Variation in developmental time affects mating success and Allee effects

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A fundamental question in biological conservation and invasion biology is why do some populations go extinct? Allee effects, notably those caused by mate location failure, are potentially key factors leading to the extinction of sparse populations. Several previous studies have focused on the inability of males and females to locate each other in space when populations are at low densities but here we investigate the effects of differences in the timing of male and female maturation on mating success. We develop a generalized model to clarify the role of protandry (the appearance of males before female emergence) and variability in adult maturation times. We show that temporal asynchrony can substantially reduce the probability of successful mating. We then apply this generalized model to estimate mating success in invading populations of the gypsy moth in North America in relation to local climate and its associated seasonality. Considerable geographic heterogeneity was observed in simulated mating success and this variability was not correlated with previous evaluations of bioclimatic requirements and habitat suitability. Furthermore, we found that the generalized model of temporal asynchrony provided reliable predictions and that detailed modeling of gypsy moth development was not necessary.
frequent cause of Allee dynamics. A few studies have investigated how mate-location failure may contribute to Allee dynamics but these studies have largely focused on the spatial isolation of individuals: widely dispersed males and females are typically limited in their abilities to locate each other for mating (Hopper and Roush 1993, Wells et al. 1998, Berec et al. 2001, South and Kenward 2001). However, in addition to dispersion in space, mates may be segregated in time (seasonally) and this could greatly impact mating success. The impact of temporal dispersion may be particularly acute in species where reproduction occurs during a discrete season. In particular, for many animal and plant species, the timing of female adult maturation occurs slightly after peak male maturation (Fagerström and Wiklund 1982). This sexual dimorphism in the timing of maturation is referred to as “protandry” (Morbey and Ydenberg 2001) and it may limit mating success, particularly at low population densities.

Calabrese and Fagan (2004) provide the most extensive investigation to date of the role of temporal dispersion and protandry in generating Allee dynamics. They developed a model of mating probability based upon two parameters that characterize the temporal overlap between males and females: the length of the population-level breeding period and the duration of the individual reproductive period. The potential for asynchrony was used to calculate the probability that a female was mated and its consequence on the growth rate of the population. They demonstrated that sexual asynchrony and protandry generated a loss of reproductive potential and contributed to an Allee effect, resulting in population decline at low densities.

While Calabrese and Fagan (2004) broke new ground in demonstrating the potential for developmental dispersion and protandry to affect mating success and contribute to Allee dynamics, they did not explore the underlying mechanism that governs this temporal variability. While both dispersion in maturation time and protandry are life history attributes, they may be quite plastic and subject to environmental influences. In particular, the development of poikilothermic organisms is largely determined by temperature. Therefore, weather conditions affect the timing of sexual maturation, consequently influencing mating success and expression of an Allee effect. Since climate varies geographically and temporally, this variability could impose limitations in the establishment of invading species.

In this paper, we model mating success as a function of both variation in developmental time (within each sex) and protandry to demonstrate that both strongly influence mating success. In contrast to the approach of Calabrese and Fagan (2004), we explicitly modeled mating probabilities in discrete time steps instead of considering mating probability as a function of the temporal overlap. We consider a generalized model and then apply it to populations of the gypsy moth, Lymantria dispar, (Lepidoptera: Lymantriidae), a native of Eurasia that is currently in the process of invading North America. We examine how variation in climate throughout North America influences gypsy moth mating success and speculate on how this might impact its ability to invade new habitats. We report that the generalized model successfully predicts gypsy moth mating success given appropriate values of protandry and dispersion of maturation times. We also found that mating failure is likely to contribute to the occurrence of Allee effects, even in relatively dense populations.

Methods
To explore the effects of temporal variation in maturation on mating success, we developed a generalized model and a gypsy moth-specific model. Both were deterministic, but they differed in the way that maturation times were calculated. The general model represented the distribution of developmental times using a Gaussian function with known variance but the gypsy moth model utilized a previously developed phenological model that predicted the distribution of maturation times based upon daily temperatures. Simulations were performed using the R language (Anonymous 2006).

