



Ultra-low Gossypol Cottonseed



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Global Cotton Lint Production in 2011 = 26.1 MMT

Global Cottonseed Production in 2011 = 48.8 MMT (~11 MMT Protein)

Can meet the protein requirements of 600 million people/year (50 g/day)



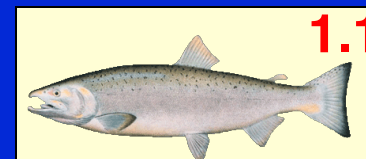
Presence of toxic gossypol prevents the use of cottonseed as food & also limits its use as feed

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Feed Conversion Ratio (FCR) for Various Animals



Animal	FCR	Reference
Beef	5.8 ^a	Klopfenstein et al., 1991
Pig	3.3 ^a	Losinger, 1998; Cromwell, 1991
Duck	2.705 ^b	Shalev and Pasternak, 1989
Turkey	2.102 ^b	Shalev and Pasternak, 1989
Chicken	2.047 ^b	Shalev and Pasternak, 1989
Shrimp	2.0 ^c	Tacon, 2002
Tilapia	1.5 ^d	McGinty and Rakocy, 1989; Boyd et al., 2005
Salmon	1.1 ^d	Villamar, 2002
Channel Catfish	1.0 ^d	Lovell, 1990

^aFeed grain/weight gain ratio

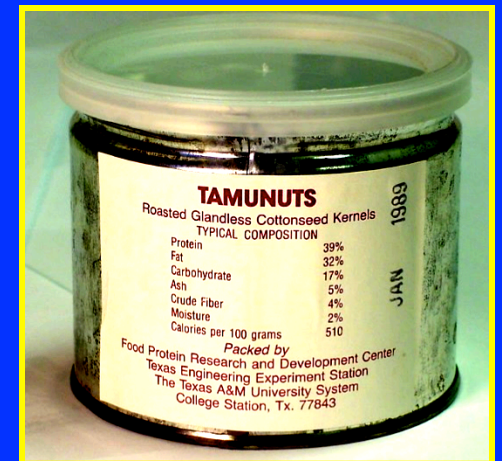
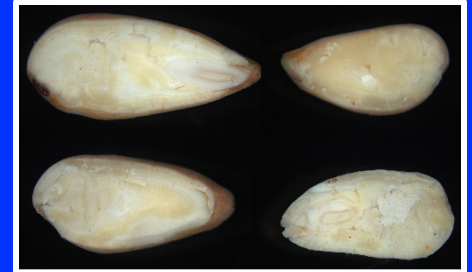
^bTotal feed utilization of live weight for seven week-old broilers

^cDry aquafeed/weight gain ratio

^dFeed/weight gain ratio

Attempts Using Plant Breeding

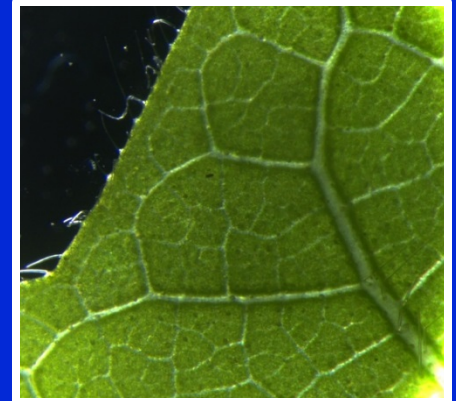
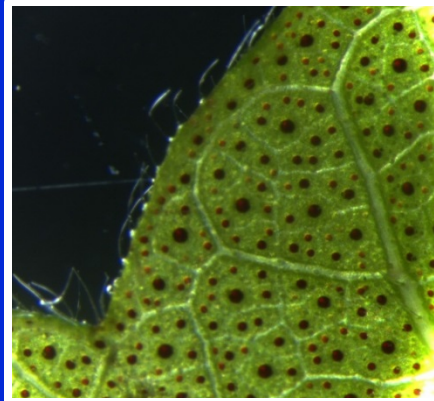
- McMichael discovered a **GLANDLESS** mutant of cotton in 1954 (was grown by Hopi Indians of Arizona)
- Seed from glandless cotton was gossypol-free
- National and International breeding programs launched to transfer the trait to commercial varieties
- Animal nutrition studies showed glandless cottonseed to be a relatively good source of feed for pigs, chicken, and shrimp
- Glandless cottonseed was even considered fit for human consumption



