

# **Reniform Nematode Resistance from *Gossypium longicalyx* - Cytogenetics, Breeding & Molecular Marker Development**

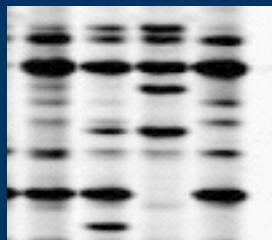
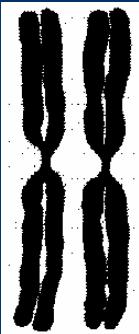
**Nilesh Dighe, David Stelly,  
Alois Bell, Forest Robinson and Monica Menz**

**Texas A&M University  
&  
USDA-ARS, College Station**

# Objectives

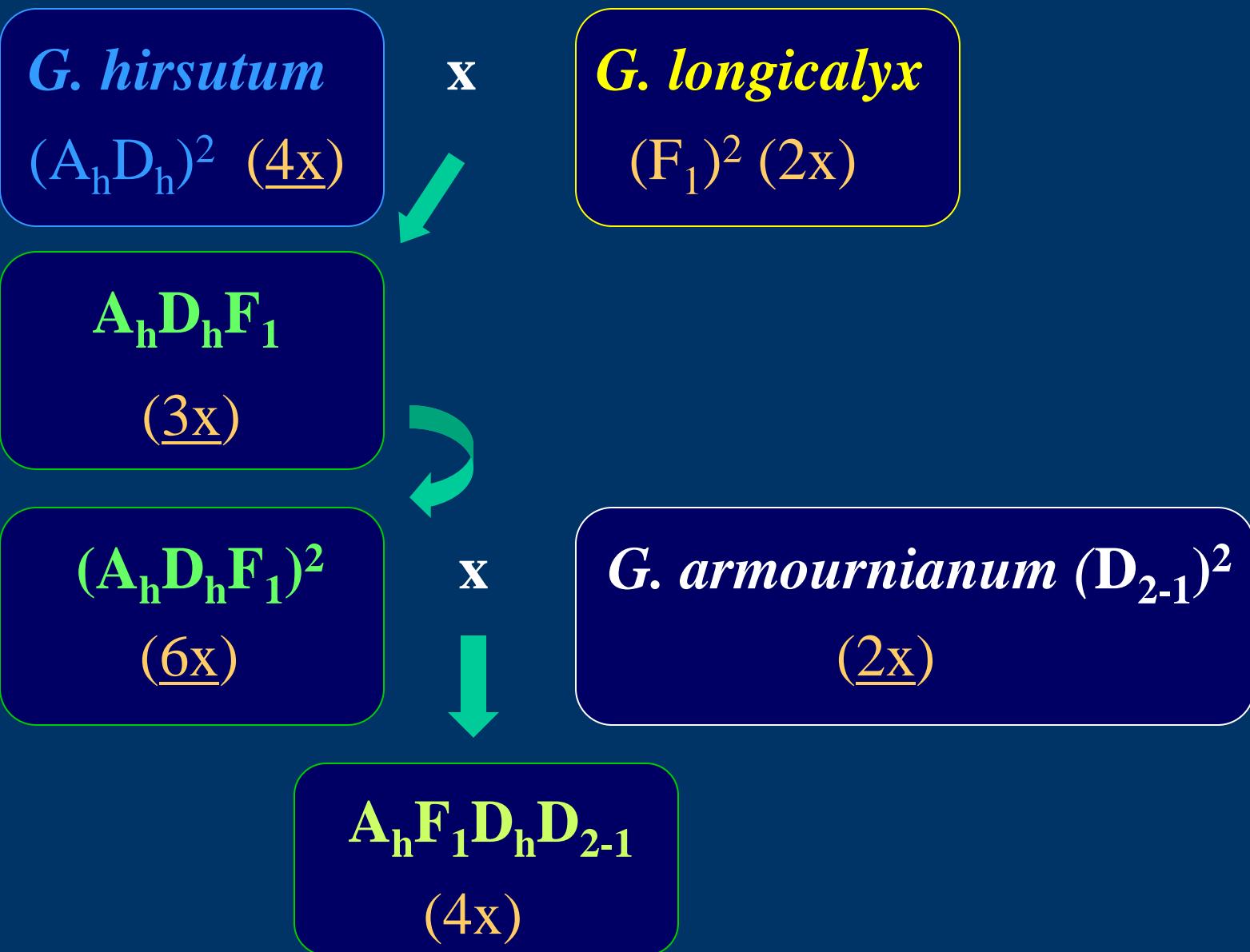


- **INTROGRESSION (selective):**
  - Resistance to reniform
  - NOT undesirable traits
- **CYTOGENETICS:**
  - Chromosomal constitutions
  - Numbers and kinds of aberrations
  - Recombination
  - Alien segment size
- **MARKERS:**
  - Analyses
  - MAS