Generalized model
The failure of opposite sexes to locate each other for mating at sparse population levels is affected by both the temporal and spatial isolation of males and females. For simplification, we did not explicitly model spatial isolation and instead focused on the temporal separation, though spatial isolation was implicitly accounted for in our model (Instantaneous mating success, below).

Distribution of maturation times
Our model was generalized to represent a univoltine species in which mating occurs during a period that follows adult sexual maturation. To model this, we generated synthetic maturation distributions, using Gaussian density functions with varying standard deviations (SD) to simulate the dispersion of maturation of each sex, and varying differences between the mean time of male and female maturation (AT) to simulate protandry. Hereafter, parameters $em_d$ and $ef_d$ refer to the rate at which males and females become sexually mature on day $d$.

Individual longevity
A second factor included in the model was adult longevity. We arbitrarily assumed 50% daily survival
of each sex though we experimentally tested how variation in longevity impacts population-level mating success. Hereafter survm and survf refer respectively to the daily survival rates of males and females. We also assumed that the maximum longevity of males and females was three days. Adult longevity varies considerably among species and for many species it may greatly exceed three days. However, in these cases, we do not expect that our simplification of longevity would qualitatively alter the relationships explored here.

**Instantaneous mating success**

The third component was a model to calculate the probability of mating given the instantaneous abundance of sexually mature males and females. This model was not spatially explicit, but it accounted for the ability of mates to locate each other in space at a given abundance. Various models of mating success have previously been applied: negative exponential models, rectangular hyperbolic models and power complement models (reviewed by Boukal and Berec 2002). Since these models exhibit very similar behavior (with slight differences in biological representation), we focused only on the negative exponential, assuring no loss of generality. This model assumes that males and females are randomly distributed in space, so the probability of mate encounter is derived from the Poisson probability distribution (Boukal and Berec 2002). Thus, the probability that a female is mated on day \( d \) is given by:

\[
P_d = 1 - \exp(-se_{Md})
\]

where \( s \) is the daily probability that a female is mated in the presence of a single male and \( M_d \) is the number of males potentially able to mate the female on day \( d \), or in other words: the number of males becoming sexually mature on day \( d \), plus the number of males becoming sexually mature on day \( d-1 \) which survived one more day, plus the number of males becoming sexually mature on day \( d-2 \) which survived two more days. We calculated the abundance of mature males \( M_d \) available on each day based upon the total number of males \( E \), the distribution of sexual maturation times \( (em_d) \) and male longevity \( \text{survm} \).

**Combining instantaneous mating success with longevity and maturation times**

When a female lives from day \( d \) to day \( d+t \), we sum the number of mature males it can encounter on each day from \( d \) to \( d+t \). Thus, its mating probability is given by:

\[
P_{dt} = 1 - \exp(-s(M_d + \ldots + M_{d+t}))
\]

For each day \( d \), we calculated the mating success of a female which became sexually mature on day \( d \): either it will not survive to day \( d+1 \) (the proportion of such females is \( e_{fd} (1 - \text{survf}) \)), or it will survive only to day \( d+1 \) (the proportion of these females is \( e_{fd} \text{survf} (1 - \text{survf}) \)), or it will survive to day \( d+2 \) (the proportion of these females is \( e_{fd} \text{survf}^2 \)). The global probability of mating for the entire population is finally given by:

\[
MS = \sum_d e_{fd}((1 - \text{survf})P_{d,0} + \text{survf}(1 - \text{survf})P_{d,1} + \text{survf}^2P_{d,2})
\]

**Effect of asynchrony on the mating success**

We generated synthetic data in which distributions of male and female sexual maturation dates were based on normal distributions with SD varying across 100 levels between 0.1 and 10 days and with differences in mean male and female emergence times (\( \Delta T \)) varying across 11 levels between 0 and 10 days. Then, we used the deterministic model previously described to evaluate the population-level mating success and investigate the effects of both protandry and dispersion in maturation time. We assumed a total number of males \( E = 100 \) and we tested different values of \( s \) (\( s=0.1 \) and \( s=0.5 \) corresponding to low and high mate-finding efficiency respectively).

