

GENETIC ANALYSIS OF RYE (*SECALE CEREALE* L.)

II. LEAF PEROXIDASE ISOENZYMES IN TRISOMIC AND TELOTRISOMICS OF CHROMOSOME OF 1R<sup>1</sup>

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**Summary.** Leaf peroxidase activity was analysed in trisomics and telotrisomics of the rye variety "Esto". Trisomic 1R showed a typical alteration in peroxidase pattern in comparison with the variety "Esto". The same difference was observed in telotrisomic 1RS; trisomic 1RL was comparable with the standard. This means, that at least one gene for peroxidase activity (PER 1) is located on the chromosome arm 1RS.

Only few genetic analyses were carried out in rye in comparison with other cereals. They concerned the inheritance of characters and their localization on specific chromosomes (Schlegel, Mettin 1982). However a successful use of modern breeding methods demands the instruction of important characters. At the same time it is necessary to find markers which make the breeding-process more efficient. The trisomic analysis is one of the methods to localize genes on chromosomes. Using trisomics and their specific segregation after crossing in  $F_2$ -like progenies it was possible to find several markers on chromosomes (Sturm, Engel 1980, Sturm et al. 1981, Sturm, Müller 1982, Melz et al. in press). Until now biochemical markers of rye were localized only by analysing wheat-rye hybrids (Irani, Bhatia 1972, Tang, Hart 1975, Höhler et al. 1979, Rao, Rao 1980, Artyomova 1982, Hsam et al. 1982). The aim of our work was to find a possibility to localize biochemical markers of rye by trisomics as well.

MATERIAL AND METHODS

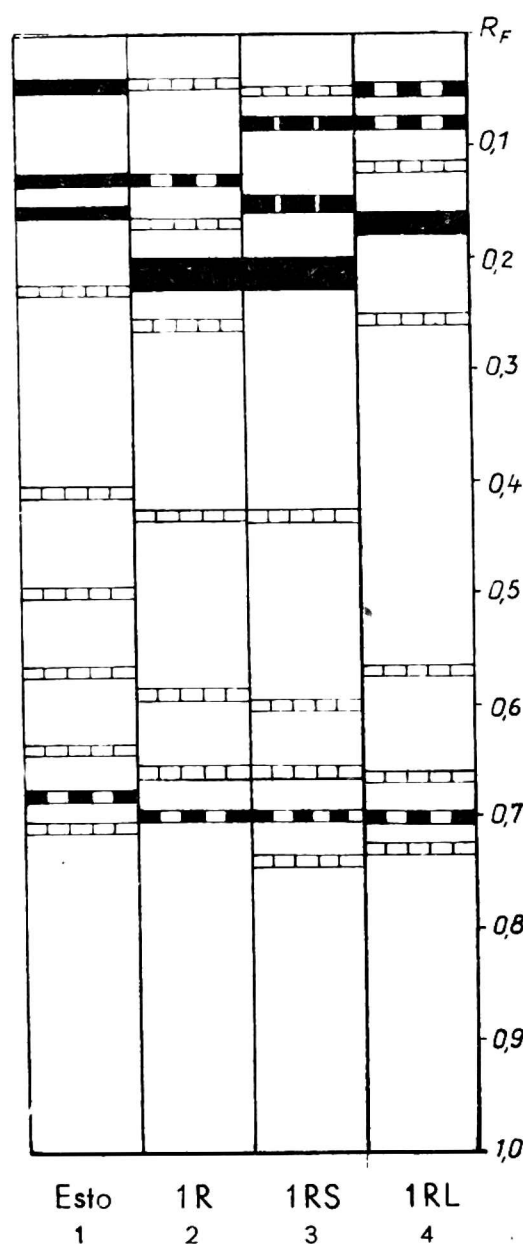
The analysis was done by the help of a set of trisomics derived from the rye variety "Esto" (Sturm 1978). The trisomics were named following the nomenclature of the 1st Workshop on Rye Cytogenetics in Wageningen (1982), which means that the

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chromosomes were designated as  $1R$  to  $7R$ . In addition to this, a set of telotrisomics (Sturm, Melz 1982) was used for analysis. Trisomic and telotrisomic plants were cloned and the youngest leaves of the clones were taken for preparation with five repetitions. The clones grew under the same environmental conditions. The leaves were extracted in 0.1 M Tris-HCl-buffer pH 0.8 (Staples, Stahmann 1964) and then a disk-electrophoresis was carried out in basic standard system No. 1 (Maurer 1968) with cylindric gels ( $5 \times 70$  mm). Multiple forms of peroxidases (EC 1.11.1.7) were detected by staining (Rychter, Lewark 1971). Zymograms of trisomics and telotrisomics were compared with the zymogram of the standard variety "Esto".

