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A stochastic growth model of grapevine with full interaction between environment, trophic competition and plant development.

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Abstract

Grapevine development is mainly determined by environmental factors whose effects are modulated by its complex topological structure. The trophic relationships between all the organs of the different axes appear to be the main underlying process which drive axis organogenesis in fluctuating environment. A new modelling approach is proposed based on GreenLab formalism in which axis organogenesis is controlled by stochastic processes according to trophic competition between the different axes. In this model, a water budget was also implemented to account for the effects of water depletion. The model was validated at organ and axis scales on a large range of environmental conditions in terms of photosynthetic active radiation, temperature and soil water supply. The efficiency of the model to simulate plant development at a detailed scale proved its ability to further analyse of the retroactions between plant development and the different environmental variables in order to improve crop management.

1 Introduction

Grapevine development is mainly determined by environmental factors such as temperature (through thermal time [1]) and abiotic stresses (e.g. soil water deficit [2]). Moreover, the source-sink relationships during plant development also modulate the vegetative and reproductive development [1], in relation with changes in trophic competition inside the plant [3]. The effects of all these factors on plant development have resulted in the development of diverse agronomic practices, such as pruning [4] or irrigation [5]. However, to improve these practices and increase our understanding of plant development in contrasting environments, modeling approaches appear useful and complementary to experiments.

Grapevine shoot is a modular branching system, with one primary axis and many secondary axes organized into a structure of three successive phytomer types (P0, P1 and P2). P0 phytomer bear no tendril or cluster. Phytomers bearing one tendril or cluster are classified as P1 or P2 phytomers depending on their positions with respect to the previous P0 phytomer [1]. Temperature determines the potential organogenesis of the axes [1], and trophic competition modulates their development [1]. The impact of trophic competition on axis organogenesis varies according to the type of axis. Primary axis appears almost insensitive to trophic competition. Secondary axes born by P0 phytomers are less affected than the ones born by P1 or P2 phytomer [1]. Soil water deficit reduces the photosynthetic activity of the plant [2] and in this way tends to have the same effect on plant development as an increase in trophic competition. Furthermore, soil water deficit also induces a signalic regulation, which leads to reduce the potential organogenesis of the axes [6].

A model taking into account the effects of temperature and soil water deficit on shoot development have been proposed [2]. Besides, a first attempt to describe biomass fluxes in grapevine was made with GreenLab model [3]. This model proved effective for analyzing the relationships between plant functioning and plant development. However, in this approach, plant organogenesis was fixed and defined with no retroaction between trophic competition and organogenesis. Thus, it did not allow simulating the plant development in fluctuating environments. A formalization of the interactions between plant organogenesis and plant functioning in GreenLab model has been already done [7]. But these studies were purely theoretical with no confrontation with plant measurements. We propose in the present study, a new version of the grapevine model with stochastic organogenetic rules and probability distribution linked to temperature, trophic competition and water plant status. To assess pluriannual simulations this model includes reserve mobilization and storage. The results of the simulation of the model were compared to experimental data collected on two cultivars ‘Syrah’

and ‘Grenache N.’ submitted to different levels of photosynthetic active radiation (PAR) and soil water status (FTSW).

2 Materials and methods

2.1 Model description

General structure. GreenLab is a model which aims to calculate at each time step of the model total assimilate supply (Q) and organ demand (D). In GreenLab the Q/D ratio was observed to be an adequate indicator of plant trophic status [7]. The aims of this present version was to input in the model relationships between Q/D values at each growth cycle and organogenesis processes. Moreover, the influence of the plant water status on both organ demand and assimilate supply was also implemented in this version (Fig. 1).

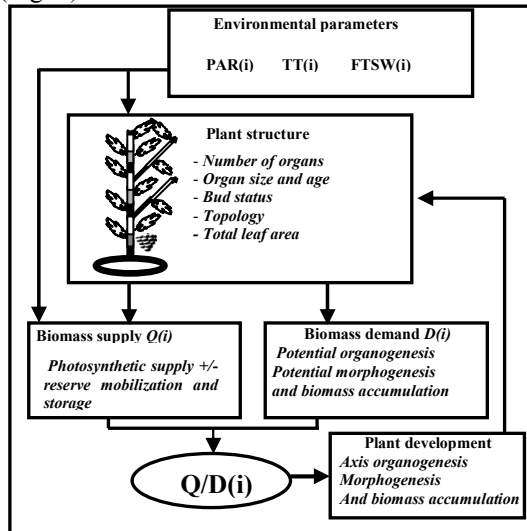


Figure 1 : General structure of the model and method of calculation of Q and D .

