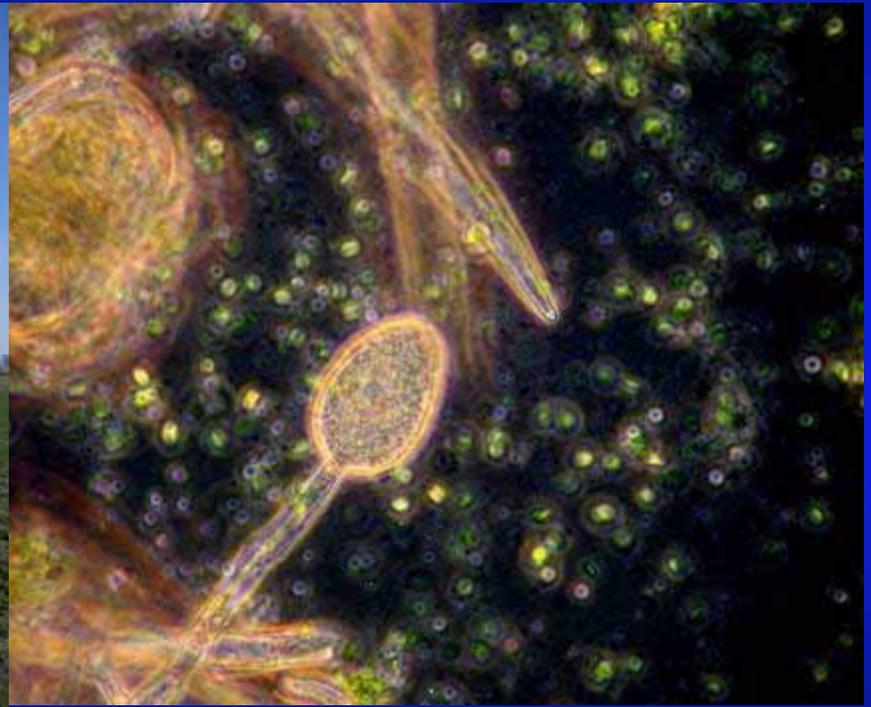


Plant Diseases of Local and National Concern

What's wrong with my plants?!



