

Spatio-temporal variation of male sterile frequencies in two natural populations of *Beta maritima*

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The spatio-temporal variation of sex phenotype frequencies is investigated within two gynodioecious populations of *Beta maritima* located along the English Channel, in which the proportions of females differ significantly: 0.19 (Population A) and 0.62 (Population B). A genetical analysis of maternal progenies obtained from *in situ* open-pollinated plants (G_1 generation) allows us to define two types of parents: segregating plants (females, intermediate-females and some hermaphrodites) which yield different types of progenies, and non-segregating plants, hermaphrodites which yield only hermaphrodites. Molecular analysis (Saumitou-Laprade, 1989) confirms the results of a preliminary study (Boutin *et al.*, 1987), in which it is shown that cytoplasmic type is related to segregation in the maternal progeny.

A differentiation in space has been pointed out by the comparison between the sex phenotype frequencies in the two populations. This differentiation is not due to a variation of cytoplasmic frequencies but to a variation in the nuclear genetic components of these populations: the level of restoration is higher in population A than in population B.

A comparison of the G_0 and G_1 generations did not allow to predict any detectable temporal change in population A, whereas the frequency of females in population B has been predicted to decrease quickly, probably due to a rapid invasion of restorer genes. We suggest that pollen flow is an important factor which determines the rate of the dynamics of male sterility during the life-span of a population.

INTRODUCTION

In natural populations of several wild plant species, nucleo-cytoplasmic determination of male sterility appears to be much more common than purely nuclear determination (*e.g.*, *Origanum vulgare*, Kheyr-Pour, 1980; *Plantago lanceolata*, Van Damme, 1983; *Thymus vulgaris*, Dommée, 1973). The intraspecific variation in the frequency of female (= male sterile) plants results from the interaction of hereditary information from two sources, nucleus and cytoplasm, leading to the formation of either female or hermaphrodite phenotypes. Several models (see Ross, 1978, for review; Gregorius *et al.*, 1982) based on purely nuclear control of gynodioecy have failed to explain the diversity of observed situations (particularly, female frequencies higher than 50 per cent and only a slightly higher female fitness of females than hermaphrodites), whereas they can be explained by nucleo-cytoplasmic models (Delannay *et al.*, 1981; Gouyon and Couvet, 1985; Couvet *et al.*, 1986). In self-compatible species, a

stable equilibrium of the frequency of females, if it exists, is reached slowly with oscillations of the frequency of females (Charlesworth, 1981). In self-incompatible species, stable limit cycles between the frequency of cytoplasmic and nuclear genes are usually found (Gouyon *et al.*, in preparation; Boutin-Stadler *et al.*, in preparation). Thus, proportions of sex phenotypes have been expected to change over time. The occurrence of genetic differentiation among populations can be explained both by isolation among populations and by the interactions between nuclear and cytoplasmic genes determining male-sterility: when a new site is founded by hermaphrodites and females, female frequency in later generations will depend on whether or not the hermaphrodites possess the restorer genes of the cytoplasm of the female plants.

The variation of sex phenotype frequencies has often been studied indirectly by comparing populations situated in different successional stages of the vegetation (Dommée *et al.*, 1983; Gouyon *et al.*, 1983; Dommée and Jacquard, 1985; Balhassen

et al., in press). Less frequently, variation has been followed over time within populations. For example, in such a study, marked decrease in female frequencies was observed by Krohne *et al.* (1980) in *Plantago lanceolata* from one year to the next.

In *Beta maritima*, the available data indicate that the male sterility system, although similar to that studied by Owen (1942) in cultivated sugar beet, has different nuclear and cytoplasmic components (Boutin *et al.*, 1987; Boutin *et al.*, 1988a; Halldén *et al.*, 1988). In the populations studied a strong correlation exists between the type of cytoplasm and the presence or the absence of segregation of females in the progenies (Boutin-Stadler, 1987; Boutin *et al.*, 1988b; Saumitou-Laprade, 1989).

In this paper, we first study the variation of sexual phenotype frequencies in space by comparing two populations of *Beta maritima*. The differences observed among the segregations in the progenies of plants from these populations give us an insight in their genetic structure and enable a rough estimate of the level of restoration. Secondly, we study the variation of sex phenotype in time, that is whether these populations are in an equilibrium stage by comparing the observed sex phenotype frequencies *in situ* (referred to as the G_0 generation) with the expected ones in the G_1 generation.

