A NOVEL NONHEME HALOPEROXIDASE GENE FOR DISEASE RESISTANCE IN TRANSGENIC COTTON AND TOBACCO Kanniah Rajasekaran, Jeffrey W. Cary, Thomas J. Jacks and Thomas E. Cleveland USDA, ARS, Southern Regional Research Center New Orleans, LA Kurt Stromberg University of Southwestern Louisiana Lafayette, LA

## **Abstract**

Genetic transformation of cotton and tobacco were carried out with a novel gene, chloroperoxidase (CPO), from *Pseudomonas pyrrocinia*. Chloroperoxidases and other haloperoxidases convert hydrogen peroxide to much more potent antimicrobial compounds, hypochlorous acid and peracetic acid.

We tested the potential of chloroperoxidase as an antifungal agent using transgenic tobacco model system. A large number of transgenic tobacco plants producing chloroperoxidase were obtained by Agrobacteriummediated transformation. Successful transformation was shown by npt II ELISA, PCR, Southern, northern and Using the substrate, western blot analyses. monochlorodimedon, the halogenating activity of the chloroperoxidase enzyme was also demonstrated in transgenic tobacco leaf extracts. Plant extracts from tobacco plants transformed with the chloroperoxidase gene significantly reduced the number of fungal colonies arising from germinating conidia of both Aspergillus flavus and Verticillium dahliae compared to the extracts from the nontransformed control. For example, leaf extracts from transformed tobacco inhibited the growth of germinating conidia of A. flavus by more than 90%. Leaf extracts from transgenic tobacco also reduced the number of colonies arising from germinating conidia of Fusarium moniliforme, although the results were not significant. The transformed tobacco plants showed greater levels of disease resistance in planta, against a fungal pathogen, Colletotrichum destructivum, which causes anthracnose and a bacterial phytopathogen, Pseudomonas syringae pv. tabaci, which causes the disease, fireblight.

Parallel transformation experiments with cotton have also been carried out. Analyses to date indicate successful transformation as shown by *npt* II ELISA, PCR-Southern and by the demonstration of enzymatic activity in transgenic cotton cells. We have just begun assaying the chimeric cotton callus cells for antifungal activity. Preliminary results with extracts from selected cotton callus colonies indicate significant antifungal activity against germinated spores of *A. flavus* and *V. dahliae*. Several of the transgenic callus colonies have been induced to form somatic embryos and we are in the process of regenerating plantlets for further evaluation in the greenhouse.

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