Most Common Diseases of Alfalfa In Utah

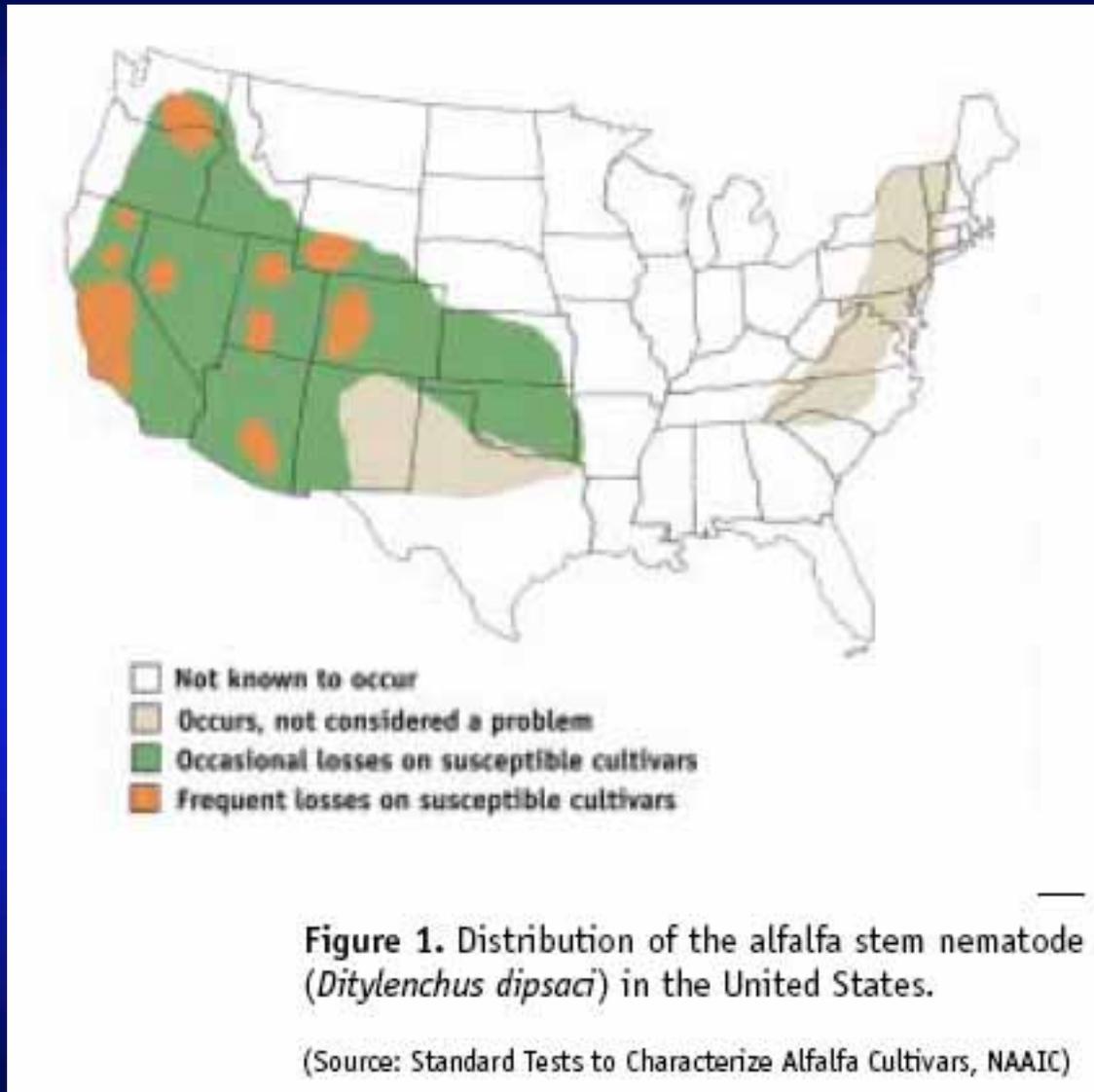
Spring Black Stem

Verticillium Wilt

Alfalfa Stem Nematode-*Ditylenchus dipsaci*



Distribution of alfalfa stem nematode in the U.S.



Alfalfa stem nematode symptomology

Stunted shoots
White flagging
Diminished stand



Chlorotic white flagged shoots



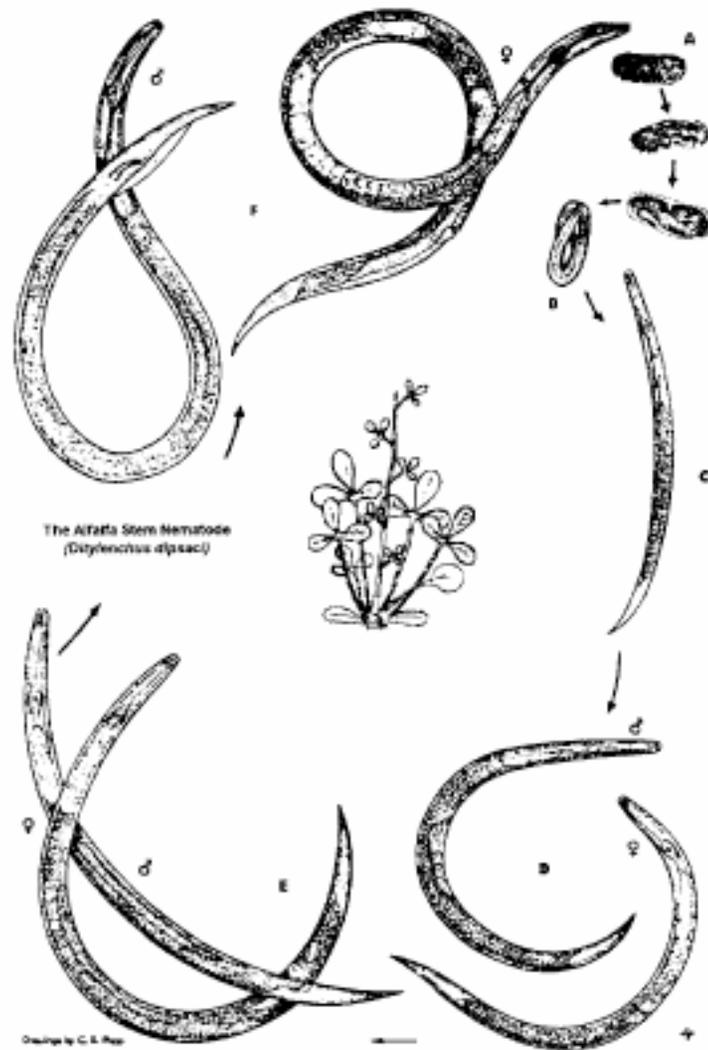


Figure 2. Life cycle of the alfalfa stem nematode (*Ditylenchus dipsaci*). A, egg; B, larvae within egg membranes; C, D, and E, second, third, and fourth stage larvae, respectively; F, mature adult male and female.

