

Determination of vegetative  
compatibility groups using molecular  
markers, and their aggressiveness of  
*Verticillium dahliae* occurring on  
sunflower

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# OUTLINE

- Introduction
- Symptoms of *V. dahliae*
- Losses
- Genetic diversity
- Vegetative compatibility groups
- Objectives
- Methods

# Major diseases of sunflower

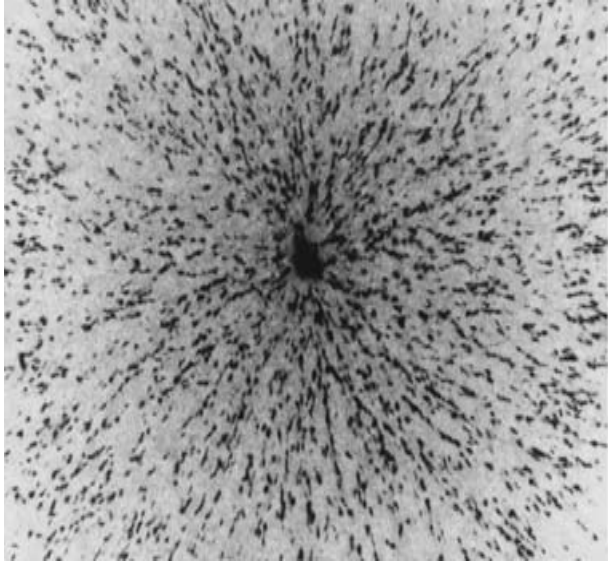
- Sclerotinia stalk and head rot
- Rust
- Verticillium wilt
- Downy mildew
- Phomopsis
- Phoma black stem and leaf spot



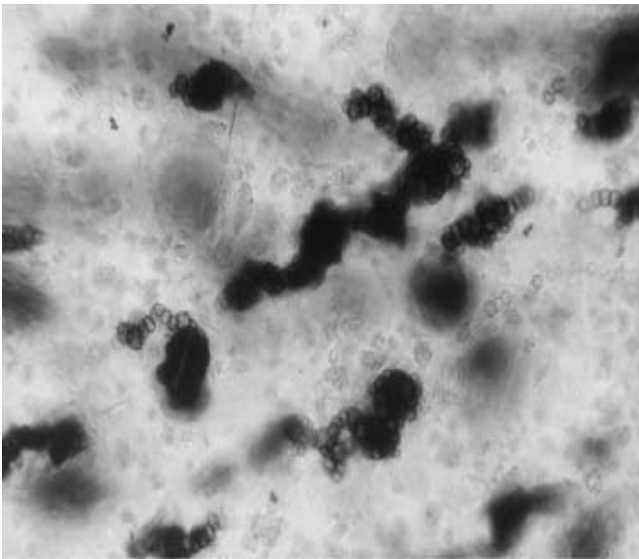
Slide obtained from Dr. Sam Markell, NDSU.



## Culture: microsclerotia



The microsclerotia are formed in a radial pattern



Globose to elongate *V. dahliae* microsclerotia



Large, globose to irregularly shaped *V. dahliae* microsclerotia with connected dark hyphae

# Losses & genetic resistance

- The disease has economic implications annually.
- Reduced head size (with severe and mild infection 42% and 18% respectively)
- Seed size, oil content, and yield (up to 66%).
- Managed by genetic resistance
  - \* V-1 gene ~ effective for 20 yrs
  - \* Vd strains not controlled by V-1 in 1985 (Argentina), 2002 (MN), 2004 (ND)

# Genetic diversity of *V. dahliae*

- No sexual stage that allows recombination of genes
- Vegetative compatibility grouping (VCG)
- Vd has 4 VCGs: VCG1, VCG2 (A,B), VCG3, VCG4 (A,B)
- On sunflower, VCGs of *V. dahliae* isolates have not been determined on wide scale for populations study (Continental & inter-continental)

