

New Approaches in Assessing Fusarium in California Cotton

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Thanks to:

- Grower and PCA Cooperators
- National Cotton Council & Cotton Inc. State Support Committee, CA Department of Food and Agriculture
- seed companies

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Fusarium in cotton at 7-9 node stage
(Fungal disease caused by *Fusarium oxysporum f. sp. vasinfectum*)

Fusarium – *recognized and researched as a disease in cotton for years, so why look at it again?*

- **What is different, new, or newly recognized?**
- First seen in finer-textured soils than typical for other FOV races assoc. with RKN – range of soil textures
- Confirmed that RKN pop'n's were very low (usually associated with damaging FOV pop'n's in CA cotton)
- Newly recognized strain for California (race 4)
- Few infected fields confirmed 2001-2004 (17 or 18)
- Mike Davis (UCD) and staff have been working on strain identification - has info on Australian strains as well as other races to identify that this is not Australian FOV

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Fusarium - Questions / Concerns

- How identify this as a newly-recognized FOV race in California cotton? Why be interested?
- Could what we are seeing be related to damaging strains of FOV isolated in Australia? *At this point nothing identified has been of either of two known serious Australian FOV strains*
- What available methods can ascertain with more certainty different races of FOV dealing with?
 - *important in trying to assess potential for damage*
 - *Any value in identifying points of origin, sources?*
 - *Improved tools to look at genetics of host plant resistance*

1st Steps: Sampling fields, Collecting Information, When to Sample?



Symptoms can develop across a range of times and eventually lead to leaf abscission – foliar damage and leaf loss can occur over period as short as 1-2 weeks or much longer

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Recognizing symptoms



Vascular staining - easily seen in lower stem & upper tap root – how differ from verticillium vascular symptoms?

1. Seen more readily in root as well as lower stem
2. timing/growth stage when first seen much earlier
3. Staining tends to be more continuous rather than "flecking" or discontinuous discoloration

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Field Scale
Symptoms:

Field 14-17a

- Typical field had affected areas about this size, with stunted "survivor" plants that produced harvestable bolls
- One 2003 field had much larger affected areas



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FIELD TRIALS: *Observations / activities summary 2002-2004*

- Evaluated sites where plants developed symptoms consistent with Fusarium (about 65 fields in 2002, >80 in 2003, >40 in 2004).
- Samples collected in >25-30 fields each year – most confirmed as Fusarium races long-recognized in CA, but samples from about 12+ cotton fields in 2002-2004 were positive for **race 4 FOV** (about $\frac{3}{4}$ Pima, rest Acala in sandy loam to clay loam soil sites w/o Root Knot Nematode)
- Any fields where growers or PCA's contacted UCCE were evaluated for Fusarium, sampled if symptoms evident and permission given

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Approaches in Identification

Field Sampling provided samples, so what is needed for ID and decision-making?

Identification / Genetic Fingerprinting:

- partial sequences of EF, B-tubulin, phosphate permase genes
- pathogenicity tests also conducted on plants grown in greenhouses
- isolates identified and stored, database maintained

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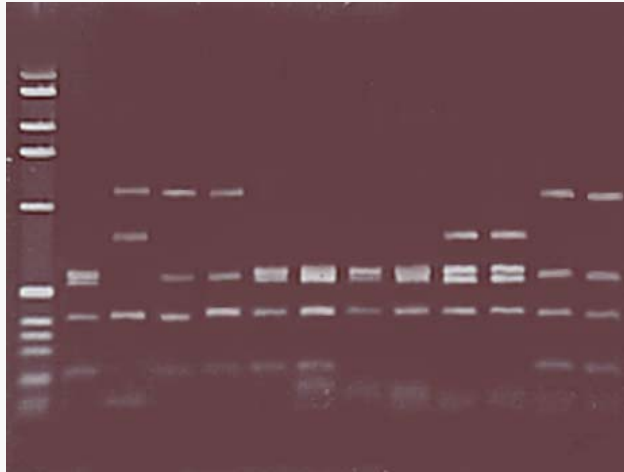
Approaches in Identification

Genetic Fingerprinting:

- IGS region of specific nuclear DNA amplified with specific primers
- amplified product digested by restriction enzymes *Scrf1* and *Rsa1*, and digested products separated by electrophoresis
- gels stained and examined to establish differences
- as develop information, may look in much more detail at base sequencing, markers and greater specificity re: sources of susceptibility and resistance

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IGS/RsaI

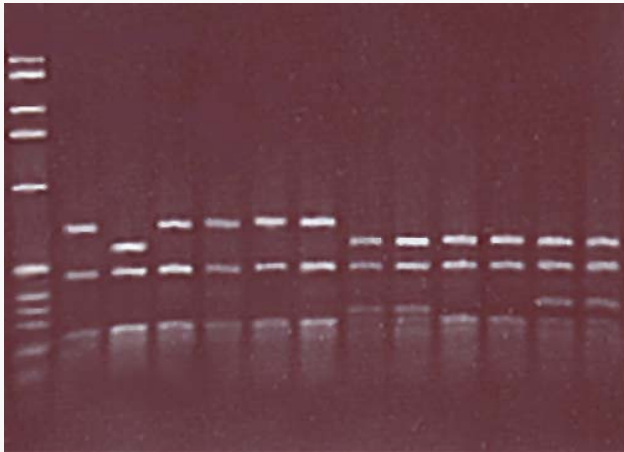


Genotype

3
17 18
5 6 19 21
1 7 8 14 15

1 3 5 6 7 8 14 15 17 18 19 21
California Strains of FOV **Aust1** **Aust2**
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IGS/ScrFI



Genotype

1 5 6 7 8
3 17 18
14 15
19 21

1 3 5 6 7 8 14 15 17 18 19 21
California Strains of FOV **Aust1** **Aust2**
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To be more certain with this analysis, can also look for more specific identification of where the genetics differ – by looking at DNA base sequence