Completes life cycle in 19-30 days under optimum conditions.

After hatching...all stages are infective.

Fourth stage juvenile survives anhydrobiosis best.

Very small...live mostly inside plants although they will exit to soil for survival if plant is dying.

A clean seed sample with sound seed
(no cracked/discolored seed, no debris)



Poorer quality seed, notice color and cracked seeds and debris.



Closer examination shows alfalfa stem nematode dried onto surface of seeds (in a state of anhydrobiosis), see arrow below.



Same seed sample, more nematodes!



Scarified seed (scratched) looks different, notice straight lines.



Alfalfa stem nematodes survive best in plant debris in poorly cleaned seed.



A closer view of nematode infected debris (leaf) in an infected seed sample.



How many nematodes are in symptomatic leaves???



Answer: More than you imagined and lots of them!



Light transmission photomicroscopy of an alfalfa stem nematode. Note the very small stylet.



Dark field microscopy can help visualize the stylet.



Nematodes enter primordial bud tissue and migrate into developing buds

QuickTime™ and a
TIFF (Uncompressed) decompressor
are needed to see this picture.



Physiologic imbalance of growth hormones

Infected stems usually become
enlarged and discolored, nodes
swell, internodes fail to elongate

Pectinase, propectinase, amylase

QuickTime™ and a
TIFF (Uncompressed) decompressor
are needed to see this picture.

Anything that moves nematode-infested soil or infested alfalfa tissue moves the nematodes...eggs can survive in hay fed to cows....the resulting cows manure can spread the pathogen when spread onto clean fields!

ASN-can infect seed (in a state of anhydrobiosis), roots, and crown tissues, even roots on occasion.

ASN-infested plants are often predisposed to infection by other diseases.

Control

Host resistance-Plant Certified Seed Of
Resistant Varieties on clean ground!!

Removal of first cutting eliminates many ASN

Cut when the top 5-8 cm of soil is dry

A 2-3 year rotation with a non-host species such as corn,
small grains or beans (DO NOT PLANT SAINFOIN!)

Chemical nematicides (follow labeled instructions)-usually
too costly, generally ineffective. Use of BIOFUMIGANTS?

