

CITRUS HUANGLONGBING (HLB) DISEASE

CURRENT USDA-ARS RESEARCH

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ARS HLB RESEARCH



- Broad in scope
- Basic / Discovery
- Applied / Development

Locations

- Ft. Pierce, FL
- Winter Haven, FL
- Beltsville, MD
- Frederick, MD
- Parlier, CA
- Riverside, CA
- Weslaco, TX

Multidisciplinary

- > Plant Pathology
- > Entomology
- > Microbiology
- > Plant Physiology
- > Plant Biochemistry
- > Molecular biology
- > Genetics
- > Omics/Bioinformatics

Collaborative



ARS HLB RESEARCH



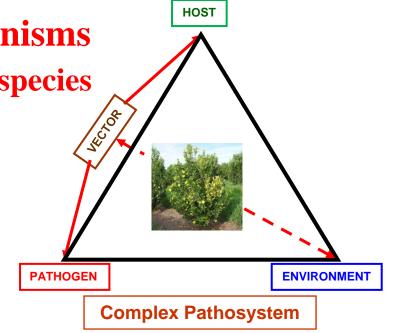
Primary Components of Complex Pathosystem to Identify Potential Targets to Mitigate Disease

- Host
 - * Responses to infection
 - * Resistance

HLB-associated microorganisms

* "Candidatus Liberibacter" species

- * Phytoplasmas
- * Other?
- Insect vector(s)
 - Citrus psyllids







BROAD RESEARCH AREAS

- **Disease detection pre-symptomatic**
- Disease diagnosis symptomatic
 - clinical samples (plant tissue)
- Candidate pathogen(s) detection & identification
 - plant tissue; insect vector

 - **Etiology Epidemiology**
 - Disease Management / Mitigation





DISEASE DIAGNOSIS / PATHOGEN(S) DETECTION & IDENTIFICATION

- PCR
 - * protocols
 - * primers
- Serological assays
- Detection arrays
- Novel platforms
- > HLB-associated microorganisms/candidate pathogens (e.g., liberibacters)
- Host response(s) to infection





ETIOLOGY

- Identification of etiological agent(s) of HLB
 - * (candidate) pathogens associated with HLB
 - > "Candidatus Liberibacter" species
 - > Phytoplasmas
 - > Other?
- Cultivation HLB-associated microorganisms in vitro
- Fulfillment of Koch's postulates





EPIDEMIOLOGY

- Diversity of HLB-associated microorganisms (e.g., Las)
- Population dynamics of HLB-associated microorganisms in citrus
- Endophytic microbial community in citrus:
 - * Characterization (qualitative; quantitative)
 - * in planta & in insecta dynamics & interactions
 - * Potential role in HLB
- Insect vector biology / ecology





INTERACTIONS BETWEEN HOST and HLB-ASSOCIATED MICROORGANISM(S)

- In planta distribution of HLB-associated microorganism(s) interactions
- Biochemical & physiological responses in HLB-affected citrus
- Gene expression in HLB-affected citrus plants
 - * Identification of genes whose expression is altered in HLB-affected plants compared to that in non-diseased plants
- Effect(s) of HLB on citrus fruit juice quality





INTERACTIONS BETWEEN INSECT VECTOR and HLB-ASSOCIATED MICROORGANISM(S)

- Acquisition & transmission parameters
- Factors that affect
 - * Acquisition & transmission of HLB-AM
 - * Vector specificity
 - * Vector competence
- Localization of HLB-associated liberibacter in various insect vector tissues





- Host resistance
 - * Screen citrus genomic resources to identify new, potential sources of resistance/tolerance to HLB
 - * New citrus scion & rootstock germplasm & cultivars with enhanced resistance/tolerance to HLB
 - * Induced resistance (pathogenesis-related proteins)
 - * Transgenics
 - > antimicrobial peptides/proteins
 - other (chimeric proteins, plant defensins)





- Cultural practices
- Therapeutic treatments
 - * Heat treatment of budwood
 - * New antibiotics
 - * New antimicrobial chemicals/compounds
 - * Other





- Novel strategies to manipulate gene expression to:
 - * Enhance innate citrus host plant resistance/tolerance to HLB
 - * Disrupt normal biological pathways involved in:
 - > Disease development
 - > Insect vector behavior





- Insect vector management
 - Improved ACP sampling
 - * Chemical insecticides
 - Biological control
 - > Parasitoids
 - > Other
 - * Volatiles
 - > Repellents or attractants
 - * Cultural practices
 - * Area-wide psyllid management





OMICS

• Sequencing, assembly & annotation of the genomes of HLB-associated "Candidatus Liberibacter species" (i.e., Las, Lam)

Genomics





Omics

- HLB-associated "Ca. Liberibacter species"
- Host responses to infection & psyllid
- Microbial community associated with HLB
 - * in planta
 - * in insecta
- Gene expression
 - Citrus host
 - Insect vector (e.g., ACP)
 - * HLB-candidate pathogens

Transciptomics
Proteomics
Metabolomics
Metagenomics
Bioinformatics