

## Comparison of Variability among Irradiated and Control Inbred Maize Lines via Morphological Descriptions and Some Quantitative Features

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### SUMMARY

*Knowledge of genetic diversity in breeding material is fundamental for hybrid selection programs and for germplasm preservation as well. Research has been done with nine irradiated (fast neutron) and four non-treated inbred lines. The aims of this study were (1) to investigate the degree of genetic variability detected with morphological description (based on CPVO TP/2/2) in these materials, (2) to compare the genetic changes among irradiated and non-irradiated maize inbred lines (based on some quantitative features). The irradiation did not change any of the characteristics clearly in positive or negative way, which can be related to the fact that the effect of induced mutation on genetic structure cannot be controlled. From the irradiated lines we have managed to select plants with earlier ripening times and better phenotypes. We could distinguish 3 main groups by the morphological features; these results match our expectations based on pedigree data. Markers distinguishable on the phenotypic level (e.g. antocyanin colouration, length of tassels) were significant in all lines.*

**Keywords:** fast neutron irradiation, genetic variability, maize (*Zea mays L.*), morphological description, quantitative features

### INTRODUCTION

Application of induced mutation in plant breeding began more than seventy years ago. Presently, there are 2252 officially recognized mutant strains generated by physical or chemical mutagenesis; more than half of which were produced during the last 20 years and likely many more strains exist bearing mutated genes built in by crosses (Maluszynski et al., 2000).

In sophisticated plant production, the most essential factors are variety, its genetic productivity, quality and yield stability. Different morphological mutants offer good possibilities for the production of modern hybrid combinations (Pepó and Pepó, 1993). The specific effects of the different radiation sources have to be investigated in detail as to their economically beneficial characteristics (Bálint, 1960). The mutant genes often build into a new strain not directly but by crosses (Szilágyi, 1978 cit. Bálint, 1996).

The recurrent selection achieved with inbred lines produced by mutational breeding can be combined with *in vitro* tissue culture and haploid techniques. In addition good quality combinations can be generated (Pepó, 2004; Pepó et al., 2004). Genetic variability – as the base of all kinds of plant breeding – is desired in maize breeding too. The determination of similarity among lines of different origin is also cannot be neglected (Rady and Nagy, 1996). Hajós

et al. (1996) studied the genetic variability in the case of maize lines produced by colchicine treatment and chronic gamma irradiation and found significant differences between the economically important phenomena of different lines.

In Sutka and Bálint's studies (1971) different quantitative characteristics responded differentially to mutagen treatments. The ear heights and the number of lateral branches were more variable. The number of grain rows is determined genetically and between certain limits can be considered as strain markers (Menyhért, 1979). In the UPOV TG 2/6 description of plant height it is also classified into the group of genetically complex, environmentally stable phenomena (Zsubori et al., 2002). Allen et al. and McKee et al. (cit. Zsubori et al., 2002) showed that the number of leaves and yield significantly correlated with the height of the plant.

The economically important features show complex mode of inheritance. Several authors published data on particular type of inheritance and values of heredity related to plant height, grain yield, etc. (Bálint, 1966; Herczegh, 1970; Schön et al., 1993). LAI (*leaf area index*), the size of assimilating surface, is one of the most important morphological markers (Berzsenyi, 2000). With the help of strain descriptions resulting from DUS studies it becomes possible to determine the most similar strains and to investigate similarity groups (Veress and Matók, 1999).

In this study we have examined the genetic variability among 9 irradiated and 4 original inbred maize lines. The detection of genetic variability has been done by morphological description (following the CPVO TP/2/2 guidelines) in the phenotypic level, supplementing with the data of the most important quantitative features of the lines. Our aim was to study the genetic variability resulted by the effect of irradiation and comparing these data to those of the control lines. Knowing the genetic diversity of these lines we can exploit their genetic potential (selecting parental lines, reducing the number of crosses) in our future hybrid breeding program.

### MATERIALS AND METHODS

Plant material: 4 original and 9 irradiated inbred maize lines (*Table 1*). For the description of the irradiation treatment see Maráz et al. (1993) and Pepó and Tóth (2004). After 12 years of inbreeding a line can be considered as homozygote. Their selection can be done by their good combining and pollen producing abilities, strength of stalks.