MATERIAL AND METHODS

Plants of the G_0 generation and their open-pollinated progenies

The two *Beta maritima* populations, called A and B, are located in the higher zone of a salt marsh in the Canche estuary (northern France). They are covered by the sea at the time of equinox spring-tides. They are separated by a distance of 1 km. Population A is located at the foot of the sand dunes; in 1984, 94 flowering plants covering an area of 80 m × 5 m grew in a herbaceous community. This population was surrounded by several other populations of *B. maritima*. Population B is located on a small sandy mound between the salt marsh and the beach and is more often covered by the sea during heavy storms; in 1984, 66 flowering plants grew in an area approximately 40 m × 10 m, among culms of *Agropyron pungens* and were isolated more than 500 metres from other *B. maritima* plants.

Three sex phenotypes were distinguished in this species by using anther and pollen characteristics defined at anthesis (Boutin *et al.*, 1987):

(i) hermaphrodites (H), with yellow stamens containing viable and functional pollen, (ii) females (Fe), with white stamens lacking pollen grains, (iii) intermediate-females (IFe), with stamens containing microspores and non-viable pollen grains. Each plant was observed three times during its flowering period in the natural populations or in the experimental garden of the university, in order to check the stability of the sex phenotypes: no difference appeared among the observations. Neither the females nor the intermediate-females transmit their genes through male gametes and so they are functionally females.

Fruits were harvested from a sample of individual plants (mother-plants) of known phenotype that had been pollinated *in situ* in each population (16 H, 11 IFe and 6 Fe in population A and 10 H, 2 IFe and 18 Fe in population B) proportional to the number of plants of each sex phenotype in the population. That explains, for instance, the very small number of IFe families studied in population B. Seeds were sown in the experimental garden of Lille University. The 63 families ranged in size from seven to 80 individuals. The total number of plants was 1446; they formed the G_1 generation and were used for the subsequent genetic analysis.

In both populations, the Fe and IFe plants always produce segregating progenies (*i.e.*, composed of two or three sexual phenotypes), whereas the H plants can be classified in two groups:

- the *segregating* ones (H segr.) which generate two or three sexual phenotypes in their progenies, like the Fe and IFe plants;
- the *non-segregating* ones (H non-segr.) which only produce H.

A joint genetical and molecular analysis (Boutin *et al.*, 1987; Boutin *et al.*, 1988a; Saumitou-Laprade, 1989) suggested the existence of two groups within these two populations, in which progenies of segregating plants carry the S cytotype and the progenies of non-segregating plants carry the N cytotype.

The variation of sex phenotype frequencies between the observed G_0 and the expected G_1 generations

The method consisted in the comparison of the sex phenotype frequencies in the *in situ* population samples with the ones expected in the next generation according to the segregations observed in the maternal progenies, in order to check if the sex frequencies are at equilibrium. In each population, the sex phenotype frequencies in the expected G_1

generation have been estimated by calculating the frequency of each sex phenotype in each progeny and taking the unweighted average over all progenies.

As the non-segregating hermaphrodites give only hermaphrodites in their offspring and as the frequency of restorer genes of the S cytoplasm is not known in the non-segregating plants, the variation in time of sex phenotype frequencies will only be estimated using segregating plants.

Statistical analysis

The *G* test of independence using William's correction (Sokal and Rohlf, 1981 p. 731) was used to test:

- the difference in sex phenotype frequency between the two natural populations (variation in space);
- the difference in frequency of the two types of hermaphrodites between the two population samples (variation of cytotype frequencies among hermaphrodites);
- the difference in cytotype frequency in the G_0 generation and the difference in the level of restoration of segregating plants (restored plants/non restored plants) between the two population samples (variation of the cytotype frequencies and variation in the level of restoration).

Within each population sample, the comparison between the sex phenotype frequencies observed *in situ* with the expected ones in the following generation (variation in time) was tested with a *G* test for goodness of fit using William's correction (Sokal and Rohlf, 1981 p. 704).

RESULTS

Sex phenotypic variation between populations A and B (variation in space)

The sex phenotype frequencies in populations A and B (table 1) appeared to be strongly different [$G_{\text{Williams}(2)} = 34.56$, $P < 0.001$]. Population A

Table 1 Distribution of sex phenotype frequencies in populations A and B from the Canche estuary

Population	No. of plants	Sex phenotypes*		
		H	IFe	Fe
A	94	0.48	0.33	0.19
B	66	0.30	0.08	0.62

* H = hermaphrodite; IFe = intermediate-female; Fe = female.

contained few Fe plants and showed a relatively high frequency of H and IFe, whereas population B contained many Fe plants and showed a low frequency of IFe.