Other Human Nutrition Studies

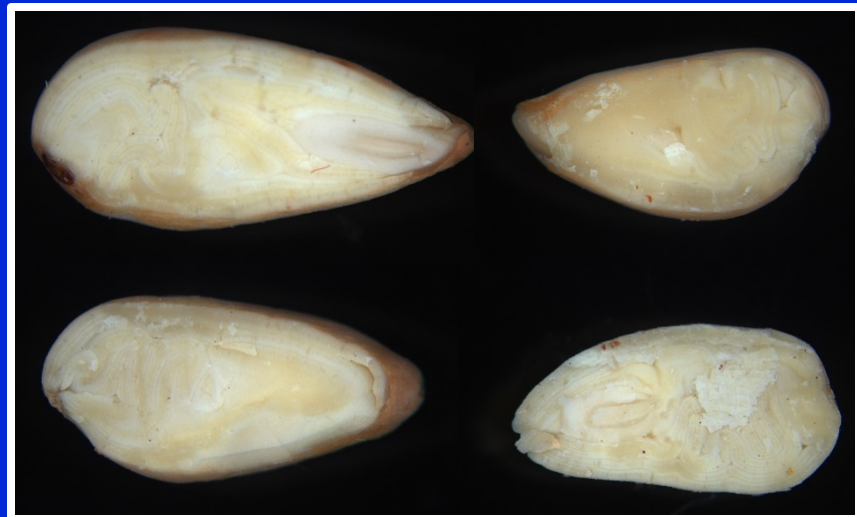
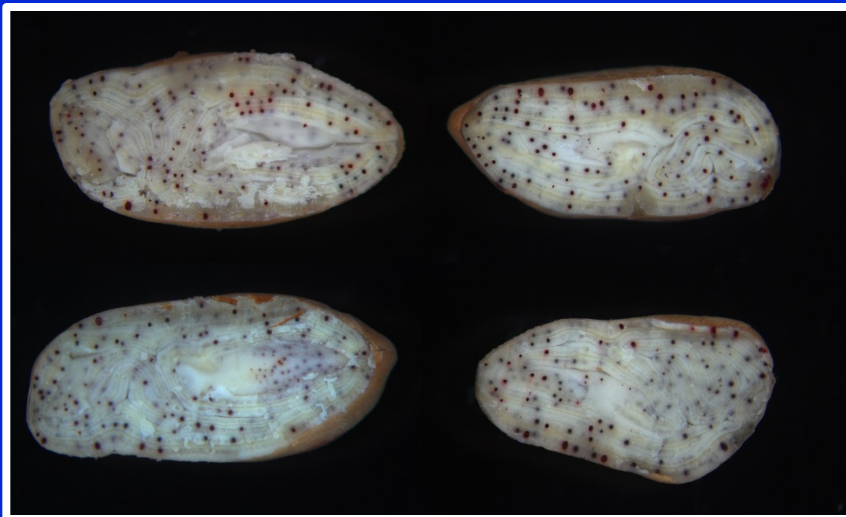
- INCAPARINA containing 38% CSF used in human nutrition trials (INCAP - a cooperative International Institute established by Central American countries and Panama in the early 1950s)
 - Favorable acceptability and no adverse effects in 1-5 year-old children
 - Satisfied protein requirement of weanling children
- CSF + millet flour blends tested in West African countries
 - Blends provided adequate protein for normal as well as malnourished infants and children
- Texas Woman's University studies (Betty Alford)
 - CSM found safe and nutritious for children
 - Cottonseed protein in the form of liquid diets or baked products maintained nitrogen balance with no change in nutrition status in young, adult women
- Texas Tech University
 - Tortillas made with 20-25% CSF improved Lysine and Tryptophan levels as well as leucine-isoleucine ratio
 - Improved shelf-life and good acceptability

