single limiting resource, DYNAGRAM is able to
- simulate the persistence of more species than essential resources. In this model we considered
- three essential resources (mineral nitrogen, water and light) but we observed long-term

- 593 persistence of more than three species. Species coexistence could partly be explained by the
- 594 cyclicity of plant growth rate due to seasonal periodic fluctuations of climatic conditions
- implemented in our model, allowing temporal turnover of species with different temperature
- 596 optimum for growth.
- Among the numerous differences with Tilman's model, we also introduced disturbances due
- 598 to grassland management (defoliation by grazing and cutting), which as such could be
- detrimental to some intolerant species and lead to species extinction. However, according to
- our fourth hypothesis, disturbances are supposed to affect more competitive species and thus
- prevent competitive exclusion, allowing more species to coexist in the plant community. This
- is confirmed by our results, as the species survival ratio increased with the defoliation
- intensity (Fig. 4A). We report the results for 21 species, but a similar relationship was
- obtained for any size of the species pool (not shown). The importance of defoliation events,
- mowing or grazing, has also been highlighted for dynamic global vegetation models: the
- 606 ORCHIDEE model better reproduces seasonal variation of LAI after integrating a
- management module for defoliation activities (Chang et al., 2013). It could be of interest to
- extend this result to other disturbances affecting grasslands, such as pest outbreaks and
- diseases, for which a lack of model development has been reported (Snow et al., 2014).
- As expected, the positive correlation between n and S tended to increase with the size of the
- species pool (Fig. 4B), suggesting that disturbances play a more important stabilizing role in
- species-rich communities, in which non-equilibrium conditions brought by disturbances and
- fluctuating environment better favour species coexistence by promoting species turnover and
- preventing competitive exclusion of subordinate species. The relatively low correlation
- obtained with DYNAGRAM2 may also be explained by the identity of the selected species,
- 616 Lolium perenne and Trifolium repens, which, as shown in Table A.2, are both tolerant to
- defoliation and trampling (Kühn et al., 2004).
- These results confirm the need of taking into account the species composition of grassland
- vegetation to accurately estimate the leaf area index (Ben Touhami and Bellocchi, 2015).
- They also add to the growing evidence that increased species richness and functional diversity
- contribute to stabilizing biomass production and enhancing the resilience of grassland
- ecosystems, as shown both theoretically and experimentally (Cottingham et al., 2001; Gross et
- 623 al., 2013).

4.2. A shift in key parameters across a diversity gradient

- Our sensitivity analysis revealed the existence of an unexpected shift in the hierarchy of the
- most influential parameters on final community structure (MRT on B_i: Table 3, Fig. 5 and 7),
- species evenness (URT and dynamic trees on B_{eve} : Table 3, Fig. 6 and 7) and species survival
- ratio (URT on S: Table 3) across a gradient of species richness: as the number of state
- variables describing the plant community increased, the influence of soil fertility parameters
- related to stress tended to decline in favour of defoliation parameters related to disturbance.
- By considering various random combinations of species in addition to the clover-ryegrass
- mixture for DYNAGRAM4 and DYNAGRAM7, we showed that this general trend was
- independent of species identity, contrary to our fifth hypothesis.
- This importance of soil nitrogen availability for a low diverse assemblage (two species) is
- consistent with several recent studies that specifically raised the need to model soil nutrient
- dynamics (Ma et al., 2014; Sándor et al., 2016; Snow et al., 2017). A sensitivity analysis of
- the biogeochemical grassland model COUPMODEL (Senapati et al., 2016) showed that more
- than half of the critical parameters, recognized as most influential on ecosystem processes,
- were related to nutrient dynamics. This importance of soil nitrogen availability for
- DYNAGRAM2 could be explained by the contrasting needs of the two species, expressed by

