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NEMA, a functional–structural model of nitrogen economy within wheat culms after flowering. I. Model description

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- **Background and Aims** Models simulating nitrogen use by plants are potentially efficient tools to optimize the use of fertilizers in agriculture. Most crop models assume that a target nitrogen concentration can be defined for plant tissues and formalize a demand for nitrogen, depending on the difference between the target and actual nitrogen concentrations. However, the teleonomic nature of the approach has been criticized. This paper proposes a mechanistic model of nitrogen economy, NEMA (Nitrogen Economy Model within plant Architecture), which links nitrogen fluxes to nitrogen concentration and physiological processes.
- **Methods** A functional–structural approach is used: plant aerial parts are described in a botanically realistic way and physiological processes are expressed at the scale of each aerial organ or root compartment as a function of local conditions (light and resources).
- **Key Results** NEMA was developed for winter wheat (*Triticum aestivum*) after flowering. The model simulates the nitrogen (N) content of each photosynthetic organ as regulated by Rubisco turnover, which depends on intercepted light and a mobile N pool shared by all organs. This pool is enriched by N acquisition from the soil and N release from vegetative organs, and is depleted by grain uptake and protein synthesis in vegetative organs; NEMA accounts for the negative feedback from circulating N on N acquisition from the soil, which is supposed to follow the activities of nitrate transport systems. Organ N content and intercepted light determine dry matter production via photosynthesis, which is distributed between organs according to a demand-driven approach.
- **Conclusions** NEMA integrates the main feedbacks known to regulate plant N economy. Other novel features are the simulation of N for all photosynthetic tissues and the use of an explicit description of the plant that allows how the local environment of tissues regulates their N content to be taken into account. We believe this represents an appropriate frame for modelling nitrogen in functional–structural plant models. A companion paper will present model evaluation and analysis.

Key words: Rubisco turnover, remobilization, functional–structural plant model, nitrogen, light acclimation, senescence, wheat, *Triticum aestivum*, root uptake, common pool.

INTRODUCTION

Nitrogen (N) availability regulates numerous aspects of plant growth. Nitrogen is involved in the expansion of organs that capture resources (meristematic activity and cell extension), as well as in their photosynthetic activity. Up to 75% of reduced N in cereal leaves is involved in photosynthetic processes, mainly as Rubisco, and the rate of light-saturated photosynthesis is linearly correlated with specific N mass (Evans, 1989a). Nitrogen is also involved in the quality of harvested organs since it is a component of proteins (Ourry *et al.*, 2001). In monocarpic species such as wheat (*Triticum aestivum*), N is remobilized from vegetative organs to fill growing grains after flowering, leading to the progressive death of the vegetative tissues.

The increasing use of N fertilizers played a considerable role in increasing yield and production quality in the last century (Tilman, 1999; Cassman *et al.*, 2003; Hirel *et al.*, 2007), but resulted in unsustainable environmental and economic costs (Wallsgrove *et al.*, 1983; Tilman, 1999; Cassman *et al.*,

2003). Reducing N inputs while maintaining high yields of high quality requires novel practices, in which N acquisition and use by plants is maximized and N losses are minimized. Optimum N management practices should closely respect plant requirements and should be combined with breeding for specific traits.

Numerical models simulating crop growth and yield from environmental variables are potentially efficient tools for designing improved agricultural practices and identifying useful traits for cultivars. Models offer the possibility to predict plant responses to changes in management practices, cultivar traits and climate. In recent decades, many crop models have been developed (for a review, see Jeuffroy *et al.*, 2002) including the model of Sinclair and Amir (1992), STICS (Brisson *et al.*, 1998), CERES (Jones and Kiniry, 1986; Gabrielle *et al.*, 1998a, b), Sirius (Jamieson and Semenov, 2000; Martre *et al.*, 2006) and ORYZA2000 (Bouman and Laar, 2006). In this type of model, a few compartments are accounted for (typically roots, reproductive organs and vegetative organs). These models capture the

main processes governing N economy of crops and are able to simulate mean plant behaviour observed under intensive conditions, for which they have been developed; however, even if several crop models have a solid mechanistic basis, they also include teleonomic components and/or equations that approximate observed crop behaviour rather than expressing the physiological processes from which this behaviour emerges. This raises difficulties when the model is used as a tool for understanding how N economy is regulated at the plant scale and when addressing current needs to simulate how novel management practices and cultivars can impact N economy.

It has been proposed that the predictive ability of these models is mainly hampered by the use of the concept of plant demand for resources in teleonomic approaches (Thornley, 1998b). Before flowering, N content of the vegetative compartment is calculated from a demand in N, that would result from the difference between a target N concentration and the real one. Defining such demand is difficult, because it is not linked to a physiological process. Consequently, the quantification of N deficiency and its effect on leaf area expansion and photosynthesis by comparing N demand and supply also raise difficulties. The calculation of N remobilization after flowering also uses a target-driven approach: most models use the approximation that all N present in the vegetative compartments at flowering will have migrated to the reproductive compartment at maturity (Sirius, ORYZA2000 and CERES). This assumption cannot account for the stay-green behaviour of some cultivars, which is characterized by a significant percentage of vegetative tissues that are still photosynthetic at maturity (Borrell *et al.*, 2001) and is often associated with high yields (Benbella and Paulsen, 1998; Borrell *et al.*, 2000; Hafsi *et al.*, 2000). Finally, N mass in vegetative organs is often considered to decrease at a constant rate during grain filling even though it has been observed first to remain stable – or even show a transient increase – in the case when N fertilization is carried out at flowering (for wheat, see Triboï and Triboï-Blondel, 2002).

We hypothesize that to use models as comprehensive tools, to make them more flexible and to enable them to elucidate what could happen under novel conditions, plant functioning should not be formalized by demand-driven rules defined at the whole plant or crop scale, but rather be simulated as the result of mechanisms described at a finer scale. Such mechanisms will then be more faithful to physiological processes, which increases the realism of simulation and paves the way for adapting the model according to ongoing progress in our knowledge of physiology. The functional–structural approach is one possible way to develop more mechanistic models (Prusinkiewicz, 2004; Godin and Sinoquet, 2005; Dingkuhn *et al.*, 2006; Luquet *et al.*, 2006; Bertheloot *et al.*, 2008a; Fourcaud *et al.*, 2008; de Reffye *et al.*, 2008). It explicitly represents plant botanical structure, so that the main functional processes can be formalized at the organ or tissue scale while accounting for the effect of the local environment (Chelle, 2005).