**Effect of longevity rates**

To examine the effect of adult survival on mating success, we assumed \( SD = 6 \) days and \( \Delta T = 6 \) days. We simulated the population-level mating probability for different survival rates of males and females \( (\text{survm}, \text{survf} = 0.2, 0.5, \text{and} 0.8) \), and also for varying numbers of males \( (\text{from} E = 10 \text{ to } 100, \text{by an increment of } 10) \) for two cases: \( s = 0.1 \) and \( s = 0.5 \).

To compare the relative importance of male and female survival rates, we considered \( E = 100 \) and \( s = 0.1 \), then we simulated the mating success for various male survival rates \( (\text{survm} = 0.1 \text{ to } 0.9 \text{ by } 0.01, \text{with} \text{survf} = 0.5) \) and for various female survival rates \( (\text{survf} = 0.1 \text{ to } 0.9 \text{ by } 0.01, \text{with} \text{survm} = 0.5) \). We applied linear regression to the results of both sets of simulations \( (MS \sim \text{survm} \text{ and } MS \sim \text{survf}) \) and compared the slopes in order to evaluate differences between how male and female survival affect mating success.

**Case study: invading North American gypsy moth**

**Study system**

The gypsy moth is native to temperate Eurasia but was introduced in North America near Boston in 1869 (Riley and Vasey 1870, Forbush and Fernald 1896). The species has considerably expanded its range but currently occupies slightly less than 30% of its suitable habitat in North America, as defined by forest
comprising (Morin et al. 2004), and less than 20% of the area with highly suitable climate (Gray 2004). New isolated colonies are frequently discovered far from its current North American distribution, mostly as a result of accidental movement of egg masses by humans (Mason and McManus 1981). Since North American gypsy moth populations have been the subject of massive amounts of research, extensive biological data exists for parameterization of the mating success model described above.

Emergence curves

Because the reproductive period of the gypsy moth begins as soon as adults emerge, we considered the distribution of emergence times to represent the distribution of maturation times. We simulated the timing of male and female emergence (emd and efd) using a previously described phenological model, Gypsy moth life stage model (GLS ver. 3.0; Gray 2004). The GLS model is a composite phenological model, which combines three separate models of egg development for prediapause (Gray et al. 1991), diapause (Gray et al. 2001) and postdiapause development (Gray et al. 1995), with models for early larval stages (Logan et al. 1991) and late larval to adult stages (Sheehan 1992). Régnière and Nealis (2002) applied GSL to weather station data and found that predicted timing of both egg hatch and adult emergence closely agreed with the timing of these events observed in the field.

The GLS model was used to predict daily emergence rates for males and females based upon temperature data (daily minimum and maximum) from the time of oviposition (summer) until the time of adult emergence the following summer. Because our simulation was designed to represent a scenario in which an isolated gypsy moth population was founded by an egg mass(s) transported from a single source location, we did not initiate the GLS model with oviposition dates specific for each site where the egg mass was moved to, but instead we used a single oviposition date that would be typical of a single source location in the infested area (1 August).

We applied the model to daily temperature data (minimum, maximum) collected at 7591 meteorological stations throughout North America from 1960–2000 (obtained from the Global Historical Climatology Network; Peterson and Vose 1997). Adult emergence curves were generated for each station in each year for which data were available. The model requires temperature data from two consecutive years starting from the date of oviposition until the date of adult emergence. If data from a given weather station were missing on more than 15 consecutive days, GLS fails to produce a prediction, and therefore emergence curves were not calculated. In more occasional instances of missing data, GLS estimates replacement temperature values through interpolation (Gray 2004).

