CITRUS ADVANCED TECHNOLOGY PROGRAM

QUARTERLY & FINAL REPORTS: Control of Citrus Greening, Canker & Emerging Diseases of Citrus

SELECT PERIOD June 2016

(●) *Quarterly Report* ( ) *Final*

Proposal Title

Engineering Citrus for Canker Resistance

Today's Date	Sponsoring Organization (drop-down)	Category (drop down)
7/15/2016	Citrus Research and Development Foundation	Transgenic/Metagenomic Mediation of Resistance

REPORT UPDATE (500 words-It is not necessary in this public report to disclose your institution's proprietary information or intellectual property.)

Objective 1: Assess canker resistance conferred by the PAMP receptors EFR and XA21
Three constructs were used for genetic transformation of Duncan grapefruit and sweet orange as part of a previous grant: EFR, EFR coexpressed with XA21, and EFR coexpressed with an XA21:EFR chimera. Five putative transgenics have survived and passed a PCR screen, and these will soon be grafted onto rootstocks. To ensure that there will be sufficient events to analyze to come to a conclusion about the effectiveness of these genes, we have initiated more transformations in Duncan grapefruit at the Core Citrus Transformation Facility at UF Lake Alfred. EFR, XA21, and XA21 + EFR constructs have been re-created with the inclusion of a GFP marker for confirmation of transformants; selection is underway. In addition, we have added the recently-identified Cold Shock Protein Receptor (CSPR) to the transformation queue.

Objective 2: Introduction of the pepper Bs2 disease resistance gene into citrus
Two constructs were created to coexpress Bs2 with other R genes that may serve as accessory factors for Bs2. These constructs have been provided to the Lake Alfred transformation facility, and selection of transformants in Duncan grapefruit is underway.

Objective 3: Development of genome editing technologies (Cas9/CRISPR) for citrus improvement
The initial target for gene editing is the citrus homolog of Bs5 of pepper. The recessive bs5 resistance allele contains a deletion of two conserved leucines. The citrus Bs5 homolog was sequenced from both Carrizo citrange and Duncan grapefruit, and conserved CRISPR targets were identified. Four CRISPR constructs are being created in the Staskawicz lab:
C1) A construct targeting two sites that will produce a 100 bp deletion in Bs5 in both Carrizo and Duncan (the bs5 transgene will be added);
C2) A construct targeting a site overlapping the two conserved leucines;
C3) C2 with the addition of a bs5 repair template for Carrizo that will not be cut; and
C4) C2 with a similar repair template for Duncan grapefruit.

The constructs have been tested by co-delivery into Nicotiana benthamiana leaves with another construct carrying the targeted DNA from Carrizo or Duncan varieties, and verified to function. To aid in the selection of positive transgenics, a GFP reporter has been added into each CRISPR construct. The constructs with GFP will be re-verified in N. benthamiana and sent for transformation.

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Form PR-12 Quarterly Reports **SUBMIT REPORT**

CITRUS RESEARCH AND DEVELOPMENT FOUNDATION, INC.