Until now, functional–structural models mainly focused on carbon (C) acquisition and distribution within the aerial parts of plants (Fournier and Andrieu, 1998, 1999; Allen *et al.*, 2005; Luquet *et al.*, 2006; de Reffye *et al.*, 2008). The first

model accounting for N fluxes was GRAAL-CN (Drouet and Pagès, 2007), which was developed for maize plants (*Zea mays*) before flowering. GRAAL-CN simulates the distribution of N and C mass within the plant which is represented as an assembly of connected modules, each characterized by a botanical type (i.e. lamina, sheath, internode and root segment) and a position within the plant. However, C and N fluxes are modelled following a demand-driven approach, which raises the difficulty of quantifying this demand, as in crop models. Moreover, remobilization rates are independent of the environment and of the plant N status. A more mechanistic approach is used for N acquisition from the soil, based on the formalization of the activities of two known major transport systems for nitrate, HATS (high affinity transport system), which functions at a low soil nitrate concentration, and LATS (low affinity transport system), which functions at a high soil nitrate concentration (for reviews, see Glass and Siddiqi, 1995; Daniel-Vedele *et al.*, 1998). GRAAL-CN accounts for an up-regulation of these transport systems by carbohydrate availability, and a down-regulation by plant N content. This is in accordance with physiological and molecular studies (Delhon *et al.*, 1996; Forde and Clarkson, 1999; Touraine *et al.*, 2001; Gojon *et al.*, 2009; Masclaux-Daubresse *et al.*, 2010) that demonstrated that the expression and/or activity of root N uptake systems is down-regulated by reduced N metabolites and up-regulated by sugar signals; besides, sugar is a source of energy required for mineral N assimilation.

We previously proposed a mechanistic alternative to the demand-driven approach for modelling the N fluxes between leaf laminae of wheat after flowering (Bertheloot *et al.*, 2008a). The N content of each lamina is regulated by the turnover of Rubisco, whose synthesis depends on light intercepted by the lamina and the availability of mobile N (representing mainly amino acids and nitrate). We demonstrated that N contents could be accurately simulated by assuming all organs share a common pool of mobile N, but differ in their light environment. The role of light intercepted by a leaf is to increase the rate of protein synthesis. The mechanism behind is the effect of light on the transpiration flux that carries both mobile N and cytokinin towards the leaf (Pons and Bergkotte, 1996; Pons *et al.*, 2001); cytokinin is a hormone involved in the synthesis of photosynthetic proteins (Kusnetsov *et al.*, 1994; Aloni *et al.*, 2005). In addition, light provides the sugar required for nitrate assimilation to leaf tissues. Phloem transport was left aside, but this simplification did not hamper calculations, probably because at the stage of development concerned, phloem transport is mostly directed toward grains. Still, the model lacked a predictive capacity because it focused on a system consisting of laminae and grains, the N mass of other organs being forced to follow experimental data.

This paper proposes a comprehensive process-based model of N economy for wheat plants after flowering, as a step forward in the development of a model spanning the full growth cycle. During the post-flowering period, N economy is driven by N acquisition from the soil and by N remobilization from vegetative tissues for grain filling, because the vegetative structure (roots and aerial parts) has mainly completed its development. The objective of the study is to propose an alternative to the demand-driven approach to model, from N

available in the soil and incident light, the N fluxes within the aerial structure of plants and their impact on dry matter acquisition and allocation to grains. The model, called NEMA (Nitrogen Economy Model within plant Architecture), extends the turnover model proposed earlier (Bertheloot *et al.*, 2008a) to all photosynthetic organs (i.e. laminae, sheaths, internodes and chaff) and links it to a model of N acquisition from the soil based on the formalization of transport systems. The present paper describes the model and discusses the reasons and expected consequences of the modelling choices. In a companion paper (Bertheloot *et al.*, 2011), we describe parameter fitting, model evaluation and a study of model behaviour involving a sensitivity analysis that aims to understand how N acquisition from the soil and the effect of N on total dry mass and N content of grains are regulated at the whole plant scale.

MODEL DESCRIPTION

After flowering, each culm has its own root system, and translocation of assimilates between the culms in the same plant is very low (Williams, 1964); consequently, in NEMA, the crop is seen as a population of individual culms of density Dens_c (culms m^{-2}). Processes are described at the scale of one culm and, for the sake of simplicity, all culms are considered to be identical. This work focuses on N distribution within aerial parts, while roots are represented by a single compartment. NEMA predicts with a daily time step, for a culm free of disease or water stress, the acquisition of N and of total dry mass as well as their distribution between leaf laminae, sheaths, internodes, peduncle, chaff, ear grains and roots. The distribution of photosynthetic area between photosynthetic modules is also predicted. The distribution of total dry mass is modelled according to GreenLab formalism, i.e. following a demand-driven approach (Kang *et al.*, 2008). Obviously N is part of the total dry mass but represents a small fraction (0.5–2 %); so, for the sake of simplicity, we keep the calculations for total dry mass distribution independent of those of N. On the other hand, the calculation of total dry mass distribution contributes to the calculation for the fluxes of N, which is partly transported with carbohydrates. The inputs are: (a) a complete plant description at flowering (i.e. for each organ: N mass, total dry mass, photosynthetic and total areas); (b) soil N concentration at flowering; and (c) the daily time course of photosynthetically active radiation (PAR) above the canopy. Culm characteristics and density enable calculation of the tissue area density and thus the vertical light profile. Soil N only changes as a function of N acquisition by roots to keep the calculation simple and to focus on the modelling of N in aerial parts. The model consists of sub-modules formalizing N and total dry mass acquisition, storage and translocation within the plant, as well as tissue death following N remobilization. Calculations are detailed below, and the general principle of the model is presented in Fig. 1.

Plant structure

Roots are represented by a single compartment, while the botanical and functional organization of the aerial part of the

culm is explicitly represented (Fig. 2). The aerial part of the culm is an assembly of phytomers, each consisting of three types of entities: from the bottom to the top, internode (in), sheath (sh) and lamina (la); the upper vegetative internode is surmounted by the peduncle (ped), which bears the ear consisting of chaff (pooled in a single compartment denoted chaff) and grains (pooled in a single compartment denoted grain). All the entities except grains are exposed to light and are photosynthetic. They are defined by their type (denoted tp), and vegetative entities are numbered acropetally according to the phytomer they belong to (denoted i).

Each photosynthetic entity is characterized by its length ($L_{tp,i}^{\text{tot}}$; m) and area ($A_{tp,i}^{\text{tot}}$; m^2). The exposed length ($L_{tp,i}^{\text{exp}}$; m) and exposed area ($A_{tp,i}^{\text{exp}}$; m^2) of sheaths and internodes are also specified, since part of these entities is hidden by the sheath of the phytomer below. Laminae are approximated as planar shapes, so that their orientation can be characterized by the two angles defining the unit vector normal to the plane: (1) the azimuth angle, i.e. the angle between the projection of the normal vector on a horizontal plane and the north; and (2) the zenith angle, i.e. the angle between the normal vector and the vertical (θ_{la} ; radians). Calculation of light intercepted by the lamina of a given rank was done using the Beer–Lambert approximation: light corresponds to the mean expected value for that rank, assuming that the zenith angle was identical for all plants and all phytomer ranks and that the azimuth angle followed a random uniform distribution. All structural characteristics were assumed to remain constant over time; only the photosynthetic active area of each entity ($A_{tp,i}^{\text{green}}$; m^2) decreased following N remobilization. Structural characteristics and the PAR above the canopy (PAR_0 ; $\text{J m}^{-2} \text{d}^{-1}$) were used to calculate the PAR incident on each entity ($\text{PAR}_{tp,i}$; $\text{J m}^{-2} \text{d}^{-1}$; calculation detailed below). Following our previous observations (Bertheloot *et al.*, 2008b), both senescent and photosynthetic tissues were taken into account to calculate PAR extinction and were assumed to have an identical effect on PAR extinction. Consequently, $\text{PAR}_{tp,i}$ was only affected by variations in PAR_0 between days.