Plant topology. Grapevine shoot is composed of primary and secondary axes. The secondary axes can be separated into two types of axis depending on their observed behaviour [1]. Thus, three orders of physiological ages are implemented in the model, respectively for primary axis, P0 secondary axes and P1-P2 secondary axes (i.e. secondary axes born respectively by P0 or P1-P2 phytomers). On all the axes, the P0 phytomers bear no tendril or cluster, and the P1-P2 phytomers bear a tendril or a cluster [1]. As previously observed, the succession sequence in all the axes is repetitive (P0-P1-P2) excepted at the base of the primary axis (corresponding to the preformed part of the shoot). In this zone the succession of the

phytomers is disturbed and a consensus sequence corresponding to the sequence with the greatest probability [8] observed in natural conditions is forced in the model for both cultivars as follows:

$$P0-P0-P0-P0-P1-P2-P0-P0-P1-P2-(P0-P1-P2)... \quad (1)$$

The clusters, if present in the plant, are located on the two first phytomers P1 and P2 on the primary axis. At most two clusters are able to develop on each shoot [1].

Primary axis organogenesis. In well watered condition, the leaf appearance rate on the primary axis was observed to be constant even in situation of severe trophic competition and equal to $0.044 \text{ phytomer} \cdot \text{Cd}^{-1}$ [1][8]. This trend allows considering, when there is no water deficit, a constant time step of the model (D_{gc}) equal to the rhythm of phytomer appearance on the primary axis (23°Cd). Preliminary studies emphasized the fact that the development does not end in absence of water deficit, even in situation of high level of trophic competition [3]. Thus, in the model, the organogenesis of the primary axis continues with the same leaf appearance rate as long as $Q(i)$ is not equal to 0.

Secondary axis organogenesis. Five development states of the axillary buds are considered in the model (Fig. 2). In state 0 the bud has not broken out, in state 0-1 the bud has broken out and is in a transition phase called ‘lag period’ during which it is not able to produce a leaf. When the bud becomes active three events can occur at each development stage, the bud may produce a leaf (state 1-1), may stay quiescent (state 1) or may die (state 1-0).

In the current version of the model, the probability of budburst occurrence (P_b) (transition from state 0 to state 0-1), is set to 1 except for the two first axillary buds for which P_b is set to 0 [8]. Note that these probabilities are fixed in the model at the beginning of plant development and do not change during plant development. The transition from state 0-1 to state 1 systematically occurs after a constant lag period (d_l). Thus, an axillary bud of insertion rank j on the primary axis, which has broken up, becomes active at the growth cycle $j+d_l$ with d_l generally equal to 6 GC. [8] When a bud becomes active, P_e corresponds to the probability of an axillary bud to die, P_p corresponds to the probability of an axillary bud to produce a leaf and $1-P_p$ to its probability to stay alive without producing new leaves (i.e. quiescent).

In previous studies [1][8], differences between leaf appearance rate of P0 secondary axes and P1-P2 secondary axes were observed. Moreover, the leaf

appearance rate of the different axes was roughly constant during their development and depended on the trophic competition at their appearance which is represented by $Q(i)/D(i)$ (called afterwards Q/D). This formalism is implemented in the model by considering that P_p depends on the type of secondary axis (i.e physiological age) (k) and on the value of Q/D when the axillary bud becomes active. The probability of a secondary axis of insertion rank j and of type k ($P_p(j,k,i)$) to produce leaf (state 1-1) at growth cycle i is thus implemented in the model.

