

Assembling and testing a generic phenological model to predict Lobesia botrana voltinism for impact studies.

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Victorine Castex, Iñaki García de Cortázar-Atauri, Pierluigi Calanca, Martin Beniston, Jérôme Moreau. Assembling and testing a generic phenological model to predict Lobesia botrana voltinism for impact studies.. Ecological Modelling, 2020, 420, pp.108946. 10.1016/j.ecolmodel.2020.108946. hal-02541785

HAL Id: hal-02541785 https://hal.science/hal-02541785v1

Submitted on 22 Feb 2024

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1 **Ecological Modelling** 2 3 Title: Assembling and testing a generic phenological model to predict Lobesia Botrana voltinism 4 for impact studies 5 6 V. Castex (1), I. García de Cortázar-Atauri (2), P. Calanca (3), M. Beniston (1), J. Moreau(4). 7 (1) Institute Environmental Sciences, of Switzerland, of University Geneva, 8 Martin.Beniston@unige.ch 9 (2) INRA, US1116 AgroClim, 84914 Avignon, France, inaki.garciadecortazar@inra.fr 10 (3) Agroscope, Agroecology and Environment, Switzerland, pierluigi.calanca@agroscope.admin.ch 11 (4) Université de Bourgogne Franche-Comté, UMR 6282 Biogéosciences, Equipe Ecologie-Evolutive, 12 France, Jerome.Moreau@u-bourgogne.fr 13 14 **Informative title :** A generic model for *L.botrana* voltinism 15 16 **Corresponding author:** Victorine Castex, victorine.castex@unige.ch 17 University of Geneva - ISE 18 Boulevard Carl-Vogt 66 19 1205 Geneva, Switzerland 20 Tel: (+41) 22 379 0768 21 22 23 24 25 26 27

Abstract:

The physiological development of insect pests is driven by temperature and photoperiod. Geographic variations in the speed of growth reflect current patterns in thermal conditions as a function of latitude and altitude. Global warming will likely lead to shifts in pests' phenology. Insects are expected to overwinter earlier and develop more generations, with implications for the risks of damage to agricultural crops. Understanding and monitoring of the voltinism of insect pests will be increasingly important to anticipate critical phases of pest development and devise options for adapting pest control measures. In this study, we describe a new generic phenological model that allows to reproduce the voltinism of *Lobesia botrana*, a major insect pest that affect grapevine production (*Vitis vinifera* L.) worldwide.

Inspired by existing models, the Lobesia Generic Model (LGM) combines two submodels representing the overwintering period and the dates of flight of subsequent generations of male insects. This model uses a unique Beta function for representing the temperature responses of both the overwintering generation as well as the adult generations. The results indicate that LGM is able to simulate *L. botrana* voltinism under climatic conditions ranging from those observed in Southern Spain to those recorded in the Alsace region (France). The LGM only requires a single set of parameters valid for all the generations with a precision of around 7 days (RMSE), allowing a simple use with parameters representative of natural behaviour and found in the existing literature.

Keywords: Generic phenological modelling, *Lobesia botrana*, voltinism, process-based model, temperature, photoperiod

Key Messages

- We present a generic model of *L. botrana* voltinism that assembles existing components
- We assume that thermal requirements are the same for all generations, and use one the same
 Beta function to represent the temperature response of all generations
 - We show that the model performs well across sites ranging from Southern Spain to the Alsace region (France)

1. Introduction

Modern agriculture relies on the massive use of pesticides to increase productivity (Tilman et al. 2002). This has caused considerable stress on biodiversity and made insect pests more resistant to pesticides (Hatt et al. 2018). Furthermore, extensive commercial trade as well as increasing

temperatures have enhanced the spread of insect pests. As a result, invasive species have started to damage crops in remote areas (Bale et al. 2002; Lin 2011; Lamichhane et al. 2015). Understanding and predicting insect pest dynamics is necessary in order to ensure productivity while reducing the use of chemicals (Bregaglio et al. 2013). In the context of global warming, there is also a need to understand how pest species dynamics will evolve in the future. In this context, phenological models are interesting tools to monitor populations dynamics and to predict their evolution, e.g., under changing environmental and climatic conditions (Steinbauer et al. 2004; Nietschke et al. 2007; Stoeckli et al. 2012). Such models are capable of anticipating insects' voltinism and allows improving treatment periods (Moravie et al. 2006).

Process-based models aim to represent biological processes through equations that describe developmental, survival, fecundity or mortality rates for different life-stages of insects (from prediapause to overwintering). Among the different processes simulated, phenology (development) is probably one of the most important, because it defines the biological cycle (Chuine et al. 2013). Insects are ectotherms and consequently their physiological development is mainly driven by temperature (Zhou et al. 1995; Bale et al. 2002; Tobin et al. 2008). They exhibit a response to thermal variation (Singer and Parmesan 2010; Bale and Hayward 2010; Brodeur et al. 2013; Moiroux et al. 2014) by adjusting life history traits like the date of overwintering and population growth (Hance et al. 2007; Moreau et al. 2017). Diapause and overwintering are further determined by photoperiod (Pavan et al. 2006; Svobodová et al. 2014). Both temperature and photoperiod thus explain how insects' life cycle respond to local climatic conditions, latitude and altitude (Honêk 1996; Ioriatti et al. 2011). Changes in the growing rate and geographical distribution of pests have already been observed in the last decades (Baumgärtner et al. 2012) as a consequence of recent changes in climate.

Among the numerous species that affect agriculture, *Lobesia botrana* (Den. & Schiff) commonly named the European grapevine moth, is a good candidate for assessing effects of climate change on the development of insect pests. It is a major grapevine insect pest that is present worldwide, including Europe. As a multivoltine species, this pest is able to complete from two (in northern parts of Europe) to four (in southern parts of Europe) generations per year according to climate and latitude (Martín-Vertedor et al. 2010). Under warmer conditions it has the potential of generating more damage than currently, as its development will begin earlier in the season and last longer (Castex et al. 2017; Marchesini and Monta 2004). The life cycle of *L. botrana* is characterized by its capacity to enter into diapause at the pupal phase when day length is less than 14 hours (12 hours is the limit for pupal overwintering). The individuals will overwinter as pupa and start developing before becoming adults and mate to start their adult life and reproduction cycle (Thiery, 2008).

Many models have been developed to monitor the adult flights of *L. botrana* (Tab.1). They all can be considered as process-based models, despite using different temperature response functions (Caffarra et al. 2012; Caffarelli and Vita 1988; Gabel and Mocko 1984; Gallardo et al. 2009; Logan et al. 1976; Milonas et al. 2001; Touzeau 1981). Other models adopt a statistical approach to represent biological responses to temperature (Moravie et al. 2006).

Existing models for *L. botrana* target different purposes. Some of them describe the growing length at different developmental phases (Touzeau 1981; Gabel and Mocko 1984; Caffarelli and Vita 1988; Briere and Pracros 1998 and 1999; Severini et al. 2005). Others focus on the timing of maximum moth flight (Moravie et al. 2006; Amo-Salas et al. 2011; Ortega-López et al. 2014), forecast the emergence timing (Arca et al. 1993; Milonas et al. 2001) and population size (Gallardo et al. 2009), the number of generations occurring before diapause (Caffarra et al. 2012) or model the overwintering process (Baumgärtner et al. 2012). Finally, very complex models such as that of Gutierrez et al. (2012) try to simulate various processes simultaneously (grapevine phenology, adult development, survival and fecundity rates).

Tab.1: Comparison between local models and the generic model described in this work. The following exhaustive literature review list the existing phenological models for *L. botrana*: (1) Gabel and Mocko 1984; (2) Caffarelli and Vita 1988; (3) Arca et al. 1993; (4) Briere and Pracros 1999; (5) Gutierrez et al. 2012; (6) Milonas et al. 2001, (7) Severini et al. 2005; (8) Moravie et al. 2006; (9) Gallardo et al. 2009; (10) Amo-Salas et al. 2011, (11) Caffarra et al. 2012; (12) Baumgärtner et al. 2012; (13) Ortega-López et al. 2014; (14) Gilioli et al. 2016

	Local models (see refs. in the legend)	Lobesia Generic Model described in this study
Input parameters and processes	-Temperature: mean, minimum, maximum, daily, hourly (1; 2; 3; 4; 9; 11; 12) -Relative humidity (10; 13) -Photoperiod (1; 4; 5; 12) -Mortality and growth rates (7)	-Temperature (mean daily) -Photoperiod
Method of calculation	-Accumulation of growing degree-days - Different thermal time requirements according to the generations (1;2;3;6;9;11)	-Accumulation of growing degree-days -Same thermal time requirements to achieve subsequent phases within one generation and for all the generations
Phases modeled	-Prediapause/ Diapause/ Post-diapause (5; 7; 12; 14) -Overwintering (12) -Egg, larval and pupa phases (3; 5; 10; 11) -1 st (8; 13), 2 nd (6) and 3rd (9) adults flights	-Prediapause/ Diapause/ Post-diapause (overwintering) -Adult flights (10% and 50% of males flights)
Beginning of calculation	-1st January (8; 10; 11; 13) -1st March (6; 9) or 5th March (1) -After diapause induction (DL<14.15h) (5) -Day of overwintering (previous year) (7) -Day of the 1st individual to enter in diapause (12) -End previous generation (chrysalides) (3)	- (d _{t0})= -183 (July 15th) corresponding to the day with > 12h of day length in Spain (the most southern site of observed data). The LGM start calculating in late summer (i.e. of the previous year n-1), so for the prediapause.