# Triple Species Hybrids

(Bell and Robinson, USDA)



# Two Triple-species Hybrids

(Bell and Robinson, USDA)

TWO resistance sources:

- 1) (*G. hirsutum* x *G. longicalyx*)<sup>2</sup> x *G. armourianum* (HLA)



- 2) (*G. hirsutum* x *G. herbaceum*)<sup>2</sup> x *G. longicalyx* (HHL)



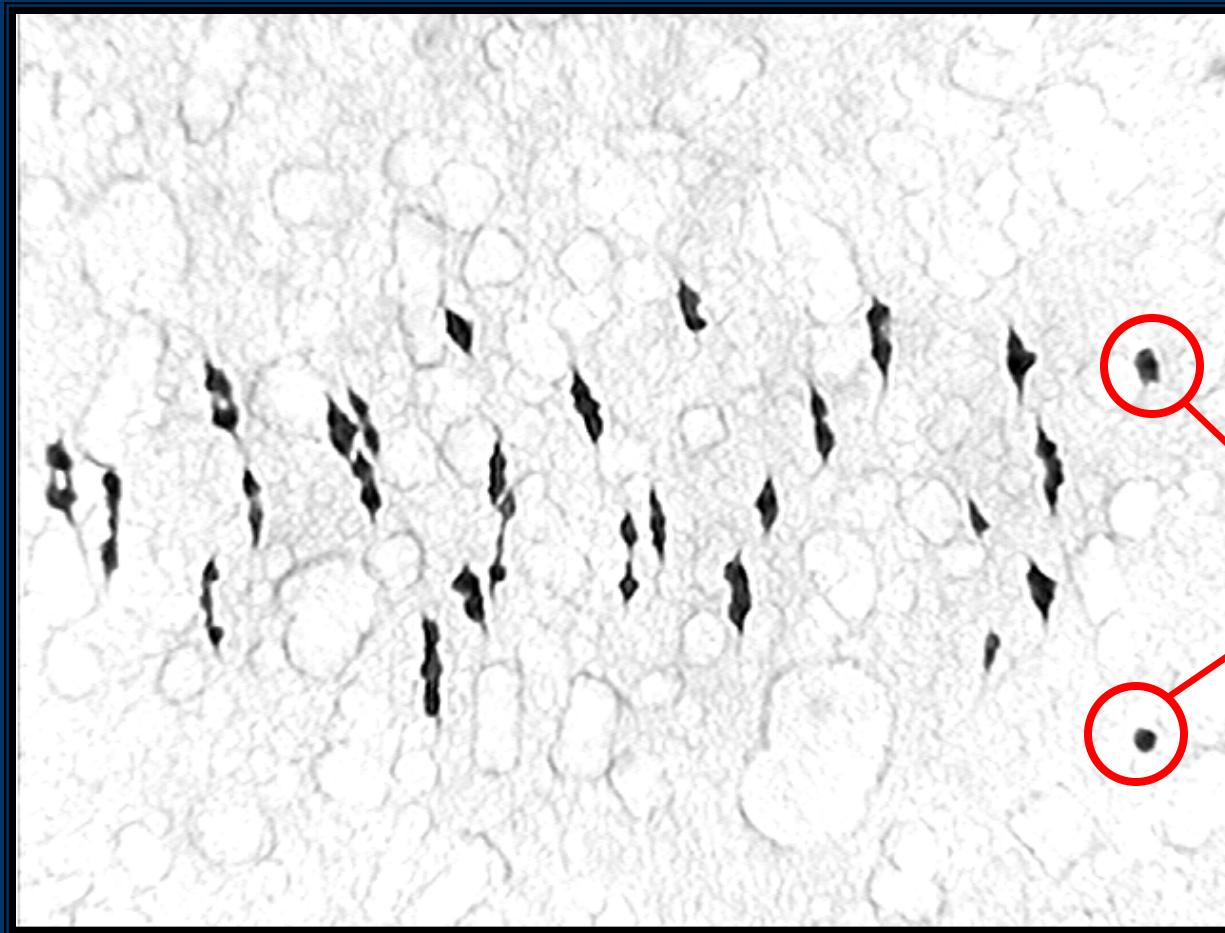
# Cytogenetics Data Summary

Generation	# Plants Analyzed
BC1F1	38
BC1S1	31
BC2F1	94
BC3F1	23
BC4F1	15
BC5F1	2
<b>Total</b>	<b>203</b>

