

Table 1. Observed and expected numbers of segregants in each of three phenotypic classes in F2 from crosses of mutant x normal (cv. 'Torsdag').

Cross	Segregation in F2			
	Observed	Expected at a 1:2:1 ratio	χ^2	P
Mutant 5 x Torsdag	34:71:40	36.2:72.5:36.2	0.56	0.9 - 0.8
Torsdag x Mutant 5	32:56:36	31:62:31	1.42	0.5 - 0.2
Mutant 66 x Torsdag	29:81:36	36.5:73.0:36.5	2.42	0.5 - 0.2
Torsdag x Mutant 66	21:41:21	20.8:41.5:20.8	0.01	1.0 - 0.9

Table 2. A morphological characterization of the original line, mutants, and heterozygotes in pea.

Variety, mutant, hybrid	Height (cm)	Node number J/	Average internode length (cm)	Stem number per plant	Length/width of leaflets (cm)	Length/width of stipules (cm)
Torsdag	77.3	18.1	4.3	1	5.4/2.8	5.8/4.0
Mutant 5	31.9*	17.3*	1.8*	1	3.9/1.8*	3.7/2.6*
Mut.5xTorsdag	51.3*	17.6	2.9*	1	4.7/2.2*	4.6/3.2*
Mutant 66	15.7*	12.5*	1.2*	3.3*	3.8/1.9*	3.8/2.4*
Mut.66xTorsdag	31.8*	16.4*	1.9*	2.0*	4.5/2.5*	4.6/3.4*

1/Measurements were made at the full flowering stage.

Table 3. Frequencies of natural and induced mutations in the original and semidominant mutant lines of pea.

	Original line		Mutant 5		Mutant 66	
	Control	Gamma-ray treated	Control	Gamma-ray treated	Control	Gamma-ray treated
Total number of families	241	221	221	199	195	161
Families with mutants (%)	0.4	7.2	4.1**	14.1*	3.6*	5.6**
Total number of plants	7993	4927	6141	3575	4457	2504
Mutants (%)	0.04	1.48	0.31***	2.40**	0.25***	0.48***
Number of mutation types	1	15	8	18	8	10

* - P=0.05; ** - P=0.01; *** - P=0.001 as compared with the original line