CSM: Cottonseed Meal; CSF: Cottonseed Flour



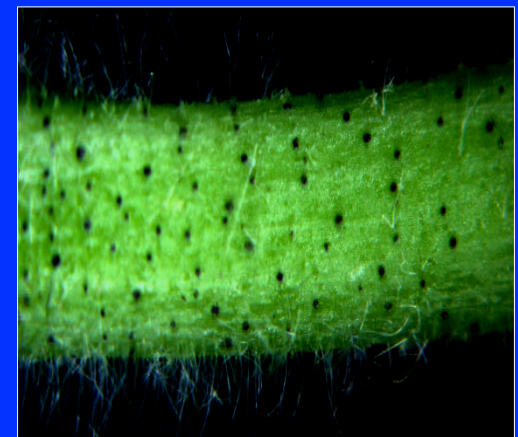
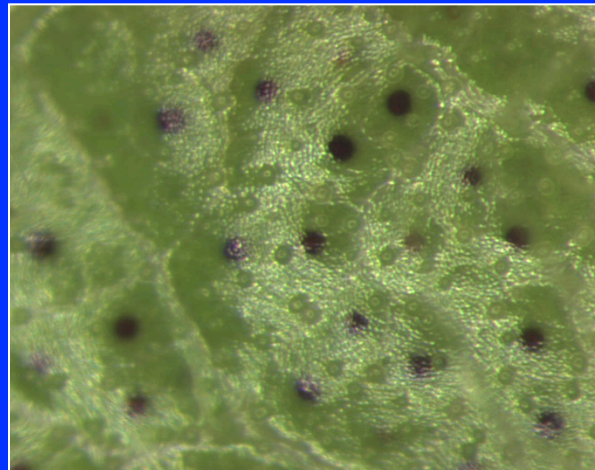
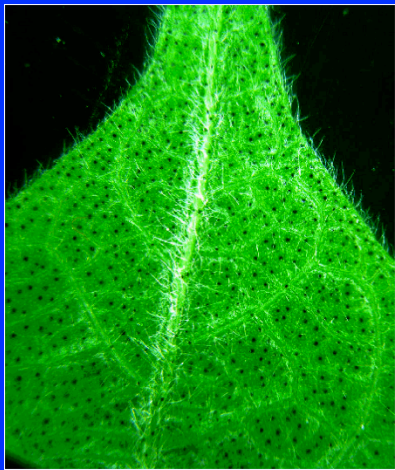
Wild-type 'Glanded' Cotton

Mutant 'Glandless' Cotton



Gossypol and other related terpenoid aldehydes present in the glands in leaves, stem, and floral parts confer resistance to insect pests

These terpenoids are also induced in response to pathogens and serve as phytoalexins



Traditional Insect Pests that Cause Greater Damage to Glandless Cotton

Cotton Bollworm (*Helicoverpa zea*)

Tobacco Budworm (*Heliothis virescens*)

Beet Armyworm (*Spodoptera exigua*)

Non-traditional Insects that Damage Glandless Cotton

Black Fleahopper (*Spanogonicus albofasciatus*)

Pillbug (*Porcellia sp.*)

Spotted Cucumber Beetle (*Diabrotica undecimpunctata*)

Grape Colaspis (*Maecolaspis flava*)

Green Dock Beetle (*Gastrophysa cyanea*)

Blister Beetle (*Epicauta vittata*)

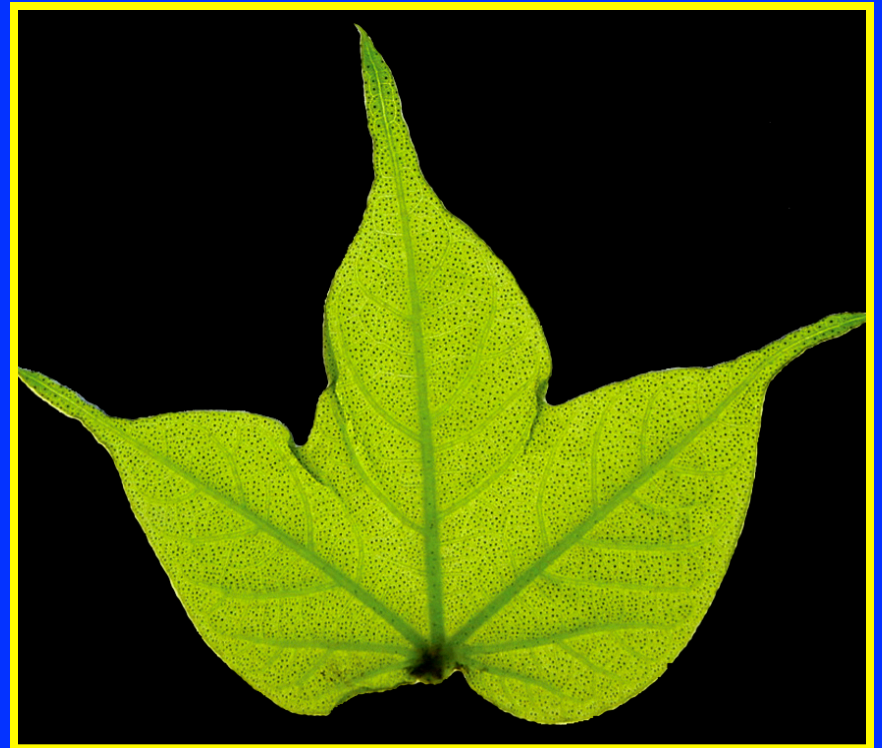
- Currently, there is no large-scale cultivation of glandless cotton.