Type of model	-Population / Rate sum model (6; 9; 12) -Empirical / Biologically Based model (1; 3; 4; 6; 10; 11; 13) -Physiologically Based Demographic Model (5; 14) -Aged structure (7)	- Voltinism Process Based Model with a statistical approach
Strength	Perform well under same conditions as considered during model development Easy to implement if local data are available Suitable for integration into Decision Support Tools	- 1 set of parameters - Suitable for large scale impact studies as climate change scenarios
Weakness	 Not necessarily reproducible at other sites with the same parameters (applicability to new situations needs to be verified) Different sets of parameters for each region, generation and between the phases (based on the analysis of the different models, ESM7). Local models are not applicable to large scale studies 	-Predictions across sites are robust even if performance quality can be low in specific cases

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Irrespective of type, structure and complexity, most of the models given in Tab.1 were developed to (a) operate at the local scale, implying that they require local sets of parameters (e.g.,: Trentino, Italian Alps, in Caffarra and Eccel 2011), and (b) to account for possible differences in thermal requirements between different generations, meaning that parameters are usually specified for each generation separately (Touzeau 1981; Gabel and Mocko 1984; Caffarelli and Vita 1988; Milonas et al. 2001; Gallardo et al. 2009; Caffarra et al. 2012). For this reason, some authors have already highlighted the difficulty of using and comparing these models within a larger range of climatic conditions than initially targeted (Flores and Azin 2015; Allen 1976; Gallardo et al. 2009; Amo-Salas et al. 2011). In fact, a major downside with local models is that they need local adjustments when applied outside the geographic area targeted in first place. To the best of our knowledge, only a few models, such as the one developed by Baumgärtner et al. (2012) or Ortega-López et al. (2014) have been applied to large geographical scales.

The overall purpose of this work is to present a generic model for L. botrana (hereafter called Lobesia Generic Model or LGM) that considers a large scale of application and a single response to temperature or thermal time requirements for all generations (Tab.1). The model is built by assembling existing model components and calibrated in such a way as to obtain a generic set of parameters. Model development and calibration were carried out using PMP, a user-friendly software providing an interface for designing, setting up, testing and running phenological models (Chuine et al. 2013).

The specific aims of this study are 1) to analyze existing models of L. botrana voltinism in order to propose a generic approach for simulating representative phase of development (10% and 50% of the male flights) of an adult population from diapause to last flight; and 2) to propose a validated standard set of parameters for applications at the large scale. We use for this purpose a large database of flight observations compiled in a wide latitudinal transect running from southern Spain to northern France (Fig.1) and representing Mediterranean and Continental climate types (Jongman et al. 2006; Beck et al. 2018).

2. Materials and Methods

2.1 Data used in the study

153 2.1.1 Phenological data of L. botrana

The observed data of *L. botrana* correspond to the number of individuals captured per week, by pheromone (sexual confusion) delta traps placed at the beginning of March. Traps are changed and counted weekly to represent the distribution of population from the first flight of the first generation (who overwintered) to the last adult flight of the last generation. Traps are considered one of the best way to capture and represent male flight activity (Milonas et al. 2001; Gallardo et al. 2009; Ortega-López et al. 2014). Data were provided by private sector enterprises, inter professional councils in the agronomical and viticulture sectors and academic institutions (Tab.2). As illustrated in the map (Fig.1), 12 sites were used to cover latitudes between Lat.36.710°N and Lat.48.213°N and an altitude range from 0 m a.s.l. (Saint Gilles, France) and 692 m a.s.l. (Requena, Spain). We have assumed that it was possible to combine all the data from different sites to calibrate and evaluate the model, even if there is no standardized protocol for the observation of adult flights in trap catches.

Tab.2. *L. botrana* adult flights data and meteorological data acquisition per site in latitudinal order.

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Lobesia Botrana **Meteorological station** Altitude Site Latitude Longitude Site (data provider) Years obs Data provider Period Number (m a.s.l) (°N) (°E) Sanlucar (Junta de 2008, 2009, 1 Jerez de la 2007-Andalucia) 2015 frontera (Junta de -6.1 56 36.7 2015 Trebujena (Junta de Andalucia) 2 2013, 2014 Andalucia) Yecla_C_abajo (O. 2010, 2011, Yecla-Pinillos 2007-38.6 3 comarcal Agraria de 2012, 2014, -1.1 602 2017 (SIAM) Alahama) 2015, 2016 Fuente de la 2012, 2013, Higuera Alforins 2010-4 2014, 2015, Caudete (SIAR) 38.7 -1.0 625 (Generalitat 2017 2016 Valencia) Requena Las Monjas 2012, 2013, Requena Cerrito -2010-5 (Generalitat 39.5 2014, 2015, -1.11 692 Arcis (SIAR) 2017 Valencia) 2016 Villar del Arzobispo 2012, 2013, Chullilla/ Villar 2010-(Generalitat 2014, 2015, 39.3 380 6 (SIAR Riegos -0.8 2017 Valencia) 2016 IVIA) Cariñena (D.O.P-2011, 2012, Calatayud 2004-7 41.3 -1.6 591 2013, 2014 2016 Vino las piedras) (Sarga) Avignon (Chambre Avignon (INRA 2003-8 2005, 2009 43.9 4.8 24 2016 Agri. Gard - France) Climatik) St-Gilles (INRA St Gilles (Chambre 2010-9 2015, 2016 43.7 4.4 0 Agri. Gard - France) Climatik) 2017 1997, 1998, Changins Changins 1997-10 (Agroscope) 1999, 2000, (Meteosuisse 455 46.4 6.2 2017 Switzerland 2001, 2010 IDAWEB) Eibindinger 2009, 2010, Staefa 2004-11 (Agroscope) 2011, 2012, (agrometeo 47.2 8.7 408 2017 Switzerland 2013, 2014 Meteosuisse) 2010, 2011, Scherwiller (Fredon 2009-Bergheim 48.2 7.4 200 12 Chambre Agri. 2012, 2013, (INRA-Climatik) 2016 Alsace - France) 2016

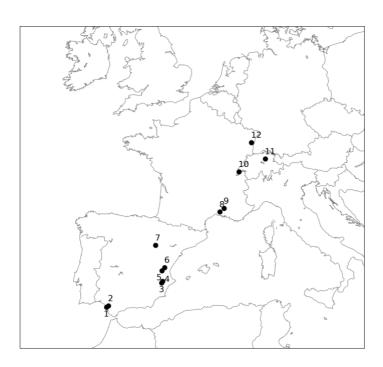


Fig.1. Map of the study sites in Europe (produced with MakeMap from an open source code). The numbers in the map represent the study sites introduced in Tab.2.

Of the initially 90 records, 51 records of (non consecutive) observations of *L. botrana* adult flight (entire cycle of generations) on an annual basis (Tab.2) were retained after quality control. From those annual male flights, we identified two main phenological observations: the 10% and 50% of the total adult flights per generation as done in Gallardo et al. (2009). To do so, we identified and separated the generations of adults as explained in ESM1 (Electronic Supplementary Material). Then, we calculated the cumulated sum of the number of individuals caught in the traps, for each generation (G1; G2; G3; G4). From this sum, we took the 10% and 50% of the cumulated sum per generation (G1-10%, G1-50%; G2-10%, G2-50%; etc.). This data was then related to its corresponding Day Of Year (DOY) of observation. The observed data are not continuous in time (from 1997 to 2016), because observations are not made systematically. Also, we only used the observed data when we could clearly identify the beginning, the peak and the end of the generations that represent around 70% of all the data we collected.

2.1.2 Meteorological data

Meteorological data were obtained from different meteorological services at each site (Tab.2). Daily minimum, maximum and mean temperatures were collected from meteorological stations in an array with a maximum distance of 10-15 km from the collection sites of *L. botrana* flights, and almost at the same altitude as the vineyards, which we considered reasonable for the representation of local meteorological conditions.

2.2. A generic phenological model for L. botrana

2.2.1 General description of the model

The LGM is a combination of two submodels allowing the simulation of at least five phases of development representing two development stages (Fig.2): 1) The overwintering submodel: It considers three phases, prediapause, diapause and post diapause from the year n-1 to n; 2) The male adult flights submodel: reproduce the 10% and 50% of the total of the population of each generation for the year n. At least two generations can be reproduced, with an option for additional generations depending on local climatic conditions and latitude.

Phase is a specific part in a physiological development process (e.g., in the overwintering stage, we identify the prediapause phase).

Baumgärtner et al. (2012) propose a complex model to simulate the overwintering stage and the adult development (at different phases) considering the speed of growth. In this study, we will only consider the first cohort (i.e., the emergence of the first individuals) of each generation (corresponding to our 10% of flight in Fig.2) considering the accumulation of thermal time.