Genetical analysis of the G_0 generation

As cytoplasmic inheritance is maternal, both the female parent and its offspring possess the same cytoplasmic type. Consequently, the structure of the samples of the two populations can be examined using both cytoplasmic and nuclear information, the latter being based on the sex phenotype of the S plants: hermaphrodites possess the nuclear restorer genes and females do not possess them. In population A, in which IFe plants were numerous, segregations in the maternal progenies between phenotypic classes are significantly different ($\chi^2_{(2)} = 41.29$, $P < 0.001$): the mean hermaphrodite frequency is higher in IFe families (0.28) than in Fe (0.16) families and is highest in segregating H families (0.47). These results of segregations (Boutin-Stadler, 1987) strongly suggest that IFe are partially restored.

The following points appear:

Firstly, *among the hermaphrodite sampled* (table 2), the frequencies of non-segregating and segregating plants, having respectively the N and the S cytoplasm, are significantly different in the two populations [$G_{\text{Williams}(1)} = 4.90$, $P < 0.05$]: the more frequent type of H is the non-segregating one in population B and the segregating one in population A.

Table 2 Frequencies of segregating and non-segregating hermaphrodites in the population samples

Population	Total no. of H	Segregating	
		Segregating	Non-segregating
A	16	0.75	0.25
B	10	0.20	0.70

Secondly, *within the sample* of each population (table 3), the frequency of each type of H can now be specified and the frequency of each type of cytoplasm can be estimated: population B contains a higher proportion of non-segregating H (0.23) than population A (0.12), but there is no significant difference in cytoplasm frequency between the two populations [$G_{\text{Williams}(1)} = 1.32$, $P > 0.05$].

Thirdly, *among the segregating plants* (table 3), the frequencies of the three sex phenotypes are significantly different between the two samples [$G_{\text{Williams}(2)} = 17.48$, $P < 0.001$]; the ratio of

Table 3 Sex phenotype frequencies according to the type of segregation and the level of restoration in the samples of populations A and B. H: hermaphrodite; IFe: intermediate-female; Fe: female

Population	Segregating plants (S)				Non-segregating plants	Total No. of plants
	Restored plants		Non-restored			
	H	IFe	Fe	Total		
A	0.36	0.33	0.18	0.88	0.12	33
B	0.10	0.07	0.60	0.77	0.23	30

restored plants (H + IFe) to the non-restored plants (Fe) is significantly higher in population A than in population B [$G_{\text{Williams}(1)} = 17.60$ $P < 0.001$], and therefore the level of restoration is higher in population A than in population B (table 4).

Table 4 Variation of the restoration level* between samples of plants from populations A and B and between generations

Population	Level of restoration* in		
	G_0 In situ generation	G_1 Expected generation	Ratio G_1/G_0
A	3.83	6.25	1.63
B	0.28	2.28	8.14

* Restoration level = (No. of restored plants: H segr. + IFe)/(No. of non-restored plants: Fe).

Comparison of the expected G_1 generation with the G_0 generation (variation in time)

The comparison of the proportions of sex phenotypes in G_0 and expected G_1 (table 5) reveals no difference for population A [$G_{\text{Williams}(2)} = 2.36$, $P > 0.05$] but a significant difference for population B [$G_{\text{Williams}(2)} = 27.84$ $P < 0.001$]. The level of restoration thus increases relatively more rapidly in population B which showed a higher frequency of females in the G_0 generation (table 4).

Table 5 Comparison of the sex phenotype frequencies among segregating plants of the population samples in the expected G_1 generation with their observed frequencies in the G_0 generation

Population	Generation	No. of plants	Sex phenotypes among S plants			$G_{\text{Williams}(2)}$ P
			H segr.	IFe	Fe	
A	G_0	29	0.41	0.38	0.21	(2.36)
	G_1		0.35	0.51	0.13	>0.05
B	G_0	23	0.13	0.09	0.78	(27.84)
	G_1		0.09	0.61	0.31	<0.001

DISCUSSION

The G_0 populations (variation in space)

Although the number of loci involved in the restoration process, their dominance, and the nature of interactions between loci are not known, it is nevertheless possible to evaluate the approximate level of nuclear restoration of the S cytoplasm. Indeed, the difference in female frequency between the two populations is not related to a difference in cytoplasm frequency; thus the main difference between the two G_0 populations may be attributed to the approximate level of restoration, which was higher in population A.

The occurrence of females in gynodioecious species (e.g., *Thymus vulgaris*) is interpreted as the consequence of a nucleo-cytoplasmic differentiation between populations (Couvét *et al.*, 1985). Study of gene flow among several populations of *Beta maritima* (Saumitou-Laprade *et al.*, in preparation) showed the existence of a genetic differentiation between populations, suggesting that populations were isolated from each other. These results can help us to understand the variation in sexual phenotype frequencies in our populations. Indeed the two populations differ only by the frequency of restorer genes. Population A, in the neighbourhood of other populations could more easily exchange nuclear genes with the adjacent populations compared with the isolated population B, in which restorer genes may have appeared recently.