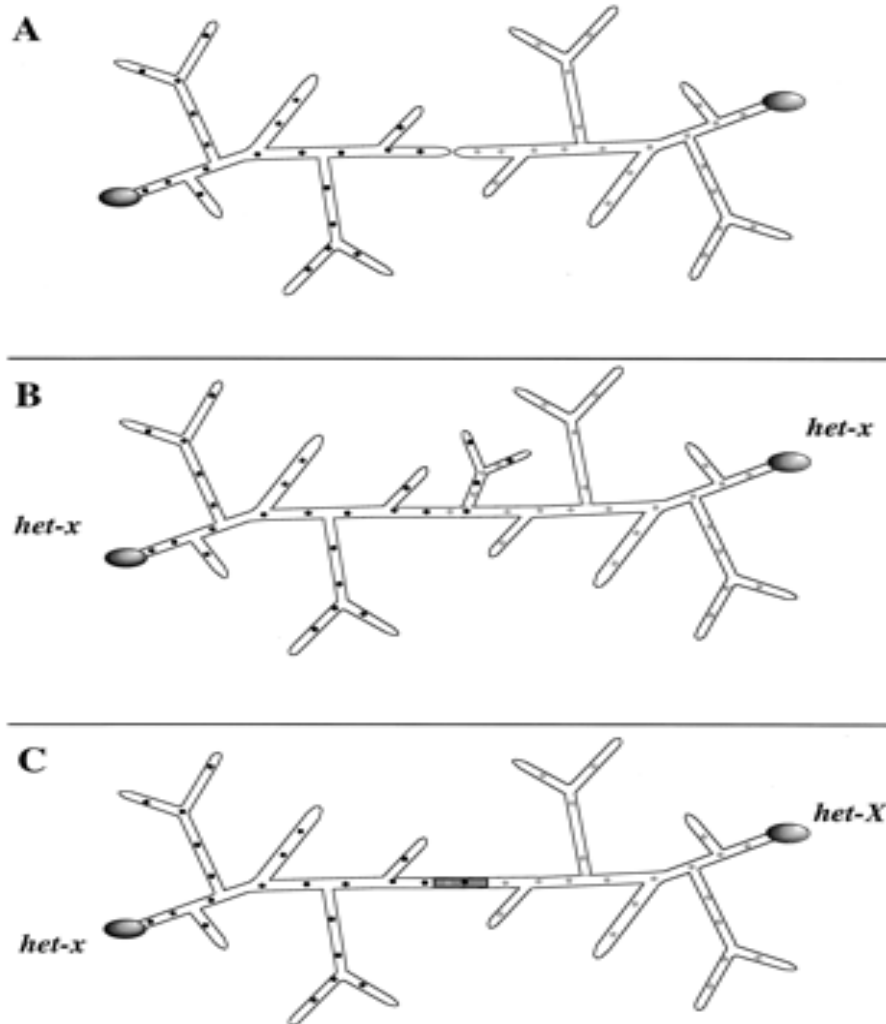
# Importance of studying VCGs

- VCGs vary in their aggressiveness, physiology, and geographic distribution.
- Breeding programs
- Management (crop rotation)
  - \*Example: potato Vs mint (Omer et al., 2008)