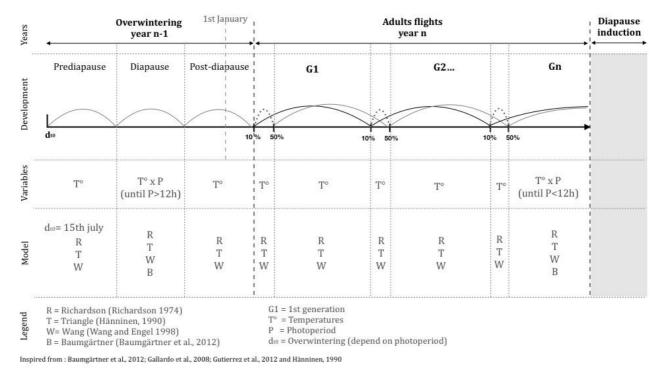


Fig. 2. LGM conceptual modelling framework. This aims at reproducing the overwintering period and the adults

flights: 10%-10% and 50%-50% of flight between generations and the 10% and 50% within a generation itself (doted line). Also the variables used (T°=Temperature and P=Photoperiod) or the functions used in the model representing the temperature response: Richardson (R), Triangle (T) and Wang (W) are represented for each physiological developmental phase of *L. botrana*.

- The overwintering of *L. botrana* is simulated following Baumgärtner et al. (2012) and Gutierrez et al. (2012) i.e. considering three phases.
- 216 1- Prediapause: development depend on temperature
- 2- Diapause: development depend on a combination of temperature and photoperiod
- 218 3- Post-diapause: development depend on temperature

- The adults flights consider two or more adults generations and depend on temperature and local climatic conditions. We propose to simulate the 10% and 50% of the total of the population of each generation. The calculation considers the adults that enter into diapause in the previous year (n-1).
- We apply the following equation to describe the temperature effect (eq. 1):

$$F_s(t_d) = \sum_{d_0}^{d_s} f(t_d) \qquad eq. 1$$

where t_d is the daily temperature, f(t) is the function describing the temperature response (described later), d_0 is the first date of the developmental phase, d_s the day of the end of the developmental phase and $F_s(t_d)$ represents the cumulated temperatures (forcing units).

According to Baumgärtner et al. (2012), d_0 (first date of the developmental phase) depends on the latitude and the photoperiod. The analysis of the equations and the parameters proposed by Baumgärtner et al. (2012) to simulate the end of diapause, allowed us to propose a simplification of the original equations (described in ESM2) using daily values and considering a photoperiod threshold set at 12h to stop the development of the last generation. The final proposed equations can be described as following:

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$$F(t_d) = \sum_{d_0}^{d_s} (f(t_d) * g(P_d)) \text{ or } F(t_d) = \sum_{d_0}^{d_s} (f(t_d) + g(P_d))$$
 eq. 2

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$$g(Pt_d) = \begin{cases} 0 \\ \frac{P_d - Plow}{Phigh - Plow} & P_d \leq Plow \\ 1 & Plow < P_d < Phigh \end{cases} eq. 3$$

where t_d , f(), d_0 , d_s and $F_s(t_d)$ represent the same variables and parameters as in (eq. 1), and g() is the function describing the photoperiod response and P_d is the daily photoperiod. The photoperiod function is varying between a low and a high threshold (Plow and Phigh respectively). We explored two possibilities of combination of the temperature and photoperiod effect: the multiplicative (usually used in other models as Wang and Engel (1998)) or the additive (as proposed by Baumgärtner et al. (2012))

2.2.2 Temperature response functions

We tested three different functions to describe the response of *L. botrana* development to temperature (Fig.3):

1) The Richardson model (R) (Richardson et al. 1974) is a step function that assumes a linear response of development to daily temperatures for temperatures in the range Tlow to Thigh:

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$$f(t_d) = \begin{cases} 0, & t_d \leq Tlow \\ t_d - Tlow, & Tlow < t_d < Thigh \\ Thigh - Tlow, & t_d \geq Thigh \end{cases} eq. 4$$

Such approaches have been used by Arca et al. (1993) to calculate the emergence of adults and larvae of *L. botrana*.

2) The Triangular model (T) (Hänninen 1990) assumes a linear, positive response to daily temperatures for t_d in the range Tmin to Topt, and an linear but negative response for t_d in the range Topt to Tmax (Maiorano et al. 2012):

$$f(t_d) = \begin{cases} 0, & t_d \leq Tmin \\ \frac{t_d - Tmin}{Topt - Tmin}, & Tmin < t_d \leq Topt \\ \frac{t_d - Tmax}{Topt - Tmax}, & Topt < t_d < Tmax \\ 0. & t_d \geq Tmax \end{cases} eq. 5$$

3) The Wang model (W) (Wang and Engel 1998) is a Beta function that considers three cardinal temperatures Tmin, Tmax and Topt:

$$f(t_d) = \begin{cases} 0, & t_d \leq Tmin \\ \frac{2(t_d - Tmin)^{\alpha}(t_d - Tmin)^{2\alpha}}{(Topt - Tmin)^{2\alpha}}, & Tmin < t_d < Tmax \\ 0, & t_d \geq Tmax \end{cases} eq. 6$$

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$$\propto = \ln(2)/\ln(\frac{Tmax-Tmin}{Topt-Tmin})$$
 eq. 7

This function is very similar to other functions described in the literature to simulate *L. botrana* response to the temperature (Wang and Engel 1998) and have been applied by Milonas et al. (2001), Briere and Pracros (1998), Logan et al. (1976) and Baumgärtner et al. (2012) for *L. botrana*.



Fig. 3. Comparison of three different approaches (Richardson, Triangle or Wang) of response to temperature with model parameters adapted to *L. botrana*. The three representations use the same parameters with $T_{min}=5^{\circ}C$, $T_{opt}=20^{\circ}C$ and $T_{max}=30^{\circ}C$, except for Richardson that only uses $T_{low}=5^{\circ}C$ and $T_{high}=20^{\circ}C$.

270 2.2.3 Modelling parameterization and evaluation framework

Overwintering submodel

- We assume that the overwintering starts (d_{t0}) when day length is more than 12 hours (Roditakis and
- Karandinos 2001; Baumgärtner et al. 2012). Individuals that enter into prediapause in the year n
- 274 (autumn) will overwinter (spring) in the following year. Diapause induction starts when the day length
- falls below <14 hours, with a critical threshold at <12hours (Fig.2), According to this assumption and
- 276 (eq.1, 2 and 3) of Baumgärtner et al. (2012), it is possible to define when the first cohort begins its
- diapause.

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- The diapause is a combination of temperature and photoperiod. The DOY of diapause induction was
- evaluated according to (eq.3) in Baumgärtner et al. (2012) calculating the day-length as a function of
- 280 the DOY and the latitude. In spite of the fact that (eq.3) predicts a critical day-length threshold
- increasing linearly with latitude, the DOY allowing diapause induction does not vary significantly
- with latitude, ranging from DOY 183 to DOY 193 across our sites. For this reason, we tested the
- sensibility of the model to different starting date (d_{t0}), July 1st for the southernest site and the July 15th
- as an average date for all the sites.
- The post-diapause depends on temperature to allow the emergence of the first generation of adults.
- The cumulation of thermal accumulation from the prediapause to the post-diapause determines the
- first adult flights. This first sub model (overwintering) was necessary to determine the start of the
- calculation of the 10% of flight of the adults from the 1st generation (G1-10%).

Submodel for the adults flights

- In order to simulate the 10% and the 50% of each generation, we made the three following assumptions:
 - 1. eggs laid by G1-10% (or G1-50%) adult will reach the adult stage at G2-10% (or G2-50%) and this hypothesis is valid for all the following generations (Gn-10% or Gn-50%);
 - 2. the same functions and set of parameters can be used to simulate the thermal time accumulation between G1-10% and G2-10% (or G(n-1)-10% and Gn-10%), G(n-1)-50% and Gn-50%, as well as Gn-10% and Gn-50%;
 - 3. the same set of parameters can be used for all sites (and generations) without significant repercussions on the quality of the simulations (if compared to results obtained with site specific parameters) (Parker et al. 2011).

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- We calibrated and tested the three models Richardson (R), Triangle (T) and Wang (W) for the 10%-
- 303 10% and 50%-50% of adult flights between the generations and the 10%-50% of adult flights within
- 304 the generation itself. Preliminary simulations were made with all parameters, called Allp, meaning that
- the model calculates alone without any restriction of parameters (ESM8).

At the same time we also tested different sets of parameters (Tmin=0°C, 5°C, 7°C; Tmax=28°C,

33°C) Gutierrez's (2012) parameters (Tmin=8.9°C, Topt=30.8°C and Tmax=33°C) and some

Baumgärtner's (2012) parameters (Tmin=4.5°C) (ESM6, 7 and 8).

