

# Rapid Diagnostic Test for Pitch Canker

Tyler J. Dreaden, Graduate Student, Forest  
Pathology, University of Florida

# Presentation outline



- Background on pitch canker
- The problem
- Seed screening project
  - ▣ ISTA culture based method
  - ▣ Intro to PCR
  - ▣ New PCR based seed screening method
  - ▣ Current status
  - ▣ Potential applications

# Background Information

- Pitch canker is a serious disease of pines
- **Causal agent:** *Fusarium circinatum*
  - (previously known as *Fusarium subglutinans* f. sp. *pin*i or *Fusarium moniliforme* var. *subglutinans*)
- **Host range:**
  - Conifers, mostly *Pinus* spp.
- **Symptoms:**
  - death of shoots, occasionally trees
  - crooked stems and forks
  - pitch bleeding from shoots, resin-soaked wood
  - Can infect at any stage in pine development
  - Pre- and post-emergence “damping off”
  - Infected seedlings are often planted
- **Signs:**
  - rarely salmon colored sporodochia
  - microconidia (oval) and macroconidia (canoe-shaped)



Sporodochia

# Pitch canker disease cycle

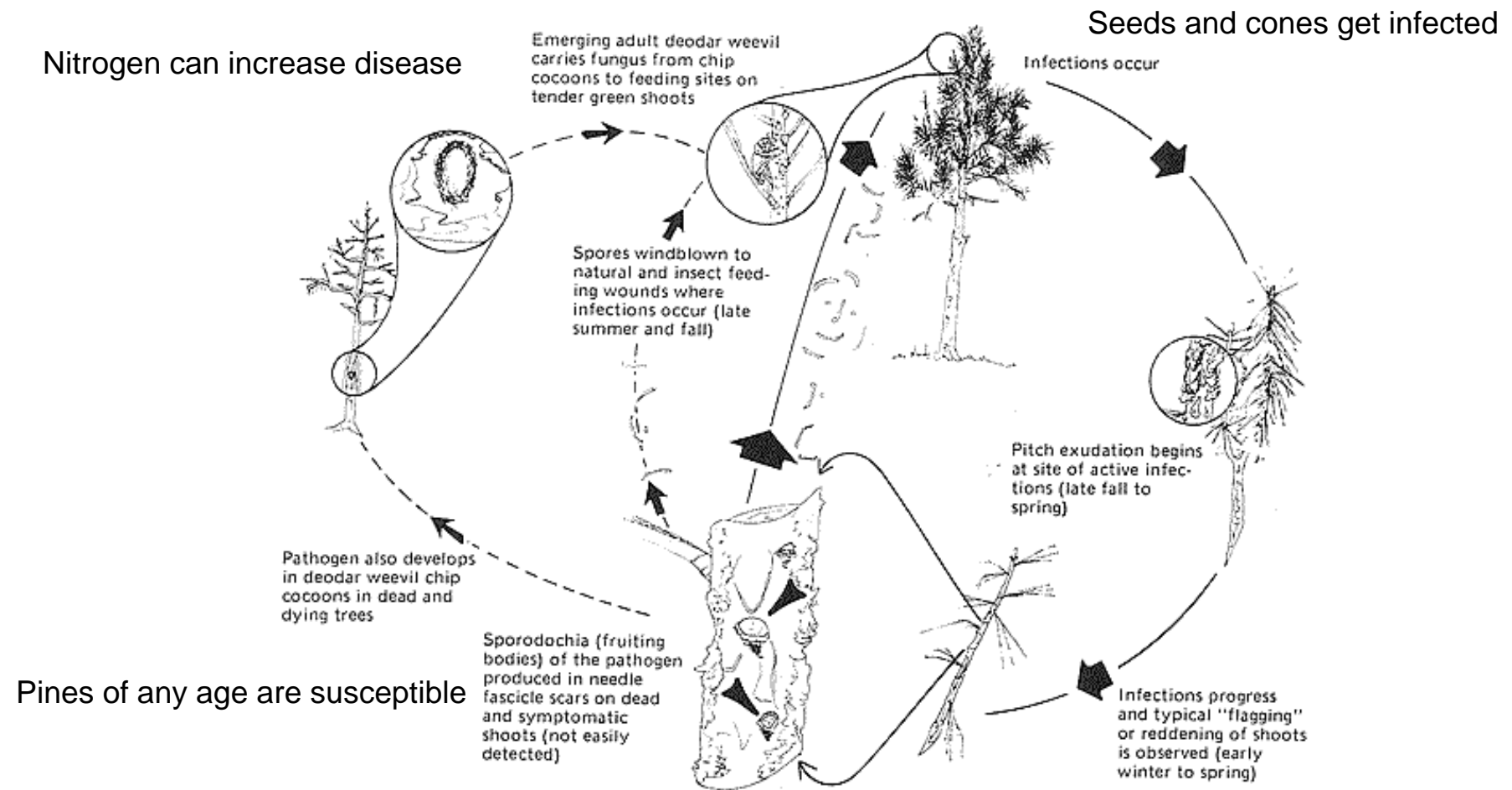


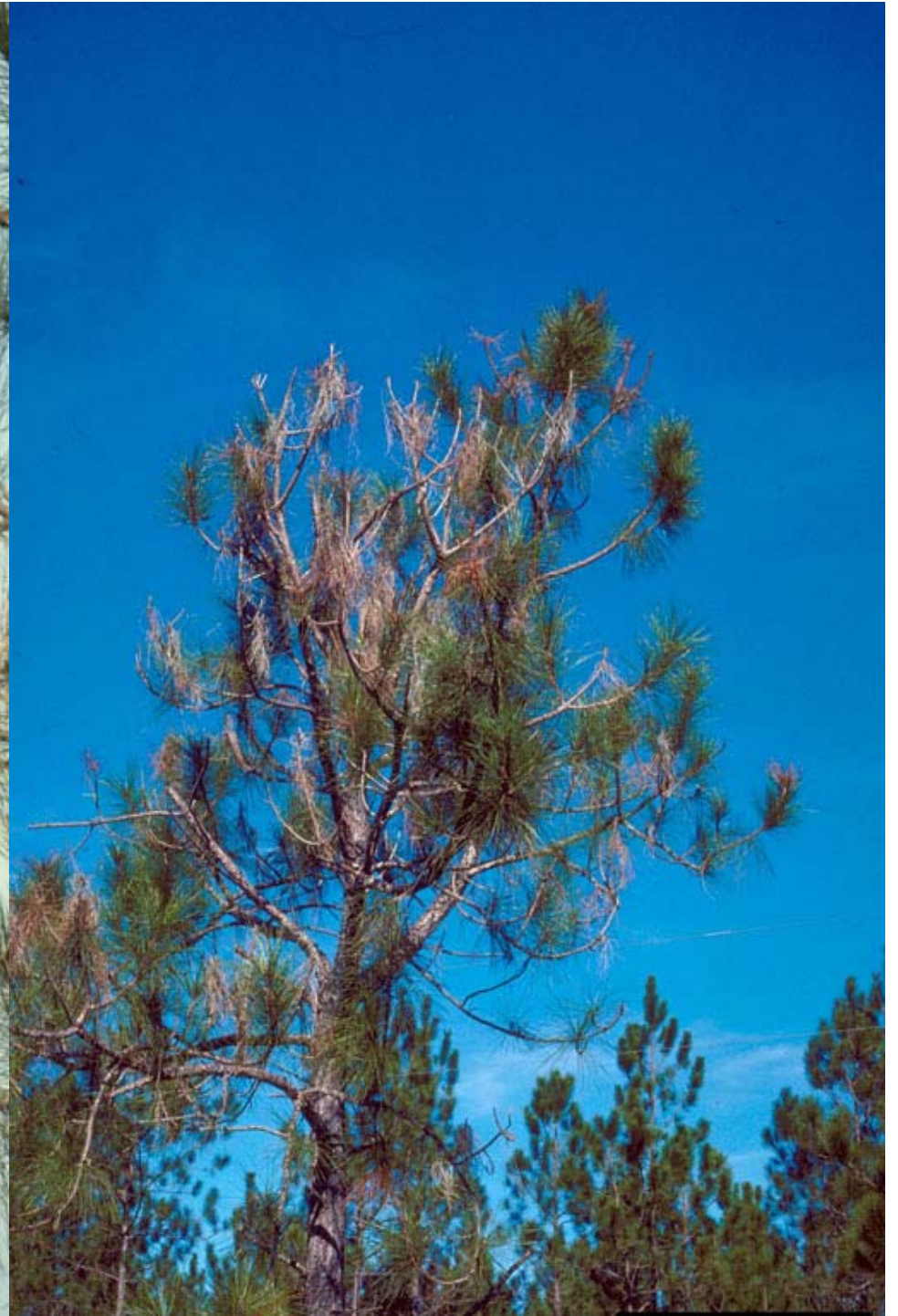
Fig. 13. Simplified life cycle of the pitch canker fungus on slash pines in Florida.

[http://www.fl-dof.com/publications/Insects\\_and\\_Diseases/td\\_cs\\_pitch\\_canker\\_fig13.html](http://www.fl-dof.com/publications/Insects_and_Diseases/td_cs_pitch_canker_fig13.html)