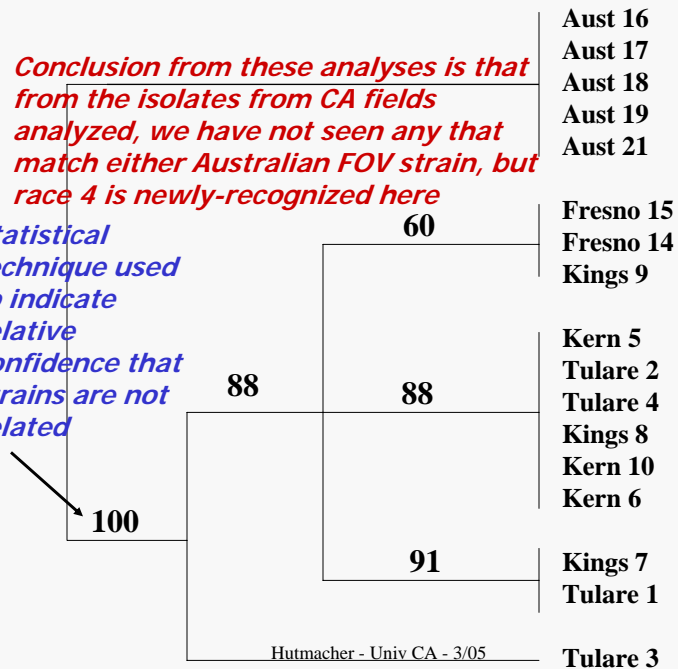
Partial Sequence of EF gene

| | | | | | |
|---------|---------|------------|------------|------------|----|
| Aus 16: | GACGGGG | CGCGTTTGCC | CTCCT.ACCA | TT...ACAAC | CT |
| Aus 17: | GACGGGG | CGCGTTTGCC | CTCCT.ACCA | TT...ACAAC | CT |
| Aus 18: | GACGGGG | CGCGTTTGCC | CTCCT.ACCA | TT...ACAAC | CT |
| Aus 19: | GACGGGG | CGCGTTTGCC | CTCCT.ACCA | TT...ACAAC | CT |
| Aus 21: | GACGGGG | CGCGTTTGCC | CTCCT.ACCA | TT...ACAAC | CT |
| Kgs 9: | GAC.GGG | AGCGTTTGCC | CTCTT.ACCA | TTCTCAGAAC | CT |
| Frs 14: | GAC.GGG | AGCGTTTGCC | CTCTT.ACCA | TTCTCAGAAC | CT |
| Tul 2: | GAA.GGG | AGCGTTTGCC | CTCTT.ACCA | TTCTCACAAC | CT |
| Krn 5: | GAA.GGG | AGCGTTTGCC | CTCTT.ACCA | TTCTCACAAC | CT |
| Kgs 8: | GAA.GGG | AGCGTTTGCC | CTCTT.ACCA | TTCTCACAAC | CT |
| Krn 10: | GAA.GGG | AGCGTTTGCC | CTCTT.ACCA | TTCTCACAAC | CT |
| Tul 1: | GAC.GGG | AGCGTTTGCC | CTCTT.ACCA | TTCTCACAAC | CT |
| Kgs 7: | GAC.GGG | AGCGTTTGCC | CTCTT.ACCA | TTCTCACAAC | CT |
| Tul 3: | GAC.GGG | AGCGTTTGCC | CTCTTAACCA | TTCTCACAAC | CT |

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Conclusion from these analyses is that from the isolates from CA fields analyzed, we have not seen any that match either Australian FOV strain, but race 4 is newly-recognized here

Statistical technique used to indicate relative confidence that strains are not related



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California Cotton FOV's
(Identified to Date) – Davis et al

| Races 1&2 <i>(Americas)</i> | Race 3 <i>(Nile Valley)</i> | Race 4 <i>(Asia)</i> | Race 8 <i>(Asia)</i> |
|--|---------------------------------------|---|--------------------------------|
| Kern Kings Tulare | Tulare | Kern Fresno Tulare | Kings Tulare |

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No. of Race 4-infested Fields

| | Fresno Co. | Tulare Co. | Kern Co. |
|---------------|-------------------|-------------------|-----------------|
| 2002-3 | 6 | 1 | 0 |
| 2004 | 6 | 1 | 4 |
| Total | 12 | 2 | 4 |

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To deal with disease issues such as FOV race 4, efforts also needed in:

- Identifying if varietal resistance / tolerance is available or can be developed
- Containment practices / management to reduce rate of spread

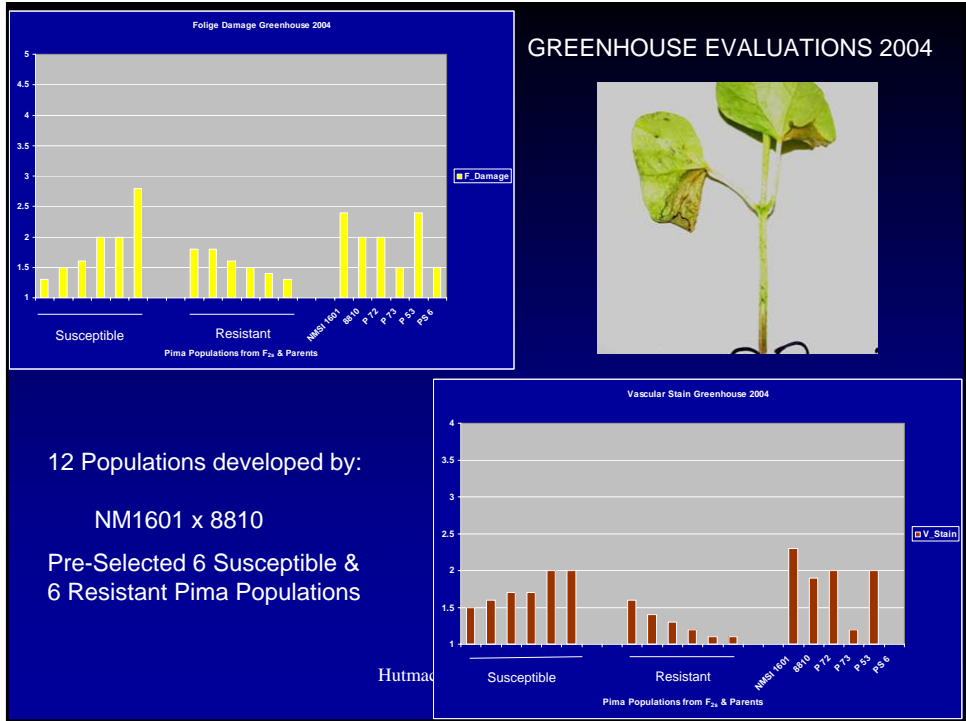
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HOST PLANT RESISTANCE EVALUATIONS FOV RACE 4 – 2003 and 2004

