because of rapid autoagglutination of the cells. None of the vertebrate erythrocytes and the trypsin-treated mammalian cultured cells were agglutinated by the SF.

Our results indicate that the SF attaches to the surfaces of cell plasma membranes of insect cultured cells. The SF-attaching sites (cellular receptors) are restricted to certain surface areas of the plasma membranes. The agglutinating capacity of the SF suggests the existence on the SF molecule of polyvalent active sites which are responsible for the binding of the SF to cellular receptors on the plasma membrane. Since the SF greatly enhances NPV infection in vitro, our present findings strongly indicate that the SF acts in vitro as an enhancer in the attachment of cell-membrane-budded nucleocapsids of NPV to cultured insect cells as in the case of NPV enhancement in vivo<sup>8-10</sup>. This phenomenon may be analogous to the antibodymediated enhancement of animal virus infections, such as the enhancement of the virus-specific antibody at a subneutralizing concentration in the attachment of the virus to Fc-receptorbearing cells15.

The NPVs are known to cause the hemagglutination of vertebrate erythrocytes<sup>16-20</sup>. Recently, Anderson et al.<sup>21</sup> found that the enveloped nucleocapsids of a GV of Plodia interpunctella agglutinated several species of vertebrate erythrocytes, in particular, rabbit erythrocytes. In our present study, the SF did not agglutinate the erythrocytes of several vertebrate species including that of the rabbit. Thus, the hemagglutinin present in enveloped nucleocapsids of P. interpunctella GV appeared not to be related to the SF.

The SF agglutinates not only lepidopterous insect cells but also those of mosquitoes and a cockroach. Since the enveloped arboviruses, i.e. Alphavirus group, multiply in the vector mosquito, the SF through its effective attachment may enhance the infection of these viruses in mosquito cells in vitro. Such a study may explain, in part, the variations in the efficiency of mosquitoes as vectors.

The basis for the attachment of the enveloped baculoviruses to the host plasma membrane to initiate fusion has not been clarified. Our previous study in vivo suggested that the phospholipids and ionic charges of the SF and viral envelopes were involved in attachment<sup>22-24</sup>. The SF appeared to enhance the attachment and/or fusion of the enveloped virion to the plasma membrane.

The question of virus receptor sites has not been resolved<sup>25</sup>. The adsorption of the SF to certain regions of the plasma membrane suggests the presence of receptor sites on the cell membrane. The attachment of the SF to the host cell membrane and its capacity to enhance the baculovirus infection provide us with a unique opportunity to further investigate this problem.

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## The effect of X-rays and EMS on the behavior of the transposing element, TE98, in Drosophila melanogaster

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Summary. The sensitivity of TE98 (carrying  $w^{+R}$  and  $rst^+$ ) to X-rays does not differ significantly from the mutability of curled and karmoisin loci. In addition no spontaneous mutants of TE98 were recovered, indicating its extreme stability. On the effect of EMS no white mutants were found supporting the view that the  $w^+$  gene of TE98 is duplicated.

Several classes of transposable genetic elements are known in Drosophila melanogaster. Similarly to procaryotic transposons, they play an important role in spontaneous and induced mutational events<sup>2</sup>. The mutagen sensitivity of a transposable DNA sequence inserted adjacent to the white locus has been demonstrated3. Few data are available on the response to mutagenic agents of other Drosophila transposons. We report on the behavior of a large transposing element TE (in a position named TE98 in 3R) under conditions of induced mutagenesis.

In the course of a cytogenetic analysis of the 87A-C region of the 3rd chromosome, we isolated a large number of deficiencies induced by X-rays. In these experiments we used the transposing element TE98 described by Ising and Block4 as a genetic marker. This transposing element is cytologically visible in polytene chromosome preparations and it carries a piece of the X chromosome with the white  $(w^{+R})$  and roughest  $(rst^+)$ genes. TE98 is a derivative of TE1 from which more than 100 new positions have been identified. The transposing elements

Table 1. Frequency of flies with mutant phenotypes in controls and after X-ray treatment

Chromosomal region examined	Number of flies examined	Number of flies with mutant phenotype	Mutation frequency
cu*	62919	45	$7.1 \times 10^{-4}$
kar*	62919	48	$7.6 \times 10^{-4}$
TE98	205975	176	$8.5 \times 10^{-4}$
Control cu* and kar*	3919	-	_
Control TE98	82 284	_	_

<sup>\*</sup>Data are taken from Gausz et al.9.

TE28 and TE98 are flanked by foldback elements of the FB family<sup>5,6</sup>.

The cytological position of TE98 is distal to the 87A4-5 band in the right arm of the 3rd chromosome (Gyurkovics, unpublished observation). In the presence of w on the X chromosome, TE98-carrying flies exhibit the wild type (red) eye color. However, when the transposing element is eliminated by a deletion, the loss of the  $w^{+R}$  allele results in white-eyed flies. If TEs are placed near to heterochromatin by inversion or translocation, this results in flies with white-variegated eyes. To obtain more data on the repsonse of TE98 to mutagens, in addition to the X-rays, we also used ethyl methane sulphonate (EMS).