UGA1501012



Canker margin





Cotyledon infection from  
contaminated seed





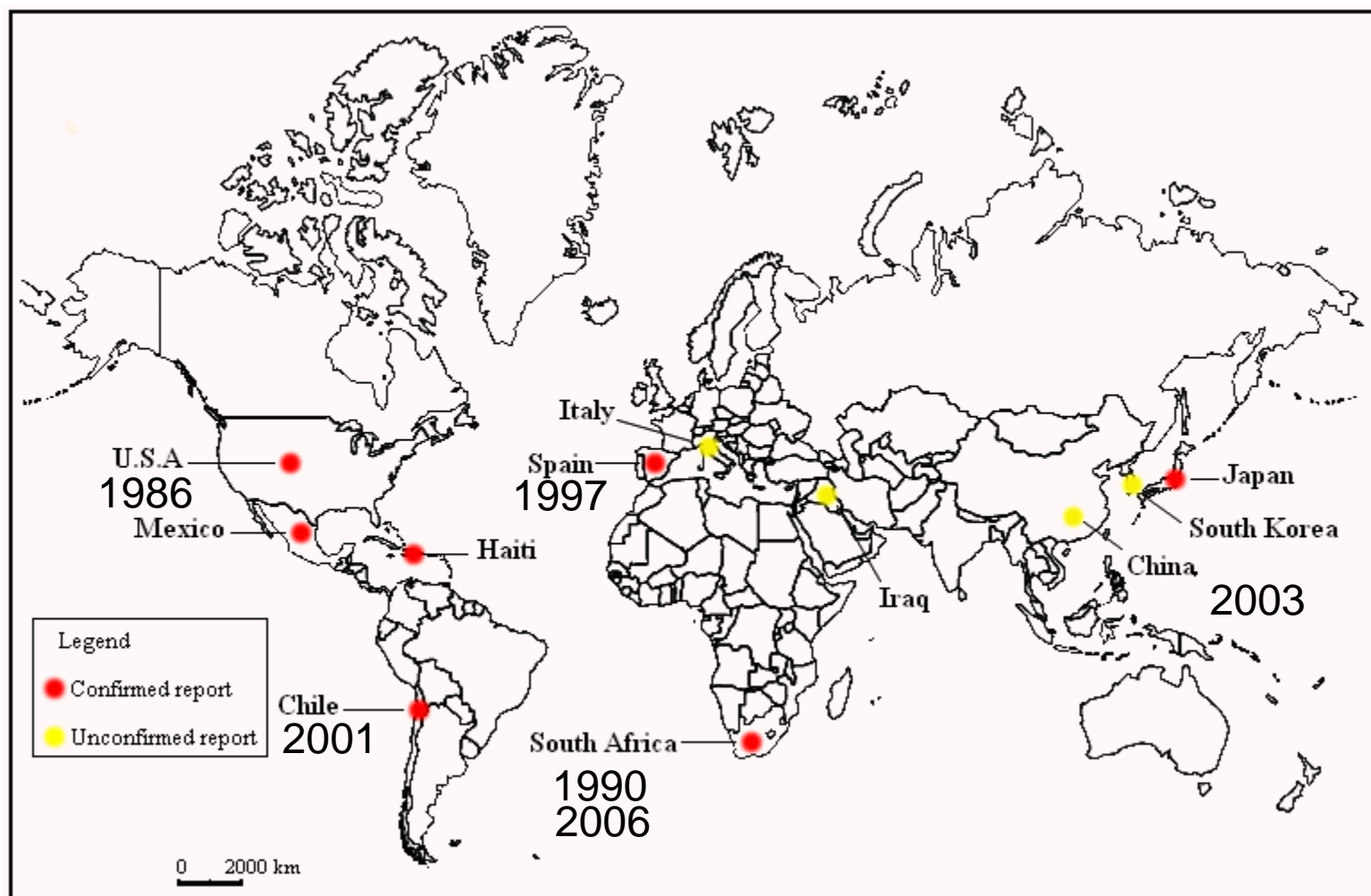


Figure 1. Worldwide distribution of pitch canker (Ganley, 2006).

# Why is regulation needed



- Massive plantations exist of susceptible pine species such as *P. radiata* and *P. taeda* in many countries
- Pitch canker does not currently exist in many areas
  - ▣ Need to keep new strains out
- Seeds are being shipped globally
- Pitch canker can be transmitted
  - ▣ Contamination of the seed coat
  - ▣ Contamination of the embryo



# The problem.....



- ❑ Currently, seeds are tested and certified using the International Seed Testing Association Guidelines
- ❑ ISTA Guidelines require testing 400 seeds per seedlot
- ❑ Seeds are tested by a “blotter paper method” that involves growing *F. circinatum* out of crushed seeds onto blotter paper
