

OBSERVATION OF LINKAGE BETWEEN *Wsp* AND THE ISOZYME LOCUS *Alat-p*

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The current linkage map (2) places the waxless mutant, *wsp*, on one end of the chromosome 7 linkage group; however, verification of this location has been difficult (G. A. Marx, pers. commun.). In my own work I have been unable to find linkage between *Wsp* and the loci T1, K, *Est-4*, or Bt, all believed to be on chromosome 7. In this communication I present evidence for linkage between *Wsp* and the isozyme locus *Alat-p*, which codes for the plastid-specific form of alanine aminotransferase (ALAT-1).

An F2 population from the cross A778-26 x A73-91 was analyzed for segregation at both *Wsp* and *Alat-p*. A778-26 was homozygous dominant for Bt, *Wsp*, and R and possessed the fast allozyme of ALAT-1. A73-91 was Wellensiek's tester (bt, *wsp*, r and displayed the slow allozyme of ALAT-1). Both lines were obtained from Dr. G. A. Marx. The ALAT-1 phenotype could be determined in young leaf extracts by horizontal starch gel electrophoresis using the buffer system consisting of 50 mM Trizma base (Sigma) neutralized to pH 8.0 with L-histidine-HCl as the electrode buffer and a 3.5 mM Trizma/histidine pH 8.0 buffer in the gel. The alanine aminotransferase assay contained 10 ml Tris-HCl pH 8.0, 20 mg L-alanine, 10 mg alpha-ketoglutarate, 15 mg NADH, 3 mg pyridoxal-5-phosphate and 200 units of lactate dehydrogenase. The assay mixture was poured on to the cut surface of the gel and allowed to sit for 10 min. The excess assay solution was poured off the gel and the gel incubated for 10-20 min at 20-30 C. The ALAT activity could be observed as dark bands against a bright fluorescent background when viewing the gel under ultraviolet light (302 nm).

Variation was present for ALAT-1 but not for the more intensely staining, cytosolic ALAT-2. Both *Wsp* and *Alat-p* segregated as monogenically inherited characters, with the alleles of *Alat-p* showing codominant expression (Table 1). Joint segregation analysis indicated that the two loci were linked, being separated by about 16 recombinant units (Table 2). These results suggest that *Alat-p* might be useful in confirming the map location of *Wsp*. However, *Alat-p* did not show linkage with T1, K, 6-pgd-p or *Pep-4*. The last two loci are linked to *Rrn-2*, the ribosomal RNA gene cluster presumably located on chromosome 7 (2). *Alat-p* also assorted independently of St, Le, Td, D, B, Fa, A, Gp, PI, *Aat-m*, *Acp-1*, and 6pgd-c (data not presented). Thus, at present, the precise location of *Alat-p* and *Wsp* remains uncertain.

1. Blixt, S. 1974. In: Handbook of Genetics, Vol. 2. K. C. King (ed.) Plenum Press, NY p. 181-221.
2. Polans, N. O., N. F. Weeden, and W. F. Thompson. 1986. Theor. Appl. Genet. 72:289-295.

Table 1. Segregation at WSP and Alat-p.

Locus	No. plants with designated phenotype*			Expected ratio	χ^2
	fast/+	H	slow/+		
WSP	62		18	3:1	0.26
Alat-p	20	43	17	1:2:1	0.68

* Designations: dominant phenotype or homozygous fast = +; heterozygous = - .
recessive phenotype or homozygous slow = - .

Table 2. Joint segregation of Wsp and Alat-p.

N	No of plants with designated Phenotype*						χ^2	Recomb.	
	+/+	+/H	+/-	-/+	-/H	-/-		Fraction	SE
80	19	33	5	1	5	12	29.0	16	4

*Designations as described in Table 1.