$$P_p(j,k,i) = c_p(k) \cdot \left(\frac{Q(j+d_i)}{D(j+d_i)} \right) \quad (2)$$

A consistent correlation between the rate of secondary axes which ended their development (T_e) and the Q/D values at the corresponding growth cycle (i) was observed in previous studies [3]. The sensitivity of secondary axes to trophic competition also depended on the number of leaves on the corresponding axis (c) and on the type of secondary axis (i.e physiological age) (k). Thus, in the model the probability of a secondary axis of type k , with number of leaves c , of insertion rank j , to die at the growth cycle i ($P_e(j,k,c,i)$) is expressed according to the predicted rate of end of development of secondary axes of type k and of number of leaves c at growth cycle i ($T_e(k,c,i)$) as follows [3]:

$$\left\{ \begin{array}{l} \text{if } (1-T_e(k,c,i)) \cdot N_{II}(k,c,i) < N_{II,p}(k,c,i) \\ P_e(j,k,c,i) = \left(\frac{N_{II,p}(k,c,i) - N_{II}(k,c,i) \cdot (1-T_e(k,c,i))}{N_{II,p}(k,c,i)} \right) \\ \text{if } (1-T_e(k,c,i)) \cdot N_{II}(k,c,i) \geq N_{II,p}(k,c,i) \\ P_e(j,k,c,i) = 0 \end{array} \right. \quad (3)$$

with $N_{II}(k,c,i)$ the total number of secondary axes of type k , with a number of leaves c at growth cycle i . $N_{II,p}(k,c,i)$ is the potential number of secondary axes calculated by considering that no new axillary bud died (state 1-0) at growth cycle i .

With [4]:

$$T_e(k,c,i) = 1 - \frac{1}{1 + \exp \left[\frac{-\frac{Q(i)}{D_p(i)} + a_e(k,c)}{b_e(k,c)} \right]} \quad (4)$$

with $D_p(i)$ the potential demand for biomass, calculated by considering that no new axillary bud died (state 1-0) at growth cycle i .

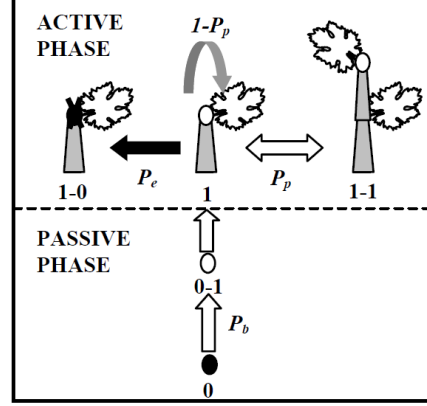


Figure 2: Different states of the axillary buds and transition probabilities.

Photosynthetic production. The photosynthetic production of biomass at each growth cycle (GC) i ($Q_p(i)$) is implemented in the model using a Beer-Lambert law production function, previously used for grapevine [3] as follows:

$$Q_p(i) = PAR(i) \cdot Sp \cdot RUE(i) \cdot \left[1 - \exp \left(\frac{-k \cdot S(i)}{Sp} \right) \right] \quad (5)$$

with $PAR(i)$, the photosynthetic active radiation at each growth cycle ($mol \cdot m^{-2} \cdot GC^{-1}$), Sp , the projected leaf area (m^2), $RUE(i)$ the radiation use efficiency at growth cycle i ($g \cdot mol^{-1} \cdot m^{-2}$), $S(i)$, the total leaf surface at growth cycle i (m^2) and k the extinction coefficient for the Beer Law.

Reserve behaviour. During the ‘heterotrophic’ phase of plant development, from budburst to flowering [9], biomass from reserve may be mobilized. The initial biomass ($Q_{r,ini}$) of reserve is an input of the model at the beginning of plant development. A parameter β_h describes the part of non-structural reserve biomass available for shoot development [10]. The time course of reserve mobilization is modelled with a beta function ($f_r(i)$), and a duration of biomass reserve mobilization ($T_r(i)$). At each growth cycle i the quantity of biomass in the reserve ($Q_r(i)$) is calculated as follows:

$$Q_r(i) = Q_r(i-1) - f_r(i) \cdot \beta_h \cdot Q_{r,ini} \quad (6)$$

During the ‘autotrophic’ phase (from flowering to maturity) reserve ‘storage’ and ‘mobilization’ can both occur [9].

Reserve ‘storage’ occurs when:

$$Q_p(i) > r \quad (7)$$

Reserve ‘mobilization’ occurs when:

$$Q_p(i) \leq r \quad (8)$$

with c ($g \cdot GC^{-1}$), an input parameter of the model.

Which could be simplified for young plant [8][10] to $r=0$.