Grains were characterized by their total dry mass ($M_{\text{grain}}^{\text{tot}}$; g) and their N mass ($N_{\text{grain}}^{\text{tot}}$; g). For photosynthetic entities and roots, two forms of dry matter and N were distinguished: structural N and dry mass ($N_{tp,i}^{\text{struct}}$, $M_{tp,i}^{\text{struct}}$ for an entity of type tp and rank i ; N_r^{struct} , M_r^{struct} for roots; g), which were assumed to be constant for the period after flowering, and remobilizable N and dry mass ($N_{tp,i}^{\text{rem}}$, $M_{tp,i}^{\text{rem}}$ for an entity of type tp and rank i ; N_r^{rem} , M_r^{rem} for roots; g), that can be synthesized or degraded. Remobilizable N mass of photosynthetic entities was assumed to consist only of N associated with the photosynthetic apparatus.

All modules are connected to a common pool of mobile N (N_c^{mob} ; g), which is the N form involved in the transport, i.e. mainly amino acids and nitrate. Total N mass of a module was calculated as the sum of its structural, remobilizable and mobile N masses. The mass of mobile N within each plant module was calculated from the mass of mobile N in the whole culm (N_c^{mob}) by assuming that the distribution of mobile N paralleled that of the remobilizable N mass, as in Bertheloot *et al.* (2008a). Total dry masses of photosynthetic entities ($M_{tp,i}^{\text{tot}}$; g) or roots (M_r^{tot} ; g) are sums of their structural and their remobilizable dry masses.

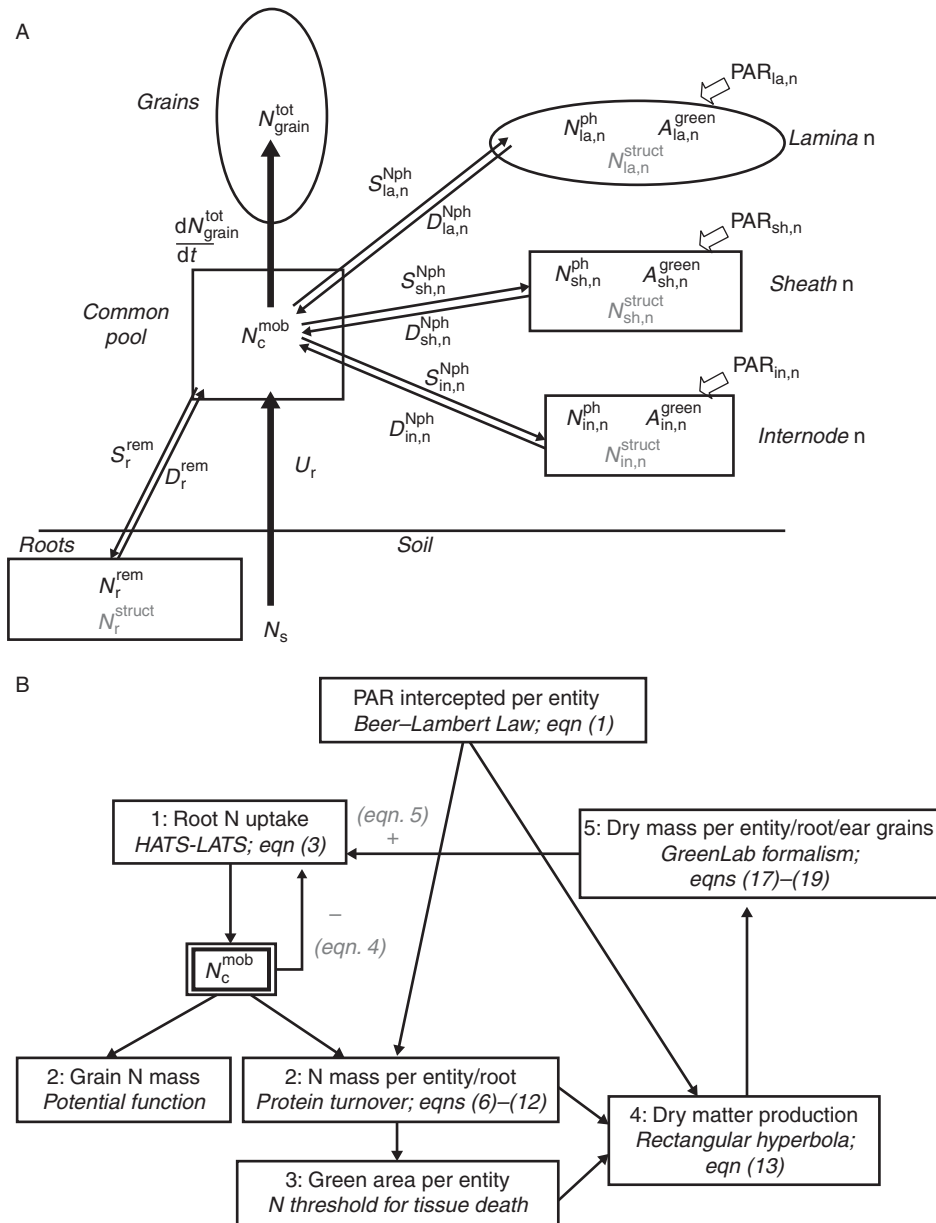


FIG. 1. Overview of the model of N economy within a wheat (*Triticum aestivum*) culm after flowering. (A) Structure and functioning of N acquisition and distribution within a culm. Each lamina (la), sheath (sh) and internode (in), all chaff, all grains (grain), all roots (r) and the peduncle are represented by different modules, but only the uppermost phytomer (n), grains and roots are shown on the figure. Symbols in black represent the N-related model variables: PAR intercepted by each entity, photosynthetic N mass (N^{ph}) and area (A^{green}), remobilizable N mass in roots ($N_{\text{r}}^{\text{rem}}$) and total N mass in grains ($N_{\text{grain}}^{\text{tot}}$); N^{struct} , the structural N mass, shown in grey, is constant throughout the grain-filling period. A supplementary module describes the mobile N pool ($N_{\text{c}}^{\text{mob}}$), which is enriched by root N uptake and assimilation (U_{r}) and N^{ph} , $N_{\text{r}}^{\text{rem}}$ degradation ($D_{\text{Nph}}^{\text{Nph}}$, $D_{\text{r}}^{\text{rem}}$), depleted by grain N filling ($dN_{\text{grain}}^{\text{tot}}/dt$) and N^{ph} , $N_{\text{r}}^{\text{rem}}$ synthesis ($S_{\text{Nph}}^{\text{Nph}}$, $S_{\text{r}}^{\text{rem}}$). N fluxes between modules are represented by arrows. (B) Calculations within one model time step. Each box represents one meta-process, with the calculation principle indicated as well as the numbers of the equations; the number in the box corresponds to the calculation step; the dependence between one meta-process and the others is indicated by arrows. In addition, the common pool of $N_{\text{c}}^{\text{mob}}$ is explicitly represented because of its central role in the regulation of N fluxes. The PAR intercepted by each entity is calculated once at flowering.