- ❑ This process is very slow, laborious, and prone to false negatives

# Objectives of study

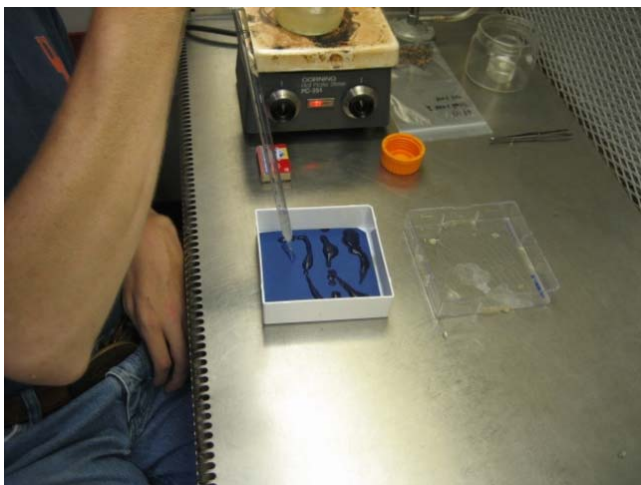
- Develop qPCR seed screening method
- Compare qPCR method to current culture-based method by screening 6 slash pine seed lots using both methods
  - Compare detection results
  - Compare time and cost associated with both methods



# ISTA culture based screening method

- Based on

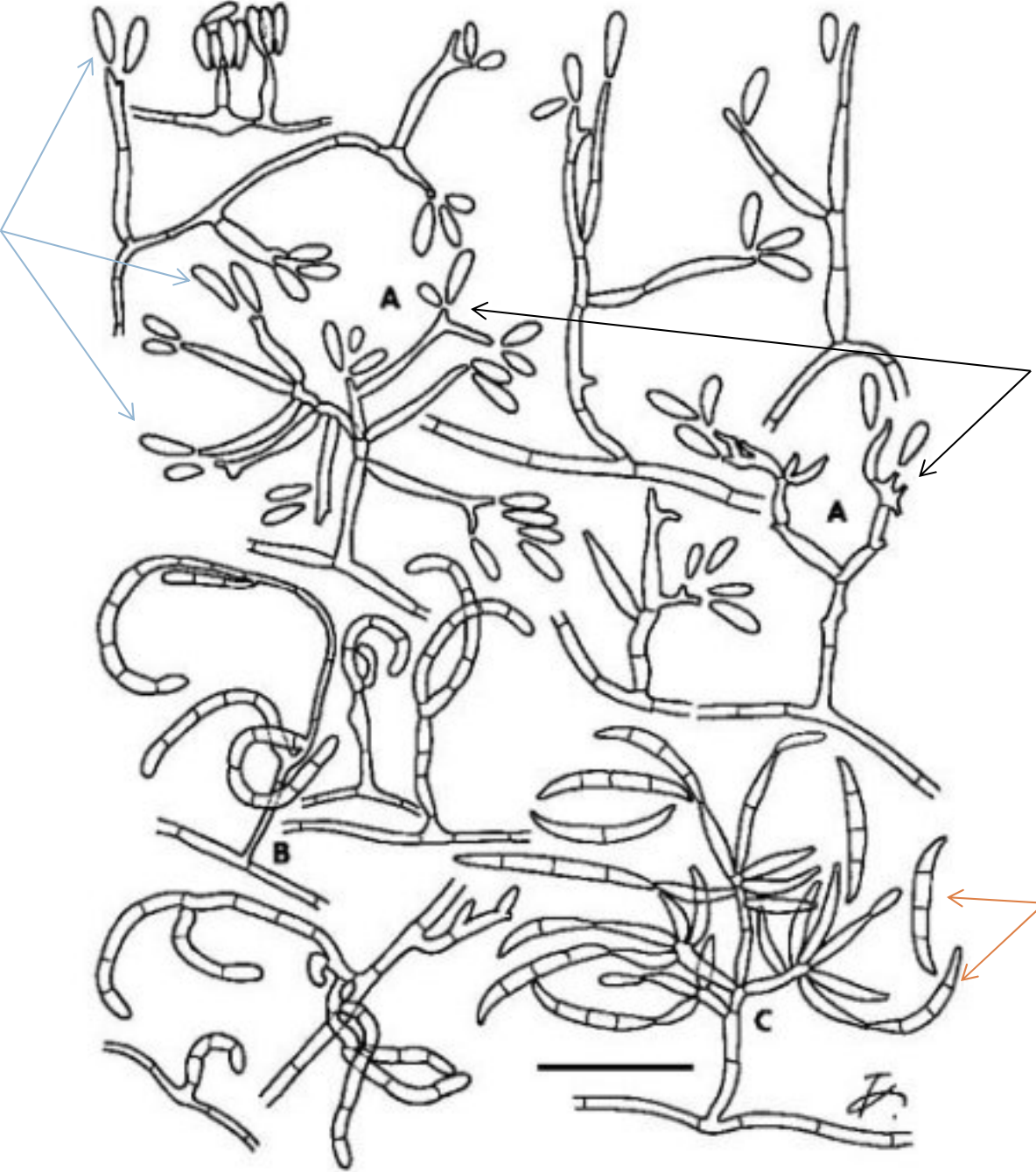
- ▣ Anderson, R.L. 1986. New method for assessing contamination of slash and loblolly pine seeds by *Fusarium moniliforme* var. *subglutinans*. Plant Disease 70: 452-453.
- Relies on placing seeds on selective media and identifying fungi that grow based on morphology under a microscope
- Also called blotter paper method