- Fresno County (50 entries – 2003), (100 entries – 2004) 25' plots. Planted July 3 (2003), May 24 (2004)
- Kern County 24 entries, 3 reps with 15' plots. Planted May 26, 2004 (grower fields)
- Greenhouse 50 entries, total 9 plants/reps & 5 as control. During Oct. 19th to Dec 7th.



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Field Variety Screens – FOV race 4 sites 2003-2004

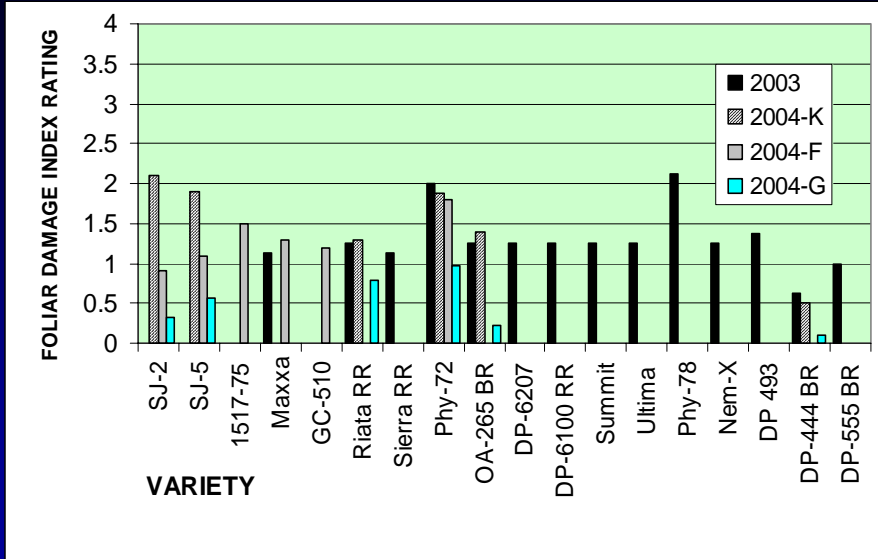
Susceptible varieties severely affected – others grow through it to varying extents even if infected

Range of measurements made – focus here on:

- root vascular staining index
- foliar symptom index
- surviving plant number
- plant height

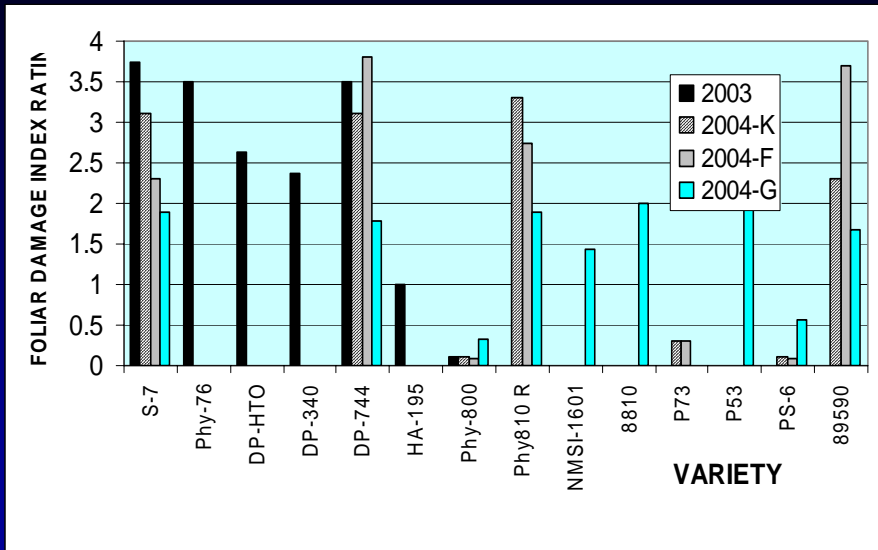
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Foliar Damage Rating – Acala/Upland



