

Identifying a gene for Southern root-knot nematode resistance



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Dept of Crop and Soil Sciences Institute for Plant Breeding, Genetics & Genomics 5th Annual Soybean Precision Genomics Workshop & Mutant Finder Field Day



August 3-4, 2016, University of Missouri- Columbia

- Presentations on soybean mutant resources
- Tutorials on cutting edge technologies in soybean
 - targeted mutagenesis/gene editing
 - gene silencing
 - other functional genomics applications
- Mutant populations & seeds
- Funding is available to help defray attendance costs
 - Contact Bing Stacey (staceym@missouri.edu)



2nd 15 Years Fine-mapping • Candidates • Identification



2 188 F5:6 RILs

17 had recombination between flanking markers Interval screened with 7 markers

235-kb with 30 gene models

Theor Appl Genet (2013) 126:1825–1838 DOI 10.1007/s00122-013-2095-8

ORIGINAL PAPER



Fine mapping and identification of candidate genes controlling the resistance to southern root-knot nematode in PI 96354

Anh-Tung Pham · Kaitlin McNally · Hussein Abdel-Haleem · H. Roger Boerma · Zenglu Li

13 Candidate genes



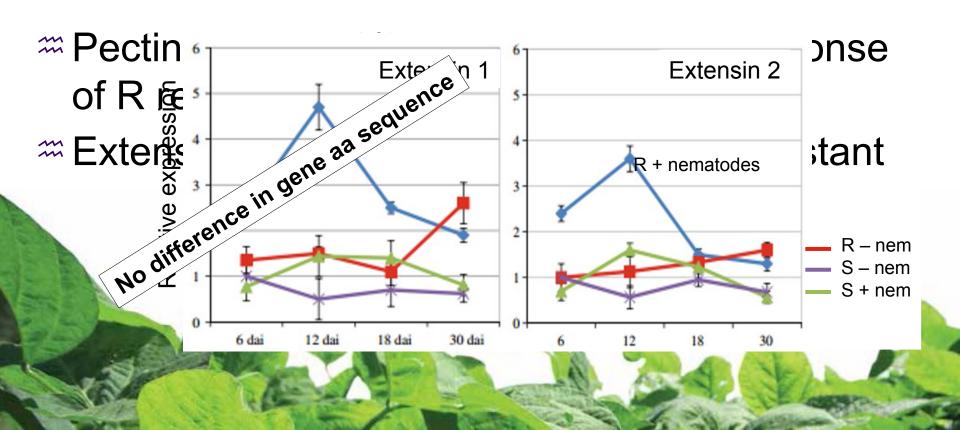
Section 34 Based on

- Root-specific expression based on Soyseg
- Upregulated in other spp in presence of nematodes
- Sequenced R & S alleles/promoters
 - 22 15 changes in S vs reference
 - 2 93 changes in R vs reference
 - 2 had no polymorphisms
 - 5 had silent mutations or in introns
 - ** 1 had a deletion not associated with resistance

