ORIGINAL PAPER



Climate changes affecting biotic interactions, phenology, and reproductive success in a savanna community over a 10-year period

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Received: 22 May 2017 / Accepted: 3 October 2017 / Published online: 16 October 2017 © Springer Science+Business Media B.V. 2017

Abstract Climatic parameters are able to influence the timing of phenological events affecting the degree of synchrony among plant species, their interactions, and reproductive success. Shrubs of Malpighiaceae family in the Brazilian Tropical Savanna present sequential flowering phenology. We verified variations in climatic factors (temperature and precipitation) over a period of 10 years (2005-2014) and correlated them with the onset of flowering of four of these Malpighiaceae species. Furthermore, we tested whether the phenological synchronization among species has changed over time affecting the herbivory and fruit set. Herbivory and fruit production were recorded during three reproductive seasons (2008/2009, 2011/2012, 2013/2014). We developed a mathematical model to estimate the flower and fruit production in response to phenological changes for the next 5 years. Results show that climatic factors changed, influencing the onset of species flowering. The degree of overlap among species also changed and the effects on species interactions were species specific. The mathematical model successfully presented a tendency on flower and fruit production contributing to the predictions of the outcomes in

Handling Editor: Heikki Hokkanen.

Electronic supplementary material The online version of this article (doi:10.1007/s11829-017-9572-y) contains supplementary material, which is available to authorized users.

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² Laboratório de Mecânica de Estruturas, Faculdade de Engenharia Mecânica, Universidade Federal de Uberlândia, Uberlândia, Minas Gerais CEP 38400-920, Brazil response to phenological changes. We confirm the effects of climate changes on plant phenological events and the importance of feature plasticity for better performance of species.

Keywords Synchrony · Flowering · Abiotic factors · Malpighiaceae · Mathematical modeling

Introduction

Flowering phenology is a critical life-history trait in angiosperms and has potential significance for plant ecology and evolution (e.g., Amasino 2010; Torres and Galetto 2011). Due to its influence on a wide range of ecological processes, including plant interactions, and reproductive success (Pilson 2000; Kawagoe and Kudoh 2010; Del-Claro et al. 2016), the pattern of flowering is one of the most investigated aspects of plant phenology (Hegland et al. 2009; Vilela et al. 2014), especially nowadays with the occurrence of noticeable climate changes (Franks et al. 2013; CaraDonna et al. 2014). Shifts in the timing of flowering have great consequences on plant developmental traits and interactions with mutualistic (pollinators and seed dispersers) and antagonistic (herbivores) partners which may affect plant success and survival rates of species (Herrera et al. 2002; Memmott et al. 2007; Torezan-Silingardi 2011; Encinas-Viso et al. 2012; Velasque and Del-Claro 2016).

The flowering pattern of individual species varies widely (Forrest et al. 2010; Forrest and Thomson 2011). In the Cerrado habitat, for some species it is staggered over a long period while others present a more pronounced peak with relatively few individuals occurring in the beginning or final of the flowering period (Ferreira and Torezan-Silingardi 2013). These variations are likely the result of a combination of abiotic, biotic, and phylogenetic factors (Mitchell et al.

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2009; Rusquist and Staton 2013; Del-Claro and Marquis 2015) and the action of these factors may have led to either the minimization or the maximization of flowering within and among plants species, inducing synchronous and asynchronous phenologies at population and community level (Van Schaik et al. 1993; Vilela et al. 2014).

Sympatric species can exhibit clumped flowering phenologies that overlap extensively, possibly reducing herbivory according to the Herbivore Satiation Hypothesis (Beattie et al. 1973), i.e., a plant-plant association in which one species could decrease the likelihood of detection or vulnerability of other species by insect herbivores (see Barbosa et al. 2009; Lamarre et al. 2014). Alternatively, less synchrony between herbivore and flowering plants (e.g., Forrest and Thomson 2011) with a mismatch in the timing of interacting organisms can also reduce the herbivore pressure (Hoye and Forchhammer 2008). Nevertheless, during periods of overlap, different plant species may attract the same pollinators which can lead to competition for these cross-pollination vectors (e.g., Tanakura et al. 2009; Muchhala and Thomson 2012). The Pollinator Competition Hypothesis introduced by Robertson (1895) assumes that pollinators are a limiting resource and flowering events should be evenly spread through time (staggered phenology) to reduce the competition for pollinators and increase the effectiveness of pollination. Thus, a balance in plant life-history strategies based on a trade-off between a staggered phenology and a more overlapped flowering is expected (see Herrera et al. 2002, Forrest et al. 2010).