- strong differences in species-specific parameters δ_i (leaf nitrogen concentration) and k_i (half-
- saturation constant for nitrogen uptake) between the grass and the legume (Table A.2). The
- inclusion in the model of the symbiotic fixation of atmospheric nitrogen N₂ by the legume
- 644 (Lazzarotto et al., 2009) would probably have still accentuated this contrast, hence the model
- sensitivity to resource parameters. Conversely, a species-rich grassland, such as the one
- modelled by DYNAGRAM21, allows functional redundancy among species, making access to
- light a key factor of plant growth expressed by a strong importance of defoliation parameters.
- One further development could consist on taking into account the uncertainty in parameter
- distribution, following a method for sensitivity analysis recently proposed (Paleari and
- 650 Confalonieri, 2016).

4.3. Grassland diversity, productivity and stability

- 652 Contrary to drivers of community structure, parameters influencing final total green biomass
- did not depend on the size of the species pool (Fig. 7C). Thus, the positive effect of species
- richness on productivity cannot be explained by changes in key drivers. For any combination
- of species, biomass production was mainly limited by nutrient and water availability, i.e.
- stress factors, while disturbances always played a minor role.
- Here we assume that the green biomass at the end of the tenth growing season is proportional
- 658 to the aboveground net primary productivity (ANPP) of the final plant community. We base
- 659 this assumption on the observation of time series from various simulations that showed strong
- periodic seasonal similarities after six or seven years of repeated climate and management
- conditions, suggesting that the system has rapidly reached a cyclic equilibrium (results not
- shown). A stability analysis based on phase or bifurcation diagrams should be carried out to
- assess the impact of key forcing parameters, e.g. the fertilisation intensity Φ and the stock
- density SD, on steady states of the system (Louie et al., 2002). As we simulated seasonal
- dynamics, cyclic steady states are expected but this stability analysis is out the scope of the
- present paper.

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- Another possible critical choice is the date at which values of biomass state variables were
- kept to compare simulation outcomes. We chose the 1st of October at year 10 of each
- simulation, just before the second cut and after all other management practices occurred
- eventually in the last year, but other choices could have been possible, such as the day before
- the first management event, i.e. the 1st of May at year 10. This second choice led to similar
- results and to the same conclusions (results not shown).

4.4. Simplicity or complexity?

- Our study highlights consequences of the chosen number of state variables on model's
- behaviour. In the case of our grassland model, we evidenced a shift in the hierarchy of key
- parameters involved in the dynamics of community structure depending on the size of the
- species pool. This should warn the modeller that mathematical grassland models that simplify
- the plant community by considering a small number of species or PFTs are likely to reveal
- 679 impacts of soil fertility, whereas more complex models that include a larger species pool will
- be more sensitive to disturbance processes. Therefore, this result suggests some conditions in
- the choice of the number of species to define the hierarchy of drivers involved in plant
- growth, such as performed in the CoSMo sub-model that ranked disturbances as the most
- important drivers and nitrogen availability as the less important one (Confalonieri, 2014).
- This shift of the most influential parameters follows a kind of continuum (Fig. 7). However,
- whereas the most influential parameters are markedly different between models with 2 and 21
- species, the difference between 7 and 21 species is relatively low. Therefore, the choice of 7