Microconidia

Polyphialides

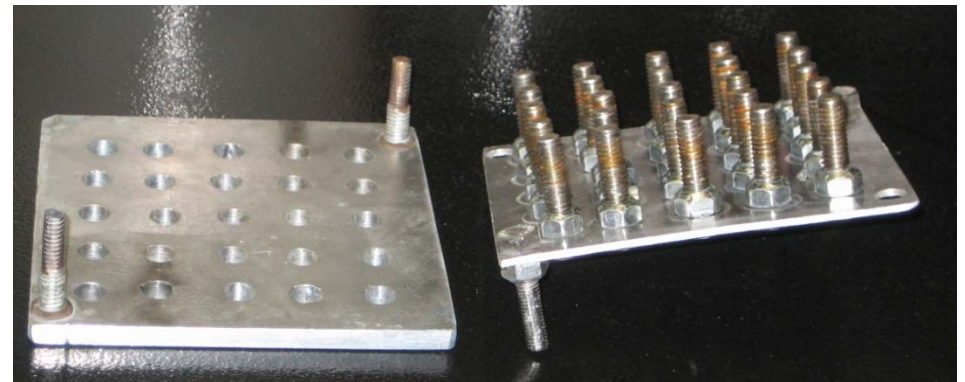
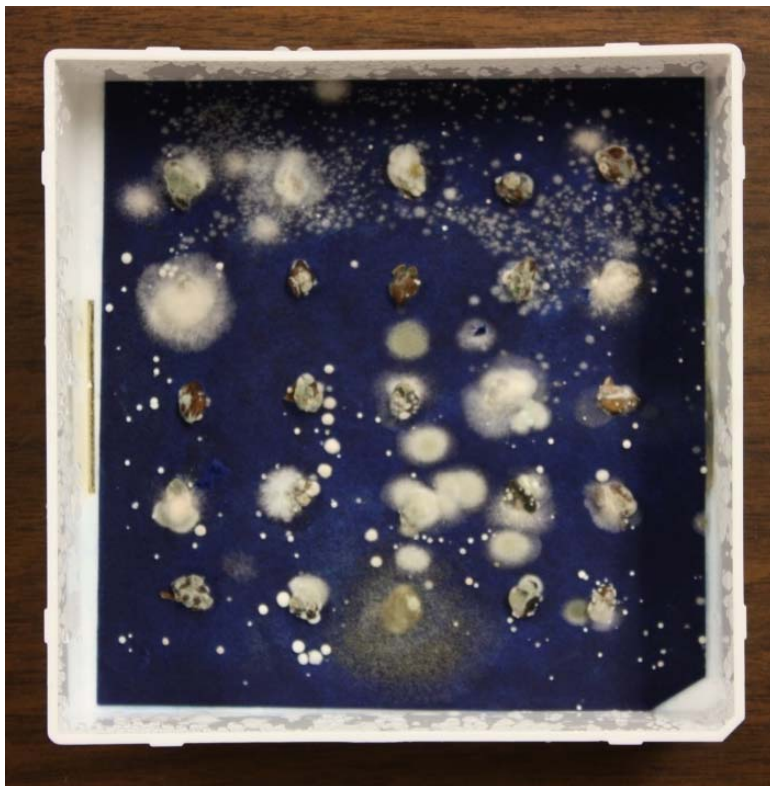
Macroconidia





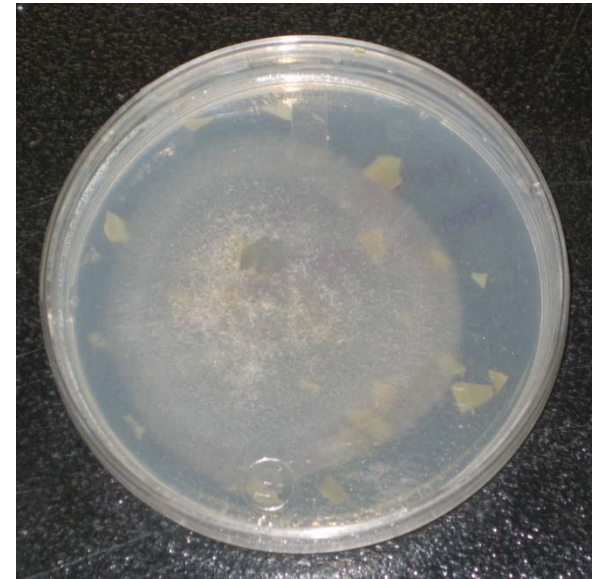
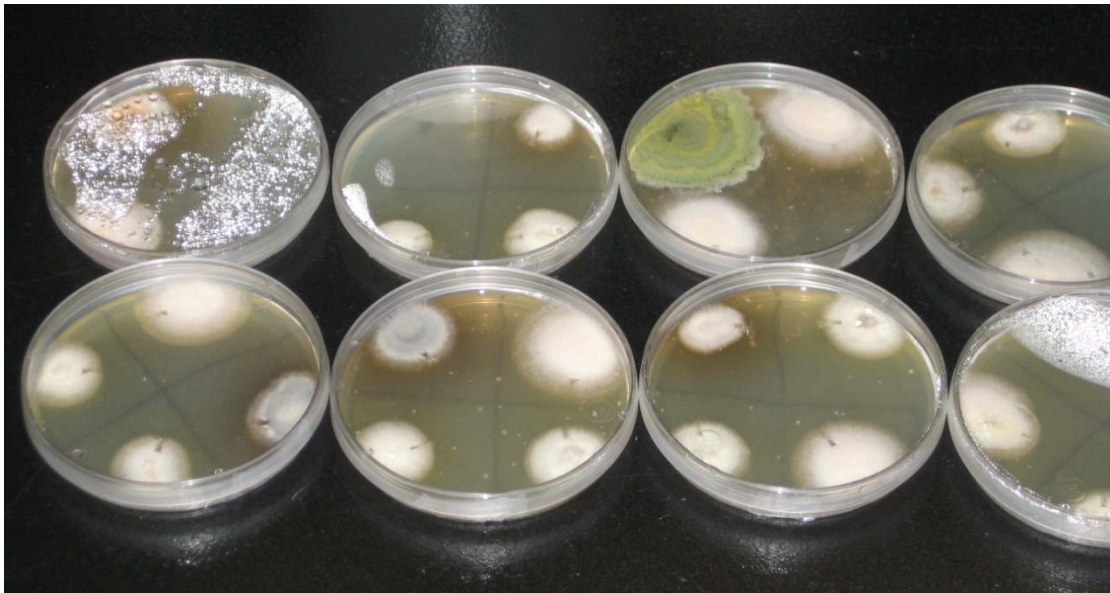
# Culture based screening method