Many plant species have presented phenological shifts in the timing of their life-history events (Rafferty and Ives 2011), affecting the strength of mutualists and antagonistic interactions (Burkle et al. 2013; Rafferty et al. 2013, 2015; Velasque and Del-Claro 2016) and consequently their fitness (see English-Loeb and Karban 1992; Thomson 2010; Lange and Del-Claro 2014). Recent studies attempt to answer why and how the phenological overlap of interacting species has been changing. Many have concluded that climate change is the main disturbance responsible for alteration of timing of life-history events in a wide array of communities (Forrest et al. 2010; Rafferty and Ive 2012; Diez et al. 2012), due to its potential to desynchronize the phenology of interdependent species (Rafferty et al. 2015) and vary the degree of overlap among related plant species (Forrest et al. 2010).

In a recent meta-analysis, Chambers et al. (2013) related phenological drivers and trends among Southern Hemisphere species and discussed how phenological information contributes to our knowledge of the adaptive capacity of species, their resilience and constraints. However, the authors concluded that our general understanding and capacity to predict phenological responses remains patchy and constrained. Furthermore, few studies have been able to directly address what are the consequences of phenological shifts for species interactions (but see van Asch et al. 2007; Fabina et al. 2010; Liu et al. 2011) and for plant fitness (but see Dominguez and Dirzo 1995; Miller-Rushing et al. 2010). Therefore, there is an urgency to translate a basic understanding of phenology into forecasts about continued phenological changes and to predict the ecological consequences of these changes (Diez et al. 2012), especially for South American species (Staggemeier and Morellato 2011; Morellato et al. 2013).

Integrating life-history theory and mathematical models into ecological studies should help in forecasting changes and in understanding what may be the consequences for plant interaction and reproductive success (Fabina et al. 2010). Mathematical models could be used to generate testable predictions that address how phenological shifts can be generated to investigate consequences for temporal overlap of interacting species (Memmott et al. 2007; Gilman et al. 2012) and how it could affect plant fitness.

Some empirical models have historically been proposed (Porter et al. 1993) but, being based only on experimental data, these are unable to predict plant responses to new scenarios. Only recently some *mechanistic* (methodologies that take into account *cause–effect* relations based on plant phenology and biology) models have been introduced (Hoogenboom 2015; Jones et al. 2003), having success in predicting tendencies of growth.

Cerrado (the Brazilian Tropical Savanna) is the second largest South American ecosystem and presents great seasonal variation as a remarkable characteristic (Oliveira-Filho and Ratter 2002). Initially, we (a) tested whether the climatic parameters (temperature and precipitation) have changed over 10 years (2005-2014) and then we used empirical data of four Malpighiaceae species of Cerrado, which present previously described sequential flowering (Torezan-Silingardi 2007; Vilela et al. 2014), to (b) answer whether the onset of flowering is correlated with those climatic parameters, and (c) whether the phenological synchrony among species has therefore changed over time. We also aimed to (d) detect the effects of a sequential flowering and a higher overlap phenology for herbivory and fruit set of species. We developed a mathematical model, CSVM program (Claro 2015), to (e) estimate the flower and fruit production of each analyzed plant species for the next 5 years in a scenario with continued phenological changes. We hypothesized that temperature and precipitation rates have presented variation over the years analyzed and the onsets of flowering have also changed affecting the degree of overlap among Malpighiaceae species. Furthermore, we expected that the degree of plant phenological overlap (staggered or synchronized) influences the herbivory and fruit set of species.

Materials and methods

Study site

Fieldwork was carried out from May 2008 to March of 2014, spanning three reproductive seasons for each species studied (2008/2009; 2012/2013; 2013/2014), at the *Reserva Ecológica do Clube Caça e Pesca Itororó de Uberlândia* (CCPIU) (18°59'S, 48°18'W), Uberlândia, Minas Gerais State, Brazil. We used a 400-ha Cerrado site consisting of a dense scrub of shrubs and trees, known as Cerrado sensu stricto (Oliveira-Filho and Ratter 2002).