- species or PFTs could be a good compromise to take into account functional plant diversity
- with keeping the number of state variables as lower as possible.
- This issue relates to a more general problem in ecological modelling: how to keep the model
- as simple as possible without missing key variables and processes? DYNAGRAM is simpler
- than many other grassland models but more complex than other ones. As compared to more
- parsimonious mechanistic models based on ODE systems (Louie et al., 2002; Siehoff et al.,
- 693 2011; Tilman, 1985), DYNAGRAM includes a consumption-degradation loop for nitrogen, the
- main limiting soil nutrient, and takes into account the seasonality of plant growth and resource
- competition (Lazzarotto et al., 2009). This added complexity was necessary to assess the
- 696 model's sensitivity to soil fertility, climatic conditions and defoliation.
- As both knowledge of grassland ecosystems and computer performance progress, the
- 698 implementation of more and more details and processes is a general trend in grassland
- modelling. Our choice of a minimal model complexity implies some strong assumptions. As
- 700 compared to more detailed models, such as MODVEGE, GEMINI or PROGRASS (Jouven et al.,
- 701 2006; Lazzarotto et al., 2009; Soussana et al., 2012), DYNAGRAM considers only one single
- biomass compartment for each species without distinguishing aboveground plant structural
- components (i.e. vegetative or reproductive, green or dead) and ignores the belowground
- biomass (roots, rhizomes and storage organs) as well as many associated physiological
- processes. In accordance with most grassland growth models, DYNAGRAM only simulates
- vegetative growth and does not include seed reproduction or emergence of new species from
- 707 the soil seed bank or by seed dispersal.
- Model degradation techniques can be applied to assess consequences on model performance
- 709 (simulation time, prediction accuracy) of reducing the number of variables and parameters in
- a complex model. Applied to grassland models, they often conclude to the necessity of
- keeping the full complexity to ensure an acceptable goodness of fit with empirical data
- 712 (Lohier et al., 2016). In this paper, our goal was more strategical than tactical: we developed
- 713 DYNAGRAM to address a specific theoretical question about the response of a grassland
- model to climatic, edaphic and management forcings in function of the number and identity of
- 715 the state variables describing the plant community. The next step will be to calibrate and to
- validate the model against experimental data so as to use it for the simulation of climatic and
- 717 management scenarios applied to real systems. Its performance in terms of computation time
- as compared to more complex models makes it possible to implement DynaGraM as a
- submodel in a spatially explicit, mosaic landscape model, such as WoodPaM (Gillet, 2008;
- 720 Peringer et al., 2013).
- Alternatives to the development of new models may rely on re-usable components or
- submodels that could be coupled with generic crop simulators to simulate an assemblage of
- several species and reproduce plant community dynamics (Confalonieri, 2014). Another
- 724 promising approach consists in comparing the performance of multiple grassland models so as
- 725 to implement a multi-model ensemble prediction system in grasslands (Palareti et al., 2016;
- 726 Sándor et al., 2017).

5. Conclusion

- 728 Our simulations with DYNAGRAM confirmed most of our expectations but provided new
- 729 insights on the diversity-productivity-stability relationship. After reaching a cyclic
- equilibrium, the final structure of the grassland plant community, described by green biomass
- of a given set of species, depends on the size of species pool. In accordance with the insurance
- hypothesis (Yachi and Loreau, 1999), increasing species richness insures the ecosystem
- against declines in its productivity caused by climatic fluctuations and periodic anthropogenic
- disturbances. However, depending on species identity and environmental conditions, the

- stability of the community, measured by the proportion of surviving species after 10-year
- simulations, and the evenness component of its diversity tend to decrease with the size of the
- species pool, due to environmental filtering and competitive exclusion.
- Moreover, results of our tree-based sensitivity analysis revealed an unexpected shift in the key
- drivers of the final community structure, but not of its total biomass, when increasing the size
- of the species pool from two to 21 species. These findings create new challenges in ecological
- modelling to be raised through mathematical and simulation studies.

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Appendix A: Variables and parameters of DynaGraM

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Table A.1 Summary of all variables and parameters of DYNAGRAM. Values of the species-specific parameters are given in Table A.2.
 1013