Desired Trait

Seeds:
Gossypol < 450 ppm



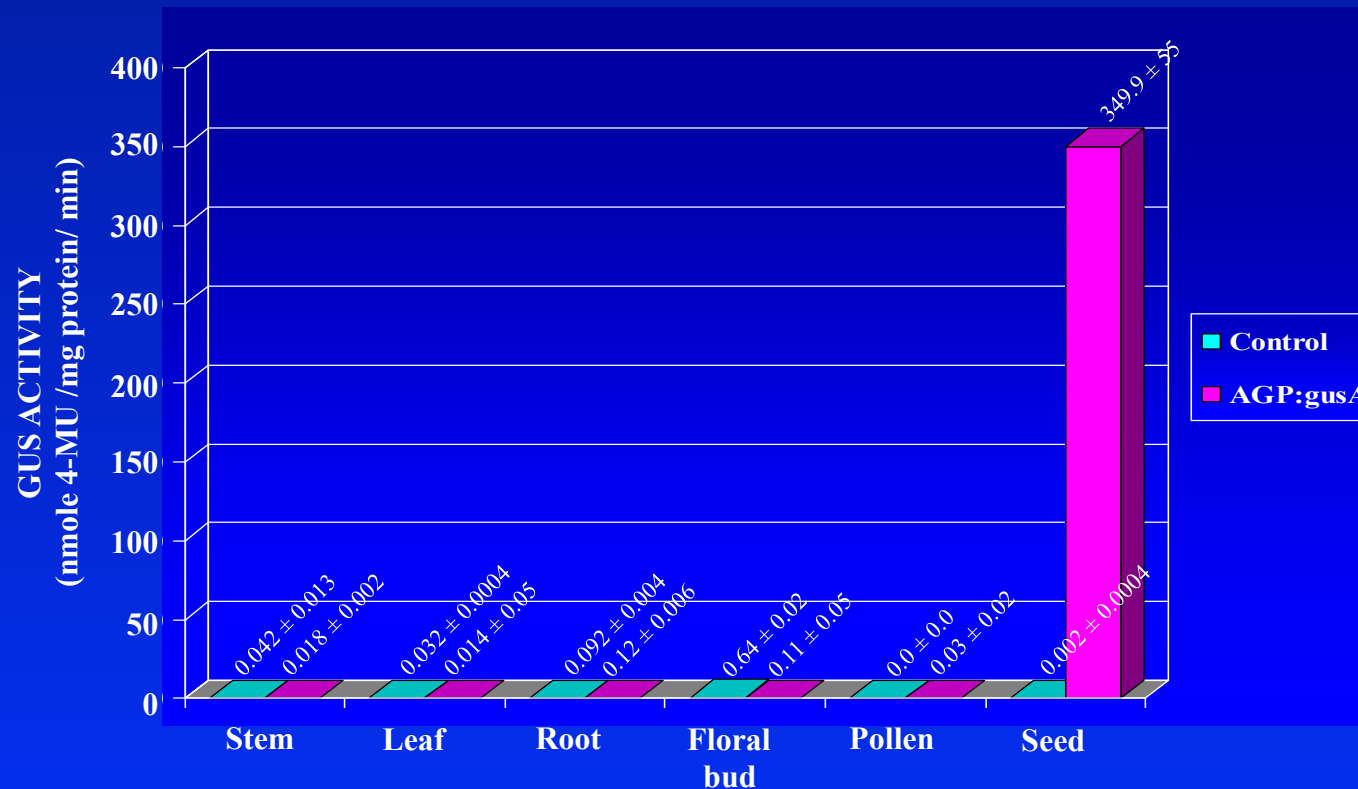
Leaf & other Organs:
Full Compliment of Gossypol
and Related Terpenoids
for Defense