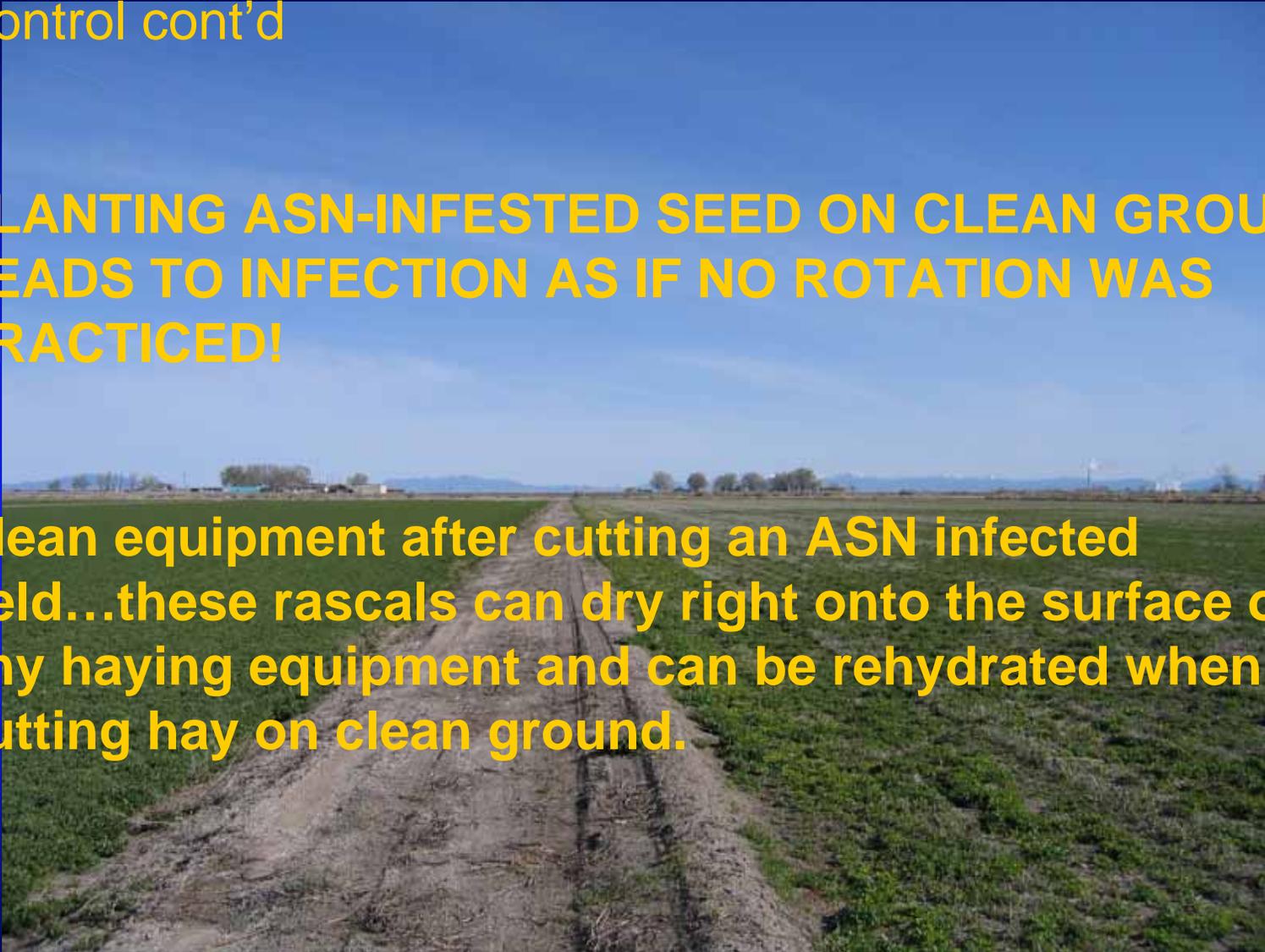
Fumigating infected seed before planting on clean ground helps
to begin with fewer or non-infected seed? Not effective?

Planting clean seed on ASN infested ground leads to infection

Control cont'd

**PLANTING ASN-INFESTED SEED ON CLEAN GROUND
LEADS TO INFECTION AS IF NO ROTATION WAS
PRACTICED!**

**Clean equipment after cutting an ASN infected
field...these rascals can dry right onto the surface of
any haying equipment and can be rehydrated when
cutting hay on clean ground.**



New Disease and Old Disease (first the old)



Phytophthora infestans on tomato-an old disease that is occurring in some greenhouses around Utah (it's wet this year!).



Sporangia on underside of leaf-enlarged view.