Symbol	Definition	Unit	Value
_	State variables		
B_i	Green biomass of species or PFT i	kg DM ha ⁻¹	$B_i(0) = 200/n$
No	Organic nitrogen, stored in the soil organic matter	kg N ha ⁻¹	No(0) = 5800
Nm	Mineral nitrogen (nitrate and ammonium) in the soil solution, available for plant uptake	kg N ha ⁻¹	Nm(0) = 20
WR	Water reserve in the soil, available for plant uptake	mm	WR(0) = WHC
	Climatic forcing variables		
Τ	Surface temperature	°C	function
P	Daily precipitation	mm d ⁻¹	function
PAR	Photosynthetically active radiation	$MJ m^{-2} d^{-1}$	function
PET	Potential evapotranspiration	mm d ⁻¹	function
	Plant species-specific parameters		
n	Number of species or PFTs	[count]	$\{2, 4, 7, 21\}$
SLA_i	Specific leaf area	$m^2 g^{-1}$	species-specific
λ_i	Proportion of biomass removed by a mowing event	d^{-1}	species-specific
ρ_i	Appetence for feeding regime of cattle	[-]	species-specific
σ_i	Utilization indicator value for the trampling tolerance	[-]	species-specific
S_i	Leaf nitrogen content	kg N (kg DM)-1	species-specific
k_i	Half-saturation constant: resource level at which growth rate is half its maximum	kg N ha ⁻¹	species-specific
u_i	Basic senescence rate of leaves	d^{-1}	species-specific
$T_{1, i}$	Lower limit of optimum of daily temperature for photosynthesis	°C	species-specific
η_i	Water stress tolerance exponent	[-]	species-specific
	Plant growth processes and tolerance parameters		
T_0	Base temperature for light utilization and development	°C	3
T_2	Higher limit of optimum of daily temperature for photosynthesis	°C	20
T_3	Maximal temperature for light utilization and development	°C	40
RUE_{max}	Maximum radiation use efficiency	g DM MJ ⁻¹	3
LAM	Proportion of laminae in green biomass	[-]	0.68
α	Extinction coefficient	[-]	0.6
v	Holling coefficient for mineral nitrogen consumption	[-]	2
ng	Holling coefficient for green biomass removal by grazing	[-]	2
T_{tol}	Amplification exponent of the control function $f(X)$ of trampling	[-]	0.5
	Soil parameters		
θ	Nitrogen mineralization rate	d ⁻¹	7 10-4
Vm_{max}	Maximal amount of Nm stored in the soil before lixiviation	kg N ha ⁻¹	120
WHC	Volumetric water content at field capacity	mm	150
PWP	Volumetric soil water content at permanent wilting point	mm	7.5
Tm_1	Critical temperature for maximal mineralization rate of No to Nm	°C	40
Tm_2	Empirical parameters describing effect of temperature on mineralization	°C	31.79
g_T	Empirical parameters describing effect of temperature on mineralization	[-]	3.36

g_{W1}	Empirical parameters describing effect of soil water on mineralization	[-]	1
g_{W2}	Empirical parameter describing effect of soil water on mineralization	[-]	6.63
g_{W3}	Empirical parameter describing effect of soil water on mineralization	[-]	-5.69
PET_{max}	Maximal measured value of PET	mm d ⁻¹	3.75
β_1	Empirical parameter for fitting f_W function of Jouven <i>et al.</i>	[-]	6.467
β_2	Empirical parameter for fitting f_W function of Jouven <i>et al.</i>	[-]	7.623 10-8
	Management parameters and auxiliary variable	les	
SD	Livestock density, number of adult bovine units per hectare	LU ha ⁻¹	$[0, SD_{max}]$
SD_{max}	Maximal possible livestock density	LU ha ⁻¹	3
κ	Daily consumption of 1 adult bovine unit	kg DM LU ⁻¹ d ⁻¹	18
Φ	Yearly input of mineral nitrogen fertilizer in the soil	kg N ha ⁻¹ a ⁻¹	0
Φ_u	Rate of mineral nitrogen restitution by cattle urine deposition	kg N ha ⁻¹ d ⁻¹	0.315
Φ_d	Rate of organic nitrogen restitution by cattle dung deposition	kg N ha ⁻¹ d ⁻¹	0.236
ncut	Number of mowing events per year	[-]	$\{0, 1, 2, 3\}$
	Climatic parameters		
ω	Seasonal oscillation period, conversion of radian in days	d^{-1}	$2\pi/365$
Tm	Mean annual values of temperature T	°C	8.517
Pm	Mean annual values of sum of monthly precipitations P	mm d ⁻¹	2.056
PARm	Mean annual values of PAR	$MJ m^{-2} d^{-1}$	5.434
PETm	Mean annual values of PET	mm d ⁻¹	1.93
Ta	Amplitude of <i>T</i>	°C	14.81
Pa	Amplitude of <i>P</i>	mm d ⁻¹	1.168
PARa	Amplitude of <i>PAR</i>	$MJ m^{-2} d^{-1}$	7.344
PETa	Amplitude of <i>PET</i>	mm d ⁻¹	3.542
Ψ_T	Origin phase of oscillations for T	d	106.5
Ψ_P	Origin phase of oscillations for <i>P</i>	d	110.5
Ψ_{PAR}	Origin phase of oscillations for <i>PAR</i>	d	80.8

