

## RFLP Report

# Chromosomal location and RFLP utility in wheat and barley of a wheat gene with homology to a 7S storage-globulin sequence

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Source of the probe

A cDNA library was constructed using poly(A<sup>+</sup>) RNA extracted from abscisic acid (ABA)-treated immature wheat embryos as described by Williamson et al. (1985). A cDNA clone (PSP511) was isolated which included a 704-bp insert containing sequences that are similar to regions of the legume 7S storage-globulin genes (Quatrano et al. 1986).

#### **Chromosomal location**

Hybridization of PSP511 to genomic DNA from 21 nullisomic-tetrasomic (NT) lines, digested with EcoRV. DraI and HindIII, and ditelosomic (DT) lines, digested with EcoRV, of the Triticum aestivum cultivar Chinese Spring (CS) revealed homoeologous sequences only on chromosome arms 4AL, 4BS and 4DS. Six hybridizing fragments were detected with all three restriction enzymes (Fig. 1), indicating the probable presence of two homologous copies of the globulin genes (Glo) in each of the three wheat genomes. Similarly, evidence was obtained for two gene copies in rye and barley, located respectively on 4H (barley chromosome 4) and 4R as shown by analysis of the single-chromosome disomic addition lines. Both copies of XGlo were polymorphic in all five genomes investigated. Segregation of the alleles on 4AL, 4BS and 4DS in wheat, 4H in barley and 4R in rye, showed no recombinants, indicating that both copies are tightly linked in each of the genomes. Each population consisted of 120 F<sub>2</sub> individuals and thus the upper limit of distance between the two genes, at 95% confidence, can be calculated as 0.25 cM. Hence, XGlo can be considered as a single locus for most marker applications. The map locations of XGlo-4A, -4B and -4D have been determined in the cross CS × Synthetic to lie 19 cM, 23 cM and 39 cM from the midpoints of the map intervals (3, 2 and 5 cM respectively) spanning the centromeres on 4AL, 4BS and 4DS. The larger map distance obtained in chromosome 4D is consistent with previous observations that the D genome chromosomes in this cross recombine more frequently than their A and B homoeologues (Devos et al. 1992).

The chromosomal locations of the XGlo loci correspond closely with those of the 7S globulin loci on the 4D<sup>t</sup> genetic map constructed by Lagudah et al. (1991) in the wild diploid T. tauschii. However, the four copies in the D<sup>t</sup> genome and the dispersal of the three copies which map on 4D<sup>t</sup> are at variance with the results described here. This may be explained by the use of pEG4.6 (P. Chandler, personal communication), a large genomic clone isolated by hybridization with PSP511 which may carry sequences in addition to the globulin genes.

Locus symbol

XGlo(XGlo-4A, -4B, -4D, -4H, -4R)

#### **Polymorphisms**

Estimates of levels of polymorphism, based on allele frequencies obtained in a panel of 15 wheat varieties

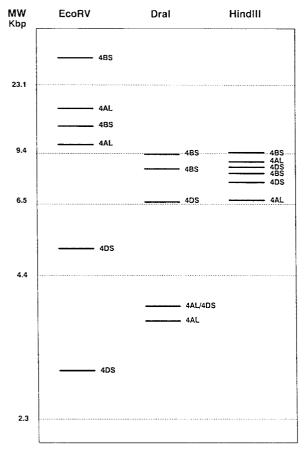


Fig. 1. Chromosome assignment of Chinese Spring restriction fragments hybridizing with PSP511

digested with three restriction enzymes, showed that XGlo-4A was highly polymorphic with all restriction enzymes (H = 51%), XGlo-4B was moderately variable (H = 39%), while at XGlo-4D the only variant allele was found in Sears' Synthetic. XGlo-4H revealed three alleles in 13 varieties digested with three enzymes, giving a relatively high potential heterozygosity index of 57%.

### Presence of related sequences in alien relatives of wheat

Hybridization of PSP511 to *DraI*-restricted genomic DNA of accessions of 21 alien relatives of wheat, including *Triticum*, *Hordeum*, *Secale*, *Aegilops*, *Agropyron* 

and Dasypyrum species, and of the Gramineae species Lolium perenne, Zea mays and Pennisetum glaucum, revealed the presence of at least one, but mostly two, hybridizing fragments in the species analysed, indicating that the globulin gene sequence is highly conserved among the Gramineae.

#### Other studies of wheat Glo genes

Expression of the wheat globulin genes is characterized by the appearance of mRNA transcripts of 2.5 kb in RNA isolated from developing wheat grains (Quatrano et al. 1986), ABA-treated wheat embryos (Williamson and Quatrano 1988), and in drought-stressed wheat leaves (Berge et al. 1989).

Probe availability

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