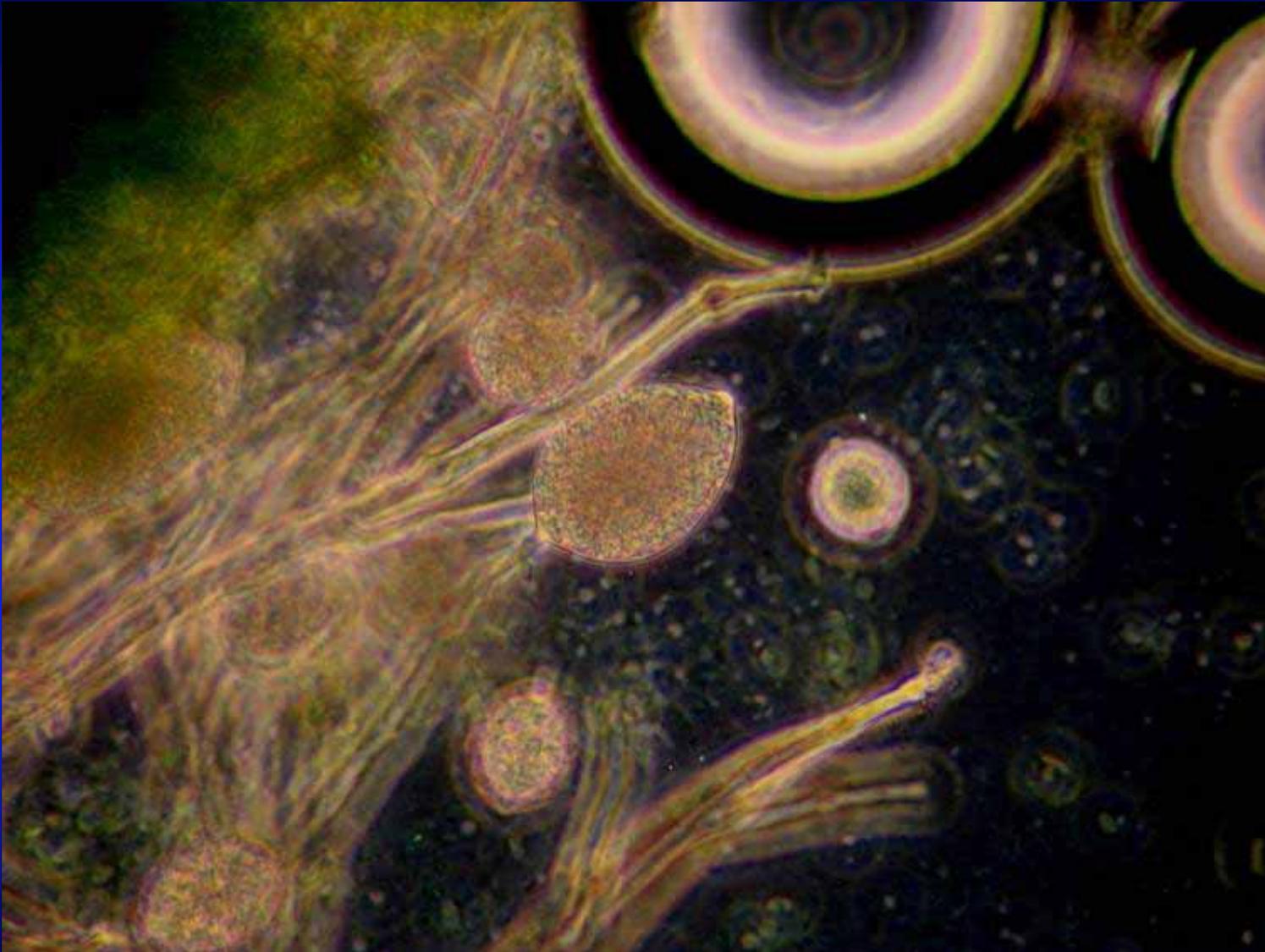
Sporangiophores and sporangia beneath the dissecting scope
(on the underside of the tomato leaf).



Spores can move by wind, water, or by mechanical means. They can germinate to directly penetrate the host. Also, we should remember that a sporangium can encyst, then differentiate, and can then germinate releasing motile zoospores (5 to 8 of them)! The zoospores swim to their new host in water!



A closer view of undifferentiated sporangia of *P. infestans*.



Just a reminder:

Some natural plant appendages such as this glandular trichome, are not spore bearing structures but are natural appendages of the plant.



Control

Sanitation.....remove infected plants and destroy them (no composting or leaving in cull pile).

Good spacing among plants....provides good air movement in the greenhouse. No overhead watering! Minimize or eliminate leaf wetness. Do not sit plants in common watering source or trays or flats.

Remove nightshade family host plants (egg plants etc.).

Chemical control may be limited for the greenhouse? If the label does not explicitly rule-out greenhouse application, then it should be OK to use it.

Cucurbit Yellow Vine Disease (CYVD)

Serratia marcescens



Identification, Phylogenetic Analysis, and Biological Characterization of *Serratia marcescens* Strains Causing Cucurbit Yellow Vine Disease

J. Rascoe, M. Berg, U. Melcher, F. L. Mitchell, B. D. Bruton, S. D. Pair, and J. Fletcher

First, second, and seventh authors: Department of E Molecular Biology, Oklahoma State University, Stillwater, Oklahoma 74078; and fifth and sixth authors: U.S. Department of Agriculture, Agricultural Research Service, Current address of J. Rascoe: Molecular Diagnostics, Current address of M. Berg: Department of Botany and Accepted for publication 23 April 2003.

Rascoe, J., Berg, M., Melcher, U., Mitchell, F. L., Bruton, B. D., and Fletcher, J. 2003. Identification, phylogenetic and biological characterization of *Serratia marcescens* strains causing yellow vine disease. *Phytopathology* 93:1233-1239.