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Table A.2 List of the 21 herbaceous species to be selected in the alternative versions of DYNAGRAM, with their plant functional type (PFT), CSR strategy (Pierce et al., 2017) and species-specific parameter values: SLA specific leaf area (Perronne et al., 2014); λ proportion of biomass removed by a mowing event, estimated from canopy height (Perronne et al., 2014); ρ appetence for cattle (Kühn et al., 2004); σ trampling tolerance (Kühn et al., 2004); δ leaf nitrogen content (Soussana et al., 2012); k half-saturation constant of the nitrogen resource, estimated from N requirement (Julve, 2017); µ leaf senescence rate (Ryser and Urbas, 2000); T_1 lower limit of optimal growth temperature (Al Haj Khaled et al., 2005); η water requirement (Julve, 2017). In bold, the two species considered for DYNAGRAM2 and kept for defining the assemblages of DYNAGRAM4 and DYNAGRAM7. PFTs are defined by four groups of grasses and three life forms of dicots (Ansquer et al., 2004; Cruz et al., 2002): A, early and fast-growing grasses in fertile and frequently disturbed grasslands; B, competitive and productive grasses in fertile and unfrequently disturbed grasslands; C, late and slow-growing grasses in infertile and frequently disturbed grasslands; D, late and slowgrowing grasses in infertile and unfrequently disturbed grasslands; E, erect forbs; R, rosette forbs; L, legumes.

