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Brief Communication

HALOTHANE SENSITIVITY, THE H BLOOD GROUP SYSTEM AND PHOSPHOHEXOSE ISOMERASE (PHI) IN PIGS A LINKAGE STUDY OF PHYSIOLOGICAL IMPORTANCE*

Previous investigations have clearly shown the existence of associations between halothane sensitivity, the H blood group system and the PHI enzyme system in pigs (*Rasmusen & Christian* 1976, *Jørgensen et al.* 1976). These associations which have considerable practical interest are most probably linkage phenomenons (*Jørgensen* 1977, *Andresen & Jensen* 1977). The major recessive locus for halothane sensitivity (HAL) comprises the two alleles N and n, n being responsible for halothane sensitivity. The distances between this locus and the loci for H and PHI are still not known exactly. This communication aims at clarifying these problems.

Fourteen families (127 pigs), informative regarding the study of linkage, were selected from previously published investigations (Jørgensen et al., Jørgensen, Jørgensen & Wäfler 1978). In five families representing double backcross matings with respect to H and HAL (Table 1), two recombinants were observed. Lod score values (Morton 1955) for linkage between H and HAL are shown in Fig. 1. As each family consists of eight or more individuals no correction due to truncated selection has been made. Fig. 1 also shows lod score values from nine backcross-intercross matings. The overall lod score has maximum for $\theta = 0.038$ with no evidence of heterogeneity among families $(\chi^2_{14df} = 10.65, P > 0.70)$. Inclusion of two families (14 pigs) representing double backcross matings with known parental linkage phase gives a maximum lod score (17.72) for $\theta = 0.029 \pm$ 0.025, with an upper 95 % confidence limit of 0.07. This recombination fraction equals the distance between H and PHI ($\theta =$ 0.026, Andresen 1970), emphasizing the close linkage between H and HAL and the very close position of HAL and PHI.

Table 2 shows segregation of HAL and PHI in four families. Maximum lod score (12.64) was obtained for θ approaching O. Addition of the value published by others (*Andresen & Jensen* 1978) gives a total of 23.67, which makes the probability against linkage between HAL and PHI extremely small.

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Recombination frequency

Figure 1. Lod score values for linkage between halothane sensitivity and the H blood group system. A: Double backcross matings. B: Intercross-backcross matings. C: Overall lod score.

Table 1.	Segregation	in double	backcross matings with	respect to
locus for	r halothane	sensitivity	and the H blood group	locus.

Mating type	Family		Progeny phenotype		
		N, aa	N, ac	n, aa n, a-	n, ac
Nn,ac x nn,a-	1a	6	1	· 0	2
	1b	6	0	0	5
		N, aa ac	$N, \frac{a}{c}$	n, aa ac	n a- c-
	2	0	4	5	1
Nn,a- x nn,ac	3	0	4	5	0
	4	0	8	2	0
		N,aa	N,a-	n,aa	n,a-
Nn,a- x nn,aa	5	0	7	1	0

Four families gave information on linkage between HAL and PHI, HAL and H, H and PHI. One recombinant between H and PHI was observed in family 1a, and this animal was also a re-

	Family	Progeny phenotype			
Mating type		N,BB	N,AB	n,BB	n,AB
Nn,AB x nn,BB	1a 1b 3 4 5	0 0 0 0 0	7 6 4 8 7	2 5 5 2 1	0 0 0 0 0

Table 2. Segregation in double backcross matings with respect to locus for halothane sensitivity and the PHI locus.

combinant between H and HAL, but not between HAL and PHI. This finding further underlines the very close or possibly even identical positions of HAL and PHI.

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