

**PI: Herbert Ohm****Project ID: 0405-OH-021****Research Area: BIO****Project Title: Mapping of Novel FHB Resistance Transferred from *Lophopyrum ponticum* to Wheat.****PI's E-mail: hohm@purdue.edu****FY03 ARS Agreement #: 59-0790-9-057****Duration of Award: 1 Year**

**PROJECT 1 ABSTRACT**  
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The objective of this research is to map novel Fusarium head blight (FHB) resistance QTL of major effect located on the long arm of chromosome 7el<sub>2</sub>, and identify DNA markers closely linked to the resistance gene(s). A new source of type II resistance against FHB was identified in a hexaploid wheat line that contains a chromosome from *Lophopyrum ponticum* (genome el<sub>2</sub>) and its derived translocation lines. The resistance was localized on the long arm of chromosome 7el<sub>2</sub>. Understanding the genetic basis of this trait is prerequisite to utilizing this source of resistance in wheat breeding. One difficulty in studying the genetics of useful traits derived from wild relatives is that the respective wheat chromosome rarely recombines with the introgressed homoeologous alien chromosome. This project is designed to map the FHB resistance gene(s) in a population derived from a cross between two substitution lines. The two alien chromosomes, 7el<sub>1</sub> and 7el<sub>2</sub>, have been shown to be homoeologous and have a high frequency of pairing. An F<sub>2</sub> population from a cross between line K2620 (FHB resistant) in which chromosome 7el<sub>2</sub> replaced 7D of wheat cv. Thatcher, and line K11463 (FHB susceptible) in which 7el<sub>1</sub> replaced 7D of Thatcher, will be available in spring 2004 for disease evaluation and DNA extraction. We are confident that we can phenotype single F<sub>2</sub> plants for this highly effective FHB resistance from 7el<sub>2</sub> based on our experience in previous tests (Shen and Ohm, 2003). However, we intend to characterize certain F<sub>3</sub> families in fall 2004. We have identified two SSR markers and one CAPS marker on the long arm of chromosome 7 of Lophopyrum that are polymorphic between 7el<sub>1</sub> and 7el<sub>2</sub>. We will also identify AFLP markers that are polymorphic between 7el<sub>1</sub> and 7el<sub>2</sub> to locate the FHB resistance of 7el<sub>2</sub>. DNA markers developed from this research will be useful in future research to reduce the length of the 7el<sub>2</sub> chromosome segment on which this resistance is located using a ph1B deletion line and also to incorporate this FHB resistance into adapted wheat lines.