In X-ray experiments males carrying w on their X-chromosome as well as TE98 on their 3rd chromosome  $(w/Y; TE98 w^{+R} rst^+/TM3)$  were irradiated with a 4000 rad (1000 rad/min) dose. The treated males were crossed to w spl/w spl; Sb/Ser virgin females (for an explanation of genetic symbols see Lindsley and Grell<sup>7</sup>). In the  $F_1$  progeny all the flies carrying the transposon exhibited a wild-type eye color, but in flies where the  $w^{+R}$  gene was mutated, excised or translocated to heterochromatin, the eyes were white or white-variegated, respectively.

EMŚ was used at a concentration of 0.025 M according to the method of Lewis and Bacher<sup>8</sup>. The crosses were identical with those described for the X-ray experiment.

For a comparison, we use our previous data obtained for the mutability of the *curled (cu)* and *karmoisin (kar)* loci<sup>9,10</sup>.

The results of the X-ray experiments are summarized in table 1. The X-ray treatment induced changes affecting the  $w^{+R}$  marker with a frequency of  $8.5 \times 10^{-4}$ , whereas flies with cu and kar phenotype appeared with a frequency of  $7.1 \times 10^{-4}$  and  $7.6 \times 10^{-4}$ , respectively. These frequencies do not differ significantly from each other indicating that the sensitivity of TE98 to X-rays is not higher than the value for induced visible recessive mutations. Of the 176 X-ray induced mutations 95 showed white and 81, variegated eyes. 32 of the white-eyed ones were identified as deletions in crosses with lethal mutations from the surroudning region. The remaining 63 mutants were lost due to sterility or reduced viability. 14 out of the 32 deletions were examined cytologically and 7 of them had one breakpoint in the 87A4-5 band, which is the insertion site of TE98 (Gyurkovics, unpublished results).

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Table 2. Frequencies of flies with mutant phenotypes in controls and after EMS treatment

Chromosomal region examined	Number of flies examined	Number of flies with mutant phenotype	Mutation frequency
cu*	8 3 0 3	7	$8.4 \times 10^{-4}$
kar*	14029	13	$9.2 \times 10^{-4}$
TE98	34495	2	$5.7 \times 10^{-5}$
Control TE98	82 284		

<sup>\*</sup>Data are taken from Gausz et al.8.

After EMS treatment (table 2) mutations at the cu and kar loci were recovered with almost identical frequencies,  $8.4 \times 10^{-4}$ and  $9.3 \times 10^{-4}$ , respectively. Changes involving TE98 appeared at a significantly lower frequency. Only 2 flies with white-variegated eyes were found among 34,495 progeny, which is a frequency of  $5.8 \times 10^{-5}$ , significantly lower than that for cu and kar loci. The variegated eye color indicates that these 2 mutations are probably the result of chromosome rearrangements. No w point mutations or deletions were isolated. Our inability to induce point mutations by EMS treatment supports the hypothesis of Ising and Block<sup>4</sup> that TE1, from which TE98 is originated, contains a duplication for  $w^{+R}$ . Recently<sup>5</sup> it has been shown by whole genome Southern analysis of TE98 that there are 3 bands of hybridization to single copy white sequences: 1 from the original location on the X chromosome, and 2 from TE98. Therefore it can be concluded that TE98 carries a duplication of the white locus region.

The frequencies of spontaneous and induced loss of 2 other transposing elements (TE6 and TE30) have been reported by Ising and Block<sup>4</sup> and by Ising and Ramel<sup>11</sup>. The red form of TE30 is lost spontaneously with a frequency of  $2 \times 10^{-4}$ . The values for TE6 vary between  $3 \times 10^{-4}$  and  $8.3 \times 10^{-3}$  depending on the genetic background. TE98 is very stable; we did not find white-eyed flies among 82,284 tested F<sub>1</sub> progeny (frequency is less than  $1.2 \times 10^{-5}$ ). In spite of the fact that TE6 and TE30 are less stable than TE98 Ising and Ramel<sup>11</sup> were not able to influence the frequency of loss by X-rays, temperature shock or interchromosomal effects of inversions on recombination. The sensitivity of the transposing elements to these external influences is different compared to the results of Rasmussen et al.<sup>3</sup>. However, the fact that 7 out of the 14 cytologically analyzed deletions induced on TE98 carrying chromosomes had one breakpoint adjacent to the transposing element, indicates that in spite of its relative stability TE98 is a 'hot spot' for induced chromosome breaks. The other breakpoints are more randomly distributed. It is not excluded that some of these breakpoints may be connected with insertion sites of dispersed repeated elements and in this case the FB-NOF sequences in TE98 may serve as portable regions of homology between the 2 breakpoints<sup>12</sup>. However, to test this possibility further studies are needed.

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