The GLS model accounts for variability in thermal requirements for development within a population. As such, it tracks the development of individuals that complete development to the adult stage, as well as those that do not. Because the model predicted that significant proportions of populations at some locations would not complete development, we transformed predictions of daily percentages of starting populations into proportions of males and females that summed to one for each sex over all emergence dates. These proportions defined the daily emergence probabilities emd and efd specific for gypsy moth development under each set of temperature conditions. Gray (2004) showed that northern regions of North America are climatically marginal for gypsy moth establishment because late oviposition, slow larval development, and early winter temperatures combine to arrest development in the non cold-hardy prediapause phase. Southern regions of North America are climatically marginal because high winter temperatures are less suitable for diapause development. When the GLS model predicted <5% of the population would successfully emerge, we rounded this to zero and assumed that there was no emergence. We found that inclusion of these marginal populations resulted, in some cases, in a gross overestimation of mating success (artificially generated by the model).

Individual longevity

Based upon previous experimental studies, we assumed that maximum male longevity was three days and their daily survival rate was survm = 0.19 (Elkinton and Cardé 1980), and that maximum female longevity was three days with a daily survival rate of survf = 0.48 (Sharov et al. 1995). The contribution of females >3 days old is negligible since fertility declines drastically in females older than three days (Richerson et al. 1976).

Model of mating success

We used the negative exponential model (Eq. 1) to calculate daily mating success based upon male abundance derived from male emergence curves predicted by the GLS model. This instantaneous mating success model was fit (s = 0.15) by Sharov et al. (1995) using daily field measurements of female mating success and male abundance from several locations. At each location, mating success was measured by dissection of 18 deployed females and male abundance was measured by the average capture of four pheromone traps. Pheromone-baited traps are a standard method for assessing sparse gypsy moth populations; they are very sensitive at low densities (Elkinton and Cardé 1981), and at these densities they are closely related to population density.
Thus, we expressed gypsy moth abundance throughout this paper in terms of males per trap. While the daily mating success model (Eq. 1) uses male abundance from a single day, the overall objective of this case study was to estimate the relationship between mating success and abundance as measured by season-long trap capture.

Combining instantaneous mating success with longevity and emergence times
Abundance of males on each day \((M_d)\) was derived from the season-long trap capture \((TC)\):

\[
TC = M_1 + \ldots + M_n + M_{n+1} + M_{n+2}
\]

where \(d = 1\) is the first day of emergence, and \(d = n\) is the last day.

But these \(M_d\) daily male abundances (measured as daily trap capture), will themselves be composed both of males that emerged that day, as well as surviving males from previous days:

\[
M_d = \text{em}_d\ E + \text{survm}_{d-1}\ E + \text{survm}_{d-2}\ E,
\]

where \(\text{em}_d\) is the rate of males emerging on day \(d\) and \(E\) the total number of males. Even though \(E\) is expressed in terms of trap capture, it differs from season-long trap capture \((TC)\) because it represents the total number of males that emerged during the entire generation. In contrast, \(TC\) includes \(E\) plus males that survived to subsequent days. Since \(\text{em}_1 + \ldots + \text{em}_n = 1\), we can deduce that \(TC = (1 + \text{survm} + \text{survm}^2)\ E\) and more precisely, \(TC = 1.23\ E\) in our case. The various parameters required in the gypsy moth model as well as the generalized model are listed in Table 1.

Simulations were carried-out for various values of season-long trap capture \((TC = 10\) to \(1000\) by \(10)\), to estimate the relationship between abundance and population-level mating success. These simulations were repeated using temperature data from four locations: Farmville, Virginia and Lakewood, Wisconsin, were located within the current range of the gypsy moth; and Santa Barbara, California and Jackson, Wyoming, were located outside the range distribution. For each location, we calculated the mean mating success over 1960–2000. Also, we fixed season-long trap capture at \(TC = 100\) and calculated the mean mating success using temperature data from each of the 7591 North American meteorological stations from 1960–2000 to examine the geographical variation in mating success, both in areas that the gypsy moth has successfully colonized and in areas not yet invaded.