The effect of temperature on the rate of processes leading to the distributions of N and dry mass is accounted for by expressing time as thermal time with a base temperature of 0°C , as widely accepted for wheat. The number of degree days ($^{\circ}\text{Cd}$) within day t is denoted $\Delta(t)$. Variables are listed in Table 1 and parameters are listed in Table 2.

Photosynthetically active radiation intercepted by a photosynthetic entity

The PAR intercepted by an entity was estimated by multiplying the area of the entity by the PAR intercepted per unit area, the latter being calculated for the position corresponding

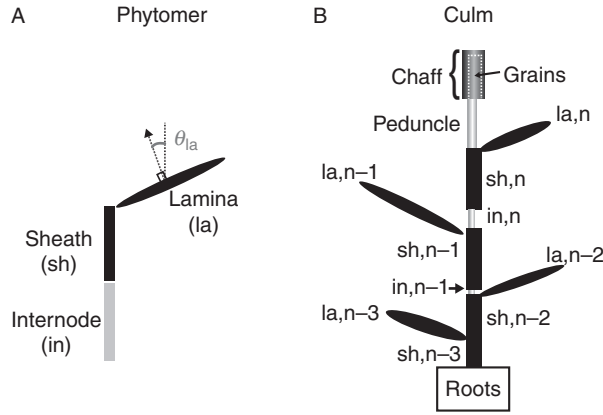


FIG. 2. Botanical structures of a phytomer (A) and a culm (B) in wheat after flowering as implemented in the model. Laminae are defined by planar shapes characterized by their orientation, θ_{la} ; sheaths and internodes are represented by vertical cylinders. Parts of internodes are surrounded by sheaths, and grains are surrounded by chaff; enclosed parts are represented by dotted lines. Phytomers are numbered according to their rank from the top of the culm, n being the uppermost phytomer.

to the middle height of the entity. For entities which are partly hidden, such as sheaths, this calculation was done for the exposed part of the entity. PAR intercepted by an entity was estimated from PAR above the canopy using Beer–Lambert’s law (Monsi and Saeki, 2005), as:

$$\text{PAR}_{tp,i}(t) = k_{tp} \times \text{PAR}_0(t) \times \exp(-k_{\text{canopy}} \times \text{AI}_{tp,i}) \quad (1)$$

where k_{tp} ($\text{m}^2 \text{m}^{-2}$) is the PAR extinction coefficient associated with entities of type tp , k_{canopy} is the mean extinction coefficient of the canopy, and $\text{AI}_{tp,i}$ ($\text{m}^2 \text{m}^{-2}$) is the cumulated area index above the middle height of the entity exposed part.

The effect of entity orientation on PAR interception was accounted for by varying the extinction coefficient from 1 for horizontal entities to k_{vertical} for vertical entities. The lamina extinction coefficient k_{la} was calculated with the following function of lamina orientation:

$$k_{la} = k_{\text{vertical}} + (1 - k_{\text{vertical}}) \times \cos(\theta_{la}) \quad (2)$$

k_{canopy} was calculated as the mean of the extinction coefficients of all entities weighted by their exposed areas.

$\text{AI}_{tp,i}$ was calculated as the product between culm density, Dens_c , and the exposed areas above the middle of the height of the entity tp,i exposed part, whose distance from the top of the canopy was calculated from the plant structure described in Fig. 2. Following the usual calculation of PAR by Beer–Lambert’s law, only half the exposed areas of the cylinders representing sheaths and internodes were considered.

Nitrogen uptake by roots and assimilation

The calculation of N acquisition by the culm is based on that of GRAAL-CN. Processes of N uptake by transport systems and of N assimilation into reduced N (that enriches the common pool) are not distinguished in that formalization. Nitrogen acquisition within a time step is modelled as the

TABLE 1. Model variables: their symbols, definitions and units.

Symbol	Definition	Unit
$\Delta(t)$	Thermal time within 1 d	$^{\circ}\text{Cd}$
t	Time	days (d)
tp,i	Indices for an entity of type tp and rank i	
PAR interception		
$A_{tp,i}^{\text{exp}}$	Exposed area of entity tp,i	m^2
$A_{tp,i}^{\text{tot}}$	Total area of entity tp,i	m^2
$\text{AI}_{tp,i}$	Cumulative area index above entity tp,i	$\text{m}^2 \text{m}^{-2}$
Dens_c	Density of culms in the field	Culm m^{-2}
k_{canopy}	Canopy extinction coefficient	$\text{m}^2 \text{m}^{-2}$
k_{tp}	Extinction coefficient of entity tp	$\text{m}^2 \text{m}^{-2}$
$L_{tp,i}^{\text{exp}}$	Exposed length of entity tp,i	m
$L_{tp,i}^{\text{tot}}$	Total length of entity tp,i	m
PAR_0	Photosynthetic active radiation above the canopy	$\text{J m}^{-2} \text{d}^{-1}$
$\text{PAR}_{tp,i}$	Photosynthetic active radiation incident on entity tp,i	$\text{J m}^{-2} \text{d}^{-1}$
N mass, dry mass and photosynthetic active area		
$A_{tp,i}^{\text{green}}$	Photosynthetic area of entity tp,i	m^2
N_c^{mob}	Mobile N mass in the common pool	g
$[N]_c^{\text{mob}}$	Mobile N mass concentration	g g^{-1}
$N_{tp,i}^{\text{ph}}$	Photosynthetic N mass of entity tp,i	g
N_r^{rem}	Root remobilizable N mass	g
N_r^{struct}	Structural N mass of roots, entity tp,i	g
N_r^{tot}	Total N mass of grains, roots, entity tp,i	g
$M_{tp,i}^{\text{green}}$	Photosynthetic tissues dry mass of entity tp,i	g
$M_{tp,i}^{\text{rem}}$	Remobilizable dry mass of roots, entity tp,i	g
$M_{tp,i}^{\text{struct}}$	Structural dry mass of roots, entity tp,i	g
$M_{tp,i}^{\text{tot}}$	Total dry mass of grains, roots, entity tp,i	g
N fluxes, dry mass fluxes and tissue death		
D_r^{Mrem}	Remobilizable dry matter degradation rate for roots, entity tp,i	g d^{-1}
$D_{tp,i}^{\text{Nph}}$	Photosynthetic N degradation rate for entity tp,i	g d^{-1}
D_r^{Nrem}	Remobilizable N degradation rate for roots	g d^{-1}
$E_C(t)$	Effect of C and N availability within the culm on root N uptake	Dimensionless
$[N]_s$	Soil N concentration	g m^{-3}
$P_{tp,i}$	Dry mass production by entity tp,i	g d^{-1}
$q_r, q_g, q_{tp,i}$	Sink strengths of grains, roots, entity tp,i	Dimensionless
$Q_{tp,0}$	Normalizing factor for the beta function	Dimensionless
S_r^{Mrem}	Remobilizable dry matter synthesis rate for roots, entity tp,i	g d^{-1}
S_r^{Nrem}	Remobilizable N synthesis rate for roots	g d^{-1}
$S_{tp,i}^{\text{Nph,phloem}}$	Photosynthetic N synthesis rate from phloem N for entity tp,i	g d^{-1}
$S_{tp,i}^{\text{Nph,xylem}}$	Photosynthetic N synthesis rate from xylem N for entity tp,i	g d^{-1}
$t_{tp,i}^{\text{init}}$	Thermal time at which grains, roots, entity tp,i begin to grow	$^{\circ}\text{Cd}$
U_r	Root N uptake rate per unit root mass	$\text{g g}^{-1} \text{d}^{-1}$