Plant species

In the Cerrado of central Brazil, shrubs of Malpighiaceae family are diverse and abundant (Gates 1982; Anderson 1990; Ferreira and Torezan-Silingardi 2013). Several species exhibit a staggered phenological development, in which individuals of different species resprout, bloom, and set fruit sequentially over time (Munhoz and Felfili 2005; Torezan-Silingardi 2007; Mendes et al. 2011). Four of these Malpighiaceae species were selected for the study because they present a sequential flowering phenology previously described by Torezan-Silingardi (2007) and Vilela et al. (2014): Peixotoa tomentosa A. Juss., Banisteriopsis laevifolia (A. Juss) B. Gates, Banisteriopsis campestris (A. Juss.) Little and *Banisteriopsis malifolia* (Ness and Mart) B. Gates. These species are deciduous, hermaphroditic, bee-pollinated, self-compatible, and share both pollinators and herbivores. The sequential resprouting and flowering of these species in the Cerrado also provides an uninterrupted food supply to a diverse herbivore guild. The first to blossom is P. tomentosa in June, followed by B. laevifolia, B. campestris, and B. malifolia. The greatest similarity in herbivore composition occurs between P. tomentosa and B. laevifolia (70%). Between these two species and B. campestris the similarity is almost 65%. *B. malifolia* shared almost 50% of its herbivores with the other plant species (Vilela et al. 2014). The immature structures (e.g., young leaves, buds, and flowers) have low structural resistance to physical damage, making the shoots, and inflorescences especially attractive to chewing and sucking insects (see Torezan-Silingardi 2007; Vilela et al. 2014).

Climatic analysis

The climate at the study site is markedly seasonal, with dry/ cold season from April to September (rainfall 22 ± 20 mm; temperature $19^{\circ} \pm 3 ^{\circ}$ C) and a wet/warm season occurring from October to March (rainfall, 270 ± 50 mm; temperature, $23^{\circ} \pm 5^{\circ}$ C) (additional details in Réu and Del-Claro 2005). The annual mean of temperature and total precipitation were provided by the Climate Station of the Geography Institute of Federal University of Uberlandia (Fig. 1). Initially, we performed the Repeated Measures ANOVA to compare the monthly means of temperature and monthly precipitations of each year from 2005 to 2014. We considered climatic factors since 2005 based on the sequential phenological results provided by Torezan-Silingardi (2007) for the same Malpighiaceae species at same place during 2005/2006. Further, we conducted Spearman's correlations between climatic factors (monthly means of temperature and monthly precipitations) and phenological parameters (the number of individuals which begin their flowering in each month) (e.g., Morellato et al. 2010; Staggemeier and Morellato 2011).

Data collection

Individual plants (N = 30) of each plant species were monitored every 2 weeks during their reproductive period (see Vilela et al. 2014). Plant phenology (intensity and number of buds, flowers, and fruits) was recorded according to Torezan-Silingardi and Oliveira (2004) and Vilela et al.



Fig. 1 Annual mean temperature (°C, line) and precipitation (mm, bar) from 2005 to 2014 in Uberlândia, Minas Gerais State, Brazil

(2014) and fruit production was expressed as the proportion of fruit produced per buds (e.g., Del-Claro et al. 2013). At the end of the breeding season, we also quantified herbivory (leaf area loss). We recorded data from nine leaves per plant (three from the most apical stem, three from a middle stem, and three from the most basal stem) in order to determine the mean herbivory rate per species. This procedure was done without leaf removal. Measurements of herbivory rates on leaves were assessed by placing leaves on a transparent grid (divided into millimeters). An index of herbivory from each leaf was calculated as the proportion of points in the grid falling within damaged and undamaged areas of the leaf blade (see Moreira and Del-Claro 2005).

Statistical analysis

Species flowering variation

Circular statistical analyses were performed using the date of first flowering of each species separately for each year studied. To calculate the circular statistic parameters, months were converted to angles from 0 =January to 345 =December at intervals of 15 because the measurements were made every 2 weeks (e.g., Cardoso et al. 2012). The frequency of individuals at the onset of flowering for each species was considered for calculating the parameters: the mean vector (μ) , length of mean vector (r), median, circular standard deviation, Rayleigh test (z). The mean date for each phenophase is determined by converting the mean angular directions to corresponding mean dates (see Morellato et al. 2000, 2010; Staggemeier et al. 2010). When the mean angle was significant, we performed the two-sample Watson-Williams tests (F) to determine differences among the mean data of the onset flowering of each species over time (Zar 1996).

Community flowering variation

Circular statistical analyses were also performed considering all species studied to analyze the degree of overlap in each reproductive period to verify for possible community flowering variation. In this case, we used more phenological variables: date of first bud, first flowering and first fruiting, date of peak of bud, flowering and fruiting. The phenological data were analyzed with the statistical software Oriana 4.0.