- Crush seeds on blotter paper with selective media



# Culture based screening method

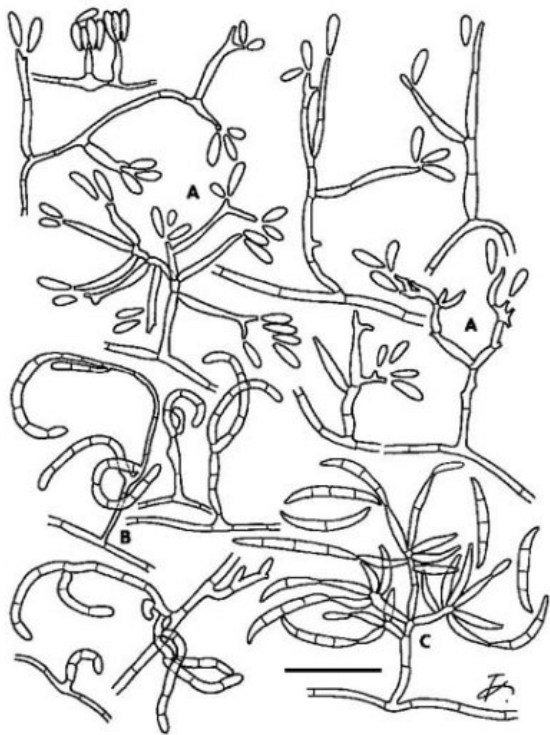
- Subculture on carnation water agar
  - ▣ Contamination





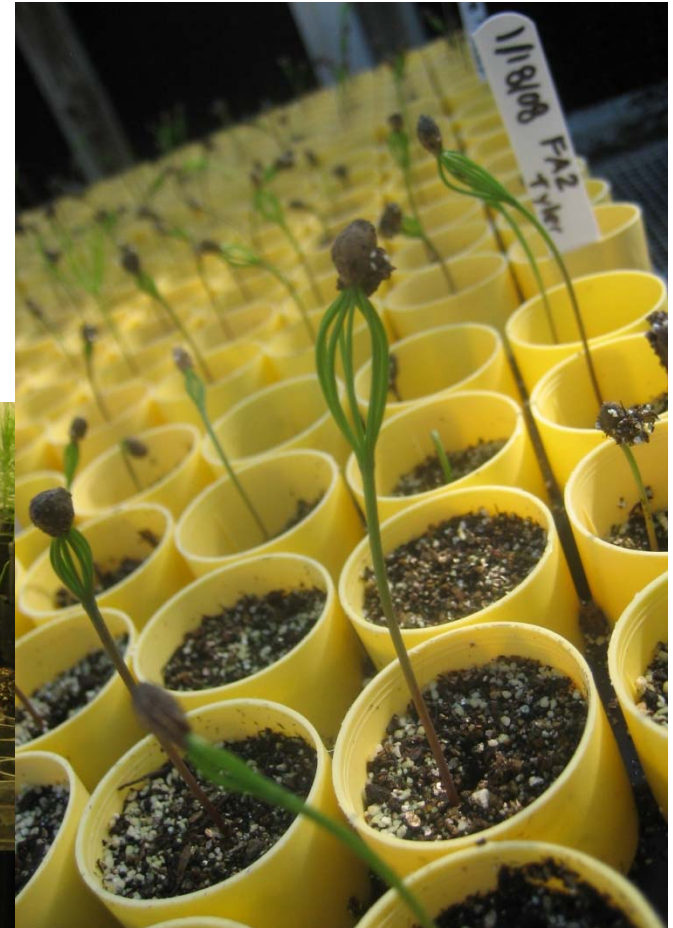
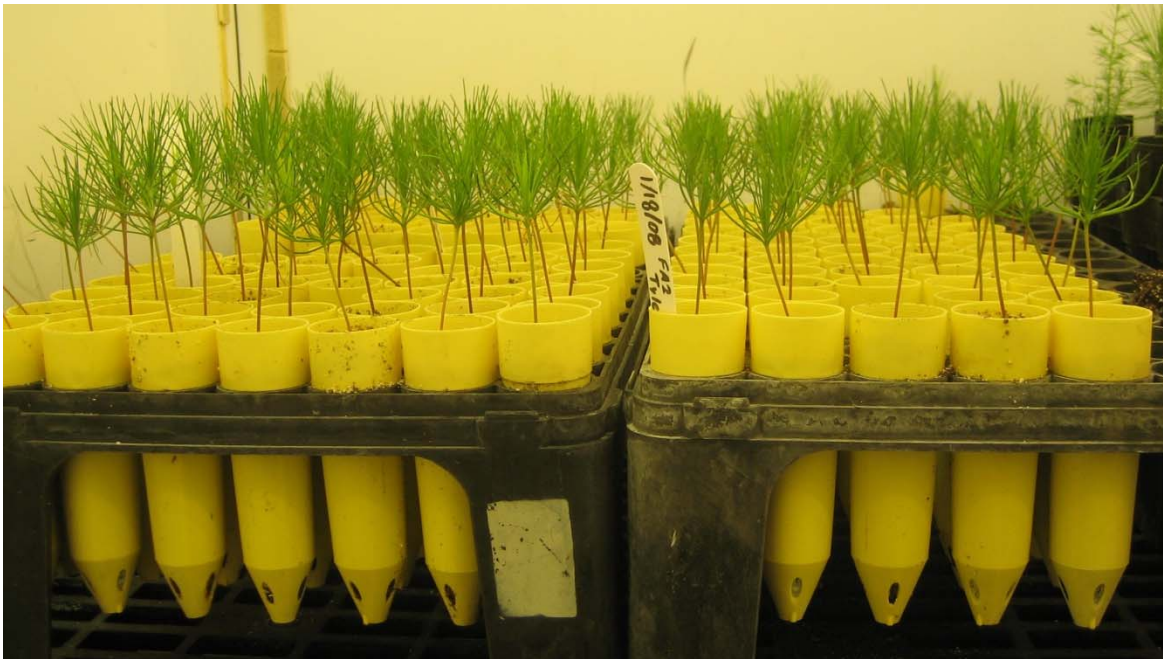
# Culture based screening method