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Symbol	Name	PFT	CSR	SLA	λ	ρ	σ	δ	k	μ	T_1	η
Lol.per	Lolium perenne	A	R/CR	0.0286	0.40	9	8	0.0182	13.00	0.0303	7.5	1.000
Ely.rep	Elymus repens	A	CSR	0.0233	0.70	6	7	0.0106	12.00	0.0178	9.5	1.000
Ant.odo	Anthoxanthum odoratum	В	S/SR	0.0283	0.70	4	5	0.0173	12.00	0.0192	11.0	1.000
Arr.ela	Arrhenatherum elatius	В	SR/CSR	0.0299	0.80	8	3	0.0238	11.00	0.0250	8.5	1.000
Dac.glo	Dactylis glomerata	В	CSR	0.0253	0.50	8	6	0.0094	11.00	0.0230	10.0	1.000
Poa.tri	Poa trivialis	В	CSR	0.0331	0.50	7	6	0.0224	11.00	0.0250	8.0	2.000
Agr.cap	Agrostis capillaris	C	SR	0.0286	0.80	5	5	0.0205	7.50	0.0205	16.0	1.000
Bro.ere	Bromopsis erecta	C	S/SR	0.0155	0.90	12	1	0.0105	5.15	0.0160	15.3	0.500
Fes.rub	Festuca rubra	C	SR	0.0164	0.90	5	6	0.0137	7.00	0.0138	13.0	1.000
Nar.str	Nardus stricta	C	S	0.0110	0.58	9	2	0.0175	4.85	0.0115	15.0	1.000
Tri.fla	Trisetum flavescens	C	SR/CSR	0.0205	0.91	5	4	0.0130	6.00	0.0212	14.0	1.000
Bra.pin	Brachypodium pinnatum	D	SR/CSR	0.0234	0.92	12	1	0.0202	6.25	0.0167	16.5	1.000
Poa.cha	Poa chaixii	D	S/CSR	0.0207	0.40	3	4	0.0175	10.00	0.0250	8.0	1.000
Ger.syl	Geranium sylvaticum	E	CSR	0.0315	0.82	8	3	0.0175	10.50	0.0220	8.5	2.000
Her.sph	Heracleum sphondylium	E	C	0.0230	0.75	6	3	0.0175	11.00	0.0176	9.0	1.000
Pla.lan	Plantago lanceolata	R	C/CR	0.0196	0.69	3	9	0.0191	9.00	0.0159	12.0	1.000
Tar.off	Taraxacum officinale	R	CR	0.0324	0.92	7	8	0.0175	9.00	0.0224	12.5	1.000
Ver.cha	Veronica chamaedrys	R	SR/CSR	0.0277	0.80	3	7	0.0196	9.00	0.0220	13.0	1.000
Lot.cor	Lotus corniculatus	L	R/SR	0.0224	0.95	9	1	0.0175	7.75	0.0173	11.5	0.707
Tri.pra	Trifolium pratense	L	R/CSR	0.0228	0.82	8	3	0.0196	8.00	0.0175	11.5	1.000
Tri.rep	Trifolium repens	L	R/CR	0.0343	0.90	9	8	0.0195	8.50	0.0234	13.0	1.000

1035	Online Supplementary Material
1036 1037	The following supplementary material is available for this article.
1038 1039 1040	Appendix B Preliminary identification stage of the sensitivity analysis of DYNAGRAM. This appendix details all results of the preliminary sensitivity analysis we performed to select the six key control parameters. Results stay in the three following tables:
1041 1042 1043	Table B.1 Parameter sensitivity of total green biomass B_{tot} to deviations by \pm 20% from the default parameter values for DYNAGRAM2 and DYNAGRAM21. Changes were applied independently for each parameter, all other parameters keeping their default values.
1044 1045 1046	Table B.2 Parameter sensitivity of species evenness B_{eve} to deviations by \pm 20% from the default parameter values for DYNAGRAM2 and DYNAGRAM21. Changes were applied independently for each parameter, all other parameters keeping their default values.
1047 1048 1049 1050 1051	Table B.3 Ranking of parameters according to the sensitivity of total green biomass B_{tot} and of species evenness B_{eve} to variations in the default parameters for DYNAGRAM2 and DYNAGRAM21. Values were computed by both increasing and decreasing the default value of each parameter. Parameters were ranked according to the mean values of the 2 and 21 species.
1052 1053 1054	Appendix C Main effects of key control parameters on biomass and species survival ratio. This appendix contains two figures that detail the main effects of the six parameters on total green biomass B_{tot} and species survival ratio S as revealed by dynamic trees.
1055 1056 1057 1058 1059	Figure C.1 Main effects of six key parameters on total green biomass B_{tot} , from dynamic trees performed on DYNAGRAM2 (A) and DYNAGRAM21 (B) outputs. Climate parameters: Tm (temperature), Pm (precipitation); soil fertility parameters: Φ (fertiliser input), θ (mineralization rate); defoliation parameters: SD (livestock density), $ncut$ (mowing frequency).
1060 1061	Figure C.2 Main effects of six key parameters on species survival ratio <i>S</i> , from dynamic trees performed on DynaGraM2 (A) and DynaGraM21 (B) outputs. Climate parameters: <i>Tm</i>

(temperature), Pm (precipitation); soil fertility parameters: Φ (fertiliser input), θ

(mineralization rate); defoliation parameters: SD (livestock density), ncut (mowing

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frequency).