When reserve ‘storage’ occurs, the part of biomass allocated to the reserve is estimated from an allometric relationship combining the photosynthetic production to the reserve biomass. A parabolic function is implemented.

$$Q_r(i) = Q_r(i-1) + a.Q_p(i)^2 + b.Q_p(i) \quad (9)$$

When reserve ‘mobilization’ occurs, a parameter β_a , accounts for the part of biomass that may be mobilizable from the reserve [10]. This parameter allows to estimate the initial available biomass of reserve when the mobilization of reserve begins (during the autotrophic phase), at growth cycle z . The fraction of biomass (μ) that may be mobilized from reserve is considered as constant at each growth cycle. Consequently, we can estimate the quantity of available ($Q_{r,a}(i)$) and total ($Q_r(i)$) reserve biomass as follows:

$$Q_{r,a}(i) = \begin{cases} \beta_a.Q_r(z).(1-\mu.(i-z+1)) & \text{if } Q_{r,a}(i-1) > 0 \\ 0 & \text{else} \end{cases} \quad (10)$$

$$Q_r(i) = \begin{cases} Q_r(i-1) - \mu.\beta_a.Q_r(z) & \text{if } Q_{r,a}(i-1) > 0 \\ Q_r(i-1) & \text{else} \end{cases} \quad (11)$$

with z , the growth cycle when reserve mobilization begins.

Biomass allocated to the non-perennial organs (Q).

The combination between photosynthetic production and reserve behaviour allows to calculate the part of biomass allocated to the shoot ($Q(i)$). To take into account the part of reserve biomass mobilized from the reserve and really converted into biomass for shoot development a parameter α is implemented in the model [11]

$$\text{In the ‘heterotrophic’ phase of plant development:} \\ Q(i) = Q_p(i) + \alpha.f_r(i).\beta_h.Q_{r,mi} \quad (12)$$

In the ‘autotrophic’ phase of plant development.

$$\left\{ \begin{array}{l} \text{if } Q_p(i) > r \\ \quad Q(i) = (1-b)Q_p(i) - aQ_p(i)^2 \\ \\ \text{if } Q_p(i) \leq r \\ \\ Q(i) = \begin{cases} Q_p(i) + \alpha.\mu.\beta_a.Q_r(z) & \text{if } Q_{r,a}(i-1) > 0 \\ 0 & \text{else} \end{cases} \end{array} \right. \quad (13)$$

Biomass allocation between non-perennial organs. In the model biomass allocation processes are separated in two phases, the first phase is a phase of biomass allocation for each growth process (organ expansion, and secondary growth) and the second phase is a phase of allocation between the different organs.

Demand for organ expansion is calculated as already proposed in the GreenLab model [7]. Each organ (o) for each physiological age (k) is defined using a sink strenght ($P_o(k)$), and a beta law function depending on its chronological age (n) ($f_o(n)$) and identical for all the physiological ages. For clusters, two phases of development exist, one from flowering to veraison and the other one from veraison to maturity [12]. Therefore two sink strenghts with their own associate beta function are implemented in the model. The total demand for organ expansion at the growth cycle i ($D_{exp}(i)$) is thus defined by:

$$D_{exp}(i) = \sum_k \sum_0 P_o(k) . \sum_n f_o(n) . N_o(n, k, i) \quad (14)$$

where $N_o(n, k, i)$ is the total number of organs of type o , physiological age k and chronological age n in the plant at growth cycle i .

The total demand for secondary growth at growth cycle i ($D_s(i)$) is implemented in the model, according to the total number of leaves ($N_l(i)$) and using a constant sink strenght (P_s).

$$D_s(i) = P_s.N_l(i) \quad (15)$$

Thus the total biomass demand ($D(i)$) of the shoot at each growth cycle i can be expressed by [13]:

$$D(i) = D_{exp}(i) + D_s(i) \quad (16)$$

The part of biomass allocated to organ expansion (Q_{exp}) and secondary growth (Q_s) is calculated as a function of the total demand:

$$Q_{exp}(i) = D_{exp}(i) . \frac{Q(i)}{D(i)} \quad \text{and} \quad Q_s(i) = D_s(i) . \frac{Q(i)}{D(i)} \quad (17)$$

The biomass allocated to each organ for its expansion ($\Delta q_o(n, k, i)$) is given by:

$$\Delta q_o(n, k, i) = P_o(k) . f_o(n) . \frac{Q_{exp}(i)}{D_{exp}(i)} \quad (18)$$

The biomass allocated for secondary growth of the different internode is implemented according to the formalism previously proposed [13], using a flexible submodel of allocation with two modes of allocation, one based on the pipe model theory and the second

based on a uniform allocation. In the first mode, the biomass allocated for secondary growth of each internode is calculated according to the surface of active leaves above it in the plant topology. In the second mode the surface above the internode is not considered and all the leaves participate equally to secondary growth. The two models are combined with a parameter λ [13].