Table 1

Origin of the tested lines

Inbred lines	Origin	Type of irradiation	Dose (Gray)
GK Mo 17 [control]	Cereal Research Non-Profit Company, Szeged	-	-
UD 507	GK Mo 17	Fast neutron	5
UD 355	GK Mo 17	Fast neutron	5
GK F2 [control]	Cereal Research Non-Profit Company, Szeged	-	-
UD 105	GK F2	Fast neutron	10
UD 201	GK F2	Fast neutron	12.5
GK 13 [control]	Cereal Research Non-Profit Company, Szeged	-	-
UD 123	GK 13	Fast neutron	5
UD 433	GK 13	Fast neutron	15
GK 41 [control]	Cereal Research Non-Profit Company, Szeged	-	-
UD 70	GK 41	Fast neutron	10
UD 126	GK 41	Fast neutron	5
UD 267	GK 41	Fast neutron	20

Morphological description: lines were sown in two areas in randomized block design in biennial, small plot experiments, in 4 replications. The measured characteristics were determined by the average of 15 plants/plot following the CPVO TP/2/2 guidelines (using 25 features) and other quantitative characteristics. For the calculation of leaf area we have used Montgomery's formula (cit. Anda and Tóbiás, 1999). Data were evaluated by one-way variance analysis.

RESULTS

Variability of quantitative features

Data of plant height were taken up 1 week after flowering. Heights were reduced by the effect of irradiation in almost all cases (Table 2). The only exception was UD 105 line, which significantly overgrew the starting control line. Stalk diameter, an important marker of stalk strength, showed both negative and positive alterations comparing to the original line. The ear height was the feature that

changed in the highest degree. The height did not influence the position of the upper ear. There was about a 20 cm difference in height between the original GK 13 and UD 433 lines. In these cases there were no striking differences between the control and treated lines, significant changes took place in both positive and negative ways. The number of leaves/plant showed significant negative changes only in the cases of GK 41 and its irradiated lines, except UD 126. Leaf area index was significantly increased in the cases of UD 105 (compared to GK F2) and UD 126 (compared to GK 41). A possible explanation can be found in the notable raise of the interval to mid-silking – higher with minimum of one FAO maturity group –, which results in the rise of their vegetation period. The time of mid-silking from planting showed significant decrease in each groups compared to line GK Mo 17, while significant increase was observed in the case of other groups. The more than two-week shift of flowering of line UD 267 compared to the control is conspicuous. Possibly the high irradiation dose – 20 Gy – triggered such alteration.

Table 2

Variability of some quantitative features in irradiated and control maize lines (Debrecen, 2005)

Lines	Plant height (cm)	Diameter of maize-stalk (mm)	Ear height (cm)	Number of leaves per plant	Leaf area index (m <sup>2</sup> /m <sup>2</sup> )	1000 kernel weight (g)	Number of days from planting to mid-silking
GK Mo 17	228.1	22.1	79	12	2.29	366.25	-
UD 507	138.4***	20.3 <sup>+</sup>	55***	11 <sup>+</sup>	2.07*	310***	-3**
UD 355	132***	20 <sup>+</sup>	40***	10 <sup>+</sup>	1.45***	309.75***	-12***
GK F2	121.2	13.2	35	9	0.72	204	-
UD 105	186***	16**	72***	11 <sup>+</sup>	3.1***	373***	+8***
UD 201	132.4*	9.8**	33.2***	8 <sup>+</sup>	1.02**	185.25***	+6***
GK 13	162.6	13.2 <sup>+</sup>	33.4	11	1.64	350.75	-
UD 123	151.3***	14.4 <sup>+</sup>	39.7**	10 <sup>+</sup>	1.67 <sup>+</sup>	329.75***	+5*
UD 433	149.5***	13 <sup>+</sup>	53.3***	9.5 <sup>+</sup>	1.16**	269.5***	+4*
GK 41	188.9	20.7	58.5	12	2.67	313	-
UD 70	154.7***	22.1 <sup>+</sup>	48.2***	10.3**	1.38***	281.75***	+9***
UD 126	175***	18 <sup>+</sup>	51.2***	11.2 <sup>+</sup>	1.71***	344.75***	+13***
UD 267	165.2***	17.1**	42.7***	10**	1.72***	310**	+15***