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2.2.4 Calibration of the model

- 311 For the purpose of calibration and testing, we split the original data into a calibration dataset (75% of
- 312 the total sample selected randomly CAL dataset) and a validation dataset (25% of the data VAL
- dataset) (Picard and Cook 1984). The calibration of the parameters for the overwintering stage and the
- adult generations were made separately. Once the models and parameters were determined, we joined
- 315 the two submodels to run the simulations with our calibration data set and validate it with the
- 316 validation dataset. Finally in order to provide the best set of parameters, we recalibrated the entire
- 317 model using all the database (ALL dataset) using a cross validation method (Cleland et al. 2007). The
- 318 best parameters were selected according to the results of the statistical criteria described in Tab.3. The
- performance criteria were defined using the variance global of the observed data (STOT), the Sum of
- 320 Squares Residuals (SRES), the Root Mean Square Error (RMSE), the efficiency of the model (EFF)
- and the Akaike test (AIC). For more details on the description and the formulas, see the ESM3 and 8.

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323 *2.2.5 Software*

- 324 To facilitate the development and the parameterization of this phenological model, the PMP software
- 325 (Phenology Modelling Platform PMP 5.5) has been used (downloaded from:
- 326 http://www.cefe.cnrs.fr/en/logiciels/ressources-documentaires). PMP allows constructing, fitting and
- running simulations of a phenological model using meteorological data and phenological observations
- 328 (Chuine et al. 2013).

329

330 **3. Results**

- In this section we describe the results obtained from the calibration and the validation of both sub-
- 332 models: 1) the first sub model describing the overwintering period; 2) the second sub-model which
- simulates the 10-10%, 50-50% and 10-50% of the different generations of adults.

334335

336

3.1 Overwintering submodel

Model three phases testing R, T and W models with CAL, VAL and ALL data set

- 337 As described previously we tested different versions of the submodel for the three phases of
- overwintering (Fig.2): 1) different functions for temperature responses (R,T,W); 2) different starting
- dates of prediapause (d_{t0}) (July 1st and July 15th), 3) different ways to combine temperature (T) and
- 340 photoperiod (P) responses during the diapause period, using an additive (Sum) and multiplicative
- (Mult) factor. The best models obtained used the T and W functions, starting on 15th July (d₁₀) and
- multiplying the temperature and photoperiod effects. Tab.3 describes only results obtained with CAL

data set with a start date (d_{t0}) on July 15th and using a multiplicative (Mult) method for combining photoperiod and temperature (for further results concerning all dates and methods, see ESM4). The best results were obtained with the Triangle and Wang approaches showing EFF values of 0.77 and 0.74 respectively (Tab.3). The combination method of the functions had more impact on the results than the starting dates named (d_{t0}). Moreover, the date of entry in prediapause was coherent with Baumgärnter's values (Baumgärtner et al. 2012).

Tab.3. Performance statistics with the CAL data set for three different choices of the temperature response function and the model version that combines temperature and photoperiod responses as multiplicative factors. Results presented here refer to simulation with parameter values taken form the literature. The statistical analysis use the RMSE (Root Mean Square Error) EFF (Efficiency of the model) and AIC (Akaike test).

Ctatistics

	Simulations		3	Statistics	
Function	(d _{t0})	Nb data	RMSE	EFF	AIC
R	July 15th	39	9.47	0.58	98.71
Т	July 15th	39	7.02	0.77	118.28
W	July 15th	39	7.49	0.74	123.22

Cimeriations

Tab.4. Comparison of the best fitting of T (Triangle) and W (Wang) simulations to represent the overwintering with CAL data and using a multiplicative factor. Parameters in **bold** are fixed parameters. Parameters with an * are from the literature (Baumgärtner, et al. 2012; Brière et al., 1999; Gutierrez et al. 2012). The other one are free. The models 1 and 2 have free parameters. The models 3 and 4 have the parameters fixed from the models 1 and 2 and a fixed Tmin at 4.5°C as in Baumgärtner (2012) or BAUM. The models 5 and 6 consider Gutierrez's parameters (GUT) with free day length (DL). The models 7 and 8 have fixed parameters from GUT but the DL and Tmin for the 3rd phase of GUT are free. Gutierrez's or GUT parameters are Tmin=8.9°C, Topt=30.8°C and Tmax=33°C.

Model		TAllp (1)	WAllp (2)	TAllp_Tmin 4.5(BAUM) (3)	WAllpT_min 4.5(BAUM) (4)	T GUT (5)	W GUT (6)	T Mult (7)	W Mult (8)
Over- wintering phases	Parameter	All free	All free	ParTAllp +Tmin= 4.5(BAUM)	ParWAllp +Tmin= 4.5 (BAUM)	GUT+Tm in= 4.5 (BAUM)+ DL Free	GUT+Tm in= 4.5 (BAUM)+ DL Free	GUT+ (Tmin & DL Free)	GUT+ (Tmin & DL Free)
	(d_{t0})	July 15th	July 15th	July 15th	July 15th	July 15th	July 15th	July 15th	July 15th
1st phase	Tmin	4.94	2.45	4.94	2.45	8.9*	8.9*	8.9*	8.9*
Prediapa use	Topt	18.76	14.36	18.76	14.36	30.8*	30.8*	30.8*	30.8*
	Tmax	28.19	27.69	28.19	27.69	33*	33*	33*	33*
	Tmin	5.29	7.66	5.29	7.66	7.1*	7.1*	7.1*	7.1*
2nd	Topt	16.23	15.04	16.23	15.04	24.1*	24.1*	24.1*	24.1*
phase	Tmax	21.89	22.87	21.89	22.87	28.5*	28.5*	28.5*	28.5*
Diapause	Plow	9.35	9.17	9.35	9.17	12.12	8.78	12.13	3.58
	Popt	13.05	14.25	13.05	14.25	16.23	13.04	16.46	7.72
3rd phase	Tmin	10.47	11.39	4.5*	4.5*	4.5*	4.5*	4.4	16.18
Post-	Topt	10.48	11.42	10.48	11.42	30.8*	30.8*	30.8*	30.8*
diapause	Tmax	20	20	20	20	33*	33*	33*	33*
Statistics	EFF	0.77	0.74	0.67	0.69	0.30	-29.81	0.30	-5.92
Statistics	AIC	118.28	123.22	161.80	158.68	185.79	329.76	183.50	270.79

To evaluate parameter sensitivity, we compare the two best models (models 1 and 2 in Tab.4), with various models using the same functions but fixing some of their parameters from the literature (Tab.4). Because Tmin and Topt values for the post-diapause phase were close to one another, we explored the impact of fixing Tmin value to 4.5°C according to Baumgärtner et al.(2012) keeping all the other parameters from the best models. Results obtained for both models (models 3 and 4 in Tab.4) show that this parameter does not degrade the quality substantially (values of efficiency of 0.67 to 0.69).

In order to explore other values of parameters from the literature, we tested Gutierrez et al. (2012) parameters (GUT parameters are Tmin=8.9°C, Topt=30.8°C and Tmax=33°C) for all the temperature thresholds with free day length (DL) and limiting the post-diapause with BAUM Tmin=4.5°C (models 5 and 6 in Tab.4), and with the Tmin free for the post-diapause phase (models 7 and 8 in Tab.4). Results obtained show that these parameters combination decrease significantly the quality of the model, having a higher effect when using the W function.

Tab.5. Validation of the overwintering models ford both Triangle and Wang models with (d_{t0}) =15th July. Best fitting for Validation tests (VAL) were made with a part of the calibration (CAL) data set. The entire data set was then tested with the ALL the data set (CAL+VAL) using cross validation. The statistical analysis use the STOT (Total sum of the data squared) RMSE (Root Mean Square Error) and EFF (efficiency of the model) (ESM3).

		STASTISTICS							
Model	Data set used	SRES	STOT	RMSE	EFF	Nb Obs			
	CAL	1874.23	8067.08	7.02	0.77	39			
T Allp	VAL	969.45	1072.69	8.99	0.1	12			
	ALL	3839.42	9410.59	8.68	0.59	51			
Model	Data set used	SRES	STOT	RMSE	EFF	Nb Obs			
	CAL	2134.56	8067.08	7.49	0.74	39			
W Allp	VAL	1293.39	1072.69	9.07	0.08	12			
	ALL	2845.7	9410.59	7.47	0.7	51			

In order to provide the best set of parameters we validated the defined best models (T and W) using the validation (VAL) data set (Tab.5). In contrast to CAL (39 data points), the validation data (VAL) include only 12, randomly chosen observations. For this reason, apart for a low value of EFF (0.1 and 0.08) the performance statistics for VAL are more than satisfactory, with RMSE=8.99 and AIC=52.7. After a first screening the different model configurations, we calibrated T and W using all available data (dataset ALL). For both response functions EFF > 0.5 (0.59 for T, 0.70 for W, respectively). Satisfactory were also the values of the RMSE (8.68 for T, and 7.47 for W, respectively) (Fig.4). We also selected W to simulate the first flight in our model as it seems to be the more stable with all the simulations made, with fixed or free parameterization.

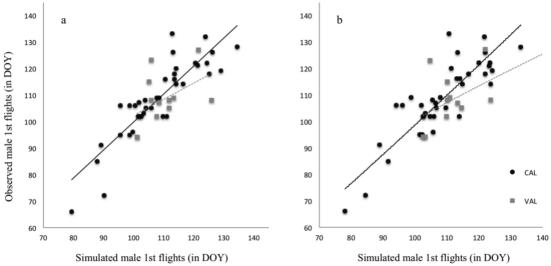


Fig.4: Representation of the day of the first flight of *L. botrana* male with a $(d_{t0})=15^{th}$ of July and a multiplicative factor: a) Triangle model b) Wang model. Simulated and observed data in DOY (Julian days) show the quality of the simulation for each data set where CAL are in black circle and the results from the validation model VAL are in grey squares.