A serious vine decline of cucurbits known as cucurbit yellow vine disease (CYVD) is caused by rod-shaped bacteria that produce phloem elements. Sequence analysis of a CYVD-specific chain reaction (PCR)-amplified 16S rDNA product showed to be a γ -proteobacterium related to the genus *Serratia*. To characterize the bacteria, one strain each from watermelon and several noncucurbit-derived reference strains were subjected to sequence analysis and biological function assays. Taxonomic placement was investigated by analysis of the 16S rDNA regions, which were amplified by PCR and directly compared, eight other bacterial strains identified by other

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Genotyping of *Serratia marcescens* Strains Associated with Cucurbit Yellow Vine Disease by Repetitive Elements-Based Polymerase Chain Reaction and DNA-DNA Hybridization

Q. Zhang, R. Weyant, A. G. Steigerwalt, L. A. White, U. Melcher, B. D. Bruton, S. D. Pair, F. L. Mitchell, and J. Fletcher

Zhang, Q., Weyant, R., Steigerwalt, A. G., White, L. A., Melcher, U., Bruton, B. D., Pair, S. D., Mitchell, F. L., and Fletcher, J. 2003. Genotyping of *Serratia marcescens* strains associated with cucurbit yellow vine disease by repetitive elements-based polymerase chain reaction and DNA-DNA hybridization. *Phytopathology* 93:1240-1246.

The bacterium that causes cucurbit yellow vine disease (CYVD) has been placed in the species *Serratia marcescens* based on 16S rDNA and *groE* sequence analysis. However, phenotypic comparison of the organism with *S. marcescens* strains isolated from a variety of ecological niches showed significant heterogeneity. In this study, we compared the genomic DNA of *S. marcescens* strains from different niches as well as type strains of other *Serratia* spp. through repetitive elements-based polymerase chain reaction (rep-PCR) and DNA-DNA hybridization. With the former, CYVD strains showed identical banding patterns despite their

ABSTRACT

Bruton, B. D., Mitchell, F., Fletcher, J., Pair, S. D., Wayadande, A., Melcher, U., Brady, J., Bextine, B., and Popham, T. W. 2003. *Serratia marcescens*, a phloem-colonizing, squash bug-transmitted bacterium: Causal agent of cucurbit yellow vine disease. *Plant Dis.* 87:937-944.

Cucurbit yellow vine disease (CYVD), which can inflict heavy losses to watermelon, pumpkin, cantaloupe, and squash in U.S. production areas from the midwest to northeastern states, causes phloem discoloration, foliar yellowing, wilting, and plant decline. Bacteria were cultured from the phloem of crown sections of symptomatic plants of *Citrullus lanatus* and *Cucurbita pepo*. Those bacteria testing positive in CYVD-specific polymerase chain reaction (PCR) were all gram negative and appeared morphologically identical, producing creamy white, smooth, entire, convex colonies on Luria-Bertani or nutrient agar. Characterized cucurbit-derived strains of *Serratia marcescens* were introduced into greenhouse-grown squash plants by puncture inoculation and into field-grown squash plants by enclosure with *S. marcescens*-fed squash bugs, *Anasa tristis*. Up to 60% of the bacteria-inoculated plants in the greenhouse and up to 17% of field plants caged with inoculative squash bugs developed phloem discoloration and tested positive for *S. marcescens* by CYVD-specific PCR. None of the controls developed phloem discoloration or tested positive by PCR. Of the diseased field plants, 12% (2 of 35) also yellowed, wilted, and collapsed, exhibiting full symptom development of CYVD. However, neither plant collapse nor decline was observed in the greenhouse-grown, puncture-inoculated plants. The morphology, growth habit, and PCR reaction of bacteria cultured from crown tissue of a subset of plants in each experimental group were indistinguishable from those of the inoculum bacteria. Evidence presented from our studies confirms that the squash bug can transmit *S. marcescens*, the CYVD causal bacterium. The *S. marcescens*-*A. tristis* relationship described here is the first instance in which the squash bug has been identified as a vector of a plant pathogen. Our experiments represent a completion of the steps of Koch's postulates, demonstrating that *S. marcescens* is the causal agent of CYVD and that the squash bug, *A. tristis*, is a vector of the pathogen.