Results

Generalized model

Simulations using Eq. 1, 2 and 3 with Gaussian distributions for sexual maturation times illustrate how temporal asynchrony between male and female maturation affects female mating success (Fig. 1a). The daily mating probability \((P_d)\) is not in complete phase with the distribution of male maturation times; the curve is extended to the right because of male longevity. The distribution of mated females is shifted quite strongly to the right of the daily mating probability distribution because it is strongly affected by the distribution of emerging females. Earlier emerging females are much more likely to be mated than late emerging females due to a decline in available males. In the case of strong protandry, mainly the first emerging cohort of females contributes to the subsequent generation.

Effect of longevity

Male survival rates affect the number of males available to mate each day, and consequently the daily mating probability \((P_d\) calculated from Eq. 1). Daily mating probability is significantly depressed when male survival is reduced (Fig. 1a–b). The population-level mating success \((\text{MS})\) increased from \(0.46\) to \(0.66\) as a result of the increase in \(\text{survm}\) from 0.2 to 0.8 (with \(\text{survf} = 0.5\), Table 2). This increase in mating success can be explained by both the increased overlap with females and the elevated availability of males each day. Female survival rates affect the number of days during which they can potentially encounter mature

<table>
<thead>
<tr>
<th>Parameters</th>
<th>Description</th>
<th>Models</th>
</tr>
</thead>
<tbody>
<tr>
<td>(s)</td>
<td>Instantaneous mating probability in presence of one mature male.</td>
<td>G &amp; GM</td>
</tr>
<tr>
<td>(\text{survm}) ((\text{survf}))</td>
<td>Daily survival rate of males (females) during the reproductive period.</td>
<td>G &amp; GM</td>
</tr>
<tr>
<td>(\text{em}_d) ((\text{efd}))</td>
<td>Daily rate of males (females) becoming sexually mature.</td>
<td>G &amp; GM</td>
</tr>
<tr>
<td>(\Delta T)</td>
<td>Protandry (delay between male and female mean time of sexual maturation); required to calculate (\text{em}_d) and (\text{efd}).</td>
<td>G</td>
</tr>
<tr>
<td>(SD)</td>
<td>Variability in development times (standard deviation of sexual maturation times); required to calculate (\text{em}_d) and (\text{efd}).</td>
<td>G</td>
</tr>
<tr>
<td>(E)</td>
<td>Total number of males.</td>
<td>G</td>
</tr>
<tr>
<td>(TC)</td>
<td>Season-long trap capture (cumulative number of daily abundance of mature males).</td>
<td>GM</td>
</tr>
</tbody>
</table>
males and high survival therefore leads to increased mating success. Since the daily mating probability given by Eq. 1 is independent of female abundance, the mating probability curve does not change with increased female longevity (Fig. 1a, 1c). However the probability that a female is mated during its reproductive period does increase slightly from MS = 0.49 to 0.65 when the female survival rate increases from 0.2 to 0.8 (with survm = 0.5, Table 2).

Though both male and female survival rates were positively related to mating success (Fig. 2, Table 2), the effect of male survival was 27% stronger (the slope estimate was 0.33, p < 0.001, for the linear regression MS ~ survm and 0.26, p < 0.001, for MS ~ survf). Mating success is obviously higher for greater levels of mate searching efficiency (s = 0.5 vs 0.1) and it is positively related to the abundance of mature males E (Fig. 2). When mate searching efficiency is high (s = 0.5), effects of survival rates on mating success are more important at low abundance of males. Mating success converges rapidly toward an upper limit with increasing number of males. In contrast, longevity is more important at high male abundance when mate-searching efficiency is low (s = 0.1).

**Effect of asynchrony on the mating success**

Both the dispersion in developmental times (SD) and differences between male and female development (protandry, DT) affected mating success (Fig. 3). Success declined as either value increased. The effect of protandry was most pronounced when dispersion was low as even small differences in developmental times between the sexes almost completely eliminated mating. The intensity of the effect of protandry on mating success increased as mate searching became less efficient (s = 0.1 vs s = 0.5, Fig. 3a vs 3b). Finally, the

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**Table 2. Effects of adult longevity on mating success simulated by the generalized model (with s = 0.1, E = 100 males, SD = 6 days and AT = 6 days).**