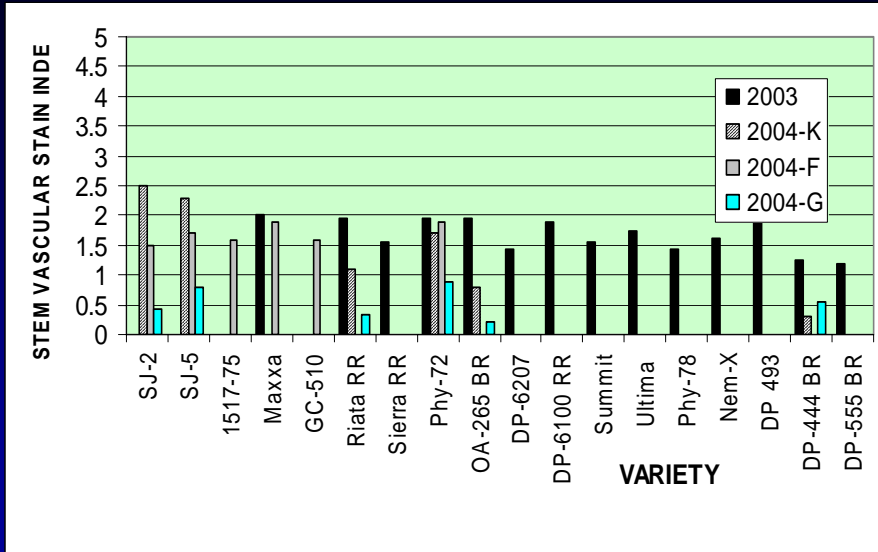
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Foliar Damage Ratings – Pima entries



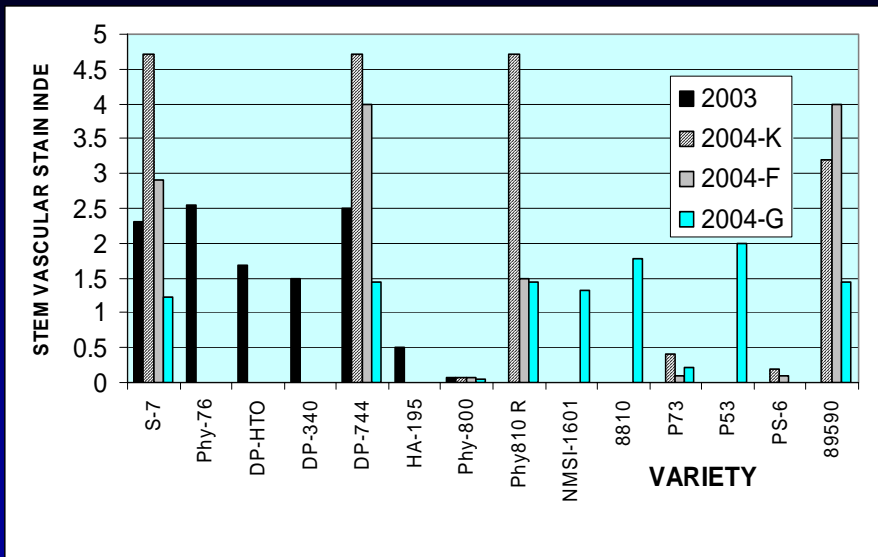
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Stem Vascular Stain Rating—Acala/Upland