Additional keywords: *Coridae*, *Heteroptera*, vine decline

The early research

Serratia marcescens, a Phloem-Colonizing, Squash Bug-Transmitted Bacterium: Causal Agent of Cucurbit Yellow Vine Disease

B. D. Bruton, United States Department of Agriculture-Agricultural Research Service (USDA-ARS), Lane, OK 74555; F. Mitchell, Texas Agricultural Experiment Station, Stephenville 76401; J. Fletcher, Department of Entomology and Plant Pathology, Oklahoma State University, Stillwater 74078; S. D. Pair, USDA-ARS, Lane, OK; A. Wayadande, Department of Entomology and Plant Pathology, Oklahoma State University; U. Melcher, Department of Biochemistry and Molecular Biology, Oklahoma State University; J. Brady, Texas Agricultural Experiment Station; B. Bextine, Department of Entomology and Plant Pathology, Oklahoma State University; and T. W. Popham, USDA-ARS, Stillwater, OK 74075

Early studies provided evidence that the disease was likely insect transmitted (8). For example, CYVD-symptomatic watermelon plants often were aggregated in small patches (12) and weekly insecticide applications significantly reduced the incidence of CYVD (8). In watermelon and cantaloupe fields of 5 ha or more, the incidence of CYVD was generally greater in perimeter rows than in the middle of the field. Bextine et al. (2) demonstrated that CYVD could be prevented in squash field plots by using mesh row-cover fabric to exclude insects. Several insect species initially were tested as potential vectors, although these preliminary studies failed to positively confirm or exclude any particular insect (25). Empirical evidence continued to implicate the squash bug *Anasa tristis* (DeGeer) as a vector of CYVD. Historically, although it never has been reported as a pathogen vector, the squash bug has been a serious pest of cucurbits, particularly squash and pumpkin. Its pest status and biology are reviewed elsewhere (21). In contrast to previous reports, squash bugs also can be an extremely serious pest of watermelon in Oklahoma and Central Texas (13,23,24,28), an area subject to severe outbreaks of CYVD (7). Pair et al.

Zhang, Q., Weyant, R., Steigerwalt, A. G., White, L. A., Melcher, U., Bruton, B. D., Pair, S. D., Mitchell, F. L., and Fletcher, J. 2003. Genotyping of *Serratia marcescens* strains associated with cucurbit yellow vine disease by repetitive elements-based polymerase chain reaction and DNA-DNA hybridization. *Phytopathology* 93:1240-1246.

Cucurbit yellow vine disease (CYVD) was first observed in squash (*Cucurbita maxima*) and pumpkin (*C. pepo*) in 1988 in Oklahoma and Texas (7), but is now known to affect other cucurbits, including watermelon and cantaloupe. The disease has been confirmed in Arkansas (J. C. Correll, *personal communication*), Tennessee (4), Massachusetts (26), Kansas (N. A. Tisserat, J. Fletcher, and B. D. Bruton, *unpublished data*), Colorado (B. D. Bruton, *unpublished data*), and Nebraska (R. M. Harveson and B. D. Bruton, *unpublished data*). Affected plants exhibit characteristic symptoms of yellowing, stunting, gradual decline, and phloem discoloration. Losses can range from less than 5 to 100% in affected fields.

Disease symptoms consistently are associated with the presence in the phloem of a rod-shaped, gram-negative bacterium, detected using transmission electron microscopy (7). The bacterium was cultured, and Koch's postulates were completed by mechanical inoculations and by transmission via an insect vector, the squash bug, *Anasa tristis*, confirming that this bacterium is the causal agent of CYVD (6).

Two CYVD pathogenic strains, W01-A and Z01-A, were originally isolated from diseased watermelon and zucchini, respectively. Sequence analysis of 16S rDNA and *groE* gene fragments of these two strains indicated that they shared more than

Infected squash will begin to yellow and suddenly collapse.



A preliminary diagnosis of a stem cut very close to the soil surface shows yellow discoloration in the vascular tissues...the symptoms diminish further up the stem away from the soil surface.



Yellow discoloration of CYVD infected pumpkin cut near soil surface.



What do we know about CYVD?

The bacterial pathogen is a rod shaped gram-negative bacteriumfor the most part it is not pathogenic to humans but strains in the same species can be pathogenic to some people (eg. HIV compromised persons). The bacterial species is known to be pathogenic to some insects.