<table>
<thead>
<tr>
<th>survf/survm</th>
<th>0.2</th>
<th>0.5</th>
<th>0.8</th>
</tr>
</thead>
<tbody>
<tr>
<td>0.2</td>
<td>0.39</td>
<td>0.49</td>
<td>0.60</td>
</tr>
<tr>
<td>0.5</td>
<td>0.46</td>
<td>0.56</td>
<td>0.66</td>
</tr>
<tr>
<td>0.8</td>
<td>0.55</td>
<td>0.65</td>
<td>0.74</td>
</tr>
</tbody>
</table>

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**Fig. 2. Effect of varying survival rates on mating success using synthetic Gaussian emergence distributions (SD = 6 days, AT = 6 days) and different values for s (s = 0.1 and 0.5 corresponding respectively to a strong and weak Allee effect), with E = 100 males.**
influence of dispersion in developmental time on mating success was relatively high in comparison to the effect of survival rates (Fig. 2).

**Case study: invading North American gypsy moth**

Simulations based on weather station data indicated that mating success was non-uniform across North America (Fig. 4). Mating success could not be simulated in extremely northern and southern locations because the GLS model predicted extremely low or no emergence of adults (locations represented by grey stars on Fig. 4). The map of mating success clearly showed a central zone stretching across the USA in which mating was expected to be optimal. In contrast, mating success was predicted to be particularly low in other parts of North America, especially in mountainous regions in the west. We examined average mating success in more details (for TC = 100 males per trap) at two locations within the current range of the gypsy moth: Farmville, Virginia and Lakewood, Wisconsin; and two locations where the gypsy moth has not yet colonized: Santa Barbara, California and Jackson, Wyoming (Fig. 4). Mating success was greatest at Farmville and lowest in Santa Barbara (Fig. 5, Table 3). These differences between locations can be explained in terms of both dispersion in developmental times and protandry (Fig. 3c). The low mating probability in Jackson was primarily due to a long delay between peak emergence of males and females (ΔT = 8.4 day average) and a

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**Fig. 3.** Mating success resulting of normal distribution of sexual maturation times, with various standard deviations SD (y-axis) and delays between males and females ΔT (x-axis): (a) s = 0.1 (low mate-finding efficiency), survm = survf = 0.50, E = 100 males; (b) s = 0.5 (high mate-finding efficiency), survm = survf = 0.50, E = 100 males; (c) s = 0.15, survm = 0.19, survf = 0.48 (values for gypsy moth), TC = 100 males. Stars symbols refer to four stations (Farmville, Virginia; Lakewood, Wisconsin; Santa Barbara, California and Jackson, Wyoming). We calculated the mean standard deviation of male emergence curves and the mean delay between male and female emergence peaks using the GLS model.
relatively low dispersion of emergence dates (SD = 6.7 days). The low mating probability in Santa Barbara was apparently the result of a different behavior: the delay between mean male and female emergence was shorter (AT = 5.2 day average), but mating apparently failed because there was so much dispersion in emergence dates (SD = 13.6 days). One consequence of the Mediterranean Santa Barbara climate is that temperatures experienced during diapause development are relatively warm and this is known to increase the variability in timing of egg hatch (Masaki 1956), which presumably would lead to highly variable adult emergence times as well. Even though this high variability allows male and female reproductive periods to overlap, it causes a severe decrease in the daily density of mature males. Since the mating probability increases exponentially with the daily number of mature males (Eq. 1), increased dispersion in the emergence curve is unfavorable for mating success.

Mating success values predicted by the generalized model are highly consistent with the values predicted by the more realistic gypsy moth-specific model. Absolute errors were below 0.01 for prediction of the mean mating probability based on the mean protandry and mean variability in males development times (Fig. 3c, Table 3). When we tested the normality of emergence times (using a Jarque Bera test which is based on skewness and kurtosis coefficients, and rejecting normality when p < 0.05), we found that these data were not normally distributed in more than 60% of the years, for each of the four locations. Despite this result, both models provided approximately the same result. This result is important because it indicates that it is not necessary to perform detailed simulations (such as performed here for the gypsy moth) to estimate mating success with an acceptable level of accuracy. Instead, one only needs estimates of the variability in development time and protandry and then mating success can be directly estimated from maps, such as those given in Fig. 3.