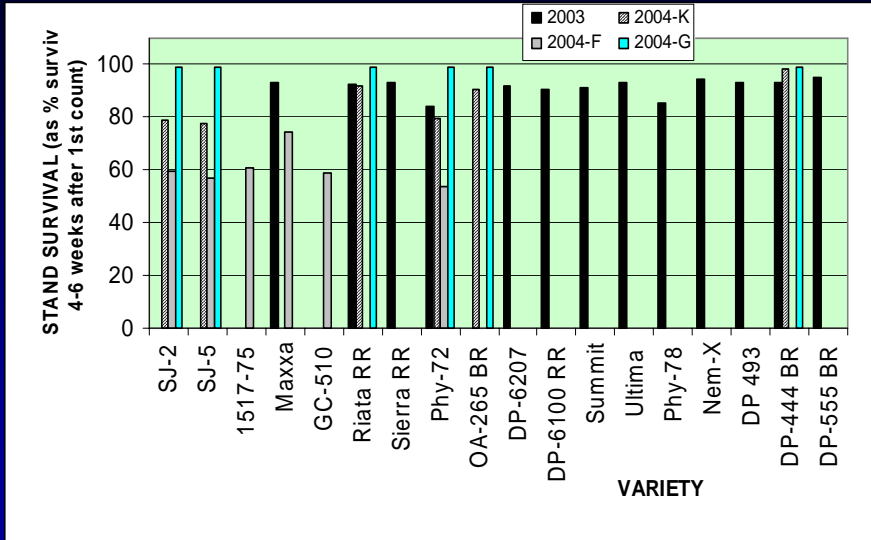
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Stem Vascular Stain Rating – Pima entries



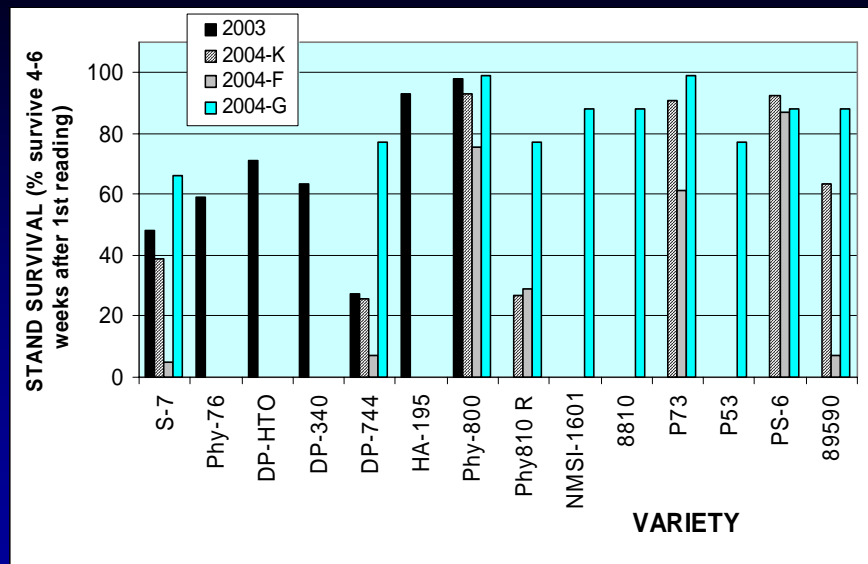
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Plant Survival Percent – Acala / Upland



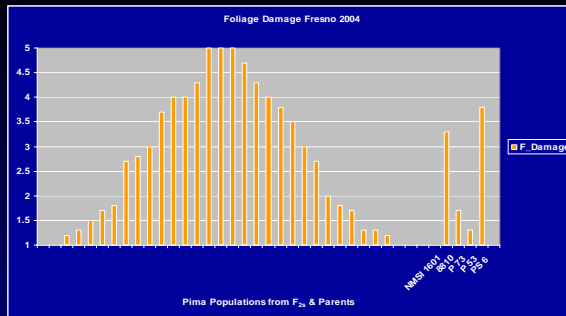
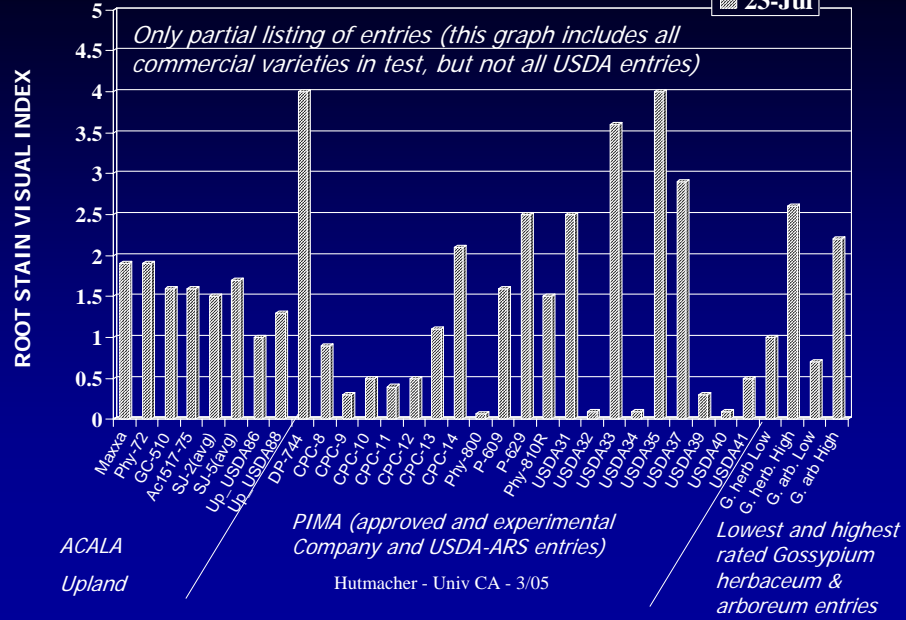
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Plant Survival Percent – Pima Entries