Pathogenic on squash, melons, cucumbers....cucurbits!

Vectored by the squash bug.....so control your insects. Squash bugs like to lay their eggs on squash plants.....being used to bait the bug away from other cucurbits to spray them there for control.

Is it here in Utah? We don't know.....I think so...probably...we need molecular diagnostic confirmation and to conduct Koch's postulates.

Diseases of National Concern (for Utah?)

Sudden Oak Death (SOD)

<http://www.suddenoakdeath.org>



The screenshot shows a web browser window with the address bar displaying <http://www.suddenoakdeath.org/>. The page title is "California Oak Mortality Task Force". The main content area features a large green banner with the text "CALIFORNIA OAK MORTALITY TASK FORCE" and a photograph of oak leaves. Below the banner, there is a paragraph of text: "The California Oak Mortality Task Force (COMTF) focuses on the plant pathogen *Phytophthora ramorum*, which can have devastating effects in the wildlands it inhabits and has had substantial impacts on the nursery industry internationally. In 14 coastal California counties and Curry County, Oregon, *P. ramorum* has caused outbreaks of Sudden Oak Death, killing tens of thousands of native oak and tan oak trees. The pathogen also infects the leaves and twigs of common ornamental nursery plants, such as rhododendrons and camellias, which serve as vectors for pathogen dispersal." To the left of the main content is a navigation menu with links: "P. ramorum Overview", "Nursery Information and Update", "Maps & Photos", "Research", "Library", "Regulations", "Management Recommendations", "Task Force Information", "COMTF Monthly Newsletters", "Training and Extension", "Upcoming Events", and "Contacts". Below the menu is a search box with a "Search" button. To the right of the main content, there are several sections: "COMTF Monthly Report: Sign up HERE" and "Current Report: May 2005"; "Host of the Month"; "New and Noteworthy" with sub-sections "SOD Outreach Survey Results" and "Training session - May 24"; "Sudden Oak Death and *Phytophthora ramorum* 2004 Summary Report"; "Press Releases and Media Assistance"; and "Guides and Resources" with a sub-section "Nursery Information: Regulatory/quarantine information from the California Department of Food and Agriculture (CDFA) and USDA Animal and Plant Health Inspection Service (APHIS); diagnostic guides; photos;".

Asian Soybean Rust

<http://www.plantpath.iastate.edu/soybeanrust/>

The screenshot shows a web browser window with the URL <http://www.plantpath.iastate.edu/soybeanrust/>. The page title is "Asian Soybean Rust | Identifying, Confirming and Managing Asian Soybean Rust". The main content area features a header with a soybean leaf image and the text "Asian Soybean Rust Identifying, Confirming and Managing Asian Soybean Rust". Below this is a paragraph describing the disease: "Asian soybean rust (*Phakopsora pachyrhizi*) is a fungal disease that can quickly defoliate plants and reduce pod set, pod fill, seed quality and yield. The disease is found in every soybean region in the world. It was recently confirmed in the southern United States and is expected to spread to other parts of North America during the upcoming growing season. This site was developed as a resource for Iowa soybean producers and agriculture professionals to help identify, confirm and manage the disease." A navigation menu on the right includes "Identify", "Confirm (Fast Track)" (with sub-links for "First detectors" and "Triage Team"), "Manage", "Resources", "Image Gallery", and "Questions and Answers". Below the menu is a search box and a "NEWSREEL" section with several news items, including "Asian Soybean Rust Found in Georgia (4/27/2005)". A world map titled "Where is Asian Soybean Rust?" shows the disease's distribution, with a callout for the United States. At the bottom, a news article titled "Asian Soybean Rust Found in Georgia" is published on 4/27/2005, reporting the discovery of the fungus in a field in Seminole County.

Common characteristics of both diseases:

Both are considered new diseases

Both pathogens have a very wide host range....and they are getting wider for SOD (Jasmine is the new host of the month)!

Both are important to the agriculture of the U.S.

Both are capable of moving from infected areas to uninfected areas

Both are known to have economic impacts but both diseases are potentially more ecological damaging than anyone is currently admitting (more realization for SOD at this time).

Can these diseases occur in Utah?

Never say never, if the conditions are right (adequate moisture, suitable host, inoculum of the pathogen) it can/will happen. I hope neither pathogen gets to Utah!

Thank You!

Drive safely!

