Group 3 Chromosome Bin Maps of Wheat and Their Relationship to Rice Chromosome 1

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Wheat deletion lines have also been used extensively

Group 3 Chromosome Bin Map



Figure

643

nificantly matched an Arabidopsis coding region. The blastX comparison of all mapped-EST unigenes against the Arabidopsis protein database returned 1799 (32%) matches for all mapped-EST unigenes and 313 (32%) for wheat group 3. The number of unigene matches per Arabidopsis chromosome did not significantly differ from that expected, on the basis of the estimated coding region content per chromosome, for wheat group 3 or total mapped-EST unigenes with either blastN or blastX analyses.

DISCUSSION

Large-scale sequencing of ESTs and generation of a high-density chromosome bin map for hexaploid wheat

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and distribution of genes among the three genomes of polyploid wheat. Genetics **168**: 701–712. Greene, R. Kantety *et al.*, 2003 Comparative DNA sequence analysis of wheat and rice genomes. Genome Res. **13**: 1818–1827. Sal ina, E., A. Borner, J5(w1A.)-212Nl ina,J5-10Korzunin**3**(Greene, R. Kantety *et al.*, 2003 Comparative DNA sequence