

## ELIMINATION OF AN INTRODUCED *B*-CHROMOSOME FROM A WILD POPULATION OF *ARCYPTERA FUSCA* (Orthoptera)

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### SUMMARY

Males from a population of the mountain grasshopper *Arcyptera fusca* possessing a persistent *B* chromosome were transferred to a population without *Bs* in two successive years. The *B* was present in the second year, but now appears to have been eliminated. A total of seven successive generations have been analysed.

### 1. INTRODUCTION

Supernumerary chromosomes are widespread in many species of plants and animals. They usually occur only in some populations and in varying frequencies (Jones and Rees, 1982). A majority of the supernumerary systems known in nature are characterised by boosting mechanisms which lead to increases in the frequency of *Bs* in the progeny of *B*-containing individuals. It seems that such boosting system can permit *Bs* which have deleterious effects to be maintained for many generations in a population. In fact, if such a boosting system exists, the critical reproductive fitness of a *B*-containing animal permitting the *B* to be maintained in a population is not 1 but some value less than 1, depending on the accumulation mechanism. Rates of transmission of *B* chromosomes have been well studied in some species of grasshoppers by comparing the karyotypes of both parents and progeny from single matings in laboratory conditions (Lucov and Nur, 1973, Hewitt, 1973) and some detailed studies concern the movement of *Bs* in natural populations (Shaw, 1983*a, b*). However, the ability of a genome to receive and incorporate extra material under natural conditions is difficult to test since unknown factors may condition the survival of this extra material.

The present report concerns the examination during seven consecutive years of two selected natural populations (with and without *Bs*) where individuals containing *Bs* were inoculated into the population without *Bs*.

### 2. METHODS

Meiotic chromosomes were analysed in squash preparations of single follicles fixed in a (1:3) glacial acetic acid: ethanol mixture.

### 3. RESULTS AND DISCUSSION

During the past seven years we have analysed the structure of the chromosome system which exists in different populations of the mountain