## RESULTS AND DISCUSSION



The standard variety "Esto" showed a typical zymogram (Fig. 1), while trisomic  $1R$  ( $1R$  = SAT chromosome) was different. The band at  $R_F = 0.23$  is of considerable intensity, while it is weak in the standard and the other trisomics. This result inclined us to test the two possible telotrisomics of chromosome  $1R$  ( $1RS$  and  $1RL$ ) as well. The zymogram of telotrisomic  $1RL$  was identical with that of the standard, while telotrisomic  $1RS$  showed the same pattern as trisomic  $1R$  (Fig. 1). This means, that at least one gene for peroxidase activity is located on chromosome arm  $1RS$ . The gene should be designated as PER 1.

These results prove, that it is possible to localize biochemical markers using trisomics and telotrisomics of rye. Moreover, findings of Höhler et al. (1979) were confirmed, but there is no qualitative effect in case of trisomics as in wheat-rye hybrids. As special crosses were not made before preparation and all

Fig. 1. Leaf peroxidase isoenzymes

1 - "Esto", 2 - trisomic  $1R$ , 3 - telotrisomic  $1RS$ , 4 - telotrisomic  $1RL$

trisomics and telotrisomics descend from "Esto", a strong band ( $R_F = 0.23$ ) is not likely to be due to specific allele combinations. On the other hand, it must be concluded, that there is a dosage effect in the case under study. This agrees well with the findings for *Petunia* (van den Berg, Wijsman 1982), barley (Nielsen, Frydenberg 1971), maize (Nielsen, Scandalios 1974), tomato (Fobes 1980) and *Lolium* (Lewis et al. 1980). The reason for the dosage effect could be the additional gene copy in the trisomic genotype  $AAA$  resulting in more gene products than with the

genes in the disomic genotype *AA*. This results finally in a higher intensity of a usually weak band at  $R_F=0.23$ .

It must be expected that there are further genes controlling leaf peroxidases, because Garcia et al. (1982) observed 13 endosperm and embryo peroxidase genes in rye. Our research work in this field has not been completed yet, but there is some reason to expect further peroxidase genes on rye chromosomes *2R*, *3R* and *7R*.

If it is possible to localize more isoenzymes on rye chromosomes, the identification of chromosomes will be more reliable. This possibility is especially important for identification of rye chromosomes and their part in substituted or translocated wheat, for recognition of structural changes in rye chromosomes after crossing with other species and for completion of a standard nomenclature of rye chromosomes.

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### ANALIZA GENETYCZNA ŻYTA (*SECALE CEREALE* L.)

#### II. IZOENZYMЫ ПЕРОКСЫДАЗ В ЛИՒՒՑИАՒИ ТРИСОМИКՒՒ И ТЕЛОТРИСОМИКՒՒ ХРОМОСОМУ 1R

##### Streszczenie

Analizowano aktywność peroksydazy w liściach trisomików i telotrisomików chromosomu 1R żyta odmiany Esto. Trisomik 1R charakteryzuje się typową zmianą w obrazie peroksydazy w porównaniu z odmianą Esto. Taką samą różnicę zaobserwowano u telotrisomika 1RS; trisomik 1RL był porównywalny ze standardem. Oznacza to, że przynajmniej jeden gen dla aktywności peroksydazy PER1 znajduje się na ramieniu chromosomu 1RS.

### ГЕНЕТИЧЕСКИЙ АНАЛИЗ РЖИ (*SECALE CEREALE* L.)

#### II. ИЗОЭНЗИМЫ ПЕРОКСИДАЗ ЛИՒՒՑЬЕВ У ТРИСОМИКОВ И ТЕЛЕТРИСОМИКОВ ХРОМОСОМЫ 1R

##### Резюме

Анализировалась активность пероксидазы листьев у трисомиков и телотрисомиков ржи сорта Esto. Трисомик 1R характеризовался типичным изменением в спектре пероксидазы по сравнению с сортом Esto. Такая же самая разница наблюдалась у телотрисомика 1RS; трисомик 1RL сравнивался со стандартом. Это значит, что по крайней мере один ген для активности пероксидазы PER1 находится на плече хромосомы 1RS.