product of a rate of acquisition per unit root mass (U_r ; $\text{g g}^{-1} \text{d}^{-1}$) and the mass of the root compartment. U_r is described as a potential rate that depends on the nitrate concentration in the soil ($[N]_s$; g m^{-3}), modulated by two dimensionless functions, $E_C(t)$ and $E_N(t)$, which express the positive and negative effects of culm C and N availability, respectively, on N acquisition. The potential rate is described as the sum

TABLE 2. Model parameters: their symbols, definitions and units

Symbol	Definition	Unit
PAR interception		
θ_{ia}	Angle between the vertical and the normal vector to the lamina plane	Radians
$k_{vertical}$	PAR extinction coefficient for vertical entities	$m^2 m^{-2}$
Root N uptake		
$\beta_C \beta_N$	Coefficient for C and N availability effect on root N uptake	Dimensionless
$k_{r,1}$	Constant of the Michaelis function reflecting HATS activity	$g m^{-3}$
$k_{r,2}$	Rate constant of the linear function reflecting LATS activity	$g m^{-3} ^\circ Cd^{-1}$
$(S_r^{Mrem})_{min}$	Minimum threshold of dry matter influx into roots to sustain root N uptake	$g d^{-1}$
$U_{r,max}$	Theoretical maximum N acquisition at saturating soil N concentration	$g m^{-3} ^\circ Cd^{-1}$
N fluxes		
$\delta_r^N \delta_{tp}^N$	Relative degradation rates of remobilizable N for roots, entities tp	$^\circ Cd^{-1}$
γ	Relative rate of potential grain N filling during cell division	$^\circ Cd^{-1}$
$k_{tp,1} k_{tp,2}$	Michaelis–Menten constants defining photosynthetic N synthesis associated with xylem influx for entities tp	$g g^{-1}, J m^{-2} d^{-1}$
$p_r p_{tp,i}$	Proportion coefficient for N influx following dry mass influx into roots, entities tp,i	Dimensionless
$\sigma_{tp,i}^{Nph}$	Relative rate of photosynthetic N synthesis associated with xylem influx for entities tp,i	$g g^{-1} ^\circ Cd^{-1}$
Death of photosynthetic active tissues and photosynthesis		
d_{tp}	Proportion of maximum specific N mass at which tissues die for entities tp	Dimensionless
ω_{tp}	Proportion coefficient linking photosynthesis at saturating PAR and N mass per unit photosynthetic area	d^{-1}
ε_{tp}	Photosynthetic efficiency	$g J^{-1}$
Dry matter fluxes		
$\alpha_{grain} - \beta_{grain}; \alpha_r - \beta_r; \alpha_{tp} - \beta_{tp}$	Two parameters determining the shape of the beta function for grains, roots, entities tp	Dimensionless
$\delta_r^M \delta_{tp}^M$	Relative degradation rates of remobilizable dry mass for roots, entities tp	$^\circ Cd^{-1}$
$\sigma_{grain}^M \sigma_r^M \sigma_{tp,i}^M$	Relative sink strength of grains, roots, entities tp,i	Dimensionless
$t_{grain}^{Macc} t_r^{Macc} t_{tp}^{Macc}$	Period during which grains, roots, entities tp can accumulate dry mass	$^\circ Cd$

of (1) a Michaelis–Menten function of $[N]_s$, corresponding to the activity of the HATS transport system, and (2) a linear function of $[N]_s$, corresponding to LATS activity:

$$U_r(t) = E_C(t) \cdot E_N(t) \cdot \left[\left(\frac{U_{r,max} \cdot \Delta(t) \cdot [N]_s(t)}{k_{r,1} + [N]_s(t)} \right) + (k_{r,2} \cdot \Delta(t) \cdot [N]_s(t)) \right] \quad (3)$$

where $U_{r,max}$ ($g g^{-1} ^\circ Cd^{-1}$) is the theoretical maximum reduced N influx per gram of roots at the soil saturation concentration of N, $k_{r,1}$ ($g m^{-3}$) is the Michaelis constant and $k_{r,2}$ ($g m^{-3} ^\circ Cd^{-1}$) is the rate constant for the linear component.

N_c^{mob} , which represents circulating amino acids and nitrate, was used to account for the known negative feedback of circulating N on N uptake by transport systems. $E_N(t)$ is assumed to be negatively related to the mobile N concentration (i.e. total mobile N mass divided by the total dry mass of green tissues in the culm; $[N]_c^{mob}$; $g g^{-1}$) following an exponential function:

$$E_N(t) = \exp\{-\beta_N \times [N]_c^{mob}(t)\} \quad (4)$$

where β_N (dimensionless) is a parameter.

Dry mass import by roots (S_r^{Mrem} ; $g d^{-1}$) was used to account for the positive effect of carbohydrate availability on

N acquisition by the culm. Below a given threshold ($(S_r^{Mrem})_{min}$ of dry mass influx into roots (calculation is detailed below), N acquisition was assumed to be null; above, $E_C(t)$ was assumed to be exponentially related to S_r^{Mrem} , as:

$$E_C(t) = 0 \quad \text{if } S_r^{Mrem} \leq (S_r^{Mrem})_{min} \\ = 1 - \exp(-\beta_C \cdot S_r^{Mrem}) \quad \text{else} \quad (5)$$

where β_C is a dimensionless parameter.

Nitrogen storage and translocation

The modelling of N accumulation in grains follows that of Bertheloot *et al.* (2008a) and Martre *et al.* (2003): at any time step, N influx into grains follows a potential rate if sufficient N_c^{mob} is available in the common pool; otherwise it corresponds to the mass of mobile N available at that time step. The potential rate is a parametric function of the thermal time, with two successive stages corresponding respectively to cell division and to rapid protein storage: the rate is an affine function of thermal time in the first stage characterized by the parameter γ , and is constant thereafter. All N entering grains is assumed to be permanently stored.