- Identified morphologically under a microscope
  - ▣ 3-7 hours



# Culture based screening method

- Inoculate seedlings to prove pathogenicity





# What is PCR?

- ❑ Polymerase Chain Reaction (PCR)
- ❑ Method of replicating DNA without living organisms
- ❑ Amplifies a specific region of DNA (can be used to identify species)
- ❑ Driven by change in heat
- ❑ Prior knowledge of organism's DNA is needed
- ❑ Can be used to detect organism presence in samples

# PCR components

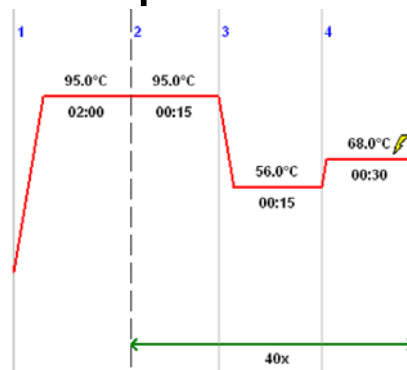


- ❑ Template DNA- DNA from sample to be replicated
- ❑ Primers- single stranded DNA that specify the end points of the PCR product (provide specificity)
- ❑ *Taq* polymerase- adds single nucleotides to the ends of the primers and makes a new DNA strand
- ❑ Free nucleotides- A, T, G, C building blocks of DNA required to make replicate strands



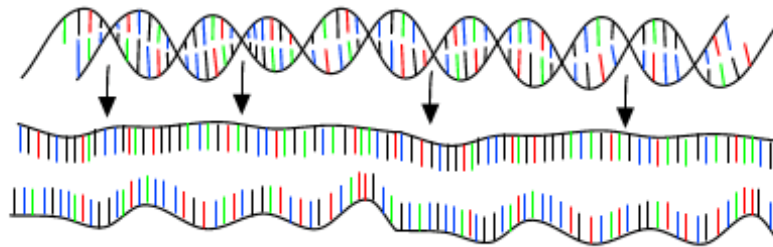
# PCR steps

- Denaturation- separation of double-stranded DNA to single strands
- Primer annealing- primers bind to a specific location on the single stranded DNA
- Extension- *Taq* polymerase, an enzyme, adds single nucleotides to the ends of the primers
- These steps are repeated 30 to 40 times. After each cycle the number of copies of the DNA doubles.



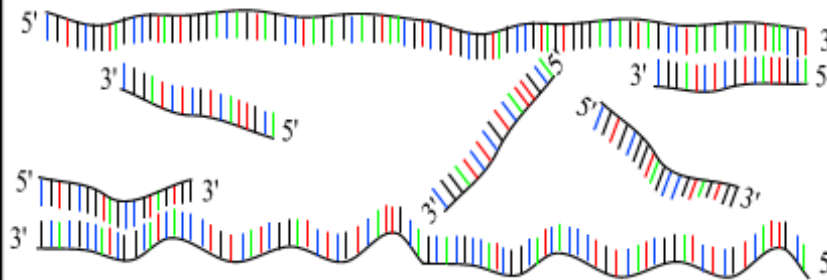
# PCR : Polymerase Chain Reaction

30 - 40 cycles of 3 steps :



**Step 1 : denaturation**

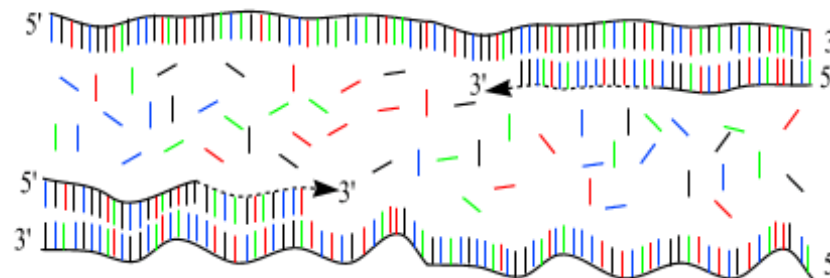
1 minut 94 °C



**Step 2 : annealing**

45 seconds 54 °C

forward and reverse  
primers !!!