# Chromosome numbers

Chrom.	51 Chr.	52 Chr.	53 Chr.
Gener.			
BC1F1	(5%) 2	(66%) 25	(29%) 11
BC1S1	(3%) 1	(84%) 26	(13%) 4
BC2F1	(22%) 21	(61%) 57	(17%) 16
BC3F1	0	(100%) 23	0
BC4F1	0	(100%) 15	0
BC5F1	0	(100%) 2	0

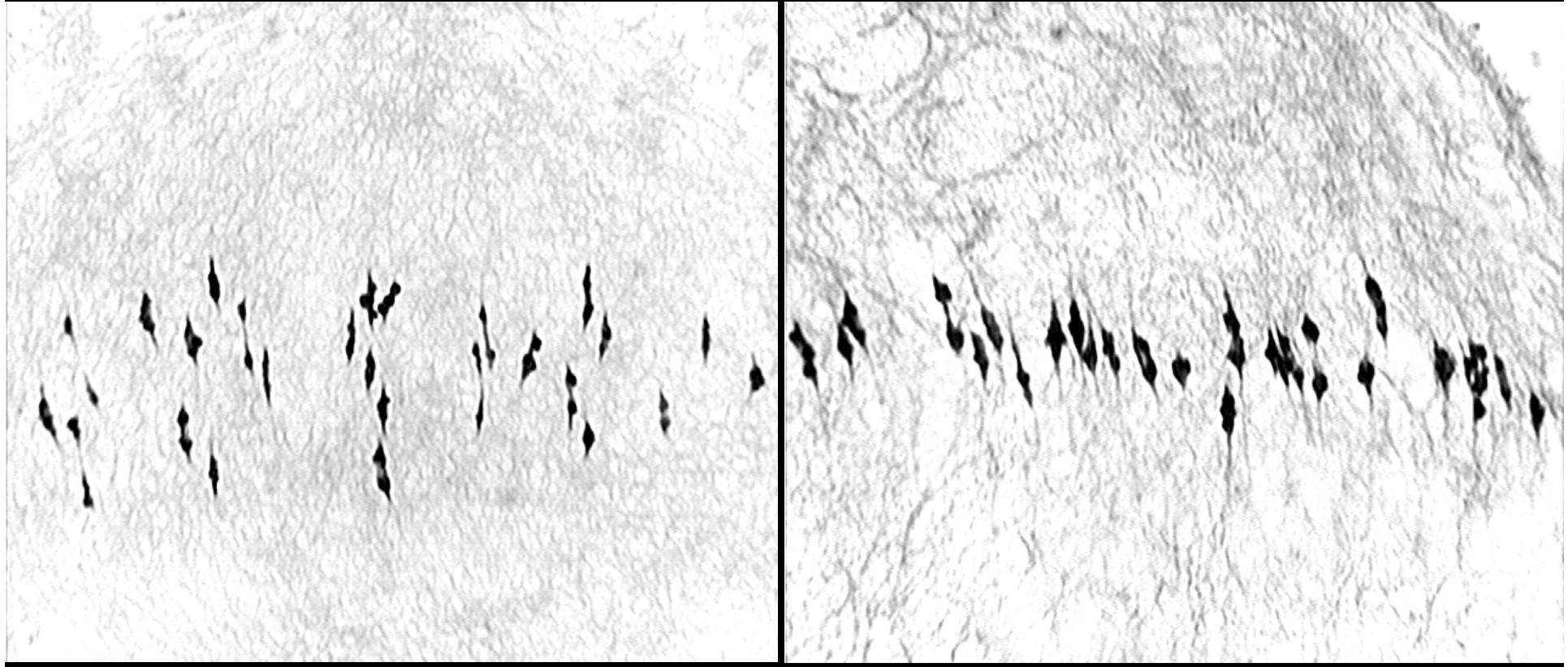
# Model Type in BC2



(Not  
recombinant)

25II+2I (Res-BC2-129)

# Modal Type in Resistant BC3, BC4 and BC5 Plants



26II

26II

# Recombination

Proximal and distal  
XOs are too distant.