Organ sizes are calculated with allometric relationships between organ masses and sizes. For leaves, individual leaf area ($S_j(i)$) at each growth cycle is calculated, using a specific leaf weight ($SLW(i)$) which was observed, in previous studies, to be highly correlated to the photosynthetic active radiation ($PAR(i)$) [14].

$$S_j(i) = q_{l,x}(i) / SLW(i) \quad (19)$$

$$\text{With } SLW(i) = a_l \cdot PAR(i) + b_l \quad (20)$$

Where $q_{l,x}(i)$ is the biomass of the leaf x at growth cycle i .

Influence of water status on plant development. The effect of soil water deficit on axis organogenesis was observed to be independent from the decrease in global photosynthetic activity [2][6], due to hormonal signalitic mainly related to ABA influx from roots [6]. However the observed decrease in axis organogenesis depended on the type of axis. Primary axis was less affected than secondary axes, and the different secondary axes (P0 or P1-P2) were affected to the same extend. Thus, as previous observed [2], the water status dependence of the leaf appearance rate is implemented in the model. For primary axis, this trend forces to change the duration of the growth cycle (d_{gc}), defined as the phyllochrone of the primary axis, and equal to 23°Cd in well-watered situation [8]:

$$d_{gc} = 23 \cdot (1 + a_{w,I} \cdot \exp(+b_{w,I} (FTSW(t) + c_{w,I}))) \quad (21)$$

with $FTSW(t)$ the fraction of transpirable soil water at date t , $a_{w,I}$, $b_{w,I}$ and $c_{w,I}$ three parameters of the sigmoid equation.

The effect of soil water deficit on $P_p(j,k,i)$ (we recall that $P_p(j,k,i)$ determines at each growth cycle i the probability to produce a leaf for a secondary axis of type k and of insertion rank j) is implemented as proposed by Lebon et al. 2006 [2]:

$$P_p(j,k,i) = \left(\frac{1}{1 + a_{w,II} \exp(+b_{w,II} (FTSW(i) + c_{w,II}))} \right) c_p(k) \cdot \left(\frac{Q(j+l)}{D(j+l)} \right) \quad (22)$$

with $a_{w,II}$, $b_{w,II}$ and $c_{w,II}$ three parameters of the sigmoid equation.

Radiation use efficiency was observed to be reduced by soil water deficit due to the limitation of CO_2 exchange [2]. We define $\alpha_w(i)$ as a function of

$FTSW(i)$ to modified RUE_{max} in situation of soil water deficit at each growth cycle i [2]:

$$RUE(i) = \alpha_w(i) \cdot RUE_{max} \quad (23)$$

$$\text{With } \alpha_w(i) = \frac{1}{1 + 6.20 \cdot \exp(-6.36(FTSW(i) + 0.14))} \quad (24)$$

2.2 Experiments used for model parametrization and evaluation.

Five experiments were carried out at the SupAgro-Montpellier-INRA campus in a greenhouse (exps. 3,4,5) and outside (exps. 1&2) on two grapevine (*Vitis vinifera* L.) cultivars ‘Syrah’ and ‘Grenache N.’ (Table 1). Experiments were managed on three-year-old plants cultivated in pots (diameter=0.3m, height=0.5m), filled with equal proportion of clay, loam and sand. Only one shoot was kept on each plant in exps. 1 to 5. This shoot was trained vertically during the experiment. All clusters were removed for plants of treatments T whereas for treatment TG (exp.1), two clusters were retained on the shoot. In these experiments water supplies and fertilization were managed in order to avoid any water or mineral deficiency. A large range of PAR conditions was observed among these experiments ranging from 0 to 53.5 mol.m⁻².d⁻¹. In exp. 5, plants were subjected to a mean value of PAR of 12.6 mol.m⁻².d⁻¹ (Table 1) from budburst to 580°Cd and were then placed below a polyester shelter in order to get a complete occultation of incident PAR (Table 1).

Table 1: Environmental conditions in the experiments.

