Managing Fire Blight by Choosing Decreased Host Susceptibility Levels and Rootstock Traits





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Fire blight bacterial infection of apple cells



Khan et al. 2013

Host resistance and fire blight management in apple orchards

- ✓ Host resistance is considered most sustainable option for disease management due to
 - Easy to deploy/implement in the orchards
 - Low input and cost-effective
 - Environment friendly
- ✓ No choice to the growers--most of the new and old cultivars are highly susceptible
- ✓ Apple breeding to develop resistant cultivars

Domestication history of the cultivated apple

45-50 Malus species-----Malus sieversii—Gene flow





Malus baccata Diameter: 1 cm



Malus sieversii Diameter: up to 8 cm







Malus sylvestris Diameter: 1-3 cm



Known sources of major/moderate resistance to fire blight to breed resistant cultivars

Source	Resistance level
Malus Robusta 5	80%
Malus Fusca	66%
Malus Arnoldiana, Evereste, Malus floribunda 821	35-55%
Fiesta, Enterprise	34-46%

- Fruit quality is the main driver for success of an apple cultivar
- Due to long juvenility of apples, it can take 20-25 years to breed resistance from wild crab apples



Genetic disease resistance in world's largest collection of apples



Evaluation of fire blight resistance of accessions from US national apple collection





- Grafted 5 replications: acquired bud-wood and rootstocks
- Inoculated with Ea273
 Erwinia amylovora strain
- Measurement of fire blight lesion length



Accessions for fire blight resistance and decreased susceptibility research and breeding



Different levels of susceptibility/resistance responses of *Malus sieversii* and *Malus x domestica* accessions

Is There a Relationship Between Root Traits and Susceptibility to Fire Blight?



Variation in root mass of a rootstock (M.7) obtained from a commercial nursery











Fire blight evaluation of scion populations and rootstock trait measurement

- Roots of M.7 rootstocks were altered to create four different root area classes (RACs) and imaged to quantify rootarea (cm²)
- Afterwards, these non-grafted rootstocks were grown in 5 gallon pots in the greenhouse
- Shoots were inoculated with Ea273 fire blight strain
- Fire blight evaluation as Percent Lesion
 Length=length of necrosis/shoot length
- At the end of experiment, fresh and dry root mass and number of primary roots





Variation in root traits of 4 different root area classes (RACs) of 'M.7' rootstocks









Fire blight susceptibility of four different root area classes (RACs) of 'M.7' rootstocks



Patterns of root area (cm²) and disease severity (%) in four different root area classes (RACs) of 'M.7' rootstocks. (A) Different root area classes (RACs) observed in M.7 rootstock, and (B) Disease severity represented as percent lesion length (%) for four M.7 RACs at 8 dai.

Root area negatively correlates with fire blight susceptibility

schematic representation of Α differential gene expression and pathway analysis for rootdependent fire blight infection in apple. (A) Numbers of differentially expressed genes (DEGs) obtained from analysis of control samples between RAC-1 and RAC-4, and between control and fire blight samples within RAC-1 and RAC-4. (B) Venn diagram numbers of unique and with shared DEGs from two different expression analyses. (C) <u>Pathways</u> showing overrepresentation from <u>common DEGs</u> that most likely correspond to effects of low root area and fire blight infection.



Take home message

 ✓ Wild *malus* species hold great genetic diversity for fire blight resistance traits but its deployment is a major bottleneck

✓ Resistance and decreased susceptibility from *Malus* sieversii and domesticated apples should be priority target for breeding

✓ Root surface area has negative correlations with fire blight susceptibility of rootstocks by impacting carbohydrate metabolism and defense pathways

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Thank you for attention!



@FruitDiseases

Research and extension to characterize mechanisms of disease resistance and pathogenesis, and development of tools for improvement of rosaceous fruits

◎ Geneva, NY *S* blogs.cornell.edu/khanlab/ III Joined December 2016

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Welcome to the Khan Lab!

HOME

We are interested in mechanisms of disease resistance and pathogenesis in rosaceous fruits, and to develop strategies to speed up variety development for improved disease resistance. Some of the tools that we use are quantitative genetics, QTL and association mapping, genome editing, bioinformatics, marker-assisted selection, and transgenics. We are also interested in developing high-throughput methods for plant resistance phenotyping

Our research focus, the rose family (Rosaceae), is culturally and commercially valuable, as many of the most popular fruits belong to it. For example, apples, pears, peaches, plums, almonds, apricots, strawberries, cherries, loguats, and raspberries belong to the rosaceae family. A number of fungal and bacterial diseases cause huge economic losses to the fruit industry globally and are a threat to sustainable production.

Currently our primary research focus is to understand and improve resistance to fire blight and apple scab-

 Understanding Underlying Mechanisms of Disease Resistance - Characterization of Mechanism of Disease Resistance

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