Proximal XO is okay.  
Distal XO is too distant.



Proximal XO is too  
distant. Distal XO is  
okay.



Proximal and distal XOs  
are okay.



# Recombination

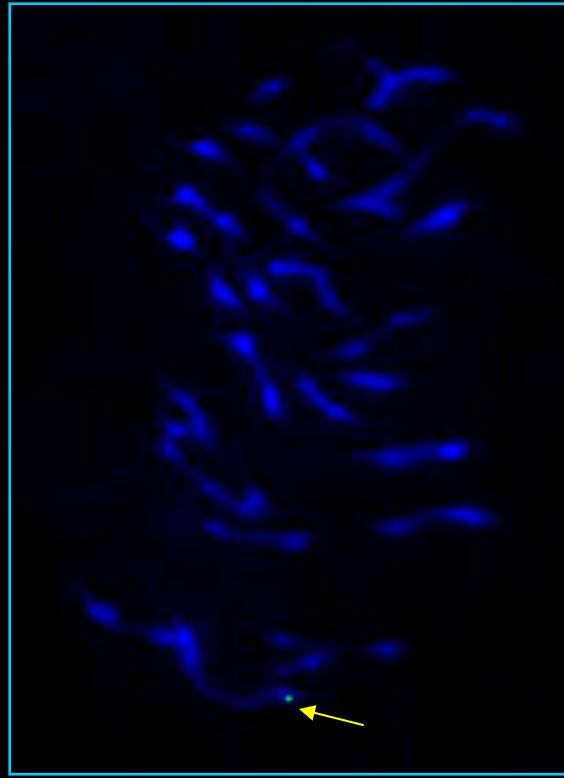
- \* Estimate of Recombination in BC2

- 2/20 Resistant Plants Analyzed form either 26II OR 1IV+24II
- At least ~10% Recombination in BC2

- \* Estimate of Recombination in BC3 and BC4

- Approx. 85% of the Resistant Plants Analyzed form 26II OR 1IV+24II

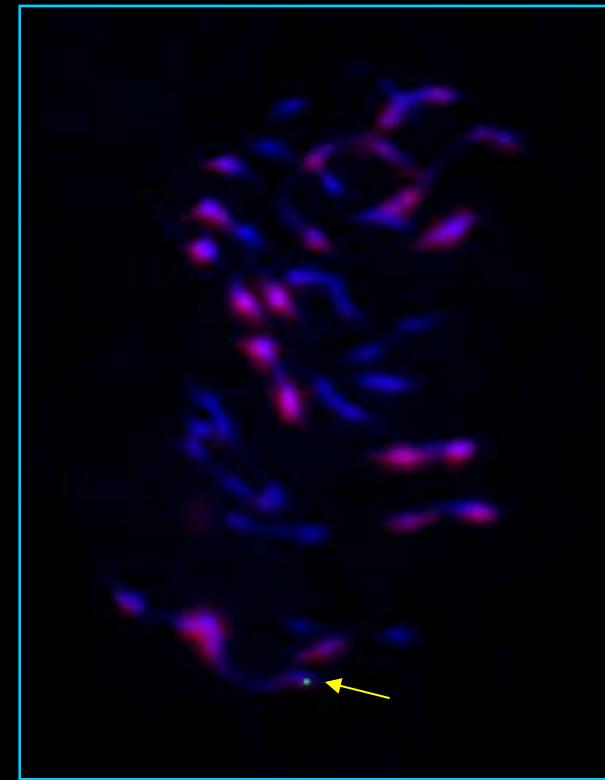
# **GISH (Genomic in situ hybridization) on BC3 1-6 (Resistant)**