Intending to confirm a segregated phenology, a null model was performed to indicate if the temporal overlap among species was less than expected by chance. Overlap was quantified via Pianka (Pianka 1973) and Czechanowski (Feinsinger et al. 1981) indices and the values were generated using a randomization algorithm (Rosario). The algorithm Rosario was designed specifically for use with interval data important (Castro-Arellano et al. 2010). Rosario maintains the shape of the empirical activity distributions for each species in the randomly generated matrices by shifting entire activity patterns a random number of intervals. For each analysis, overlap indices were calculated for 10,000 randomly generated matrices of temporal flowering phenology, creating a null distribution of overlap values. Then a one-tailed test was conducted and the p value was calculated as the proportion of randomizations that resulted in overlap that is equal to or less than the empirical overlap value (observed) (e.g., Brito et al. 2012). Simulations were conducted with the TimeOverlap program (Castro-Arellano et al. 2010).

Herbivory and productivity analyses

We compared the rates of herbivory and fruit production between the periods with minimal (2008/2009) and higher overlap (2013/2014) using Paired test t after arcsine of the square root transformation of the percentage values for herbivory data and Wilcoxon test for fruit production, used for paired nonparametric data. The data were analyzed using the software Systat 12.

Mathematical modeling

The model we developed is part of a software (Programa *CSVM*[®]—INPI—BR 51 2015 000482-4) to present a tendency of plant production in face of phenological shifts associated with climate variability in agricultural systems (Claro 2015). The software encompasses a mechanistic fuzzy logic approach, to describe the behavior of a population of plants. Considering that temporal and spatial variations on the abundance of specific partners could represent local variation on the results of flower and fruit production, the program intends to ensure a qualitative validity, by demonstrating a tendency in different scenarios. We adjusted and validated the model parameters by using Differential Evolution Optimization methods (Lobato and Steffen 2008; Vanderplaats 2005) to define some of the model's constants, by comparing the real case with the program results via optimization. This procedure was made by estimating the interval in which the variables considered may vary, normalizing the variables, and then applying an optimization procedure to achieve the best approximation of the real parameters/ variables in a specific scenario (Lobato and Steffen 2007). Initially, we considered the field data obtained by Torezan-Silingardi (2007) in 2005/2006 aiming to ensure a greater robustness for the model. In order to standardize the sample method, we employed the same methodology applied to Torezan-Silingardi (2007).

The inputs considered were rainfall, leaf sprouting and production of floral buds, flowers and fruits. Other parameters estimated by the researchers and adjusted by the optimization procedure, were the weekly rates of herbivory, pollination and autonomous self-pollination. The preliminarv data as input parameters were: area of 2500 m^2 , with 100×100 square elements of 0.5×0.5 m each, simulating for 5 years of 48 weekly time steps each (1 time step equals 1.0863 week). Containing 50 B. malifolia individuals (3×3) elements in size), 50 P. tomentosa individuals (2×1 elements in size), 30 B. laevifolia individuals (2×2 elements in size), and 40 B. campestris individuals $(1 \times 1 \text{ elements})$ in size) based on their real size and presence measured on the field. The herbivory and pollination were set to occur in 90% of the elements per repetition. The randomness of weekly herbivory considered was its species-specific base value with a random of 3%. The weekly randomness of insect pollination and autonomous self-pollination considered was also its species-specific base value with a random of 3%. We considered 13% of randomness for the baselines (rainfall, leaves sprouting, and flowering), applied both in time and geographical location. All variation parameters were estimated based on real field data.

An advance in the phenology of *B. laevifolia* was simulated, given that this phenomenon was already observed in 2012 and 2013, to investigate the consequences of a continuous shift in its phenology for the degree of overlap among species and the outcomes for plant reproduction, i.e., production of flowers and fruit set, for the next 5 years. We simulated three phenological scenarios with advancement of phenology in 2, 4, and 6 weeks on the *B. laevifolia* flowering, keeping all other parameters set as before.

Results

Climatic analysis

The monthly means of temperature and monthly precipitations varied substantially from 2005 to 2014 (Repeated-Measure One-Way ANOVA F = 3.785; df = 9; p < 0.001for temperature and F = 2.21; df = 9; p = 0.027 for precipitation) (Fig. 1). The Spearman's correlations between the onset of flowering of each species and the monthly temperature and precipitation data for each year studied (2008/2009, 2012/2013, 2013/2014) showed that the species were differently correlated with climatic factors over the years with *P. tomentosa* presenting the highest correlation results and *B. malifolia* as the species suffering less interference from climatic variations (see Table 1).