For roots and photosynthetic entities, remobilizable N mass fluxes were calculated as the difference between a synthesis rate (S_r^{Nrem} , $S_{tp,i}^{Nph}$; $g d^{-1}$) and a degradation rate (D_r^{Nrem} , $D_{tp,i}^{Nph}$ for roots and photosynthetic entities, respectively;

$g\ d^{-1}$). Synthesis and degradation rates for an entity were assumed to match the fluxes of mobile N from the common pool into the entity and from the entity into the common pool, respectively. For grains and roots, N was assumed to come only from the phloem following the osmotic gradient generated by C transport (Münch, 1930). After ligulation, laminae do not import C from other organs, so N can only come from the transpiration stream. Concerning other photosynthetic entities, no information could be found in the literature and mobile N was assumed possibly to come from both phloem and xylem. Synthesis rates were calculated differently according to the origin of the mobile N, as given in eqns 10–12. The variation in N mass of respectively the roots and the different photosynthetic modules was written as:

$$\frac{dN_r^{\text{tot}}}{dt} = S_r^{\text{Nrem,phloem}}(t) - D_r^{\text{Nrem}}(t) \quad (6)$$

$$\frac{dN_{tp,i}^{\text{tot}}}{dt} = S_{tp,i}^{\text{Nph,phloem}}(t) + S_{tp,i}^{\text{Nph,xylem}}(t) - D_{tp,i}^{\text{Nph}}(t) \quad (7)$$

The degradation of photosynthetic N produces amino acids that enrich the pool of mobile N; degradation is described by first-order kinetics as in Bertheloot *et al.* (2008a), in accordance with conclusions of Irving and Robinson (2006):

$$D_{tp,i}^{\text{Nph}}(t) = \delta_{tp}^{\text{N}} \times \Delta(t) \times N_{tp,i}^{\text{ph}}(t) \quad (8)$$

where δ_{tp}^{N} ($^{\circ}\text{Cd}^{-1}$) is the relative rate of photosynthetic N degradation for entities of type tp.

Similarly, for roots:

$$D_r^{\text{Nrem}}(t) = \delta_r^{\text{N}} \times \Delta(t) \times N_r^{\text{rem}}(t) \quad (9)$$

where δ_r^{N} ($^{\circ}\text{Cd}^{-1}$) is the relative rate of remobilizable N degradation in roots.

The N influx rate from the phloem into the roots was modelled as proportional to both $[N]_c^{\text{mob}}$ and the dry mass influx into roots S_r^{Mrem} :

$$S_r^{\text{Nrem,phloem}}(t) = [N]_c^{\text{mob}}(t) \times p_r \times S_r^{\text{Mrem}}(t) \quad (10)$$

where p_r is a dimensionless parameter.

Dry mass influx into photosynthetic entities was calculated as the difference between the rate of increase in dry mass for that entity ($S_{tp,i}^{\text{Mrem}}$; $g\ d^{-1}$) and the increase in dry mass due to its own photosynthesis ($P_{tp,i}$; $g\ d^{-1}$); consequently, the N influx rate from the phloem into an entity is written as:

$$S_{tp,i}^{\text{Nph,phloem}}(t) = [N]_c^{\text{mob}}(t) \times p_{tp} \times [S_{tp,i}^{\text{Mrem}}(t) - P_{tp,i}(t)] \quad (11)$$

$S_{tp,i}^{\text{Mrem}}$ and $P_{tp,i}$ calculations are detailed in subsequent sections.

The N influx rate through the xylem into photosynthetic entities is expressed, following Bertheloot *et al.* (2008a), as a function of the PAR intercepted, the mass of photosynthetic

tissues ($M_{tp,i}^{\text{green}}$, g) and $[N]_c^{\text{mob}}$:

$$S_{tp,i}^{\text{Nph,xylem}}(t) = \sigma_{tp}^{\text{Nph}} \cdot \Delta(t) \cdot M_{tp,i}^{\text{green}}(t) \cdot \frac{[N]_c^{\text{mob}}(t)}{[N]_c^{\text{mob}}(t) + k_{tp,1}} \cdot \frac{\text{PAR}_{tp,i}}{\text{PAR}_{tp,i} + k_{tp,2}} \quad (12)$$

where σ_{tp}^{Nph} ($g\ g^{-1}\ ^{\circ}\text{Cd}^{-1}$) is the relative rate of $N_{tp,i}^{\text{ph}}$ synthesis associated with xylem influx, and $k_{tp,1}$ ($g\ g^{-1}$) and $k_{tp,2}$ ($J\ m^{-2}\ d^{-1}$) are Michaelis constants.

Photosynthesis and dry mass production

The production of total dry mass per unit area of photosynthetic tissue was modelled as a function of the daily PAR intercepted according to a rectangular hyperbola (Thornley, 1998a). This approximation proved satisfactory when assessed using instantaneous light response curves of photosynthesis, measured for a range of leaf N statuses (data not shown).

Dry mass production of a photosynthetic entity was calculated as:

$$P_{tp,i}(t) = \frac{\varepsilon_{tp} \cdot p_{tp}^{\text{max}}(t) \cdot \text{PAR}_{tp,i}(t)}{\varepsilon_{tp} \cdot \text{PAR}_{tp,i}(t) + p_{tp}^{\text{max}}(t)} \cdot A_{tp,i}^{\text{green}}(t) \quad (13)$$

where ε_{tp} ($g\ J^{-1}$) is the photosynthetic efficiency and p_{tp}^{max} ($g\ m^{-2}\ d^{-1}$) is photosynthesis at saturating light.

Photosynthetic efficiency varies mainly with long-term adaptation to light climate and was assumed to be independent of leaf N (Hirose and Werger, 1987; Schieving *et al.*, 1992). Photosynthesis at saturating light is limited by the amount of Rubisco, which is the main form of leaf N storage (Evans, 1989b). Accordingly, p_{tp}^{max} was assumed to be proportionally related to photosynthetic N mass per unit area, with the coefficient ω_{tp} (d^{-1}) (Hirose and Werger, 1987; Thornley, 1998a):

$$p_{tp}^{\text{max}}(t) = \omega_{tp} \cdot \frac{N_{tp,i}^{\text{ph}}(t)}{A_{tp,i}^{\text{green}}(t)} \quad (14)$$

In each photosynthetic entity, a given fraction of tissues die when photosynthetic N mass per unit photosynthetic area drops below a threshold, which was expressed as a fixed fraction (d_{tp} ; dimensionless) of entity N mass per unit photosynthetic area at flowering. The photosynthetic dry mass of an entity was calculated by a cross product depending on its total dry mass, total area and photosynthetic active area

Dry mass distribution within the plant

Photosynthetic entities and roots lose dry mass by remobilization (D_r^{Mrem} , $D_{tp,i}^{\text{Mrem}}$; $g\ d^{-1}$), which was assumed to follow first-order kinetics of remobilizable dry mass (Drouet and Pagès, 2007):

$$D_{tp,i}^{\text{Mrem}}(t) = \delta_{tp,i}^{\text{M}} \times \Delta(t) \times M_{tp,i}^{\text{rem}}(t) \quad (15)$$

$$D_r^{\text{Mrem}}(t) = \delta_r^{\text{M}} \times \Delta(t) \times M_r^{\text{rem}}(t) \quad (16)$$

where δ_{tp}^M and δ_r^M ($^{\circ}\text{Cd}^{-1}$) are relative rates of remobilization for an entity of type tp and roots, respectively.