**Step 3 : extension**

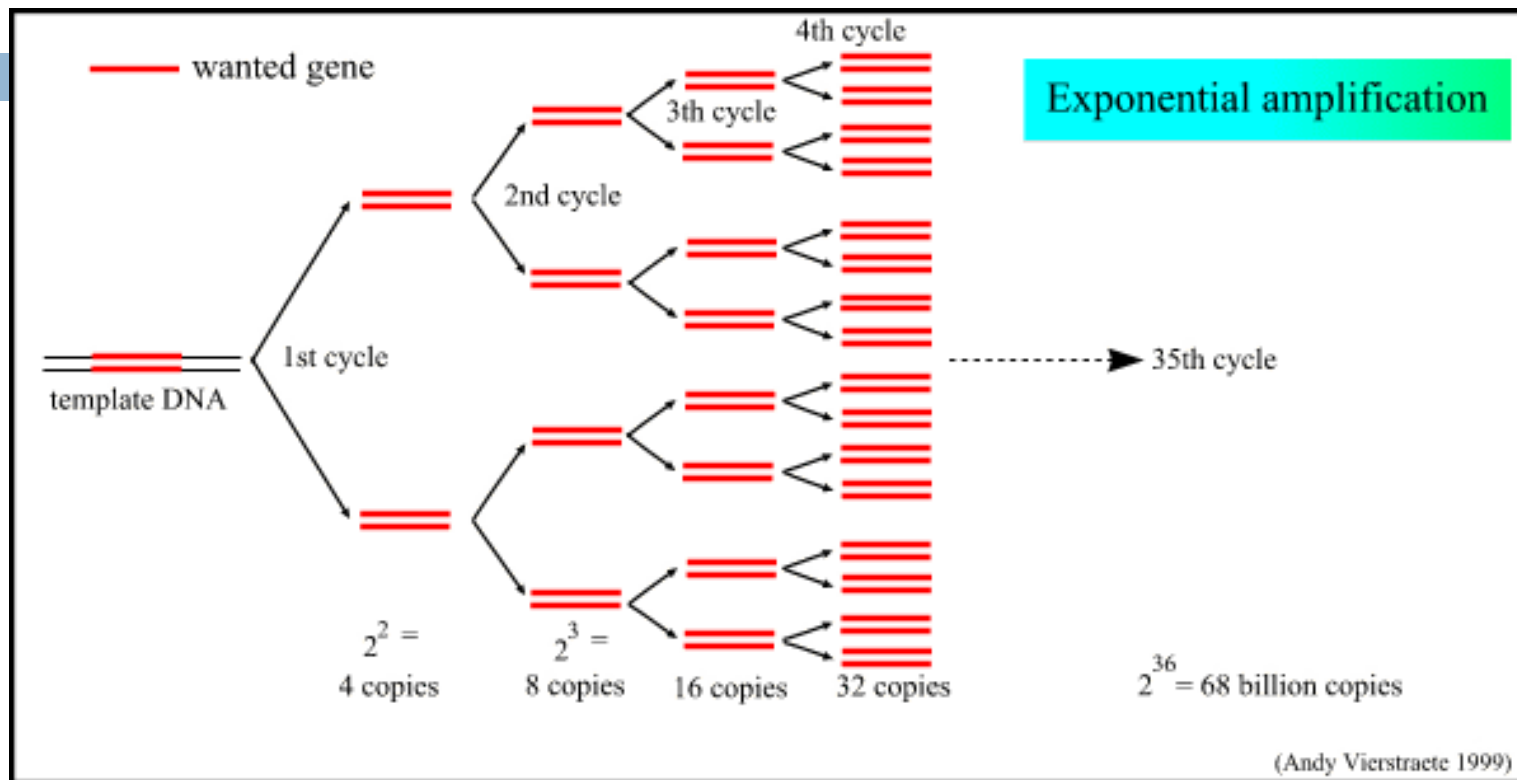
2 minutes 72 °C

only dNTP's

(Andy Vierstraete 1999)

<http://users.ugent.be/~avierstr/principles/pcr.html>





<http://users.ugent.be/~avierstr/principles/pcr.html>

# Real time PCR, qPCR

- Quantifies the initial amount of the DNA template
  - ▣ highly specific, sensitive, and reproducible
- Similar to regular PCR but the amplification is read by a computer in real-time
- Faster and more sensitive than traditional PCR
- Commonly used to test for presence of pathogens
  - ▣ Examples: Sudden oak death, soybean rust, bird flu

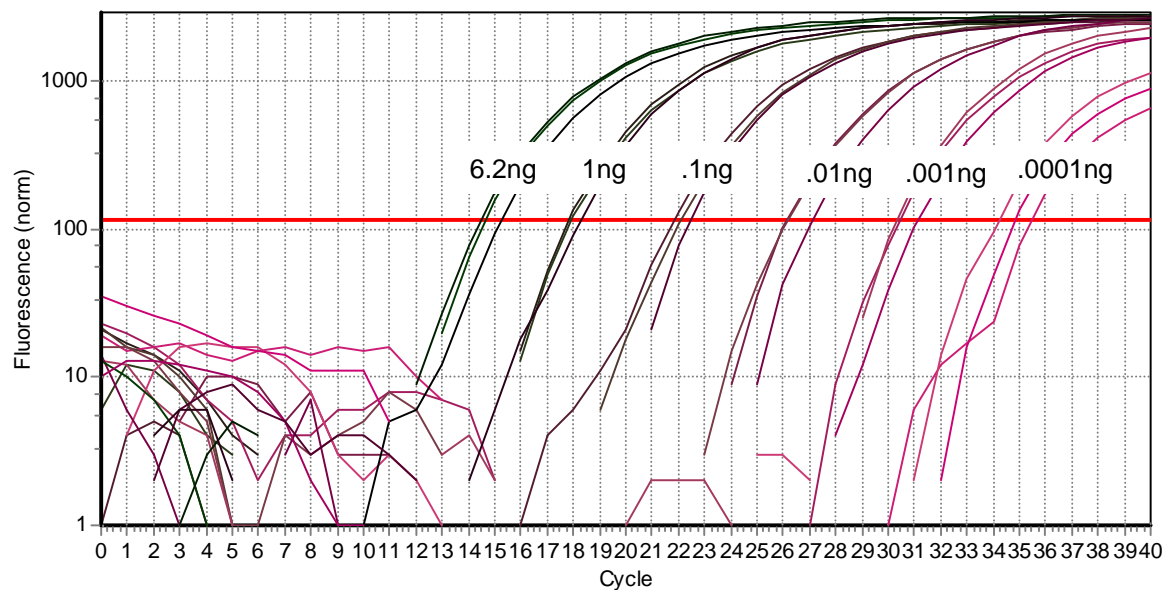
# qPCR



- Relies on the change in fluorescence due to amplification of DNA
  - ▣ Generally as amplification increases, fluorescence increases
  - ▣ Fluorescence is measured after each cycle
  - ▣ The more amplified DNA in the PCR reaction the less number of cycles needed for the fluorescence to significantly increase over the background.