**Blue---All the chromosomes**

**Red----A1-genome**

**Green—F1-genome**



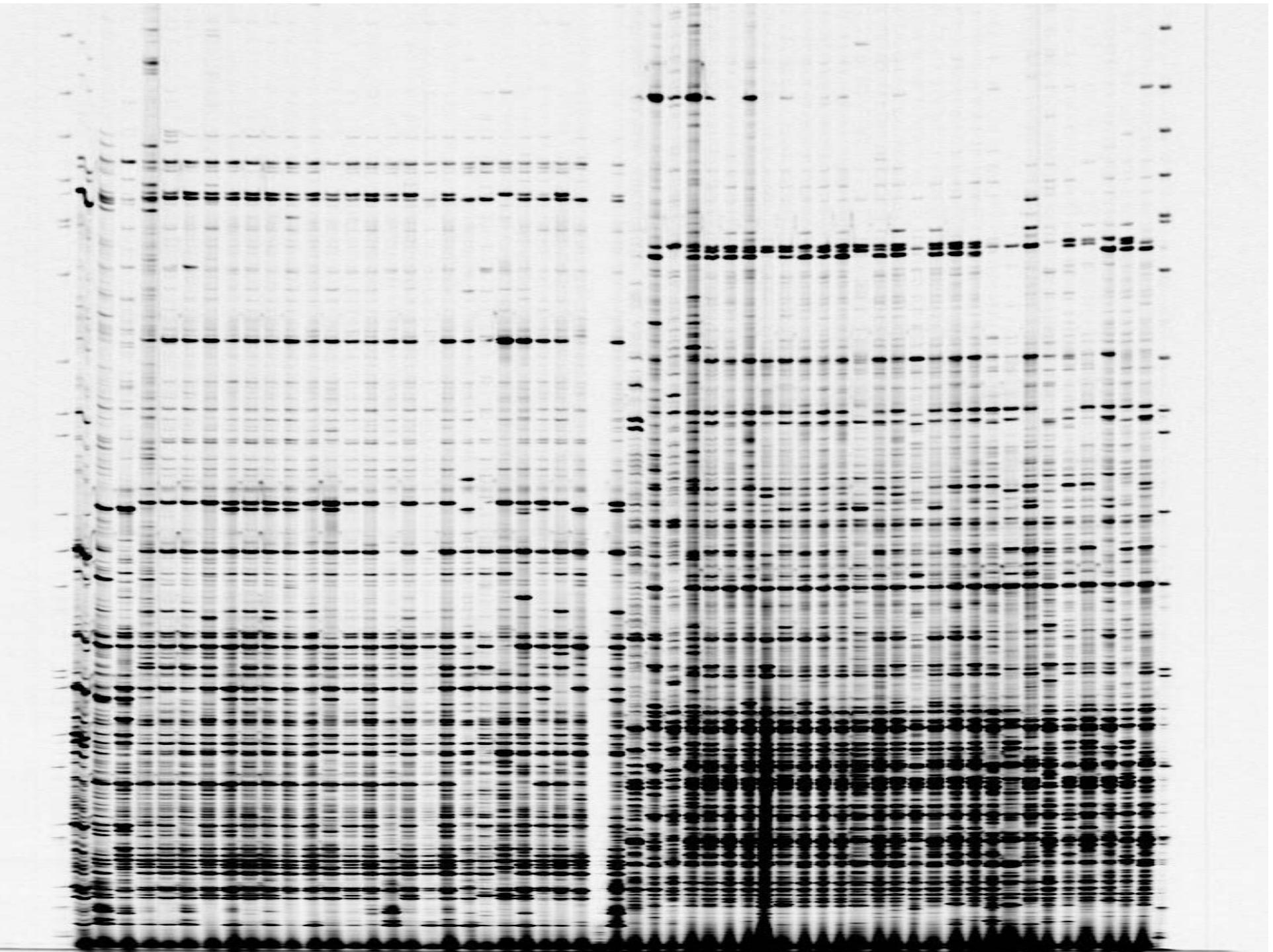
**BC3 1-6**

**Highly Resistant plant  
26II Chrom. Config**

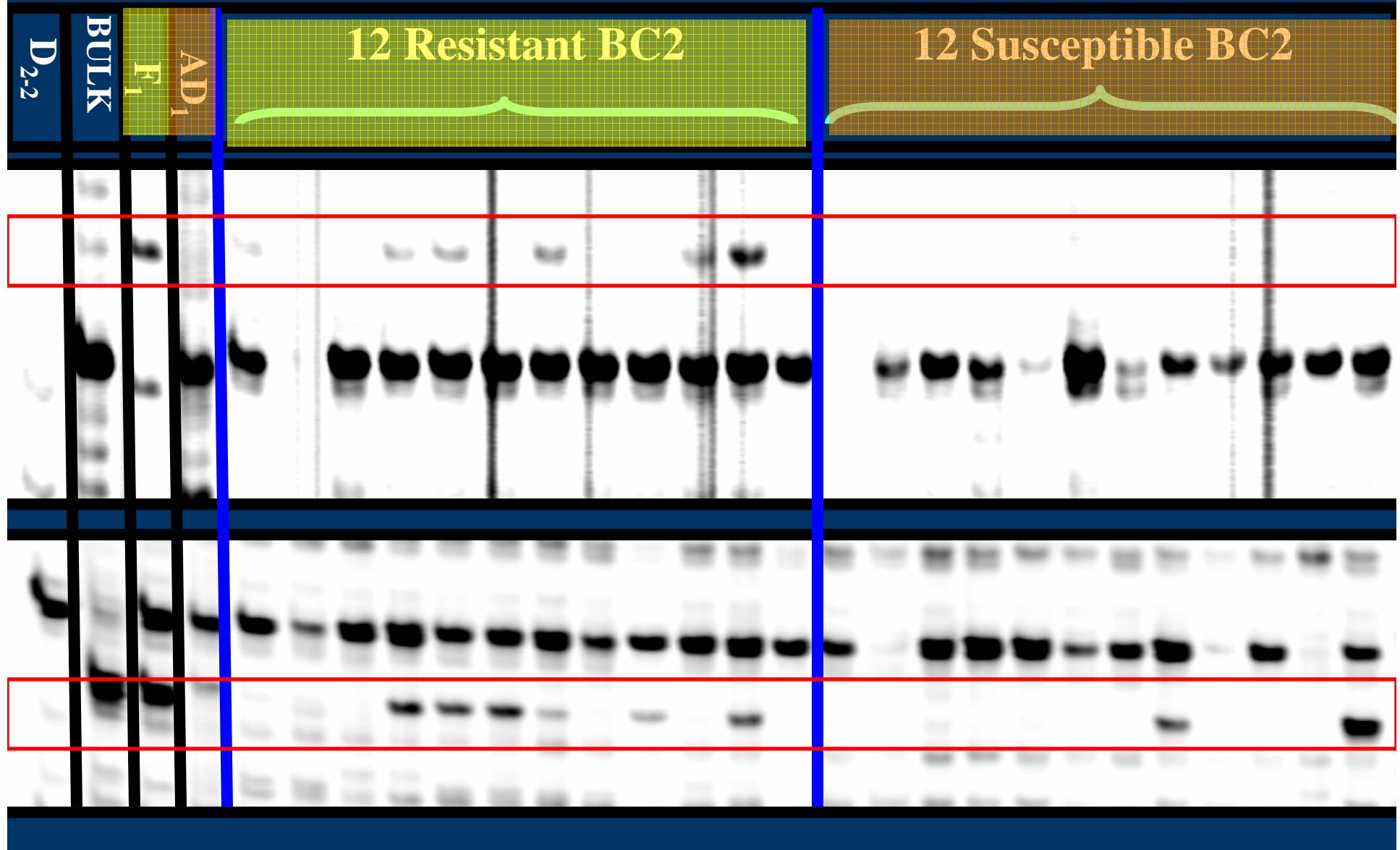
**BC4F1 Segregating 1:1 (5R:5S)**

# MARKER DEVELOPMENT

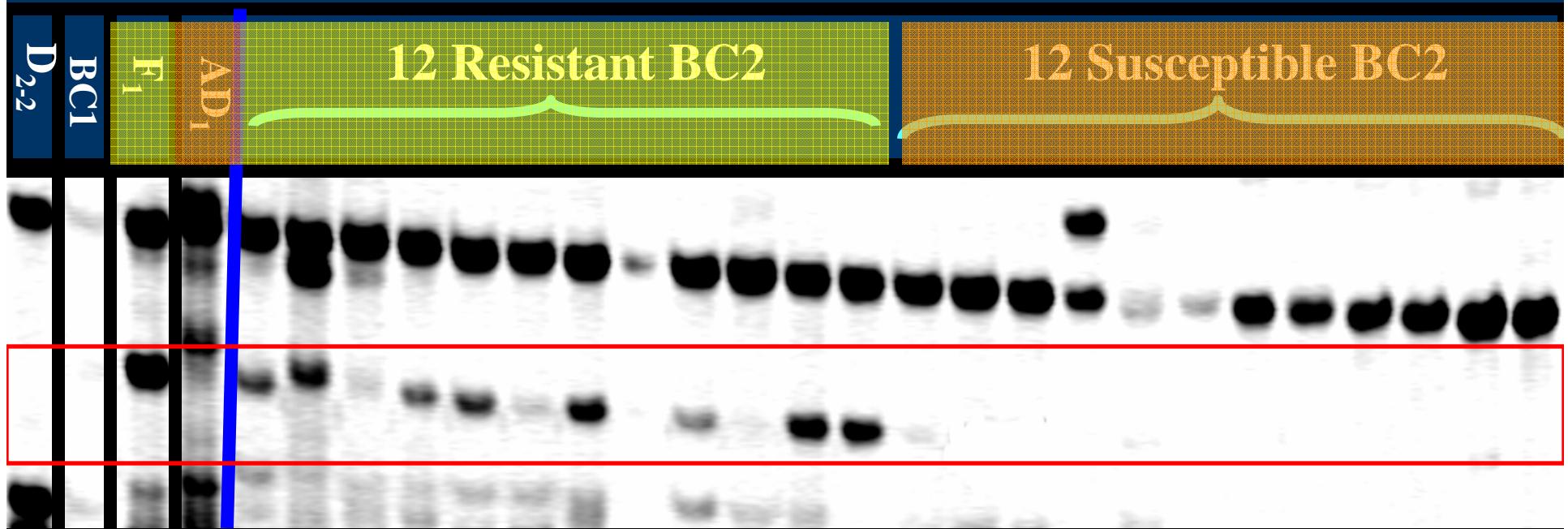
- **“PHASE-I” MARKERS:** alien AFLPs
- **SCREENING PANEL:**
  - *G. hirsutum* ( $\text{AD}_1$ )
  - *G. longicalyx* ( $F_1$ ) and *G. armourianum* ( $\text{D}_{2-2}$ )