Species flowering variation

For all species, the mean angles of onset of flowering were significantly seasonal for whole analyzed period (Rayleigh test p < 0.05) and the lengths of mean vectors (r) were nearly 1 (Table 2). The vector r varies from 0 (when phenological activity is distributed uniformly throughout the year) to 1 (when phenological activity is concentrated around one single date or mean angle) (Zar 1996). Comparisons of the mean data among the years showed that the onset of species flowering varied over time (F = 59.176, p < 0.001 for P. tomentosa; F = 48.445, p < 0.001 for B. laevifolia; F = 33.087, p < 0.001 for ; F = 60.729, p < 0.001 for B. campestrisB. malifolia, Table 2).

Community flowering variation

For the first reproductive season (2008/2009) the mean angles of onset and peak for all phenophases were not significantly seasonal (Rayleigh test p > 0.05) and the lengths of mean vectors (r) were nearly 0 (Table 3). These results show that phenological activity of the Malpighiaceae was distributed uniformly throughout the year displaying a segregated phenology. The null model simulations for this reproductive period indicated that the overlap of distributions was significantly less than expected by chance, which confirms the sequential flowering (Pianka = 0.202, p < 0.001; Czechanowski = 0.968, p < 0.001). For the second reproductive season (2012/2013) the mean angle of peak for buds and flowers and onset of fruit were not significantly seasonal (Rayleigh test p > 0.05) and the lengths of mean vectors (r) were nearly 0 (Table 3). These results show that phenological activity exhibited a degree of overlap among some phenophases. In the third year of study, the mean angles were significantly seasonal (Rayleigh test p < 0.05) and the

 Table 1
 Spearman correlations between the date of onset of flowering of *P. tomentosa*, *B. laevolia*, *B. campestris*, and *B. malifolia* with climatic parameters (temperature and precipitation) in a Brazilian Tropical Savanna of Uberlândia, Minas Gerais State, Brazil

Species	Temperature			Precipitation	Precipitation					
	2008/2009	212/2013	2013/2014	2008/2009	2012/2013	2013/2014				
P. tomentosa	- 0.57*	- 0.66*	- 0.71*	- 0.57*	- 0.67*	- 0.47				
B. laevifolia	0.27	0.1	- 0.58*	- 0.44	- 0.59*	- 0.71*				
B. campestris	0.08	0.61*	0.59*	0.76*	0.58*	0.39				
B. malifolia	- 0.59*	0.06	- 0.25	0.22	0.18	0.18				

* Means significant correlation, p < 0.05

Species	Year	N	a ± DP	Mean data	R	$Z\left(p ight)$	Watson- Wil- lian (<i>F</i>)	Р
P. tomentosa	1°	19	154.649° ± 9.775°	03/06/2008	0.986	18.455 (< 0.001)	59.176	< 0.001
	2°	14	$181.009^{\circ} \pm 11.97^{\circ}$	30/06/2012	0.978	13.402 (< 0.001)		
	3°	7	173.858° ± 19.433°	22/06/2013	0.944	6.239 (< 0.001)		
B. laevifolia	1°	30	233.995° ± 8.313°	22/08/2008	0.99	29.375 (< 0.001)	48.445	< 0.001
	2°	20	221.943° ± 13.97°	10/08/2012	0.971	18.846 (< 0.001)		
	3°	16	$207.574^{\circ} \pm 19.615^{\circ}$	26/07/2013	0.943	14.23 (< 0.001)		
B. campestris	1°	26	349.866° ± 15.99°	15/12/2008	0.962	24.052 (< 0.001)	33.087	< 0.001
	2°	13	357.807° ± 14.234°	23/12/2012	0.97	12.222 (< 0.001)		
	3°	5	$11.932^{\circ} \pm 14.772^{\circ}$	12/01/2014	0.967	4.678 (0.003)		
B. malifolia	1°	22	110.053° ± 19.451°	20/04/2008	0.944	19.605 (< 0.001)	60.729	< 0.001
	2°	22	$82.547^{\circ} \pm 9.86^{\circ}$	23/03/2012	0.985	21.358 (< 0.001)		
	3°	17	$111.088^{\circ} \pm 17.195^{\circ}$	21/04/2013	0.956	15.536 (< 0.001)		

 Table 2
 Circular statistical analysis testing for seasonality on floral phenology of *P. tomentosa, B. laevolia, B. campestris, and B. malifolia* in a Brazilian Tropical Savanna over three reproductive seasons (Years 1—2008/2009; 2—2012/2013; 3—2013/2014)

Watson-Williams tests (F) were performed to test the difference among the mean date of onset of flowering along the years