3.2 Submodel for the adults flights

We again tested different temperature response functions (R, W, T) to evaluate their performance to simulate adult flights between two consecutive generations (10-10%; 50-50%) and within a generation itself (10-50%). Following the same methodology as for the first flight, simulations were carried out by specifying all the parameters as free parameters (to be calibrated) or fixing some of them at values obtained from the literature. Summary statistics are presented in Tab.6.

3.2.1 Simulation the 10-10%-50-50% between the generations

According to the first assumption described in the modelling framework (see Material and Methods section), we explore the capacity of the model to represent with a single set of parameters all the 10%-10% and 50%-50% generations.

Tab.6. Simulation of the model hypothesis testing the 10-10%/50-50% between the *L. botrana* generations with the calibration (CAL) and validation (VAL) data set. The statistical analysis use the RMSE (Root Mean Square Error) and EFF (efficiency of the model). GUT parameters are Tmin=8.9°C, Topt=30.8°C and Tmax=33°C.

10-10	10-10%-50-50% between generations				
	Simulation	Data set	Nb data	RMSE	EFF
R Allp		CAL	140	6.97	0.95
R ParGUT	10-10 / 50-50	CAL	140	8.35	0.93
R ParAllp (VAL)	_	VAL	46	11.00	0.84
	Simulation	Data set	Nb data	RMSE	EFF
T Allp		CAL	140	6.93	0.95
T ParGUT	10-10 / 50-50 _	CAL	140	8.32	0.93
T ParAllp (VAL)		VAL	46	11.08	0.84
	Simulation	Data set	Nb data	RMSE	EFF
W Allp		CAI	140	6.93	0.95
W ParGUT	10-10 / 50-50	CAL	140	27.96	0.25
W ParAllp (VAL)	_	VAL	46	11.15	0.84

The three models in Tab.6 had high values of EFF when all the parameters are fitted (around 0.95). Values of the EFF decreased when some of the parameters were fixed (values varying between 0.25 to 0.95 depending on the model). We also validated the best model (Allp model) with the independent VAL dataset and we obtained good efficiency values for all the models (EFF = 0.84). As for the overwintering model, we propose to exclude R from our analysis because its structure to describe the development of *L. botrana* does not integrate maximum temperature (Tmax) even if the calibration did not show a significant effect of this threshold. Moreover, as in the overwintering model, the test using Gutierrez's parameters did not improve the simulation even if we obtained better results than with the overwintering models. Finally, the W function with Tmin=2.4°C, Topt=23.12°C and Tmax=32.13°C was the best one with free parameters (see ESM7) and the closer to the known biological behaviour of *L. botrana* according to the literature.

Indeed, we also explored the sensitivity of each group (10-10% and 50-50%) separately and

W ParCal Allp

W ParGUT

Tab.7. Crossed models separating the 10-10% and 50-50% between generations of *L. botrana* with CAL data set. Tests are made with free parameters and with the parameters of the best simulation from CAL Allp, testing the 10-10% /50-50% between generation. The statistical analysis use the RMSE and EFF. Gutierrez's or GUT parameters are Tmin=8.9°C, Topt=30.8°C and Tmax=33°C.

comparing the values of these parameters to the previously obtained. This analysis allowed evaluating

their robustness to simulate each phase independently (Tab.7).

10-10%-50-50% be	etween generations		Statis	stics	
R 10-10%	Simulation	Nb data	RMSE	EFF	
R Allp			8.12	0.93	
R ParCal Allp	10 - 10	93	8.26	0.93	
R ParGUT			8.81	0.92	
R 50-50%		Nb data	RMSE	EFF	
R Allp			7.77	0.94	
R ParCal Allp	50-50	93	7.87	0.94	
R ParGUT			9.06	0.92	
Т 10-10%		Nb data	RMSE	EFF	
T Allp (10-10%)			8.17	0.93	
T ParCal Allp	10 - 10	93	8.31	0.93	
T ParGUT			8.79	0.92	
Т 50-50%		Nb data	RMSE	EFF	
T Allp (50-50%)			7.72	0.94	
T ParCal Allp	50-50	93	7.81	0.94	
T ParGUT			9.03	0.92	
W 10-10%		Nb data	RMSE	EFF	
W Allp (10-10%)			8.11	0.93	
W ParCal Allp	10 - 10	93	8.33	0.92	
W ParGUT			26.14	0.26	
W 50-50%		Nb data	RMSE	EFF	
W Allp			7.74	0.94	

Results obtained in Tab.7 show that the models simulate correctly the 10-10% and the 50-50% generations when we consider parameters from the first calibration (Tab.6) and with all parameters

50-50

0.21

27.77

free but separating 10 - 10% and 50 - 50% in different datasets (for parameters values see in ESM3). The EFF was around 0.92 and 0.94 and the RMSE between 7 and 8 days for the three functions (R, T and W). Indeed, when using Gutierrez's parameters, the R and T models showed good results (EFF=0.92), which was not the case for the W model that degraded (EFF=0.21 and 0.26). This analysis confirms the robustness of the model and its parameters, allowing to use a single function to simulate all the generations of one cycle for the 10 to 10% and 50 to 50% values for each generation.

3.2.2 Simulation 10%-50% of adults flights within a generation itself

As previously reported (Fig.2) results presented in the following refer to simulations with no difference in parameter values between different generations.

Tab.8: Simulation of the 10-50% of *L. botrana* adults flights within a generation itself with CAL, VAL and ALL data set. The statistical analysis use the RMSE (Root Mean Square Error) and EFF (model efficiency). Gutierrez's or GUT parameters are Tmin=8.9°C, Topt=30.8°C and Tmax=33°C.

10%-50% within a g		Statistics			
Model R 10-50%	Nb data	Data set	RMSE	EFF	
R CAL Allp	<u></u>		5.59	0.99	
R CAL ParCAL_Allp10-10	107	CAL	5.93	0.98	
R ParGUT		CAL	6.24	0.98	
R VAL Allp	37	VAL	10.93	0.95	
R ALL	144	ALL	8.56	0.97	
Model T 10-50%	Nb data	Data set	RMSE	EFF	
T CAL Allp	<u></u>		5.53	0.99	
T CAL ParCAL_Allp10-10	107	CAI	5.91	0.98	
T ParGUT		CAL	6.24	0.98	
T VAL Allp	37	VAL	10.02	0.96	
T ALL	144	ALL	7.33	0.98	
Model W 10-50%	Nb data	Data set	RMSE	EFF	
W CAL Allp	<u></u>		5.55	0.99	
W CAL ParCAL_Allp10-10	107	CAI	5.94	0.98	
W ParGUT	144 ALL 8.56 Nb data Data set RMSE 5.53 5.91 6.24 6.24 37 VAL 10.02 144 ALL 7.33 Nb data Data set RMSE 5.55	-0.05			
W VAL Allp	37	VAL	9.84	0.96	
W ALL	144	ALL	7.15	0.98	

Results described in Tab.8 show that very high values of EFF (from 0.98 to 0.99) were obtained irrespective of the model configuration. Fixing parameters for the W model to the values suggested by Gutierrez et al. (2012) significantly degraded the results (EFF = -0.05). In line with the previous tests, the sensitivity of these models have been tested fixing some parameters (Tlow/Tmin and Tmax in ESM5 and 6). All the parameters found are close to the ones known in the literature so we kept them fixed (Tmin=8.9°C, Topt=30.8°C and Tmax=33°C in GUT or Tmin=4.5°C in Baumgärtner et al., (2012)). In the first simulations with CAL data set and using free parameters (All parameters free), W seems to achieve better performances than T and R. Indeed, with fixed parameters the results did not improve with an RMSE around 5.5 days for the 3 functions and with parameters whose values are close to those found in the literature. We also validated the best model (All parameters free model)

with the independent VAL dataset and we obtained good efficiency values for all the models (EFF=0.96) and an error of 9.84 days (RMSE). Fig.5 illustrates the distribution of the simulated and observed data for the 10-50% of male flights within the same generation for the Wang model. The figure shows the good capacity of the model to simulate this phase. In general Fig.6, the box size is proportional to the number of data and shows that the error for the simulation of all the generations for all the transect study have a small error (mean of +/- 5 days) between simulated and observed data.