  - Resistant  $\text{BC}_1$  parents (Bulked)
  - 12 most resistant plants of 186  $\text{BC}_2$  plants
  - 12 most susceptible plants of 186  $\text{BC}_2$  plants
- Li-Cor System



# Potential Markers



# Potential Markers



# Marker Development

## Research in Progress..

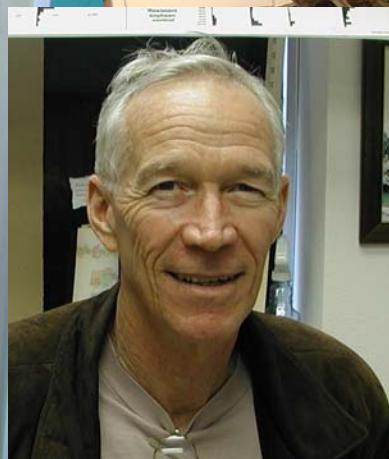
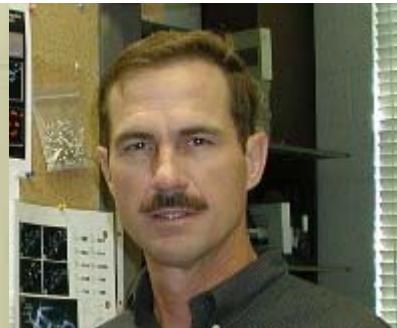
- Identify AFLP markers tightly linked to the reniform nematode resistance gene

## THEN:

- “Phase-II” Markers -- e.g. SSRs
  - Identify linkage group (*using mapped loci*)
  - MAS, especially among inbreds (“Zygosity assays”)

# STATUS SUMMARY

- The existing data indicate successful introgression!!
- NEEDS:
  - Markers for MAS
    - Backcrossing
    - Inbreeding
  - Eliminate unnecessary alien germplasm
    - GISH (physical size of alien segment)
    - Markers
    - Phenotypic screens
    - Yield and fiber assessments



**Acknowledgments**  
**Cotton Inc.**