Simulations indicated that, when season-long trap capture was below 100 males per trap, mating success rarely exceeded 60% (Fig. 5). The relative magnitude of mating success varied considerably among locations with different climates, but the same pattern of increasing mating success with increasing abundance (measured by trap capture) was evident from simulations using data from all sites. Even at an apparently favorable location, where gypsy moth populations have been established for several years (Farmville, Virginia), mating success was less than 50% when trap capture was below 60 males per trap. Mating success converged toward an upper limit when season-long trap capture reached 1000 males per trap (which corresponds to the upper bound of season-long trap capture). This limit was approximately 0.90 in Farmville, Lakewood and Santa Barbara but considerably lower, around 0.77, in Jackson. This result suggests that protandry and dispersion in development time act differently. In Santa Barbara, dispersion in development time was particularly high and mating success was quite low at low population density but increased to the same limit as in favorable sites. In contrast, emergence curves in Jackson were characterized by a very long delay between male and female mean emergence time. This protandry seems slightly more favorable than the long dispersion of developmental time at low population density, but mating success is substantially lower than in other locations at high population density.
Table 3. Estimation of the mean mating success over n years in four locations, resulting from the gypsy moth specific model, MS(GM), and from the generalized model, MS(G) using the mean standard deviation of the males emergence curve (SDM), and the mean difference in the mean emergence time between females and males (ΔT) to determine the mating success on Fig. 3c.

<table>
<thead>
<tr>
<th>Locations</th>
<th>n</th>
<th>ΔT ± SD</th>
<th>SDm ± SD</th>
<th>MS(GM) ± SD</th>
<th>MS(G)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Farmville, Virginia</td>
<td>38</td>
<td>3.5 ± 0.43</td>
<td>2.8 ± 0.5</td>
<td>0.62 ± 0.06</td>
<td>0.63</td>
</tr>
<tr>
<td>Lakewood, Wisconsin</td>
<td>26</td>
<td>4.5 ± 0.69</td>
<td>3.7 ± 0.7</td>
<td>0.56 ± 0.04</td>
<td>0.57</td>
</tr>
<tr>
<td>Jackson, Wyoming</td>
<td>35</td>
<td>8.4 ± 2.68</td>
<td>6.7 ± 2.3</td>
<td>0.42 ± 0.07</td>
<td>0.43</td>
</tr>
<tr>
<td>Santa Barbara, California</td>
<td>22</td>
<td>5.2 ± 0.43</td>
<td>13.6 ± 2.9</td>
<td>0.37 ± 0.05</td>
<td>0.36</td>
</tr>
</tbody>
</table>

To more fully understand the causes of mating failure, we illustrate a simulation from a single location (Farmville, Virginia) in a single year (1997) (Fig. 1d). Females emerging early had a very high mating probability (approaching 100%) since their emergence occurred during the period of highest male emergence. Females emerging later had very low mating success (approaching 0%). These females that emerge at particularly late dates greatly diminish the population-level mating success.

Discussion

Reproductive asynchrony has been mainly considered adaptive for populations, notably in cases of strong environmental stochasticity (Wiklund and Fagerström 1977, Waldbauer 1978, Simmons and Johnston 1997). However, this study supports the conclusion drawn by Calabrese and Fagan (2004) that asynchrony can actually generate an Allee dynamics at low densities, which could contribute to extinction. It is worth noting that the main contributor to low mating success appears to come from individuals with particularly fast or slow development (Fig. 3a–b). However, it has also been speculated that environmental stochasticity may cause selection for developmental outliers (Post et al. 2001). Thus, natural selection for this life-history trait would result from two opposing forces: (1) mate-location that could trigger selection against protandry, and (2) various factors traditionally considered as beneficial (e.g. an increase in the male mating opportunity or decrease in the waiting cost of females, Morbey and Ydenberg 2001). Since protandry is an adaptive behavior that has been maintained for thousands of years throughout biological evolution in many species, the negative effects on mate location described here must be offset by even stronger positive selection for this trait.