The dynamics of populations (variation in time)

The comparison between the *in situ* observed G_0 populations and their respective G_1 generation enables us to test whether or not the sex phenotype frequencies were at equilibrium. It is expected that:

- (i) Population A is not in an important shifting phase because there is no significant difference between generations;
- (ii) Population B is undergoing major change because, in only one generation, a major decrease of females of -47 per cent was accompanied by an increase of intermediate-females of 52 per cent.

Our data obtained from the two population samples indicate that the high proportion of females in population B may not correspond to an equilibrium state and occurs when male fertile restorer genes are rare in the population.

These data are in agreement with a model of dynamics of sex phenotype in gynodioecious populations of self-incompatible species (Boutin-Stadler *et al.*, in preparation). Three phases can be described if we suppose that female (having the S cytotype) and hermaphrodite (having the N cytotype) migrants colonize a new site:

- (i) Assuming a cost of restoration (that is, selection acting against hermaphrodites carrying the restorer genes of an another cytoplasm (Delannay *et al.*, 1981; Charlesworth, 1981; Gouyon and Couvet, 1985; Gouyon *et al.*, in preparation), in such a population, the hermaphrodites (N) rarely possess the specific restorer genes of the sterile S cytotype. Because of a cytoplasmic determination of male sterility, as long as females have even a very small advantage in seed production, the female frequency will increase in the population as well as the frequency of cytotype S.
- (ii) High female frequency in the population selects for nuclear restorer genes. Then, once introduced, restorer genes invade the S cytoplasm, very rapidly at the beginning and later on more slowly, causing a decrease of female frequency in the population.
- (iii) At this time, hermaphrodites having S cytoplasm are numerous and female frequency is low in the population. Assuming a higher female fitness of hermaphrodites having N cytotype compared to ones having S cytotype, the frequency of the former increase in the population. In the same time, due to the restoration cost, the frequency of restorer

genes decreases in the population and we come again at the initial point.

The two first phases are not observed in our study; there are no patches of unrestored females, as it has been observed in *Thymus vulgaris* (Dom-mée and Jacquard, 1985). Both populations would correspond to different stages of the second phase, population B being at an earlier stage than population A. Indeed, while the S cytoplasm has already invaded 80 per cent of both populations, restoration is also present, with a higher level of restoration in population A. The marked change in the proportion of sex phenotypes towards a lower frequency of females, detected in population B, would be due to the dynamical process of invasion by nuclear restorer genes. This is well illustrated by the variation of the level of restoration between the G_0 and G_1 generations (table 4).

The rate of decrease of female frequency in the expected $(n+1)$ th generation has been observed to be proportional to the female frequency in the preceding one. In the same way, Krohne *et al.* (1980) observed a decrease of female frequency in eight natural populations of *Plantago lanceolata* related to the female frequency in the populations, although similar results have not been found in *Thymus vulgaris* (Couvet *et al.*, 1985). We suggest that the speed of invasion of a population by restorer genes is related to pollen flow, which is probably very different between anemophilous species like *Beta* or *Plantago* and entomophilous species like *Thymus*. The minimal plant distance to maintain purity of plant varieties depends on the reproductive system (self or cross-fertilization), and within each system, there is an increase in the minimal distance from entomophilous species to anemophilous species (Levin and Kerster, 1974). In *Thymus*, pollen transfer by bees is very limited (0.50 to 1 m (Brabant *et al.*, 1980; Belhassen *et al.*, 1987)), whereas in anemophilous species such as *Beta*, pollen flow is less limited: >4.50 m (Archimowitsch, 1949) and 50 per cent of pollen is present 100 m from the source (Levin and Kerster, 1974). As a consequence, the process of invasion by restorer genes in a gynodioecious population should be faster in anemophilous species.

The distribution of sex phenotypes in the expected G_1 has been estimated using segregations obtained from the seeds produced in the G_0 generation. The study of the dynamics of change in female frequency in natural populations should take into account the life-history traits of the species: *Beta maritima* is a short-lived perennial. In our case, only six plants (four hermaphrodites and two females) in population A and three plants

(two hermaphrodites and one female) in population B were still alive the next year because the winter 1984–85 was particularly severe. The speed of the invasion of restorer genes will therefore depend on the pattern of recruitment of young individuals into the populations in the next generation (from the seed bank and the seeds of the current year).

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