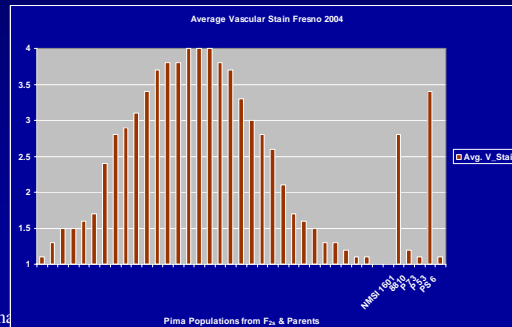
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Root Vascular Stain Index Ratings – Fresno Co. site - 2004 Fusarium race 4 field



EXPAND INFORMATION DEVELOPED ON:

- Heritability
- Potential Markers Useful in Selection and Improvement



Summary – Identification Approaches

Genetic Fingerprinting:

- As develop more information, more detailed work on heritability questions, host plant resistance in public materials, identification of markers and greater specificity re: identification of sources of susceptibility and resistance

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SUMMARY

- FOV race new or newly recognized in CA cotton
- Many evaluated Pima varieties were observed to be more susceptible to FOV race 4 (stand loss, stunting, etc) than evaluated *Gossypium hirsutum*s
- Preliminary results showed some Pima germplasm has a more complete resistance to FOV race 4.
- The impact of the disease for Acala / Upland cottons was milder than effects on most Pima, but still a problem, since those Acala / Uplands infected by FOV race 4 could reproduce & expand inoculum.

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SUMMARY

- In Pima for this study, preliminary results showed that resistance for FOV race 4 may be inherited quantitatively and controlled by several genes.
- Heritability estimates indicated that selection for FOV race 4 can be accomplished.
- Early evaluations to look for highly resistant Acala and/or Upland cottons suggests that resistance in *G. hirsutum* may be more complex.
- Evaluations will continue

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Part of dealing with any disease still involves efforts to contain problems while more permanent solutions developed:

- Identifying if varietal resistance / tolerance is available or can be developed
- Containment practices / management to reduce rate of spread

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Containment issues for growers as sites with FOV are identified ?

- *Remember that spores of this organism can be very long-lived so limit practices that expand movement*
- *What rotation crops will reduce inoculum / pop'ns most non-cotton crops will reduce inoculum levels, but unlikely any crop will eradicate it*
- *Can this strain influence other crop spp? Highly unlikely, but can impact Acalas/Uplands and Pimas*
- *Can inoculum be spread in fields with soil transport or movement of plant parts (leaves, flowers, squares?) ...yes.. By irrigation? ...yes... cultivation ...yes*

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Containment / research issues

Improvements in detection methods (could be useful in soil, and in plant tissues such as seeds – quick tests versus more in-depth methods)

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**Develop and expand methods to
provide general information
(symptoms & recommendations,
containment)**

Example: UC cotton web site

<http://cottoninfo.ucdavis.edu>

Thank you

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