Our model indicates that under virtually all situations, temporal asynchrony is likely to cause increased mating failure at lower densities. Such Allee effects have been shown to be important determinants of the establishment of newly founded populations and the spread of populations following establishment (Lewis and Kareiva 1993, Taylor and Hastings 2004). In particular, Allee effects have been shown to be critical in determining the establishment and spread of invading gypsy moth populations (Liebhold and Bascompte 2003, Johnson et al. 2006, Tobin et al. 2007). The results of our simulations thus indicate that mate location may be an important contributor to Allee effects. Of course, mating success represents just a single component of the demographics of any population; as such, quantification of Allee effects cannot be derived alone from the results of this study. However, these results do support the concept that mating failure is a major contributor to an Allee effect in the gypsy moth. In the future, it should be possible to combine the results of this study with other demographic models of gypsy moth dynamics that account for all sources of mortality and reproduction, to estimate Allee effects.

Recent analyses of historical gypsy moth invasion data indicate that geographical variation in Allee effects influence geographical variation in rates of gypsy moth spread (Tobin et al. 2007). In this study, we have shown that mating success can be expected to exhibit considerable geographical variability, with a central zone stretching across the USA where mating rates are highest (Fig. 4). Most of the current distribution of the gypsy moth in North America lies outside of this optimal area. Thus, it is likely that the climate-related geographical variation in mating success represents a component of geographical variability in Allee effects and this could have affected past rates of gypsy moth spread, as well as influence its future rate of spread.

We have demonstrated the potential for geographic variation in climate to produce geographic variation in mating success under an assumption of constant life history and thermal requirements for development across the area. These results add another dimension to our understanding of geographical variation in habitat suitability for the gypsy moth. Previous analyses characterized the geographical distribution of suitable forest host tree species (Liebhold et al. 1997, Morin et al. 2004). Gray (2004) used a network of weather station records, similar to that utilized here, to characterize geographical variation in climatic suitability for establishment based on life cycle completion (i.e. the ability of an introduced population to complete successive generations). The results presented here indicate that geographical variation in climatic
suitability for mating is yet another component of habitat suitability. Although coastal California is potentially suitable for gypsy moth development (Gray 2004) and contains suitable host trees (Liebhold et al. 1997), our results indicate that the probability of mating may be relatively low (Fig. 3c, 4, 5).

Our results provide new insights concerning the potential for establishment of new gypsy moth populations across North America, however we should moderate our interpretation. While our model accounted for geographical variability in maturation times, it assumed that other parameters (e.g. male and female longevity and survival) were constant. Substantial geographical variation in these traits could be expected to alter the model’s predictions. This study provides a preliminary understanding of mating failure in this species and while assuming various hypotheses, it rigorously describes the underlying mechanism for this failure.

As a matter of simplicity, simulations described here were intended to represent a scenario where a gypsy moth egg mass was transported from a single location and, therefore, our phenology simulations were based upon a single date of oviposition (1 August). In practice, the equilibrium oviposition date varies geographically with climate (Gray 2004). We conducted additional simulations in which the date of oviposition was varied and we found that through most of the range of equilibrium oviposition dates reported by Gray (2004), there was little variation in predicted mating success at most locations. Thus, the source location of an egg mass transported to most portions of North America could be expected to have little impact on mating success. In contrast, oviposition date does affect the probability that a population can complete development (Gray 2004) and thus the identity of the source population is likely to affect local establishment.

Our generalized model coupled with a species-specific investigation provides further evidence of the importance of mating failure due to temporal asynchrony between male and female maturation times. This mechanism alone, or in addition to other factors generating Allee effects (such as spatial isolation of individuals or other factors not associated with mate-finding), could contribute to the decline of a population, or the failure of a population to establish. As a result, the causes of Allee effects should be clearly identified to understand more precisely the dynamics of expanding or declining populations.

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