# Vegetative Compatibility Groups

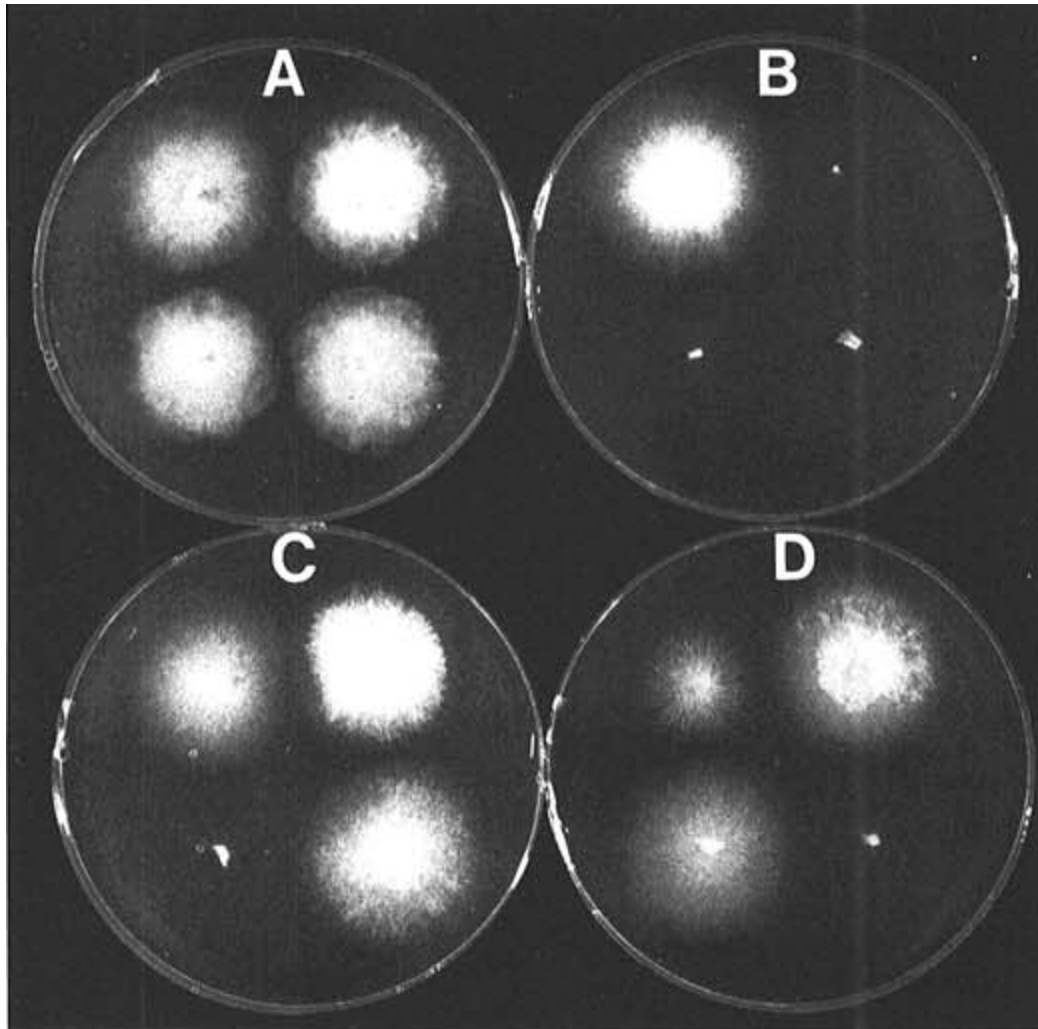
- What is vegetative compatibility ?



(A) When two different fungal individuals meet, they spontaneously undergo a cell fusion event or anastomosis.

(B) If the two individuals have the same *het* genotype, a heterokaryon is established.

(C) If the two strains differ in *het* genotype, the heterokaryotic cells are destroyed or severely inhibited in their growth