4 Remaining candidate genes

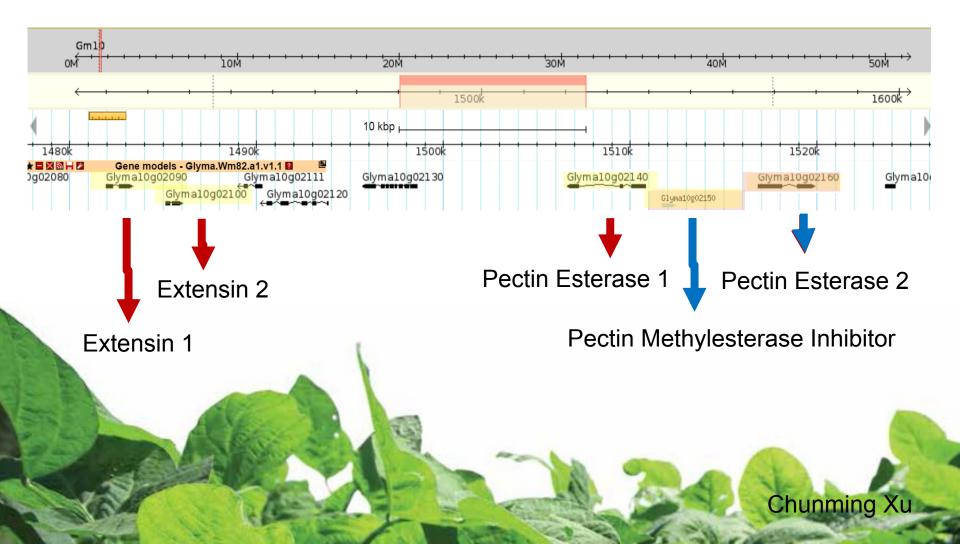


84/93 polymorphisms in these genesInvolved in cell walls



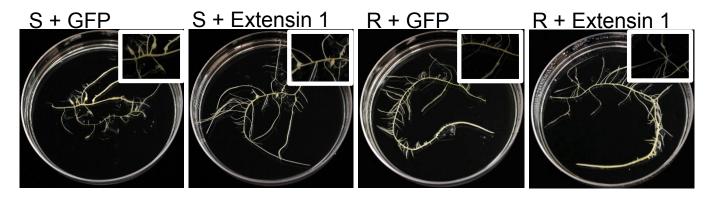
4 Candidate genes

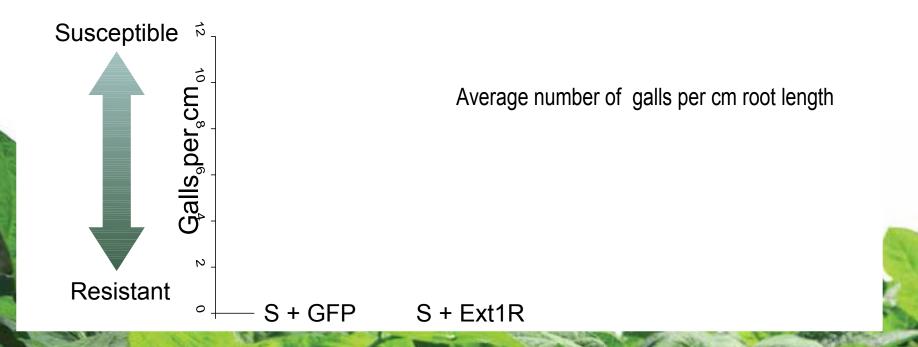




Overexpression of Extensin 1R

