LOCUS FOR THE GENE def (DEVELOPMENT FUNICULUS)

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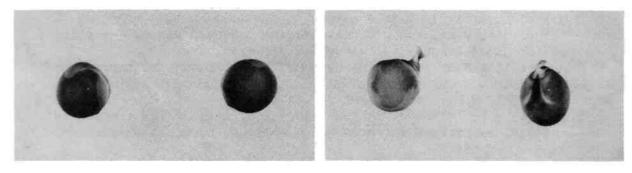
In 1966 Rozental (3) selected a spontaneous mutation, seeds of which were characterized by a strong connection of the funiculus with the hilum (Fig.1). The original test cross was carried out by Khangildin and Khangildin (2) who found monogenic, recessive inheritance and proposed the name <u>development funiculus</u> for the character and the symbol <u>def</u> for the gene. No information on the chromosomal location of the <u>def</u> locus has been found in the literature.

In 1984-88 linkage studies for the gene <u>def</u> were carried out at Wiatrowo. A line with this gene (Wt11900) was obtained from VIR/ Leningrad collection. Eleven testerlines with morphological and isozymic markers on different <u>Pisum</u> chromosomes were chosen from the Wiatrowo gene bank. The thirteen enzyme systems investigated allowed us to observe the segregation of 19 isozyme markers. Results for dihybrid segregation were analysed using a 2 x 2 contingency table to calculate the joint segregation chi-squared (to detect linkage) and the product-ratio method to calculate the recombination fraction.

 F_2 generations were analyzed beginning with cross populations with the biggest number of markers for "strategic" loci for different chromosomes, followed by populations with morphological markers for individual chromosomes as well as isozymic markers. No substantial deviations from dihybrid segregation were observed for <u>def</u> with any of the morphological markers. Thus <u>def</u> is not located in any of the well known chromosome regions which probably explains why the <u>def</u> locus was not found for over twenty years from first identification of the gene.

Isozymic markers provided the chance to investigate less known or completely unknown chromosomal regions and significant deviations from normal dihybrid segregation were revealed for <u>Def</u> with <u>Aldo</u> and <u>Gal-2</u>. (Table 1). These results indicate <u>def</u> is located on chromosome 2 (2). From the dihybrid data for these genes with <u>chi-5</u> (Table 1), one can assume that the segment Aldo Def Gal-2 is located clearly further than chi-5 (1).

The most recent studies of Weeden and Wolko (6) as well as Swiecicki (4) suggest serious changes in the Lamprecht version of the <u>Pisum</u> linkage map. For example, it is now proposed that the "upper" segment of chromosome 2 (from the <u>oh</u> locus) is located on chromosome 7. Thus on the new map the locus for def is on chromosome 7.



Def

def

Fig. 1. Normal seeds (left) without a funiculus (\underline{Def}) and mutant seeds (right) with a funiculus (\underline{def}) .

Table 1. Distribution of phenotypes in F_2 , populations segregating for def									
and chromosome 7 (chromosome 2 in Lamprecht's version) markers Aldo, Gal-2									
and Chi-5. The crosses are Wt11900 (def) x tester lines: 1) Wt8905, 2)									
Wt15063, 3) Wt15045 and 4) Wt15860									

Cross	Gene pair	Phase	Phenotype				Total	Joint	Rec.	SE
			DD	Dr	rD	rr	IOLAI	chi-sq.	frac.	5E
1)	<u>Aldo</u> :Def	С	35	3	1	6	45	22.37***	9.5	4.6
2)		С	23	1	3	3	30	8.73**	15.7	7.3
3)		С	25	1	6	9	40	16.26***	12.7	5.7
4 }		С	33	6	1	8	48	19.12***	11.8	5.0
Total			116	11	11	26	164	62.25***	15.2	3.1
2)	Aldo:Gal-2	С	22	3	2	3	30	6.00*	21.4	8.6
3)		С	25	1	5	10	40	19.12***	11.1	5.2
Total			47	4	7	13	71	25.77***	16.1	4.8
2)	Gal-2:Def	С	25	1	1	4	30	17.98***	8.1	5.2
3)		С	30	1	1	9	40	30.87***	5.0	3.6
Total			55	2	2	13	72	49.79***	6.1	2.9
3)	<u>Chi-5</u> :Gal-2	R	21	10	9	1	40	1.91	30.3	14.2
3)	<u>Chi-5</u> :Aldo	R	16	15	9	1	40	4.68*	22.8	14.8
3)	<u>Chi-5:Def</u>	R	22	9	9	1	40	1.48	32.2	14.0

*,**,*** Probability less than 0.05, 0.01 and 0.001, respectively.

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