**Is it possible to Use Seed-specific, RNAi-mediated Gene Silencing
to generate Ultra-low Gossypol Cottonseed (ULGCS)
without Compromising Plants Defense Capabilities???**

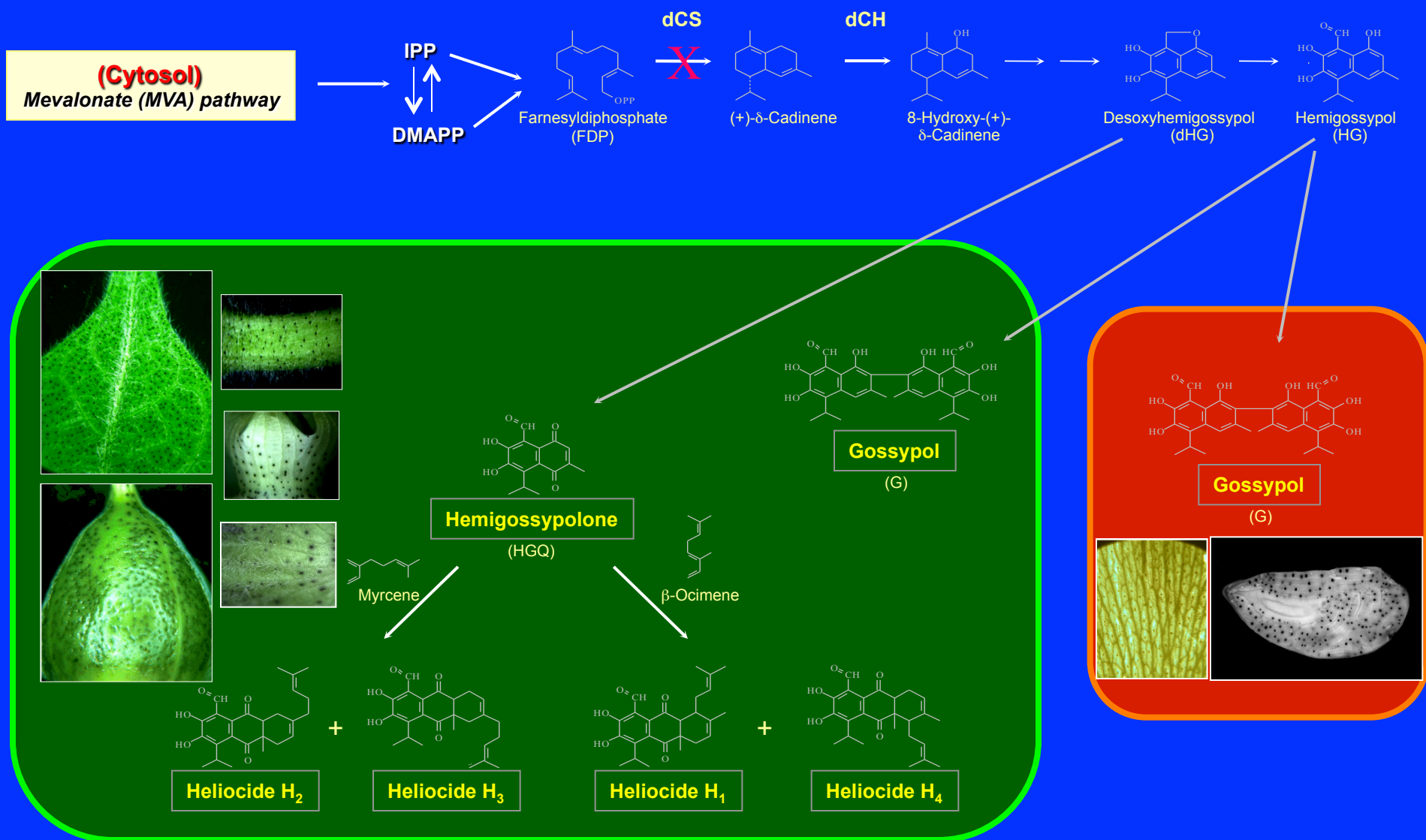
- **Efficient Transformation System for Cotton**
- **δ -cadinene Synthase Gene**
- **Seed-specific, α -globulin Promoter**
- **A Gene Silencing Mechanism (RNAi)**