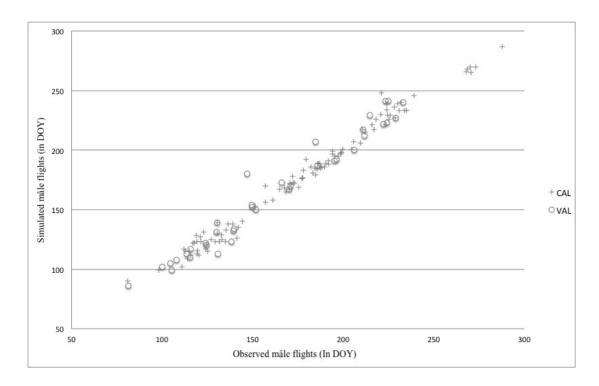
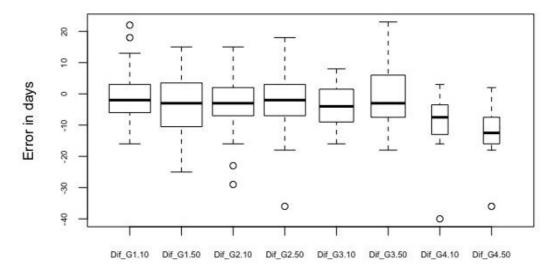


Fig. 5. Distribution of the simulated and observed data for the 10-50% of *L. botrana* male flights within the same generation with the Wang model. Simulated and observed data show here the correlation with both the results from the calibration model (CAL with crosses) and the results from the validation model (VAL with circles).



Generations at 10% and 50% of flight

Fig. 6. Error between observation and simulations in number of days of the 10% and 50% of each generation of L. botrana flights, all site and years confounded. The errors fluctuate around 10 days before or after the observation data in the boxplots, with some extreme data, also present in the observed data. The last generations (G4.10 and G4.50) show a larger difference between observation and predictions, probably due to the sparsity of data that we disposed of, as only 2 sites had a 4^{th} generation. The size of the boxes is proportional to the number of data.

4. General discussion of the results

This study examined the possibility to assemble a simple, robust, generic, yet process-based model for simulating the voltinism of *L. botrana*. This model can be useful for large scale studies and impact studies targeting the behaviour of the insect pests under future climate change. The analysis of existing model indicated that two phases (overwintering and adults) are necessary to realistically represent the life cycle of *L. botrana*. The methodology to calculate the voltinism was based on the 10% and 50% of the total male flights per generation as significant reference to make simulations.

Data used for model development and testing were obtained from a wide latitudinal transect, allowing to test the possibility to apply the model at large scales.

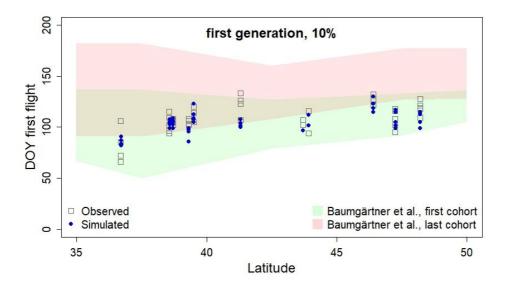
The parameterization of the model was made by analysing different published models (Tab.1). Three types of temperature response functions were examined: Richardson, Triangle and Wang. The evaluation results showed that T and W are very close in their performance. The analysis provided good results with or without the use of certain specific parameter values found in the literature. Moreover, the sensitivity of the models to different sets of parameters from the literature used by Baumgärtner et al. (2012) and Gutierrez et al. (2012) did not improved the results of the parameterization tests with free parameters (Allp). On the contrary, the tests made with the Gutierrez's parameters degraded the results (Tab.8).

Another issue raised in our study was to integrate the photoperiod in a large latitudinal transect (from Lat.36.710°N to Lat.48.213°N) and understand its interaction with temperatures (as multiplicative or additive factor). The analysis of this interaction (Tab.3 and 4) showed that the multiplicative approach was better adapted to describe these interactions than the additive approach proposed by Baumgärtner et al. (2012) (ESM2). In any case, the multiplicative approach is the one commonly implemented in complex models and usually performs accurately (Briere and Pracros 1998; Wang and Engel 1998; Baumgärtner et al. 2012). Finally, the analysis also indicated that assuming a common diapause starting date, $(d_{t0}) = July 15^{th}$, for the entire transect was a reasonable choice.

We show that LGM is able to correctly represent the life cycle of *L. botrana* in various areas with the same set of parameters, with a model performance comparable to the one of local models (Parker et al. 2011). This model is based on existing models but applied at other scale and consider less parameters that make its use easier. This approach is different to the previous one as it can be applied for large scale studies like climate change scenarios.

4.1 Proposal for a new model

The LGM uses a simplification of Baumgärtner's approach only to calculate the first cohort and not the entire population of *L. botrana* as in the original function. The overwintering submodel considers the general approach of the sum of temperatures (i.e. cumulation of temperatures as GDD or Growing Degree Days) and not the speed of growing as in Baumgärtner's original function. Then, the adults submodel uses the same parameters to cumulate heat between and within the generations. As we describe in the Fig.7, the LGM model and Baumgärtner's model have the same range of errors to simulate the first cohort of a *L. botrana* population even with a simplification of the calculation methods. This comparison between both models provided a second partial validation of the LGM model, because it is not exhaustive as the years (observations) and sites (latitude) are different in both studies.



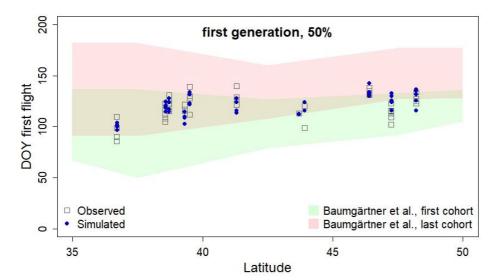


Fig.7: Comparison between LGM observed and simulated adults flights of *L. botrana* in DOY (Day of Year) and Baumgärtner's first and last cohort observations. From a geographical perspective, both models consider a wide range of sites and climates. The simulations of G1-10% and observations both fall within the spread of the first cohort appearing (in green) in Baugmärtner's model. The same observations can be made with the G1-50% that fall within the spread of the last cohort appearing (in red) in Baugmärtner's model.

In the LGM, the results of the different tests showed similar performances for all models and developmental phases studied. We nevertheless propose to use the W model to simulate different temperature responses because it better reflects the known biological response of *L. botrana* to temperature. Thus, Wang is the unique function used to simulate both overwintering (Predia, dia and Post-dia) and adult flights submodels. This is a classical approach using a first model with the calibration and validation data sets. A preliminary analysis revealed that the small divergence between the models allowed us to test another model with all the data sets. Tab.9 provides the statistical analysis with all the parameter values used for the calibration, validation and the entire data set.

It summarizes the proposed model, LGM model, showing the different parameter sets used to simulate the three phases of the overwintering period, but the single, common set of parameters used to simulate all the phases (10% and 50%) of the different generations of the adults flights.

Tab.9. Final submodels selected for their efficiency according to the calibration (CAL), validation (VAL) and the full data set (ALL) of samples and the stability of their biological response to temperature. Both set of parameters obtained using the 10% and 50% between the generations of the *L. botrana* adults flights and the 10-10%/50-50% within the generations are presented. The final results for the Wang model are close to biological thresholds reported in the literature, particularly those published by Gutierrez et al. (2012).

	Parameters								Statistics					
Models	Di	m :	T		Plow	Phigh	-	CAL				VAL		
	Phases	1 mm	торі	1 max			F	RMSE	EFF	Nb Obs	RMSE	EFF	Nb Obs	
WAllp	Predia	2,45	14,36	27,69	-	-	40,74						_	
	Dia	7,66	15,04	22,87	9,17	14,25	28,35	7,49	0,74	39	9,07	0,08	12	
	Post- dia	11,39	11,42	20,00	-	-	1,47	,	,		,	,		
W Don Alle	Predia	2,45	14,36	27,69	-	-	39,79							
W ParAllp Tmin=4.5°C	Dia	7,66	15,04	22,87	9,17	14,25	30,23	8,04	0,69	39	9,04	0,09	12	
(BAUM)	Post- dia	4,50	11,42	20,00	-	-	2,026	,	,		,	,		
W Allp 10- 10 /50-50 Between g.	Adult flight	2,38	23,13	32,31	-	-	45,61	6,93	0,95	140	11,15	0,84	46	
W Allp 10- 50 within g.	Adult flight	2,38	23,13	32,31	-	-	7,76	5,94	0,98	107	9,84	0,96	37	

Concerning the adult development stage, the results confirm our hypothesis that a standard set of parameters is sufficient to accurately simulate the voltinism of *L. botrana* from diapause to last flight in terms of the 10% and 50% of the male flights.

This finding is important because, as we described in the introduction, it is common practice to recalibrate phenological models for each generation (Beddow et al. 2010). Thus, in previous studies (Tab.1) authors proposed a different set of parameters or models for each generation considering that the biological response was different during the entire cycle. We are convinced that our approach introduces more robustness into the simulation of the life cycle of *L. botrana*. Indeed, it allows applications at large geographical scales without adapting the model to local parameters. Whether this can also be extended to other species remains to be investigated.

4.2 Advantages and limits of the generic model

To define the LGM, we considered the models with the best statistical criteria and the most representative biological processes known from the literature (ESM7). One of the major difficulties we encountered was to model the overwintering phase, as we did not dispose of data to represent the

associated intermediate phases (i.e., end of prediapause and end of diapause). The calibration of this complex submodel was hence performed targeting the date of the 10% of the first generation at each site. The lack of intermediate information is a common problem faced in developing phenology models for plants (Chuine et al. 2016). In our case, the data collection have been a long process and the quality selection of the male flights had to correspond to sites where we had meteorological data available close by (at least 15 years). Some authors like Amo-Salas et al. (2011) and Armendáriz et al. (2009) integrated relative humidity in their model, but they concluded that the value error with or without this parameter was not a significant influencing factor. Indeed, we did not consider relative humidity as in Baumgärtner et al. (2012), as we did not have such data available for all our sites.