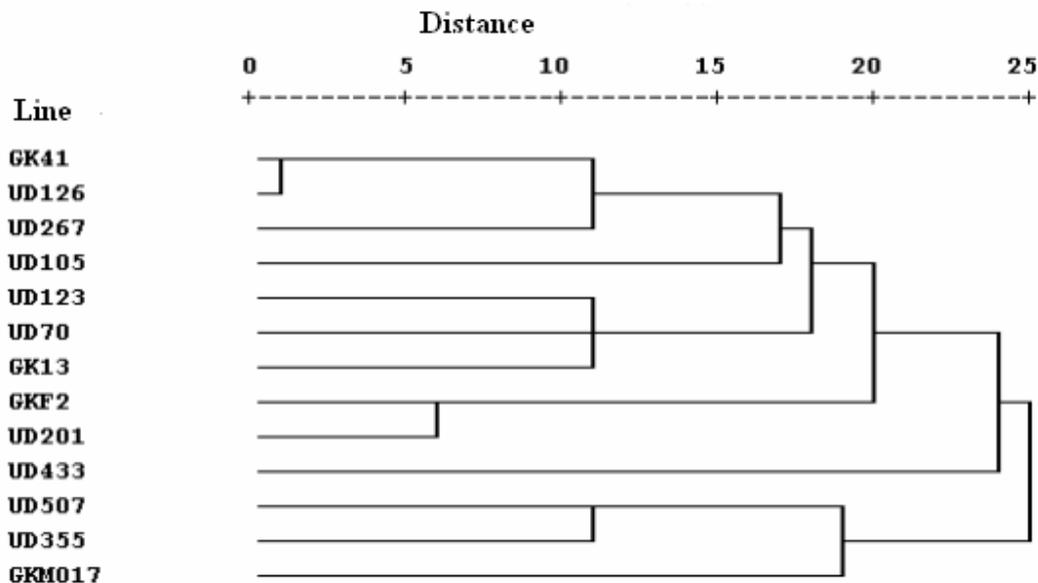
Values followed by \*, \*\*, \*\*\* are significant at 0.1; 0.05; 0.001 probability levels in comparison with control lines. Values followed by <sup>+</sup> are not significant.

### Analysis of morphological features

The dendrogram on *Figure 1* has been created by 25 examined features following the guidelines of CPVO TP/2/2. Based on the similarities among

inbred lines we could distinguish 3 major groups. Group 1: *GK 41, UD 126, UD 267; GK F2, UD 105, UD 201*. Group 2: *UD 123, UD 433, UD 70*. Group 3: *GK Mo 17, UD 507, UD 355, GK 13*.

*Figure 1: Dendrogram obtained from cluster analysis based on the morphological descriptions*



Based on the pedigree data in Group 1 *GK F2* and its irradiated lines (*UD 105* and *UD 201*) can be clearly separated from the others. Considering their kernel types these lines are not in close relationship with none of the other groups. Markers distinguishable on the phenotypic level (antocyanin colouration, length of tassels, etc.) were significant in all lines. The slightest morphological change was detected between *GK 41* and *UD 126*.

### DISCUSSION

To summarize our results we can conclude that the effects of phenomena possibly originated in the mutations caused by irradiation induced both positive and negative alterations in the plants.

Our results confirm the statement that the dose of irradiation can be different in all genotypes subject to

the aim of experiment. According to the general hypothesis (Bálint, 1967) that low doses cause micro while high doses macro mutations is correct but the barriers of micro and macro mutations can be found at different levels in each genotypes. The most effective and cost-saving methods to increase genetic variability are still the application of physical mutagens. Their importance in maize breeding can increase in the next years since the world-wide spreading of genetically manipulated maize hybrids can narrow the genetic bases either in gene centre of maize or in other maize producing areas of the world.

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