Enhanced resistance in both genotypes

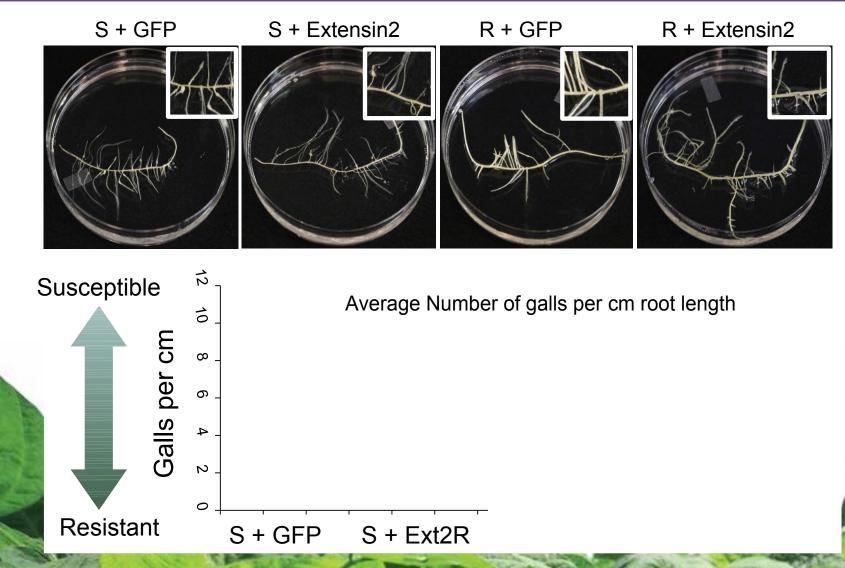




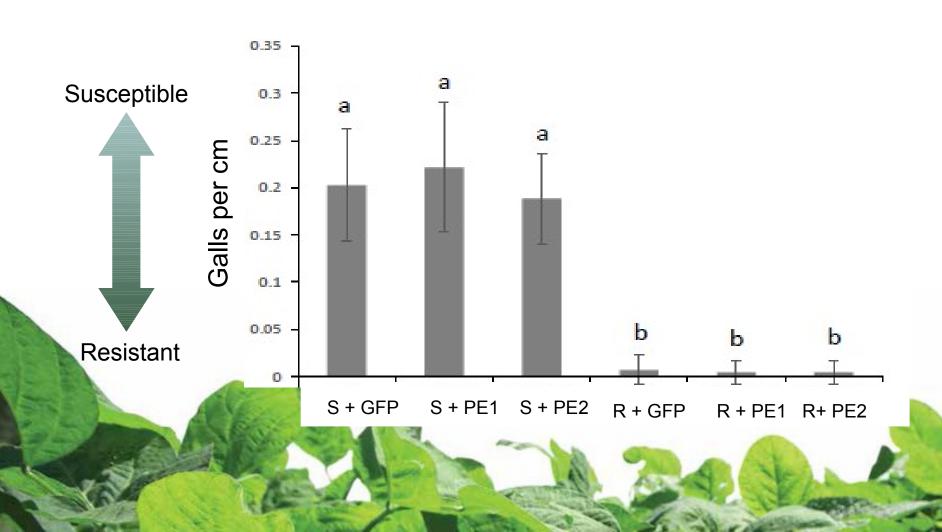
Overexpression of Extensin 2R



Enhanced resistance in both genotypes



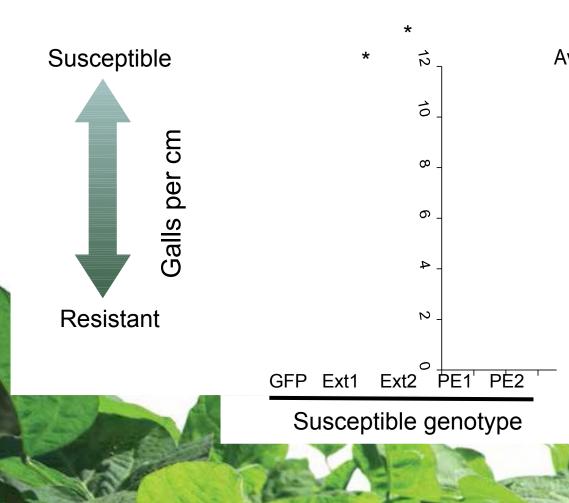
OE of Pectin Esterases from R No effect in either genotype