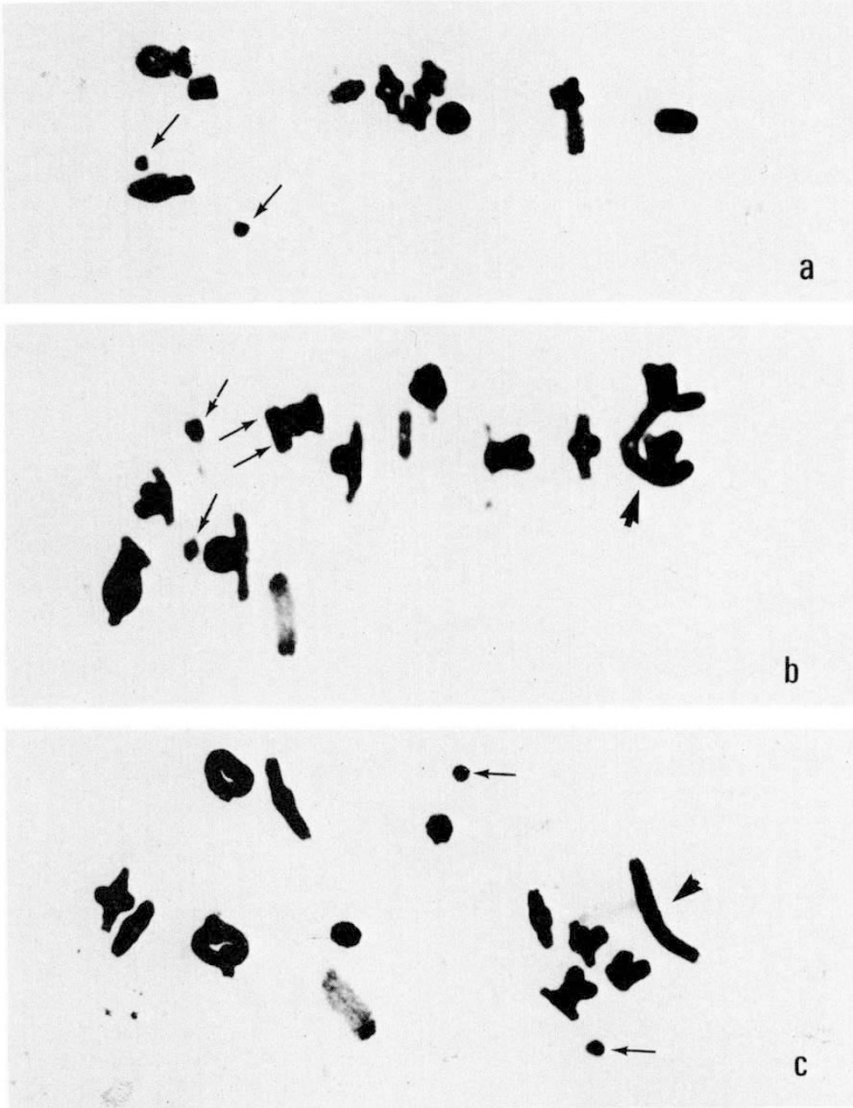


PLATE 1. Metaphase-I from a male of *A. fusca* which integrated the *B* in its genome. (a) Two unpaired *B*s. (b) Four *B*s (arrows) plus a  $L_2$  trivalent (arrow head). (c) Two unpaired *B*s (arrows) plus an extra  $L_2$  bivalent (arrow head).

TABLE I  
*Populations of Arcyptera fusca and number of individuals with and without Bs collected during seven consecutive generations in Respumoso*

References	Sample		
	0 <i>B</i>	<i>B</i> s	Total
AF.IV 1976	21	20	41
AF.IV 1977	10	9	19
AF.IV 1978	11	11	22
AF.IV 1979	21	18	39
AF.IV 1980	23	13	36
AF.IV 1981	15	10	25
AF.IV 1982	27	18	45
Total	128	99	227

Given the frequency of *B*s in Respumoso, it was expected that 50 per cent of these two inoculations would contain *B*s in their genomes. In both cases only males were used, so the incorporation of *B*s into the genepool would be through non *B* females of Pto. de la Bonaigua.

The karyotype analysis of the individuals from the 1979 generation (G79) did not reveal the existence of individuals carrying *B*s, although it does not entirely exclude the possibility that some *B* individuals were not detected since a sample of males were used. However in 1980 (G80) two out of forty males collected had *B*s with similar characteristics to those found in males from Respumoso (plate 1). Interfollicular variation in the number of *B*s, due to premeiotic accumulation (0*B*s, 1*B*, 2*B*s, 3*B*s and 4*B*s) was also observed. Samples collected during 1981 (32 individuals) and 1982 (25 individuals) did not contain *B*s and the frequency of *B*-carrying individuals at the Pto. de la Bonaigua is now probably zero.

In one of the two individuals found with *B*s at Pto. de la Bonaigua, the L<sub>2</sub> chromosome was polysomic in all cells of four of the testis follicles. The extra L<sub>2</sub> chromosome could pair and form a single bivalent, usually with a distal chiasma (plate 1c) or form multiples with the standard pair also by means of a terminal chiasma (plate 2b). The fact that such an anomaly was only found in this individual out of over four hundred individuals studied from this and other populations led us to wonder if it might be related to genetic divergence between the populations studied. Such divergence is reasonable because of the restricted mobility in the females due to the small size of the termina and wings, the unfavorability of the intervening terrain, where high mountains break the continuity of the distribution, and finally the relatively short life of the species which might contribute to a restriction in the genetic interchange between populations.

The major problem in discussing the present experiment is the partially negative results obtained. Although some of the introduced males were effective in impregnating females, it is difficult to know for certain the causes which produced the elimination in the next generation. However, the evidence that the *B* could be introduced, at least in one generation, indicates that postzygotic mechanisms of selection rather than prezygotic ones might

be effective in the elimination of the *B*. The stabilisation of a *B*-chromosome in a population could depend, as in the grasshopper *Myrmeleotettix maculatus*, on various factors such as meiotic drive in egg cells (Hewitt, 1973), selection against certain karyotypes carrying *B*s and the selective advantage of others. The genome of the recipient population could be different and not interact with the introduced *B* to produce a stable equilibrium as in the donor population. It is also possible that a relatively ineffective inoculation would permit drift alone to eliminate the *B*s rather easily.

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