Seed-specific expression of AGP::*gusA* in cotton



Sunilkumar Rathore (2002)
Transgenic Research 11: 347-359
U.S. Patent #7,626,081

Biosynthetic Pathway for Gossypol and Related Terpenoids



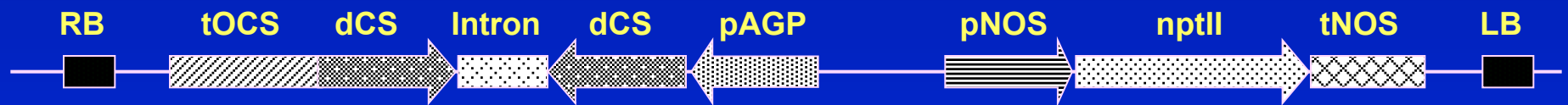
Selection of trigger region for gene silencing

5'  δ-cadinene synthase 3' 2050 bp

Trigger region  604 bp

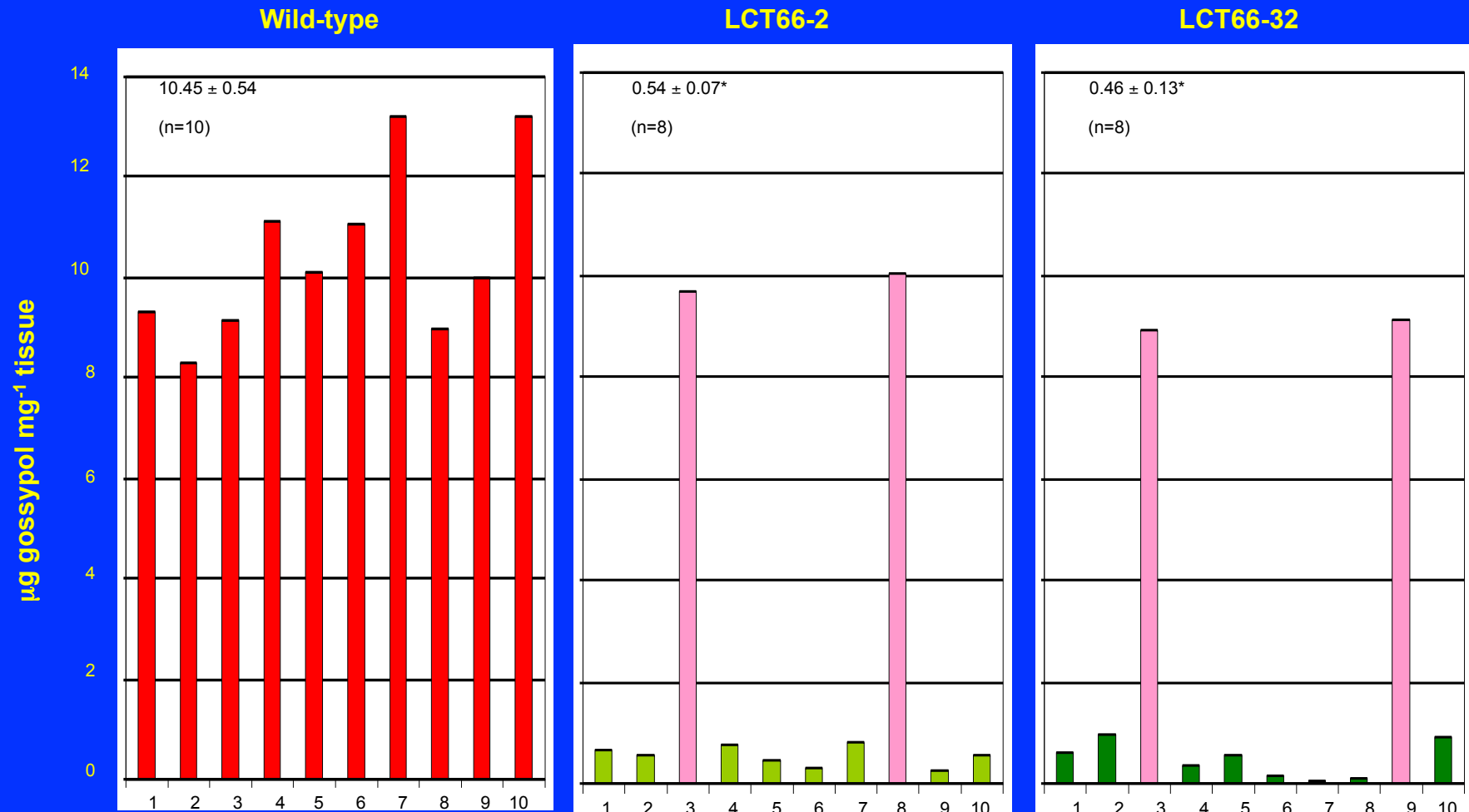
δ-cadinene synthase gene	Genbank accession no.	Plant source	Homology with the trigger sequence (%)	Homology with Complete Coding sequence (%)
Cad1-C14 (XC14)	U23205	<i>G. arboreum</i>	99.8	99.8 (1661/1665)
Cdn1-C4	AF270425	<i>G. hirsutum</i>	98.8	98.0 (1631/1665)
Cdn1	U88318	<i>G. hirsutum</i>	98.5	97.6 (1625/1665)
Cad1-C2	Y16432	<i>G. arboreum</i>	96.4	95.9 (1597/1665)
Cad1-C3	AF174294	<i>G. arboreum</i>	96.2	96.1 (1467/1527)
Cad1-C1 (XC1)	U23206	<i>G. arboreum</i>	96.0	96.0 (1599/1665)
Cad1-A	X96429	<i>G. arboreum</i>	80.9	81.4 (1348/1656)