Table 3	Circular	statistical	analyzes	testing f	for seasor	ality o	on pheno	ological	pattern	of co	ommunity	compose	ed by I	P. ton	nentosa,	B . 1	laevolia,	В.
campest	ris, and B	. malifolia	in a Braz	ilian troj	pical sava	ına du	ring thre	e repro	ductive s	seaso	ons (2008/2	2009; 202	2/201	3; 20	13/2014)		

	Flower bud		Flower		Fruit		
	Onset	Peak	Onset	Peak	Onset	Peak	
2008/2009							
Mean data							
Number of observations	110	109	97	90	87	88	
Mean vector (μ)	197,599°	166,697°	183,012°	206,112°	198,797°	219,599°	
Length of mean vector (r)	0.162	0.101	0.174	0.169	0.175	0.177	
Median	225°	225°	225°	240°	240°	255°	
Circular standard deviation	109,408°	122,594°	107,109°	108,082°	106,992°	106,622°	
Rayleigh test (Z)	2.87	1.12	2,945	2,563	2,661	2,758	
Rayleigh test (p)	0.057	0.326	0.053	0.077	0.07	0.063	
2012/2013							
Mean data	03/05/2012	14/05/2012	28/05/2012	09/06/2012	08/06/2012	01/07/2012	
Number of observations	93	93	69	69	64	64	
Mean vector (μ)	123.18°	134.554°	148.521°	160.316°	159.064°	182.271°	
Length of mean vector (r)	0.197	0.13	0.216	0.191	0.188	0.251	
Median	180°	180°	90°	210°	120°	225°	
Circular standard deviation	103.299°	115.796°	100.261°	104.28°	104.702°	95.32°	
Rayleigh test (Z)	3.604	1.565	3.228	2.513	2.269	4.02	
Rayleigh test (p)	0.027	0.209	0.04	0.081	0.103	0.018	
2013/2014							
Mean data	10/05/2013	16/05/2013	04/06/2013	17/06/2013	28/05/2013	15/06/2013	
Number of observations	73	74	45	45	32	32	
Mean vector (μ)	129.454°	135.948°	154.857°	167.452°	147.297°	165.694°	
Length of mean vector (r)	0.276	0.235	0.517	0.483	0.432	0.355	
Median	165°	105°	135°	****	127.5°	150°	
Circular standard deviation	91.88°	97.465°	65.812°	69.1°	74.219°	82.478°	
Rayleigh test (Z)	5.578	4.098	12.029	10.509	5.976	4.029	
Rayleigh test (<i>p</i>)	0.004	0.017	2.74E-06	1.54E-05	0.002	0.017	

Rayleigh test performed the significance of the mean angle and the mean data were omitted when it were not significant

lengths of mean vectors (r) were nearer 1 (Table 3). Linear graphics with the intensity of flowering of Malpighiaceae species showing the degree of overlap among the years were performed (Fig. 2).

Herbivory and productivity analyses

The periods with higher overlap of flowering accounted for lower rates of herbivory for *B. laevifolia* (t = 2.382;

df = 29; p = 0.024) and *P. tomentosa* (t = 2.883, df = 29, p = 0.007). However, there were no significant difference for *B. campestris* (t = 1.821; df = 29; p = 0.079) and *B. malifolia* (t = 0.7957, df = 29, p = 0.4327) (Fig. 3a). In relation to fruit set, the higher overlap period accounted for lower rates of productivity only for *B.laevifolia* (W = 168; n = 30; p = 0.044). There was no significant difference for either *B. malifolia* (W = 147; n = 30; p = 0.114) and *P. tomentosa* (W = 23; n = 30; p = 0.721). *B. campestris* presented an



Fig. 2 Linear graphics with the intensity of flowering of *Peixo*toa tomentosa, Banisteriopsis laevifolia, Banisteriopsis campestris, and Banisteriopsis malifolia during 2008/2009, 2012/2013, and

2013/2014 in a Brazilian Tropical Savanna (axis y means intensity of flowering phenology being 1 = 25% and 2 = 75%; n = 30)



Fig. 3 Comparative analysis of herbivory (leaf area loss) **a** and fruit production, **b** of *Peixotoa tomentosa*, *Banisteriopsis laevifolia*, *Banisteriopsis campestris* and *Banisteriopsis malifolia* in a Brazilian Tropical Savanna between the period with lower (2008/2009) and greater

Table 4 Total percentual errors given by the integral of the squared error between the real and simulated data, on a random simulation scenario

Total percen- tual squared error (%)	B. malifolia	P. tomentosa	B. laevifolia	B. camp- estris
F. buds	1.22	4.11	2.46	1.10
Flowers	1.71	3.69	3.99	5.51
Fruits	1.65	9.03	3.16	9.45

increase in productivity during the period with higher overlap, however, there was no significant difference (W = -62; n = 30; p = 0.219) (Fig. 3b).