Exp.	Treatment	Duration (°Cd)	Daily mean temperature (°C)	Daily mean PAR (mol.m ⁻² .d ⁻¹)	Daily mean FTSW
1	T	1158	24.4	53.5	-
1	TG	1158	24.4	53.5	-
2	T	1128	22.6	43.6	-
3	T	1077	21.2	12.6	-
4	T	1210	20.1	6.3	-
5	T	1166	22.4	12.6 then 0	-
6 [2]	T	803	23.4	54.8	0.74
6 [2]	T	803	23.4	54.8	0.31
6 [2]	T	803	23.4	54.8	0.18

The number of leaves on each axis was counted every three days. Axes were distinguished according to their types (primary axis or P0-P1-P2 secondary axis). Roughly, 550, 800 and 1100°Cd after budburst, four plants, per treatment were harvested. The sizes and the dry masses of all plant organs including roots were then measured. Allometric relationships were used to estimate individual leaf areas and total leaf area of the plants.

The data of exp. 6 were extracted from Lebon et al. 2006 [2]. In this experiment, one shoot without cluster was kept on each plant. Three treatments depending on the mean values of FTSW were applied (0.78, 0.31 and 0.18), on these plants (Table 1). The water deficit was applied to the plant 380°Cd after budburst. Data about the evolution of total leaf area and of the number of leaves were available whereas no data on plant dry mass were available in this experiment.

3. Results

3.1 Assessments of model simulations

The estimated values of the different parameters were obtained from the determinist version of GreenLab by model inversion, from experimental data and from literature. Model inversion of GreenLab model was used to estimate the sink strengths and the associated beta functions for all the organs, the parameters of the relationships between Q/D and axis development and the parameters of the production equation. These parameters, largely exposed in Pallas et al. 2009 [3], were obtained on an independent set of plants. Literature was used to estimate the parameters of the phyllochron [1][8], and the parameters of secondary axis development according to soil water deficit [2]. Measurements were used to estimate the parameters of reserve behaviour and the parameters of the relationship between SLW and PAR. A unique set of constant parameters was used for all experiments, and the number of clusters on each plant was an input of the model. Only the environmental data changed among the experiments (Table 1). The obtained simulations results were then confronted to the experimental measurements.

The results presented in this section are only the results of simulations made on ‘Grenache N.’, as the results of ‘Syrah’ are quite similar (data not shown). The simulations of organogenesis processes on plants are consistent with the observed values for the experiment carried out outside, in the greenhouse (Fig. 3). The model simulates well the variations of axis development, due to the modular structure of the shoot (P0-P1-P2) and due to the differences in axis behaviour related to their insertion rank [1]. Moreover the model reproduces the decrease of the number of leaves on the secondary axes in accordance with the decrease of incident PAR in the greenhouse (Fig. 3B).

At the whole plant level the model is strongly consistent with observed values for the six experiments (Fig. 4). The variable with the lowest RRMSE is the total leaf area (12.2%) and with the greatest is total dry

mass (22.1%). Note also that the dynamics of biomass growth of clusters (Fig. 4D) is quite similar with others observed dynamic in previous studies [3].

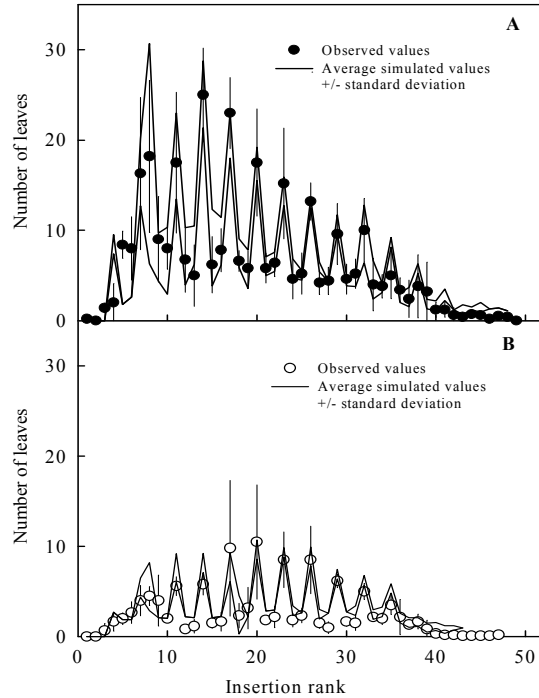


Figure 3. Observed and simulated values of the final number of leaves on each secondary axis according to their insertion rank on the primary axis. Simulation results in exp. 1 with the plants of TG treatment (A). Simulation results in exp. 3 (B). Simulation standard deviations were calculated with 50 repetitions of the adjustment procedure. Bars indicate the standard deviation of the observed values.

