LINKAGE OF REX WITH SHAKER-2 IN THE HOUSE-MOUSE

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THE dominant hair-waving gene, Rex, is one of five genes with almost identical phenotypic effects, known in the house-mouse. It arose in the fancy, and was first investigated by Crew and Auerbach (1939, 1940). Linkage tests with 10 other loci were made and no significant departure from independent segregation was obtained. The recombination frequencies, calculated from the data published by Crew and Auerbach, are set out in table 1, together with an

TABLE 1

Recombination between Rex and 10 other loci, with approximate values for the closest admissible linkage. Calculated from the data of Crew and Auerbach (1939, 1940)

Locus tested	Recombination frequency per cent.	Standard error	Closest admissible linkage (approx.) per cent.	
Albino c Pink-eye p Dilute d Brown b Non-agouti a Waltzer v Leaden ln Pied s Waved-I wa-I . Caracul Ca	52 37 50 60 51 49 45 45 47 62 41	5.5 7.6 6.5 7.6 6.5 9.8 9.8 9.8 9.8 9.8 9.8 9.8 9.8 9.8 9.8	41 22 37 45 38 29 25 28 42 28	

approximate value of the closest admissible linkage. Close linkage of Rex with any one of these ten factors was thus excluded, and at the same time the non-allemorphism of Rex with Caracul and with waved-1, two of the other hair-waving genes, was proved. The degree of independence of Rex with the agouti locus that was observed excludes the possibility of Rex being an allele of wellhaarig, a recessive hair-waving gene, since wellhaarig is linked with agouti with a recombination frequency of about 15 per cent. (Hertwig, 1942).

The present paper gives evidence which shows that Rex is fairly closely linked with shaker-2, a locus not tested by Crew and Auerbach. At the same time it is shown that Rex is not an allele of waved-2, the fifth hair-waving gene, which is also carried on this chromosome. (Snell and Law, 1939).

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The linkage of Rex with shaker-2 was revealed by number nine of the segregating lines set up in the Department of Genetics by Professor R. A. Fisher with the object, amongst others, of filling the gaps in our knowledge of linkage. The foundation matings of this line were made by Mrs V. P. Leonard in June 1944. In pursuance of the same scheme, Rex was introduced to the line by Miss M. E. Wright early in 1945, and in June of that year the first mating was made showing simultaneous segregation of Rex and shaker-2 in the same animal. The frequency of recombination between Rex and shaker-2, indicated by the data up to the time of going to press, is 20 per cent., with a standard error of $5 \cdot 1$ per cent. All the matings were coupling backcrosses of the type $\frac{\text{Re}+}{+\text{sh-2}} \times \frac{+\text{sh-2}}{+\text{sh-2}}$, and the frequencies of the phenotypic classes among the progeny are shown in table 2. The matings are separated according to the sex of the segregating parent : the sex-difference in crossing-over is not significant.

TABLE 2

Sex of segregating parent	Old comb Re+	oinations +sh-2	Recomb Re sh-2	oinants ++	Total	Recombination frequency per cent.
Male	17	14	3	4	38	18.4
Female	6	12	3	2	23	21.7
Combined	23	26	6	6	61	19.67

Coupling backcross segregation of Rex and shaker-2

The recessive hair-waving gene, waved-2, is known from the data of Snell and Law (1939) to be linked with shaker-2. The recombination frequency for males, based on 56 mice bred from backcrosses was 27 per cent., and that for females, based on 256 mice. was 25 per cent. The recombination frequency estimated from the combined data is $25\cdot3$ per cent. (S.E. = $2\cdot46$ per cent.). In order to determine the position of Rex relative to shaker-2 and waved-2, coupling backcross matings of the type $\frac{\text{Re} + +}{+ \text{sh-2 wa-2}} \Im \times \frac{+ \text{sh-2 wa-2}}{+ \text{sh-2 wa-2}} \eth$ were set up. The progeny could of course be only partially classified, since Rex and waved-2 phenotypes can be distinguished only with difficulty, if at all, and the double mutant phenotype is unknown. The appearance of a single straight-haired offspring would, however, show Rex and waved-2 to be at different loci, and the frequency of straight-haired offspring would lead to an estimate of the frequency of recombination between Rex and waved-2, and so indicate the order of the three loci on the chromosome.

Up to the time of going to press, 122 mice have been bred from these matings, and 25 were straight-haired. (10 of these lived to be classified as shaker-2 and 11 as non-shaker.) Thus Rex is shown not to be allelomorphic with waved-2, and the data indicate a frequency of 41 per cent. (S.E. = 8.9 per cent.) for recombination between Rex and waved-2, on the assumption that the double mutant is viable. Rex therefore cannot lie close to waved-2, and the order must be Re—sh-2—wa-2.

Thus the map of chromosome VII of the house-mouse becomes approximately as follows :---

Map-distance (centimorgans)	•	•		←21·5→←27·5→
Order of genes	•	•	•	Rewa-2
Recombination frequency (per	cent.)	•		$\leftarrow 20\pm5\rightarrow \leftarrow 25\pm2.5\rightarrow$

The map-distances are calculated according to Kosambi (1944) by means of the r to z transformation (Fisher and Yates, table VII). The table is entered with twice the recombination fraction as r: the corresponding value of z is then twice the map-distance.

The observed recombination frequencies for the two short segments indicate a map-distance of 49 centimorgans for the Re—wa-2 segment. This corresponds with a recombination frequency of 37.5 per cent., from which the observed frequency of 41 per cent. differs by less than the standard error of the estimate. The length of the Re—sh-2 segment is not, however, determined very exactly by the data presented here, and further work is in progress in the laboratory.

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