Mathematical modeling

The CSVM program obtained a mean squared error of 3.087% when compared to the mean field data response, achieving a close fit between the model and the real-case scenario. All mean percentage squared error achieved for each species after the optimization procedure are exhibited in Table 4. The results of advance in the phenology of *B. laevifolia* demonstrated a tendency in increasing the production of flowers, however, a proportionally lower production of fruits (Fig. 4a). For *P. tomentosa*, the model indicated a tendency in increasing amounts of both flowers and fruits (Fig. 4b). There were no tendencies of variation for *B. malifolia* and *B. campestris* (Fig. 4c, d, respectively).

Discussion

Our results present climatic variation in annual mean of temperature and precipitation from 2005 to 2014 and detected correlations with the onset of plant flowering during the three reproductive seasons studied (2008/2009, 2011/2012,



overlap of flowering (2013/2014). (Test Paired T in (**a**) and Wilcoxon test in (**b**) (*) means statistical difference; p < 0.05, Means \pm 1SE are presented)

2013/2014). Shifts on the onset of flowering resulted in a greater overlap among Malpighiaceae species during the third reproductivity period analyzed, corroborating our first hypothesis. We also demonstrated that degree of plant phenological overlap (staggered or synchronized) influenced the herbivory and fruit set of species, corroborating our second hypothesis. The mathematical model indicated a tendency in flower and fruit production for each plant species contributing to a better understanding of how the phenology may vary on a continuing climate change (see a movie produced in CSVM program as supplementary material).

The correlations between onset of plant flowering, temperature, and precipitation suggest an influence of these climatic parameters on the phenology of plant species. Plants may advance or delay their reproductive period in a seasonal climate by avoiding the sprouting of new leaves before or during unfavorable periods (e.g., water stress) or by producing new leaves to coincide with the onset of periods of favorable conditions (e.g., high radiation) (Van Shaik et al. 1993; Del-Claro and Torezan-Silingardi 2012). The impacts of climate change on phenological shifts are among the most documented process occurring during the last years, as demonstrated by several studies (e.g., Forrest et al. 2010; Forrest and Thomson 2011; Rafferty and Ive 2012; Diez et al. 2012; Liuth et al. 2013), especially the effects of rainfall and temperature in tropical climates (e.g., Morellato 2013; Bock et al. 2014). These environmental factors can affect the growth and development of the plants under different forms during the phenological phases, whereas solar radiation, and relative humidity influence many physiological processes related to flowering patterns (see Camargo 2010).

Changes in flowering patterns are likely to affect plant reproduction if competitive or facilitative interactions between plant species affect their associations with interacting animal partners (see Gross et al. 2000; Lazaro et al. 2009; Forrest et al. 2010). Annual variation on plant phenologies in subalpine meadows in Colorado, USA,



Fig. 4 Results of flower and fruit production for *Banisteriopsis laevifolia* (a) *Peixotoa tomentosa* (b), *Banisteriopsis malifolia* (c), and *Banisteriopsis campestris* (d) in four different phenological simula-



Peixotoa tomentosa

tions, considering an anticipated phenology of *B. laevifolia* in 2, 4, and 6 weeks. Default means the standard setting

produced a higher overlap among plant species leading to increased sharing of pollinators by some species (e.g., Forrest et al. 2010) and decreased fruit set for other species due to their decreased synchrony with pollinators (Thomson 2010). Rafferty and Ives (2011) showed that species experimentally induced to flower earlier in a wildflower community received more pollinator visits and Rafferty and Ives (2012) simulated changes in phenology of two perennial wildflowers and demonstrated that the seed set can vary as a result of changes in pollinator taxonomic composition and effectiveness. Therewith, flowering phenology is commonly thought to affect plant reproduction through the influence of pollinators, as well as the degree of plant dependence on this interaction (see Johnson et al. 2011).