3.2 Evolution of the trophic competition in the various experiments.

The evolution of the Q/D ratio and its absolute values are consistent with the available data previously obtained by GreenLab model inversion on grapevine [3] (Fig. 5).

In outside experiments for plants without clusters (T) (Fig. 5A, exps. 1&2), after a first decrease at the beginning of shoot development, the Q/D ratio tends to increase due to the increase of the number of autotrophic organs. Then, Q/D decreases in relation with the increase of the number of developing axes and of internode secondary growth [3].

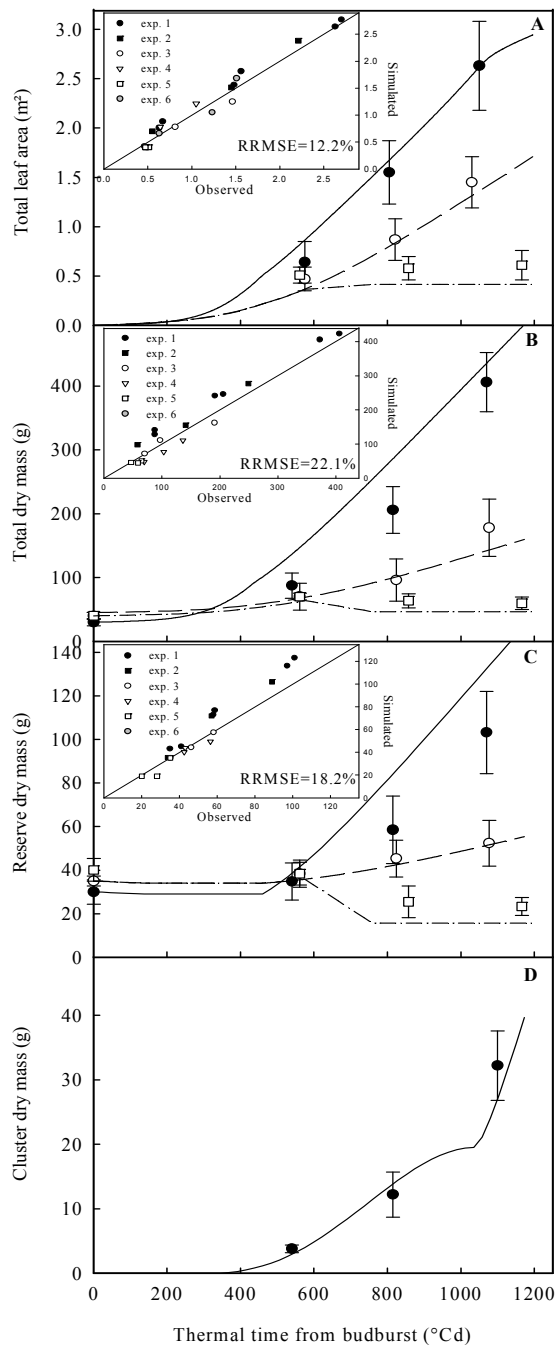


Figure 4. Evolution of total leaf area (A), total dry mass (B), reserve dry mass (C) and clusters dry mass (D) for the plants of exp. 1 (TG) of exp. 3 and 5. The symbols represent the observed values and the solid or dashed lines the simulated values. Bars indicate the standard deviation of the observed values. The inserts represent the comparison between the observed and the simulated values for the whole dataset.

The decrease of incident PAR in the greenhouse (Fig. 5A, exps. 3, 4&5) tends logically to decrease the

Q/D ratio, which appears to stabilize, with no other observed decrease after 400°Cd because the number of developing axes and secondary growth are limited. The presence of clusters tends to decrease the Q/D ratio particularly after 1100°Cd (veraison stage, Fig. 5B) corresponding to the phase of the greatest increase of cluster growth rate [12] whereas the effect of cluster development before this date is really weak.