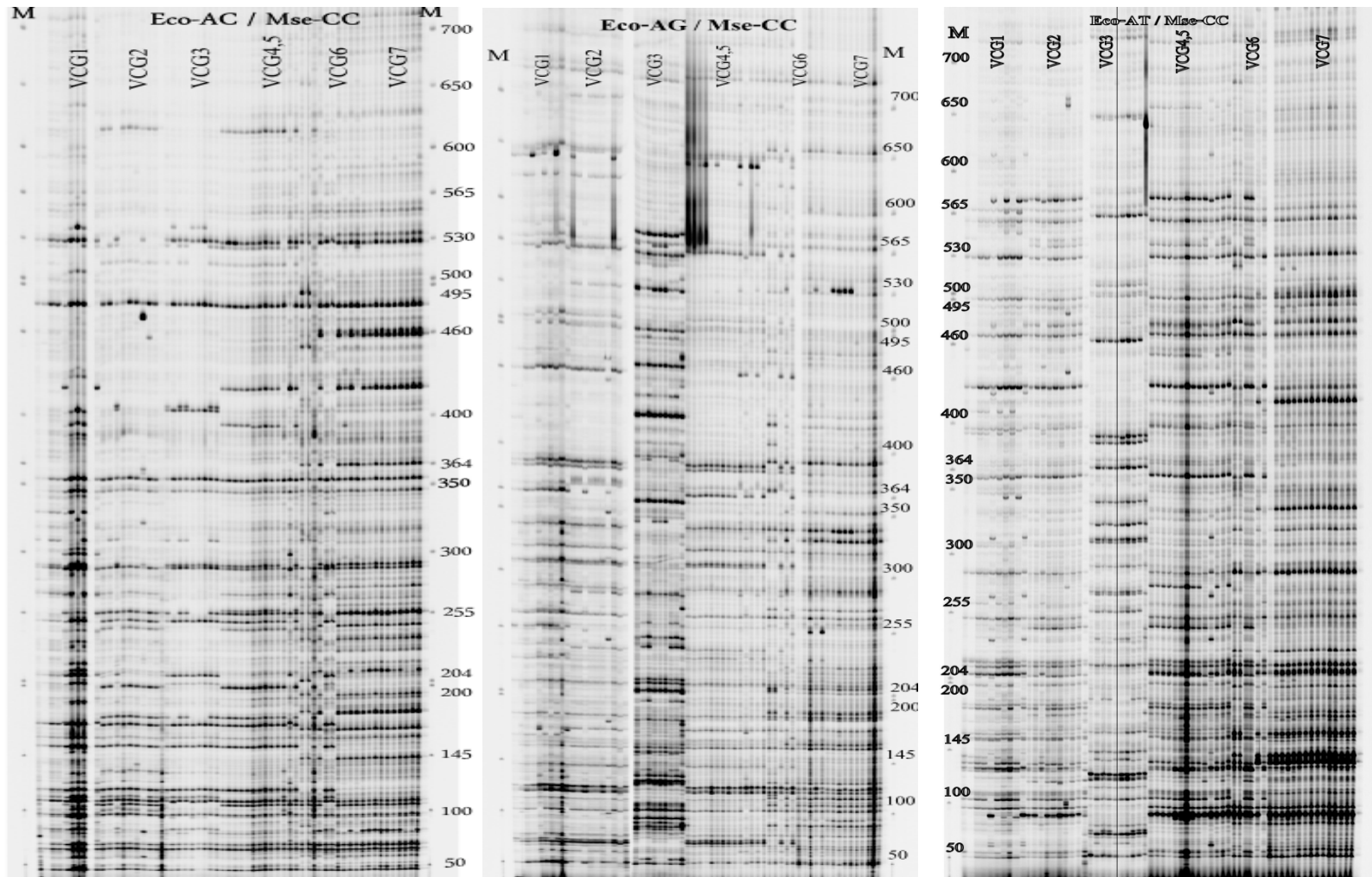
- Stable
- (*nit*) mutants
- Simple
- Many fungal plant pathogens

### Limitations

- laborious and time consuming
- mutant are not always possible
- *Vic* or *het* alleles