At any given time step, the mobile dry mass available within the culm is the dry mass produced by photosynthesis during that time step, plus the remobilized dry mass. It is distributed among modules following the GreenLab model (de Reffye *et al.*, 2008): the increase in dry mass of one entity, the root compartment or grain compartment, is the product of the dry mass available at the culm scale and the demand in dry mass of the module. This demand is defined as the sink strength of the module, which is a dimensionless function [$q_{tp,i}(t)$, $q_r(t)$ and $q_{grain}(t)$ for an entity, roots and grains, respectively], expressed relative to the sum of the sink strength of all modules. Below is the equation for an entity of type tp and rank i :

$$S_{tp,i}^{\text{Mrem}} = \frac{q_{tp,i}(t)}{\sum_{tp,i} q_{tp,i}(t) + q_r(t) + q_{grain}(t)} \cdot \left(\sum_{tp,i} P_{tp,i} + \sum_{tp,i} D_{tp,i}^{\text{Mrem}} + D_r^{\text{Mrem}} \right) \quad (17)$$

For one module, sink strength was modelled from a relative sink strength parameter (denoted $\sigma_{tp,i}^M$ for entity tp,i , σ_r^M for roots and σ_{grain}^M for grains), which corresponds to its potential sink strength, and a normalized beta function, which indicates the pattern of change in sink strength during its life. For entity tp,i , the equation is written as:

$$q_{tp,i}(t) = \sigma_{tp,i}^M \cdot \left[\left(\frac{tt - tt_{tp,i}^{\text{init}}}{tt_{tp,i}^{\text{Macc}}} \right)^{\alpha_{tp}-1} \cdot \left(\frac{tt - tt_{tp,i}^{\text{init}}}{tt_{tp,i}^{\text{Macc}}} \right)^{\beta_{tp}-1} \right] / Q_{tp,0} \quad (18)$$

where $tt_{tp,i}^{\text{init}}$ ($^{\circ}\text{Cd}$) is the thermal time at which the entity begins to grow, $tt_{tp,i}^{\text{Macc}}$ ($^{\circ}\text{Cd}$) is the duration of the period in which the module can accumulate dry mass and $Q_{tp,0}$ is a normalizing factor calculated so that the relative sink strength parameter corresponds to the maximal value of the sink strength. It is calculated as:

$$Q_{tp,0} = \left(\frac{\alpha_{tp} - 1}{\alpha_{tp} + \beta_{tp} - 2} \right)^{\alpha_{tp}-1} \cdot \left(\frac{\beta_{tp} - 1}{\alpha_{tp} + \beta_{tp} - 2} \right)^{\beta_{tp}-1} \quad (19)$$

Model implementation

The model was implemented in C++. The daily time step was sub-divided into four sub-steps in order to avoid mistakes in solving differential equations due to a too long time step. Above four sub-steps, there were no longer any significant changes in model predictions.

Model initialization method

Model initialization corresponds to a description of the state variables of the different modules at flowering. One difficulty

is to be able to quantify the different forms of N and dry matter; however, they can be indirectly estimated from measurements of total dry mass and N mass in each entity at flowering and maturity, if (a) it is assumed that the matter remaining in dead tissues at maturity corresponds to structural matter and (b) the mean percentage of mobile N in an entity is known (see Bertheloot *et al.*, 2008a). This method is used in a companion paper (Bertheloot *et al.*, 2011) to calibrate the model.

DISCUSSION

NEMA predicts the acquisition and distribution within wheat plants after flowering of both dry matter and N, using soil N concentration and the daily time courses of temperature and of PAR above the canopy as driving variables. The originality of the model is that it proposes, thanks to an explicit description of plant botanical structure, a mechanistic alternative to the conventional demand-driven approach for modelling N fluxes within the aerial parts of plants. On the other hand, like other models at the whole plant scale, NEMA uses much simplified descriptions of the processes with the aim of simulating relevant plant behaviours without multiplying the number of parameters. The choices we made were the following: N content and remobilization in each photosynthetic organ is simulated from the formalization of Rubisco degradation and synthesis, which depends on the PAR intercepted by the organ and the amount of mobile N. While individual organs differ in their PAR intercepted, they are assumed to share identical amounts of internal resources; mobile N is represented by a common pool shared by all plant organs in accordance with suggestions by several authors (Cooper and Clarkson, 1989; Kull and Kruijt, 1999; Thornley, 2004). The amount of mobile N varies according to the fluxes of N from and to plant organs: it is enriched by root uptake and assimilation or organ remobilization, and depleted by grain accumulation or protein synthesis. This modelling of N fluxes within the plant was associated with a previous model of N acquisition (Devienne-Barret *et al.*, 2000; Malagoli *et al.*, 2004; Drouet and Pagès, 2007) that integrates the activities of transport systems in response to soil availability and the feedbacks linked to plant nutrient status (for reviews, see Glass and Siddiqi, 1995; Daniel-Vedele *et al.*, 1998).

The present version of NEMA does not include a realistic description of root architecture or of soil: such developments will be required, for example, to compare cultivars differing in root architecture or to define optimum agricultural practices, since practices may affect N forms and their spatial distribution in the soil. The present implementation should, however, be sufficient to respond to the objective of testing the validity of the new concepts implemented and gain new insights into N economy regulation in the aerial parts of the plant. In contrast to N, few efforts have been made here to model C acquisition and distribution mechanistically within the plant. Nitrogen distribution, together with PAR, regulates photosynthesis, but it is not C metabolism, but the increase in total dry mass that is directly modelled. This simplification is valid since C has a much higher contribution to dry mass than N. Moreover, the modelling of total dry mass distribution between plant parts follows the conventional demand-driven

approach. A more elaborate model for C transfer to wheat grain was proposed, for instance, by Bancal and Soltani (2000) or Minchin and Lacomte (2005). Coupling this type of model with the N model proposed in this study will be considered in a future step; however, we believe that, given the determining role of N metabolism for C production at the whole plant scale and the fact that C fluxes are mainly directed towards grains after flowering, the present implementation will already provide new insights into the regulation of N economy within the plant after flowering, which is fundamental for more efficient N management practices or cultivars.

The simulations we presented in Bertheloot *et al.* (2008a) showed that it is possible to accurately simulate the gradient of photosynthetic N that exists between the top and the bottom leaves of a dense wheat canopy by considering mobile N as a pool equally accessible to all laminae and by relating the protein synthesis rate to the PAR intercepted by laminae. Functional acclimation of the lamina photosynthetic apparatus to PAR intensity has been demonstrated in different studies (Prioul *et al.*, 1980a, b; Grindlay, 1997; Terashima *et al.*, 2005). The relationship between PAR intensity and the transpiration stream is likely to be the origin of this acclimation (Pons and Bergkotte, 1996; Pons *et al.*, 2001) as the transpiration stream supplies nitrate and circulating amino acids as well as cytokinins, known to be involved in the synthesis of the photosynthetic apparatus (Aloni *et al.*, 2005). NEMA assumes that N of all photosynthetic tissues can be modelled in a unified way and extends the assumption of acclimation to PAR light to all plant organs. The observation that entities of the same type but located at different depths in the canopy followed similar patterns of N concentration depletion (Bertheloot *et al.*, 2008b) is in accordance with this assumption. Time courses of these vertical gradients were similar for leaf laminae and sheaths on the one hand, and for internodes and chaff on the other hand, but differed between the two groups. This suggests that, if the processes involved are the same, the relationship of organ N content to light climate depends on the type of organ; this would imply different parameter values for synthesis and/or degradation.