Another limit of this study concerns the data that have been selected and converted to a comparable and standardized unit, but the observed data do not consider a protocol and the heterogeneity of the observed data can alter the quality of the calibration. Observation values obtained in the laboratory (in controlled conditions, at constant temperature), where rates of development are estimated approximately (Briere and Pracros 1999; Torres-Vila et al. 1999; Picart 2009) can differ considerably from those observed in the field and between the sites or by calibrating models using local conditions (Arca et al. 1993; Torres-Vila et al. 1999; Severini et al. 2005; Moravie et al. 2006). The large latitudinal range of study is complex, as seen in Fig.5, where some extreme values, representing the 4th generations in south of Spain (not present at other sites), fall for the CAL dataset and not in the VAL dataset due to the randomization of the data selection and can alter the EFF of the tests. Nevertheless, the strength of this study has been to obtain a stable and robust model at a large spatial scale.

We believe that this work has contributed to raise new questions on scale applications for applied studies like climate change and on the methodologies used to simulate insect developmental processes. The approach developed here, making use of a simple and unique set of parameters, representative of the biological response to temperatures, has shown very interesting and encouraging results.

4.3 Future model applications

We anticipate that LGM, as a model, is appropriate to conduct impact studies that integrate climate change scenarios in order to predict the future geographical distribution of ectothermic insects and the future extent of vulnerable areas. Multivoltine species are likely to develop more generations under warmer conditions (Caffarra et al. 2012; Reineke and Thiery 2016). Sensitivity analyses with LGM, combined with phenological models for plants, as grapevine for example, will allow to identify critical shifts in the synchrony or asynchrony of different trophic levels (host plants and insect pests) and the ensuing overlap periods (Hoover and Newman 2004; Hirschi et al. 2012; Stoeckli et al. 2012; Gilioli et al. 2016). Nevertheless, more data would have been useful to validate the choice of the best models. This study could be improved by testing other regions of the world or by implementing the entire set

of data used. In concert with phenological models for crops, LGM will help raise the awareness of producers on the emergence of new threats and adaptation of future treatment practices according to geographic areas. Again, the advantages of LGM are its conceptual simplicity and generic parameterization, which allow the implementation of climate change studies at the large scale for both plants and insect pests. **Author Contribution Statement** VC and IGCA conceived and designed the research and analyzed the data. PC and JM contributed new reagents and analytical tools. MB reviewed and corrected the manuscript. All authors read and approved the manuscript. **Compliance with Ethical Standards:** All authors declare that they have no conflict of interest. This article does not contain any studies with animals or human participants performed by any of the authors. Acknowledgments A particular thanks to the Ernst and Lucie Schmidheiny Foundation for providing the funding that enabled the first author to conduct a short stay at the AgroClim Unit at INRA in Avignon (France) and Denis Thierry at INRA Bordeaux. This work would have not been possible without the help of data providers for meteorological data (MeteoSuisse, Agrometeo, IDAWEB, SIAR, SIAM, riegos.ivia, Phenoclim INRA) and for L. botrana male flight data, A. L. Espadas (Oficina Comarcal Agraria de Alhama), J. Serrano Palao (Servicio de Sanidad Vegetal de Jumilla), J. V. Badia Ballester (GVA-Generalitat Valencia) L. García de Arboleya Puerto (Junta de Andalucia), Oficina del regante de Zarragoza, J. Isiegas Mainar (D.O.P Cariñena), Mme Lauer (CA Alsace), Mr Genevet and Mr Cassarini (CA Gard), Mr Pasquier and Mr Goëlles (Agroscope) and especially Dr. Josep Armengol from the Universitat Politècnica de València for his collaboration.

640 References

Allen JC (1976) A modified sine wave method for calculating degree days. Environ Entomol 5:388–396

643

Amo-Salas M, Ortega-López V, Harman R, Alonso-González A (2011) A new model for predicting
 the flight activity of Lobesia botrana (Lepidoptera: Tortricidae). Crop Prot 30:1586–1593. doi:
 10.1016/j.cropro.2011.09.003

647

Arca B, Cossu A, Delrio G, Locci L (1993) Individuazione dei gradi giorno relativi allo sviluppo della
 Lobesia botrana(Den. et Schiff.) in Sardegna. Atti Convegno Nazionale 'Protezione delle colture:
 osservazioni, previsioni, decisioni'. 325–334

651

Armendáriz I, Pérez-Sanz A, Capilla C, et al (2009) Cinco años de seguimiento de la polilla del
 racimo de la vid (Lobesia botrana) en la DO Arribes (Castilla y León, España). Bol Sanid Veg-Plagas
 35:193–204

655

Bale JS, Hayward SAL (2010) Insect overwintering in a changing climate. J Exp Biol 213:980–994. doi: 10.1242/jeb.037911

658

Bale JS, Masters GJ, Hodkinson ID, et al (2002) Herbivory in global climate change research: direct effects of rising temperature on insect herbivores. Glob Change Biol 8:1–16. doi: 10.1046/j.1365-2486.2002.00451.x

662 663

Baumgärtner J, Gutierrez AP, Pesolillo S, Severini M (2012) A model for the overwintering process of European grapevine moth Lobesia botrana (Denis & Schiffermüller) (Lepidoptera, Tortricidae) populations. J Entomol Acarol Res 44:2. doi: 10.4081/jear.2012.e2

665 666

664

Beck HE, Zimmermann NE, McVicar TR, et al (2018) Present and future Köppen-Geiger climate classification maps at 1-km resolution. Sci Data 5:180214. doi: 10.1038/sdata.2018.214

669

Beddow JM, Kriticos D, Pardey PG, et al (2010) Potential global crop pest distributions using
 CLIMEX: HarvestChoice applications. St Paul MN Univ Minn Harvest

672673

Bregaglio S, Donatelli M, Confalonieri R (2013) Fungal infections of rice, wheat, and grape in Europe in 2030–2050. Agron Sustain Dev 33:767–776. doi: 10.1007/s13593-013-0149-6

674 675 676

Briere J-F, Pracros P (1999) A novel rate model of temperature-dependent development for arthropods. Env Entomol 28:22–29. doi: 10.1093/ee/28.1.22

677 678

679 Briere J-F, Pracros P (1998) Comparison of temperature dependant growth models with the development of Lobesia botrana (Lepidoptera:Tortricidae). Env Entomol 27:94–101. doi: 10.1093/ee/27.1.94

682 683

Brodeur J, Boivin G, Bourgeois G, et al (2013) Impact des changements climatiques sur le synchronisme entre les ravageurs et leurs ennemis naturels: conséquences sur la lutte biologique en milieu agricole au Québec

685 686

684

Caffarelli V, Vita G (1988) Heat accumulation for timing grapevine moth control measures. Bull SROP 11:24–26 ref.4

689

Caffarra A, Eccel E (2011) Projecting the impacts of climate change on the phenology of grapevine in a mountain area. Grape Wine Res 17:52–61. doi: 10.1111/j.1755-0238.2010.00118.x

692

693 Caffarra A, Rinaldi M, Eccel E, et al (2012) Modelling the impact of climate change on the interaction

between grapevine and its pests and pathogens: European grapevine moth and powdery mildew. Agric Ecosyst Environ 148:89–101. doi: 10.1016/j.agee.2011.11.017

696

Castex V, Beniston M, Calanca P, et al (2017) Pest management under climate change: The importance of understanding tritrophic relations.pdf. Sci Total Environ 616–617:397–407. doi: http://dx.doi.org/10.1016/j.scitotenv.2017.11.027

700

701 Chuine I, Bonhomme M, Legave J-M, et al (2016) Can phenological models predict tree phenology 702 accurately in the future? The unrevealed hurdle of endodormancy break. Glob Change Biol 22:3444– 703 3460. doi: 10.1111/gcb.13383

704

Chuine I, de Cortazar-Atauri IG, Kramer K, Hänninen H (2013) Plant Development Models_2013. In:
 Schwartz MD (ed) Phenology: An Integrative Environmental Science. Springer Netherlands,
 Dordrecht, pp 275–293

708

Cleland E, Chuine I, Menzel A, et al (2007) Shifting plant phenology in response to global change.
Trends Ecol Evol 22:357–365. doi: 10.1016/j.tree.2007.04.003

711

Flores C, Azin G (2015) Modelos lineales para pronosticar la actividad de vuelo de Lobesia botrana (Denis & Schiffermüller) (Lepidoptera: Tortricidae) en la localidad de Russel, Mendoza, Argentina

714

Gabel B, Mocko V (1984) Forecasting the cyclical timing of the grape vine moth, Lobesia botrana (Lepidoptera, Tortricidae). Acta Entomologica Bohemoslov.:1–14