Hairpin RNAi construct to silence δ -cadinene synthase gene



- ❖ Used pHannibal and pART27 vector systems developed by Waterhouse, CSIRO [Wesley *et al.* (2001) Plant J. 27: 581-590]
- ❖ *Agrobacterium* strain - LBA4404

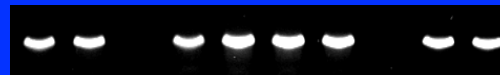
Gossypol Levels in Individual Wild-type and T1 Cottonseeds



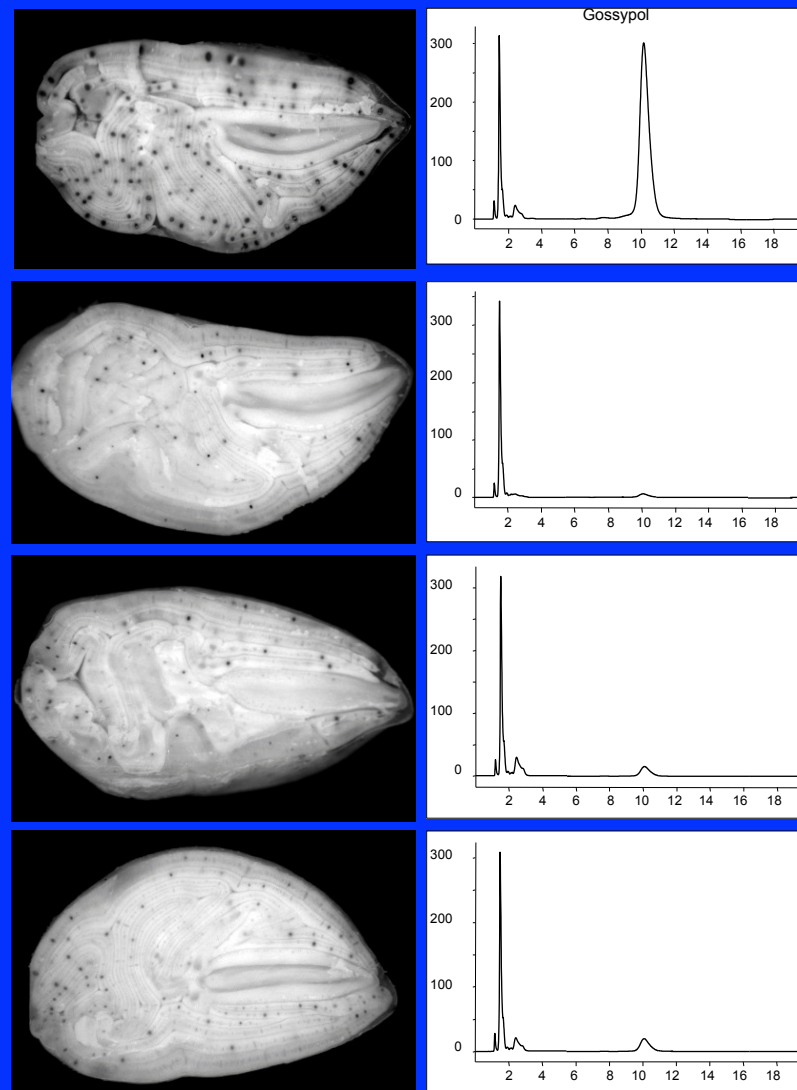
Sunilkumar Rathore (2006)