Knock out 4 genes individually

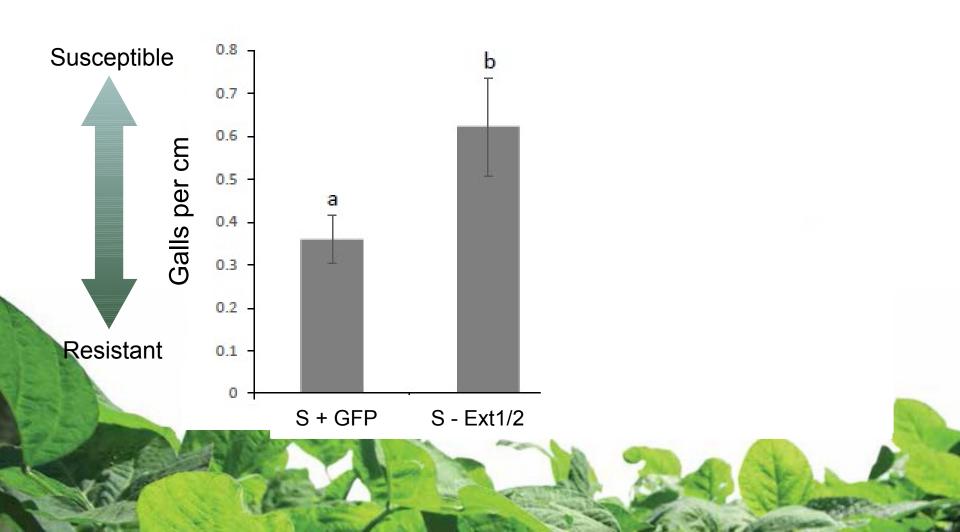


Small effects of Extensin/ no effects of pectin esterase



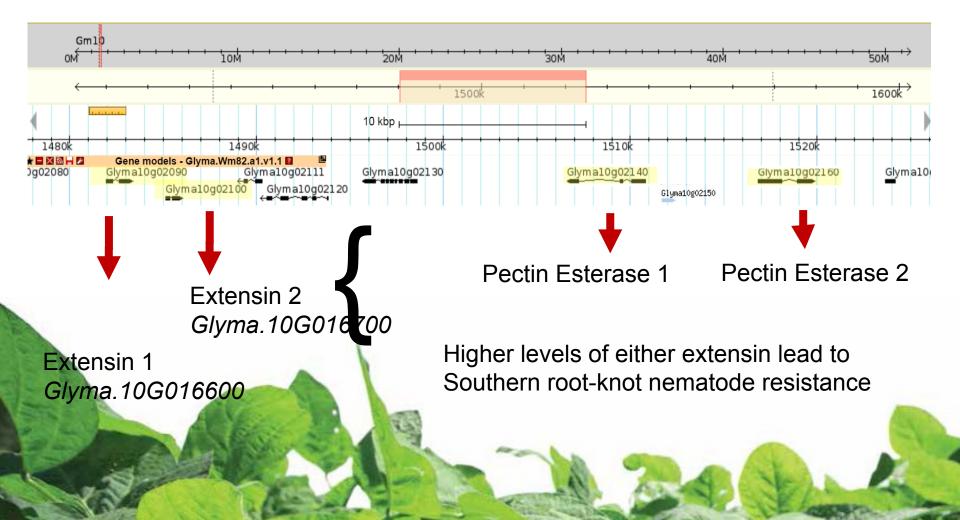
Average number of galls per cm root length

Double extensin knockouts Susceptibility increases in both genotypes



Four candidate genes





Extensin 2 promoter Relative to Williams 82



Resistance associated with expression level Related to promoter sequence

| Consensus Identity C+ 1. Bossier C+ 2. Williams C+ 3. Pl | | | | 1,100 1,200 1,300 1,400 | 1,507 |
|--|---------|------|------------|-------------------------|-------|
| | | SNPs | Insertions | Deletions | |
| | Bossier | 2 | 0 | 3 | |
| | PI96354 | 19 | 7 | 10 | |
| | | | | | |

So what is it?





Pollen Ole e I allergen Gene family, expressed in various tissues Is it really an allergen?

80mer Sliding Window Search Results

| Database | AllergenOnline Database v16 (January 27, 2016) | | |
|-------------------------------|---|--|--|
| Input Query | >query atgaaaatggcctatttccatagtacctgctttctggtttttatgcttatcgtatcactg ttggtccccggcagaggcggtgactacggtcccgaagaagaaaaacttctatcaaaaact ataggcattcaaggcattgtctattgcaaatctgcctctaaactcaccccacttgaaggg gccttgacaaggatatcatgcgaggctgttgatgaatatgggtttgaaacgacaccgttt tctttctta | | |
| Length | 249 | | |
| Number of 80 mers | 170 | | |
| Number of Sequences with hits | 0 | | |

No Matches of Greater than 35% Identity Found

So what is it?



²²² Previously cloned in soybean

DOI: 10.1007/s10535-012-0258-0

BIOLOGIA PLANTARUM 57 (1): 85-90, 2013

GmPOI gene encoding a Pollen_Ole_e_I conserved domain is involved in response of soybean to various stresses

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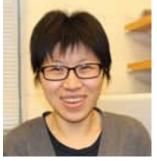
Associated with drought tolerance

- Upregulated in roots during stress
- Transgenics have higher drought tolerance
 - ☎ Higher SOD levels
 - Higher proline levels

Acknowledgements









Peter LaFayette Shi Li