717

Gallardo A, Ocete R, López MA, et al (2009) Forecasting the flight activity of Lobesia botrana (Denis and; Schiffermüller) (Lepidoptera, Tortricidae) in Southwestern Spain. J Appl Entomol 133:626–632.
 doi: 10.1111/j.1439-0418.2009.01417.x

721

Gilioli G, Pasquali S, Marchesini E (2016) A modelling framework for pest population dynamics and management: An application to the grape berry moth. Ecol Model 320:348–357. doi: 10.1016/j.ecolmodel.2015.10.018

725

Gutierrez AP, Ponti L, Cooper ML, et al (2012) Prospective analysis of the invasive potential of the European grapevine moth Lobesia botrana (Den. & Schiff.) in California. Agric For Entomol 14:225–238. doi: 10.1111/j.1461-9563.2011.00566.x

729 730

Hance T, van Baaren J, Vernon P, Boivin G (2007) Impact of Extreme Temperatures on Parasitoids in a Climate Change Perspective. Annu Rev Entomol 52:107–126. doi: 10.1146/annurev.ento.52.110405.091333

732 733

731

Hänninen H (1990) Modelling bud dormancy release in trees from cool and temperate regions. ACTA For Fenn 312

736

Hatt S, Boeraeve F, Artru S, et al (2018) Spatial diversification of agroecosystems to enhance biological control and other regulating services: An agroecological perspective. Sci Total Environ 621:600–611. doi: 10.1016/j.scitotenv.2017.11.296

740

Hirschi M, Stoeckli S, Dubrovsky M, et al (2012) Downscaling climate change scenarios for apple pest and disease modeling in Switzerland. Earth Syst Dyn 3:33–47. doi: 10.5194/esd-3-33-2012

743

Honěk A (1996) Geographical variation in thermal requirement for insect development. Eur J Entomol
 93:303–312

746

Hoover JK, Newman JA (2004) Tritrophic interactions in the context of climate change: a model of grasses, cereal Aphids and their parasitoids. Glob Change Biol 10:1197–1208. doi: 10.1111/j.1529-

749 8817.2003.00796.x

Ioriatti C, Anfora G, Tasin M, et al (2011) Chemical Ecology and Management of Lobesia botrana (Lepidoptera: Tortricidae). J Econ Entomol 104:1125–1137. doi: 10.1603/EC10443

Jongman RHG, Bunce RGH, Metzger MJ, et al (2006) Objectives and Applications of a Statistical Environmental Stratification of Europe. Landsc Ecol 21:409–419. doi: 10.1007/s10980-005-6428-0

Lamichhane JR, Barzman M, Booij K, et al (2015) Robust cropping systems to tackle pests under climate change. A review. Agron Sustain Dev 35:443–459. doi: 10.1007/s13593-014-0275-9

Lin BB (2011) Resilience in Agriculture through Crop Diversification: Adaptive Management for Environmental Change. BioScience 61:183–193. doi: 10.1525/bio.2011.61.3.4

Logan JA, Wollkind DJ, Hoyt SC, Tanigoshi LK (1976) An analytic model for description of temperature dependent rate phenomena in arthropods. Environ Entomol 5:1133–1140

Marchesini, E., & Monta, L. D. (2004). Nel Veneto quattro generazioni di tignoletta della vite. *Informatore Agrario*, 60(4), 75-78.

Maiorano A, Bregaglio S, Donatelli M, et al (2012) Comparison of modelling approaches to simulate the phenology of the European corn borer under future climate scenarios. Ecol Model 245:65–74. doi: 10.1016/j.ecolmodel.2012.03.034

Martín-Vertedor D, Ferrero-García JJ, Torres-Vila LM (2010) Global warming affects phenology and voltinism of Lobesia botrana in Spain. Agric For Entomol 12:169–176. doi: 10.1111/j.1461-9563.2009.00465.x

Milonas PG, Savopoulou-Soultani M, Stavridis DG (2001) Day-degree models for predicting the generation time and flight activity of local populations of Lobesia botrana (Den. & Schiff.) (Lep., Tortricidae) in Greece. J Appl Entomol 125:515–518. doi: 10.1046/j.1439-0418.2001.00594.x

Moiroux J, Bourgeois G, Boivin G, Brodeur J (2014) Impact différentiel du réchanffement climatique sur les insectes ravageurs des cultures et leur ennemis naturels: implications en agriculture

Moravie M-A, Davison AC, Pasquier D, Charmillot P-J (2006) Bayesian forecasting of grape moth emergence. Ecol Model 197:478–489. doi: 10.1016/j.ecolmodel.2006.03.030

Moreau J, Desouhant E, Louâpre P, et al (2017) How Host Plant and Fluctuating Environments Affect Insect Reproductive Strategies? In: Advances in Botanical Research. Elsevier, pp 259–287

Nietschke BS, Magarey RD, Borchert DM, et al (2007) A developmental database to support insect phenology models. Crop Prot 26:1444–1448. doi: 10.1016/j.cropro.2006.12.006

Ortega-López V, Amo-Salas M, Ortiz-Barredo A, Díez-Navajas AM (2014) Male flight phenology of the European grapevine moth Lobesia botrana (Lepidoptera: Tortricidae) in different wine-growing regions in Spain. Bull Entomol Res 104:566–575. doi: 10.1017/S0007485314000339

Parker AK, De CortáZar-Atauri IG, Van Leeuwen C, Chuine I (2011) General phenological model to characterise the timing of flowering and veraison of Vitis vinifera L.: Grapevine flowering and veraison model. Aust J Grape Wine Res 17:206–216. doi: 10.1111/j.1755-0238.2011.00140.x

Pavan F, Zandigiacomo P, Dalla Montà L (2006) Influence of the grape-growing area on the phenology of Lobesia botrana second generation. Bull Insectology 59:105–109

Picard RR, Cook RD (1984) Cross-Validation of Regression Models. J Am Stat Assoc 79:10

805

Picart D (2009) Modélisation et estimation des paramètres liés au succès reproducteur d'un ravageur de la vigne (Lobesia botrana DEN. & SCHIFF.). Université Sciences et Technologies-Bordeaux I

808 809

Reineke A, Thiery D (2016) Grapevine insect pests and their natural enemies in the age of global warming. 12. doi: 10.1007/s10340-016-0761-8

810 811

Richardson EA, Seeley, Schuyler D, Walker DR (1974) A model for estimating the completion of Rest for "Redhaven" and "Elberta" Peach trees. HortSciences 9:331–332

814

Roditakis NE, Karandinos MG (2001) Effects of photoperiod and temperature on pupal diapause induction of grape berry moth Lobesia botrana. Physiol Entomol 26:329–340. doi: 10.1046/j.0307-6962.2001.00253.x

818

Severini M, Alilla R, Pesolillo S, Baumgärtner J (2005) Fenologia della vite, e della Lobesia botrana (Lep. Tortricidae) nella zona dei Castelli Romani. Riv Ital Agrometeorol 3:34–39

821 822

Singer MS, Parmesan C (2010) Phenological asynchrony between herbivorous insects and their hosts: signal of climate change or pre-existing adaptive strategy? Philos Trans Biol Sci 365:3161–3176

823 824

Steinbauer MJ, Kriticos DJ, Lukacs Z, Clarke AR (2004) Modelling a forest lepidopteran: phenological plasticity determines voltinism which influences population dynamics. For Ecol Manag 198:117–131. doi: 10.1016/j.foreco.2004.03.041

828

Stoeckli S, Hirschi M, Spirig C, et al (2012) Impact of Climate Change on Voltinism and Prospective
 Diapause Induction of a Global Pest Insect – Cydia pomonella (L.). PLoS ONE 7:e35723. doi:
 10.1371/journal.pone.0035723

832

Svobodová E, Trnka M, Dubrovský M, et al (2014) Determination of areas with the most significant shift in persistence of pests in Europe under climate change. Pest Manag Sci 70:708–715. doi: 10.1002/ps.3622

836

Thiery, Denis. 2008. "Les Tordeuses Nuisibles à La Vigne." *Féret, Bordeaux*, les ravageurs de la vigne, 54.

839

Tilman D, Cassman KG, Matson PA, et al (2002) Agricultural sustainability and intensive production practices. Nature 418:671–677. doi: 10.1038/nature01014

842 843

Tobin PC, Nagarkatti S, Loeb G, Saunders MC (2008) Historical and projected interactions between climate change and insect voltinism in a multivoltine species. Glob Change Biol 14:951–957. doi: 10.1111/j.1365-2486.2008.01561.x

845 846

844

Torres-Vila, LM, Rodriguez-Molina MC, Roehrich R, Stockel J (1999) Vine phenological stage
 during larval feeding affects male and female reproductive output of Lobesia botrana (Lepidoptera:
 Tortricidae). Bull Entomol Res 89:549–556

850

Touzeau J (1981) Modélisation de l'évolution de l'Eudemis de la vigne pour la region Midi-Pyrénées.
 Boll Zool Agrar Bachic Ser II 16:26

853

Wang E, Engel T (1998) Simulation of phenological development of wheat crops. Agric Syst 58:1–24

855

Zhou X, Harrington R, Woiwod IP, et al (1995) Effects of temperature on aphid phenology. Glob Change Biol 1:303–313