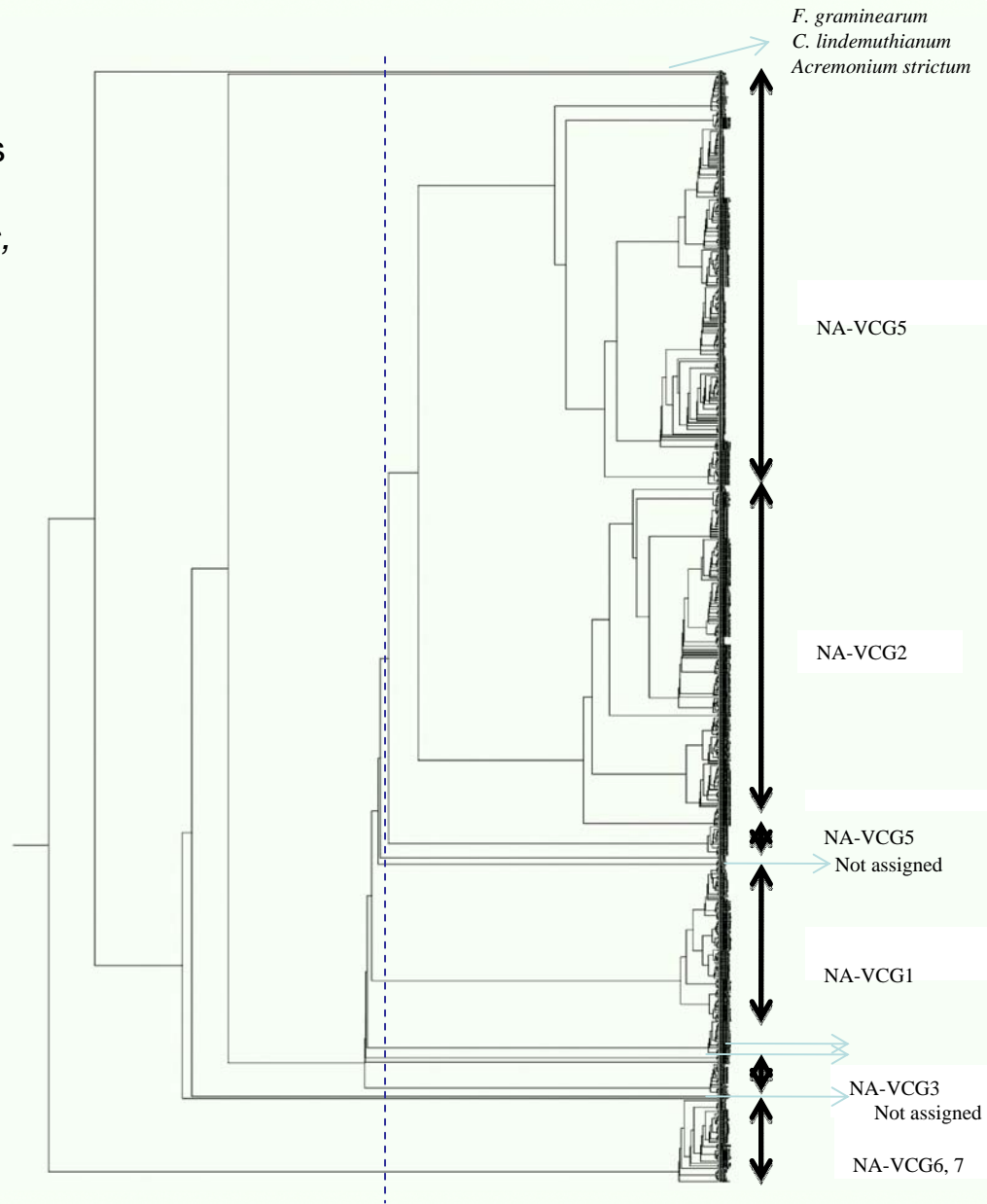
Nitrate nonutilizing (*nit*) mutants phenotypes of *Fusarium oxysporum* on media with four different nitrogen sources

# Amplified fragment length polymorphism (AFLP)



Amplified fragment length polymorphism electrophoresis of *C. coccodes* isolates for the seven NA-VCGs. The three primer sets, Eco-AC/Mse-CC, Eco-AG/Mse-CC, and Eco-AT/Mse-CC were used for selective polymerase chain (PCR) amplification reaction. PCR products were separated on 6% polyacrylamide/7 urea gel using LI-COR 4300 model. The specific AFLP bands for NA-VCGs can be detected. M: is a standardized, IRdye-700 labeled size marker of 50-bp.

Dendrogram generated by 1000 bootstrap reiterations using PHYLIP for AFLP data of 855 isolates of *C. coccodes*, and 3 outgroups including *C. lindemuthianum*, *F. graminearum*, and *A. strictum*.



# Project objectives

- Determine the VCG(s) of *V. dahliae* on Sunflower
  - \* AFLP method
  - \* Simple sequence repeats (SSR)
- Aggressiveness of VCGs on sunflower
- SCAR marker for *V. dahliae* to assess a pathogen in plants

# Isolates

- Collect *V. dahliae* isolates from US
- Dr. Gulya (USDA) and Dr. Rashid's isolates (MB)
- Other parts of the world (Argentina and Europe)
- Tester isolates from Dr. Randy Rowe (PS)

# Methods

- Isolates will be cultured, purified, and DNA extracted
- AFLP protocol, will follow Collado-Romero et al., 2008
- Fluorescent DNA sequencer (Li-COR)
- SSR markers (Atallah et al., 2009)
- Aggressiveness: root dip method, hypodermic needle inoculation

# Acknowledgements

- **National Sunflower Association**
- **Dr. Samuel Markell**



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