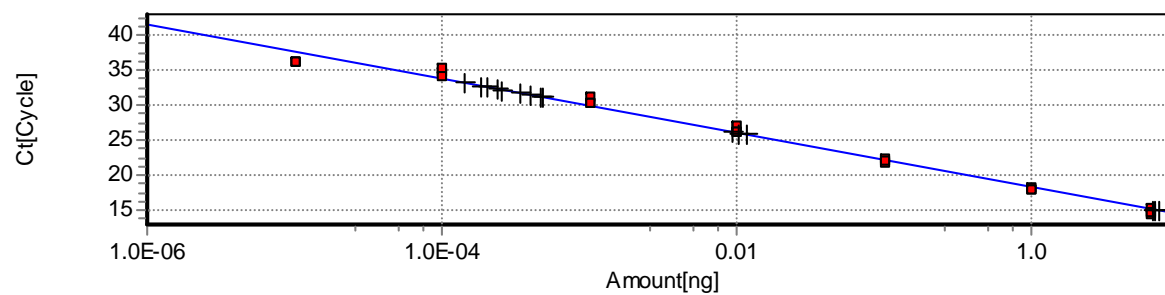


# qPCR



Cycle threshold (Ct)

Threshold: 117 (Noiseband)  
Baseline settings: automatic, Drift correction OFF



Slope: -3.883  
Y-Intercept: 18.32  
Efficiency: 0.81  
R<sup>2</sup>: 0.988

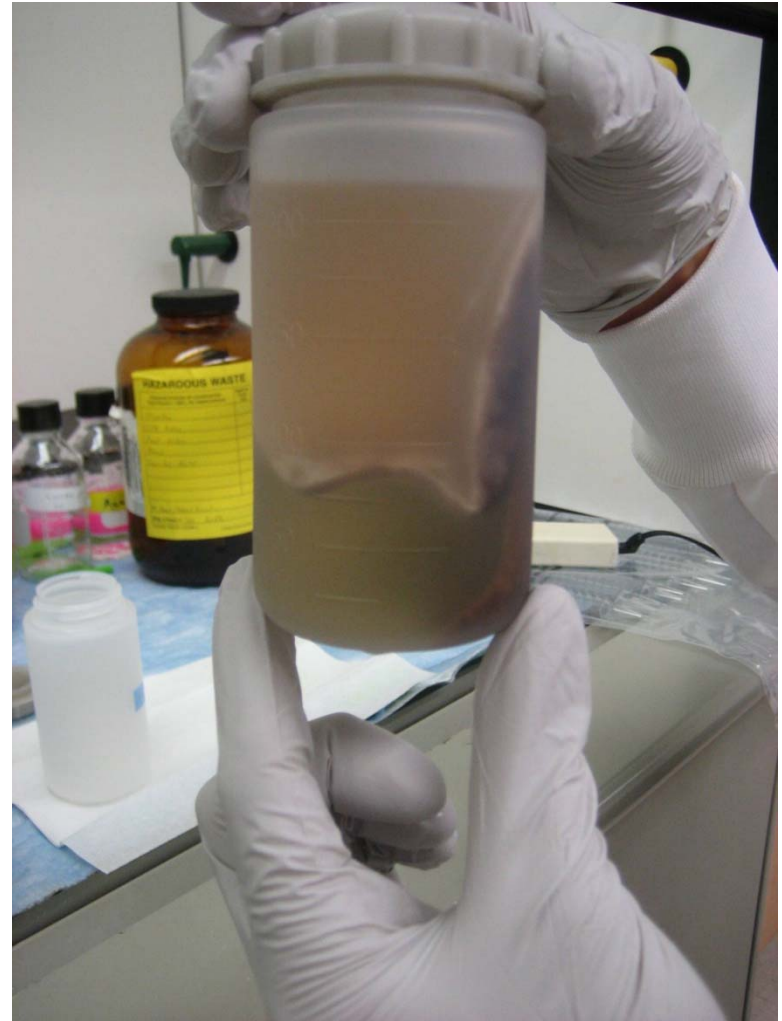
# Main point



- qPCR provides a potential method to obtain accurate, quantitative detection of *F. circinatum* from different substrates.

# qPCR screening procedures

- Extract DNA from 400 slash pine seeds
  - ▣ 6 hours for 6 DNA extractions





# qPCR method development



## □ Primers based on

- Schweigkoffer, W., O'Donnell, K. and Garbelotto, M. 2004. Detection and Quantification of *Fusarium circinatum*, the Causal Agent of Pine Pitch Canker, from Two California Sites by Using Real-Time PCR Approach Combined with a Simple Spore trapping Method. Applied and Environmental Microbiology. Vol. 70 No. 6 3512-3520.
- Produce a 360 bp product only in the presence of *F. circinatum* DNA
- 5 hours to screen 6 seed lots

# Time requirements

- ❑ qPCR can screen seeds in 2 days
- ❑ Culture-based method requires 2-3 weeks



qPCR machine

# Current status



- Blotter Method
  - ▣ ID fungi from two seed lots
  - ▣ Inoculate seedlings with *F. circinatum* obtained from seeds
- qPCR method
  - ▣ Standardize DNA extractions
  - ▣ Run qPCR reactions on 6 seed lots
- All research should be completed by the end of summer



# Potential applications



- Screen seeds before shipping internationally
- Screen seeds for potential damping off problems caused by *F. circinatum*
- Rapidly detect *F. circinatum* in diseased seedlings
- Compare cone/seed infections/contamination from different sources
- Use in epidemiology studies of *F. circinatum*

# Thanks



- Committee Members Dr. Ed Barnard, Dr. George Blakeslee, and Dr. Jason Smith
- Wayne Bell (International Forest Seed Company)
- Dr. Claire Anderson for technical assistance

# Questions