The central role of PAR light interception in N dynamics within the plant implies being able to estimate it accurately. In the present version, Beer's law was implemented and is used in a companion paper to calibrate the model and study its behaviour (Bertheloot *et al.*, 2011). This law has the advantage of allowing rapid calculations and of only requiring a simple description of plant structure. However, for a detailed investigation on how plant architecture impacts N economy or for analysing plant–plant interactions, the next step will be to couple it with a model of radiative transfer based on surface approaches (Chelle and Andrieu, 1998), as in Evers *et al.* (2010). To investigate these questions, one interesting feature of NEMA is to formalize N economy for an individual plant. Moreover, structure is an input of the model, so that it would be easy to include variability in the structure of culms constituting the field.

NEMA also extends our previous model (Bertheloot *et al.*, 2008a) in that organ N is assumed to come not only from the xylem but also from the phloem. The two transport paths involve different driving forces: N influx into an entity follows the transpiration stream when it originates from the

xylem, while, when it originates from the phloem, it is driven by a mass flow mechanism following the carbohydrate gradient that exists between sources and sinks for C (Münch, 1930). Thus, the formalization of N influx into an entity and N synthesis rate differ depending on the transport path. On the other hand, the two synthesis paths are assumed to have access to the same circulating N, which is considered as a single pool. Consequently, the formalization implicitly assumes that N concentrations in xylem and phloem are correlated with each other. This is probably because transfers of amino acids between xylem and phloem take place throughout the vascular system (for reviews, see Feller and Fischer, 1994; Delrot *et al.*, 2001).

It is likely that, similar to what is known for leaves (Bregard and Allard, 1999; Turgeon, 2006), other mature photosynthetic tissues cannot import C from other plant parts, so that the role of phloem in N import is negligible for such tissues. By allowing N to be imported through both the xylem and phloem for non-leaf tissues, the model will enable analysis of their possible contributions. In contrast, growth of young organs still hidden in the whorl or having a small transpiration area depends on phloem for import of both C and N. It will thus be important to take into account the role of phloem when adapting the model to the pre-flowering period.

The response of grain N accumulation to N fertilization at flowering was simulated with the simple approximation that the N flux from the common pool of mobile N to the grains follows a potential function of thermal time, but occurs only when mobile N is available in the common pool. Thus, the process by which N influx towards grains is driven by the osmotic gradient created by the differences in carbohydrate concentration (Münch, 1930) is not explicitly formalized. This choice was made because of its simplicity and the good predictions obtained with contrasted treatments (Bertheloot *et al.*, 2008a). It appeared to be sufficient in usual agronomic conditions, when N availability, rather than carbohydrate transport and sink limitation, limits the import of N into the ear (Dingkuhn *et al.*, 2007; Bancal, 2009); a more exact formulation may be required in a context of strong carbohydrate limitation, due to light starvation, for example.

Photosynthetic tissues were assumed to die when photosynthetic N mass per unit area fell below a critical value. The threshold value is known to vary according to N fertilization, for example (Hirel *et al.*, 2005). In a preliminary version, we implemented the frequently used assumption (Ackerly and Bazzaz, 1995; Ackerly, 1999) that tissue death occurs when C net assimilation by tissues is null. This however led to the prediction of an earlier tissue death for the lowest leaves (data not shown) compared with that observed, which may be due to inaccuracy in the calculation of net photosynthesis. Using this assumption would probably require an appropriate account of respiration costs under conditions of low light and low N. Consequently, the N threshold for tissue death was defined empirically as a fraction of entity N content at flowering, similar to that in the original model (Bertheloot *et al.*, 2008a). When this threshold is reached, leaves have a low impact on plant C and N economy, so the lack of a mechanistic approach is unlikely to hamper the use of the model.

The formalization of N acquisition by roots follows that proposed earlier based on nitrate uptake by transport systems and its subsequent assimilation (Devienne-Barret *et al.*, 2000; Malagoli *et al.*, 2004; Drouet and Pagès, 2007). Nitrate is the mineral form brought by fertilization and the main form assimilated in wheat. Changes would be required for the case of a wider range of N forms. The rate of N acquisition by roots is expressed as the sum of a Michaelis–Menten function of soil N concentration, which reflects the saturable character of HATS, and a linear function of soil N concentration, which reflects the non-saturable character of LATS [Cacco *et al.* (2002) for wheat; Malagoli *et al.* (2004) for oilseed rape; Siddiqi *et al.* (1990) for barley]. The well-known positive dependence of N acquisition, mainly N assimilation into reduced N, on carbohydrate availability (e.g. Delhon *et al.*, 1996) as well as the negative feedback of reduced N (Forde and Clarkson, 1999; Touraine *et al.*, 2001) on transport system activities were also accounted for by the use of specific functions. Regulation by C availability was based on dry mass influx into roots, which may not accurately reflect carbohydrate availability. This difficulty could probably be overcome by developing a more mechanistic module for C metabolism. In contrast, to account for N negative feedback, the formalization of a common pool of circulating N is an original and interesting feature in the model. In a previous study (Bertheloot *et al.*, 2008a), it proved to be a reliable variable to regulate leaf N dynamics and grain N filling. For regulation of root N uptake, it provides a physiological state variable, whereas previous models used indicators that do not have a causal effect on physiological activities. The functions implemented in the model are adapted from Drouet and Pagès (2007) and Cardenas-Navarro *et al.* (1999). However, there is no strong support for the choice made: more work is needed to define mathematical functions that actually reflect the regulation of N acquisition by mobile N.

In conclusion, a functional–structural model of N economy for wheat after flowering is described in the present paper which accounts for physiological activities governing N acquisition and distribution within the aerial part of the plant and, in a simplified manner, for C–N interactions. This model is a mechanistic – yet still simple – alternative to the teleonomic approaches usually used to model N economy, based on the idea expressed by different authors that Rubisco turnover plays a central role in the regulation of N fluxes within the aerial part of the plant (Thornley, 1998a, 2004; Gallais *et al.*, 2006; Hirel and Gallais, 2006). Our motivation comes from the assumption that the integration of processes governing N metabolism at the whole plant scale will provide insights into the regulation of N acquisition and the effect of N on yield by environment (light and soil N) and genotype, which is fundamental for the design of more efficient cultivars and practices as mentioned by Hammer *et al.* (2010). The use of the model as a comprehensive tool is investigated in a companion paper (Bertheloot *et al.*, 2011). The model proposed here illustrates the potentialities offered by the functional–structural approach to integrate physiological knowledge at the scales of the whole plant and field, and thus to link physiology, eco-physiology and agronomy. The generality of the processes formalized should make the model adaptable to species other than wheat and, as an individual plant type model, suitable for

plurispecific crops provided it is coupled with an appropriate model of radiative transfer. This will be one of the next steps. Another step will be to use the model proposed here as the basis for a model dealing with the more complex pre-flowering period. This will require the mechanistic formalization of both C and N metabolism and of their effects on the plant structure dynamic.

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