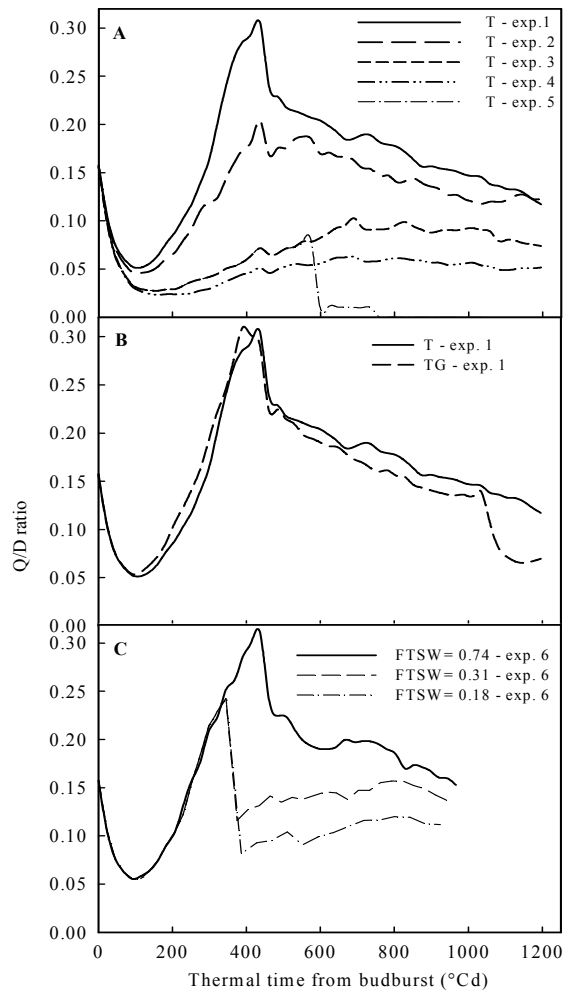


Figure 5. Evolutions of the Q/D ratio according to thermal time. in exps. 1 to 5 with T treatments (A), in exp. 1 with T and TG treatments (B) and in 6 exp. 6 (C).

Soil water deficit also decreases the values of Q/D , but no decrease of Q/D ratio is observed after 450°Cd contrary to what is observed for well-watered plants (Fig 5C). Even a slight increase in Q/D ratio is observed after 450°Cd for the plants in soil water deficit. This observation is consistent, with a decrease of the competition among the vegetative sinks in situation of water deficit [2].

3.3 Sensitivity analysis

The model was extensively tested for its sensitivity to input variables, in order to explore its behaviour and evaluate the biological coherence of its response. In the present study, one sensitivity analysis illustrates the impact of FTSW (Fig. 6) variations on the model outputs. Concerning the effect of FTSW, the model reproduces an observed pattern with potential repercussions on agronomic practices. Cluster dry mass appears to be the least affected by soil water deficit. A significant decrease of cluster dry mass is only observed for FTSW values lower than 0.35, whereas for the other variables, this decrease is observed for values lower than 0.60 [15]. This observation probably results from the decrease of the number of vegetative 'sinks' due to the signalic regulation of vegetative development, which leads to promote cluster development in situations of moderate soil water deficit.

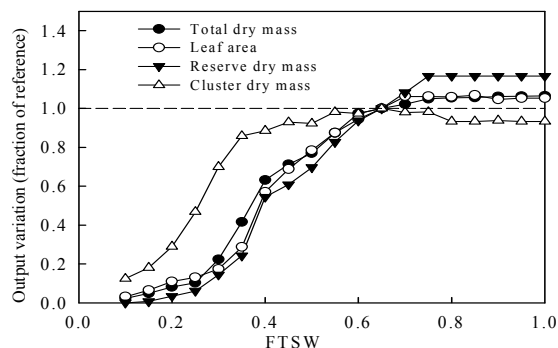


Figure 6. Sensitivity analysis of various model outputs to different values of FTSW. All model outputs were normalized according to the values observed at FTSW=0.7, 1200°Cd after budburst with PAR=40mol.m⁻².d⁻¹.

4. Conclusions-perspectives

This study is a first attempt to simulate real plant development using the principles of retroaction between plant organogenesis, morphogenesis and trophic competition developed with the GreenLab model [7][13]. The Q/D ratio appears as a relevant variable to drive plant development in fluctuating environment, including soil water deficit. This new version of the model using the main principles of GreenLab model with stochastic rules of axis development according to trophic competition appears promising to further simulate plant population development in various environments.

5. References

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