In our system, all species are self-compatible, they can require animal pollinators to move pollen or reproduce via spontaneous autogamy, however, the rate of autonomous self-pollination varies greatly among species (Torezan-Silingardi et al. in preparation). Fruit set via spontaneous self-pollination in *Banisteriopsis laevifolia* and *B. campestris* is about only 2%, while in *P. tomentosa* this value can range up to 43%, and in *B. malifolia* accounts for about 30% of the fruit set. In this sense, the species of Malpighiaceae were affected differently by the greater overlap of flowering period with *B. laevifolia*, the only significantly affected in a negative way. Based on these results, we can suggest that *P. tomentosa* and *B. malifolia* presented a more robust response to a greater overlap of phenology due to their relative independence of pollinators. *Banisteriopsis campestris* cannot be considered in this line of thought because its flowering separated from the core of other species. Therefore, the staggered strategy is more essential for *B. laevifolia* due to its greater dependence in relation to pollinators. Bishop and Schemske (1998) and Moeller (2006) have demonstrated and discussed the importance of self-compatibility by increasing fruit and seed production as an important driver of plant mating system evolution.

Our results also showed that the herbivory rate was significantly reduced for P. tomentosa and B. laevifolia and not significant for B. malifolia and B.campestris in a scenario with greater overlap. Therefore, the effects of overlap of plants in reducing herbivory were corroborated and it is likely that the effect on *B. malifolia* and *B.campestris* was lower due to its less overlapping than *P. tomentosa* and *B.* laevifolia. In addition, these species flourish in the rainy season, the period with the greatest abundance and activity of herbivore insects (e.g., Alves-Silva et al. 2013, 2014; Alves-Silva and Del-Claro 2013; Vilela et al. 2014). Thus, we suggest that greater synchrony between P. tomentosa and B. laevifolia may have allowed the occurrence of dilution of herbivory (Herbivore Satiation Hypothesis) (e.g., Clark and Clark 1991; English-Loeb and Karban 1992). A classic study of Aide (1991) investigated this effect for leaves growing synchronously within the population of the sub-canopy tree Gustavia superba (Lecythidaceae) and found out that individuals out of synchrony with their conspecifics suffered significantly more herbivore damage. It is also possible that herbivory rate diminished due to less synchrony between herbivores and flowering time. The importance of synchrony for effective herbivore consumption was already evidenced for larva of Lepidoptera feeding on leaf flush from experimental studies (e.g., Hunter and Elkinton 2000; Van Asch et al. 2007).

The mathematical model we developed was able to demonstrate with great precision (mean squared error $\approx 3\%$) any given scenario with variations in the rainfall or herbivory for the species considered. In scenario extrapolations with input not directly linked to the species themselves (i.e., rainfall, herbivory, etc.) the model is also able to make very precise *predictions*, which has been optimized and validated by field measured data. In the opposite way, when the variations imposed as entries are directly related to the species (i.e., flowering), the results should be interpreted as a *tendency* with an unknown error, due to unavailability of real field data for comparisons and model adjustment. We considered variations in the onset of plant flowering and, therefore, we have to interpret the results as a valid tendency. The simulations showed a species-specific response, also demonstrated by Diez et al. (2012) to three North American plant communities in response to climate change.

In our study, the most pollinator-dependent species, B. laevifolia, presented a tendency in increasing the production of flowers possibly as a result of the dilution in herbivore pressure (e.g., Bishop and Schemske 1998) or lowering of herbivore synchrony with plant flowering (e.g., Forrest and Thomson 2011). However, the greater synchronization with P. tomentosa and possible sharing of pollinators associated with its dependence on these vectors resulted in a tendency of reduced production of fruits. On the other hand, plants possessing the combination of simultaneously autonomous self-pollination and escape from herbivores, as demonstrated by P. tomentosa, were able to keep a disproportionate fitness advantage over plants possessing any of the other possibilities (see Herrera et al. 2002). Thus, within communities, variation in species responses may determine communitylevel patterns of phenology and resilience of species (Memmott et al. 2007; Kaiser-Bunbury et al. 2010).

We demonstrated variations in climate parameters over the recent years and the manifestation of these changes on plant phenology and biotic interactions of different plant species from the Southern Hemisphere. There is a concern that climate change is altering the timing of life-history events in a wide array of species, especially for species from Northern Hemisphere. However, few papers have related the effects of climatic change on plant phenology and interactions of South American species (Morellato et al. 2013). Our findings of significant variability in species responses to their interacting animal partners suggest that plasticity in these traits will be important for the survival and reproduction of these plant species in the future. Thus, integrating phenological investigations in the field with the existing theories and mathematical modeling allow us to interpret different phenological synchronization degrees and enable us to describe the possible maintenance and evolution of life histories in response to climate change.

Acknowledgements We thank the three anonymous reviewers for reviewing the manuscript. We also thank CNPq and FAPEMIG for funding this research (KDC/HMTS/VTSDC) and FAPEMIG for award-ing fellowships to AAV.

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