PNAS 103:18054-18059

U.S. Patent #7999148



T1 Cottonseeds from Line LCT66-32 and Individual HPLC Chromatograms for each Showing Gossypol Peaks

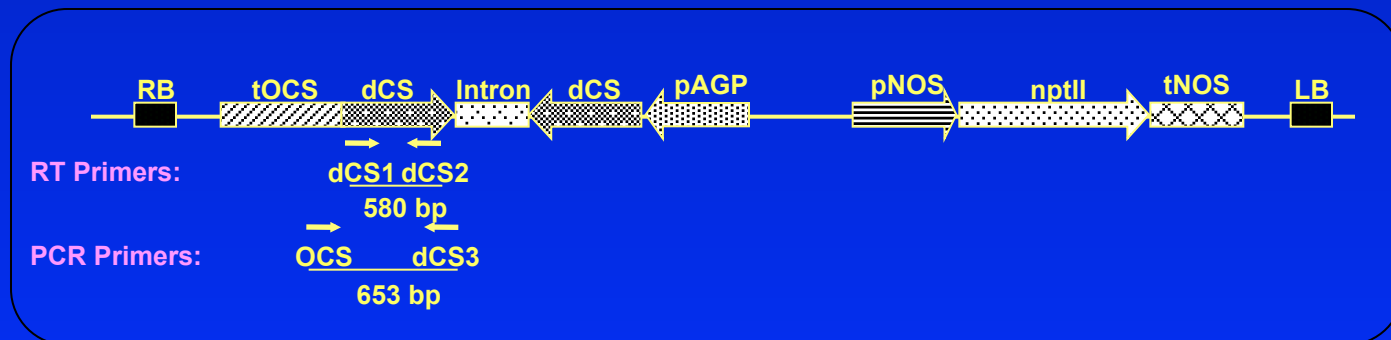
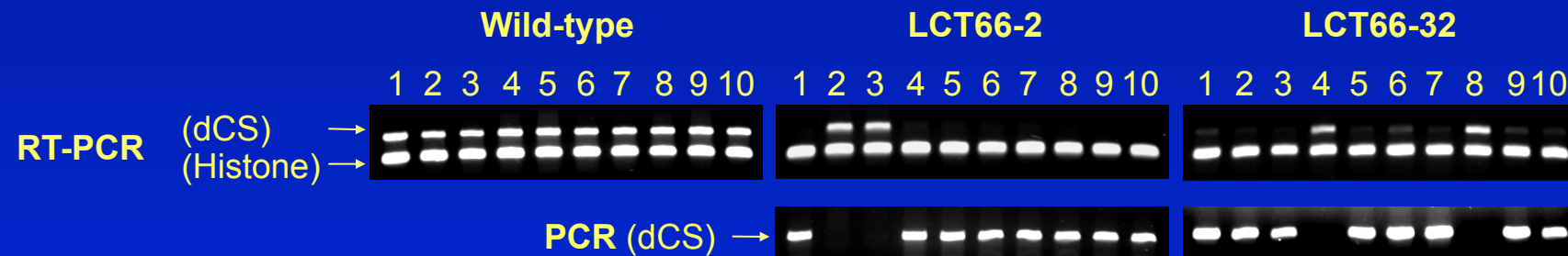


Sunilkumar Rathore (2006)

PNAS 103:18054-18059

U.S. Patent #7999148

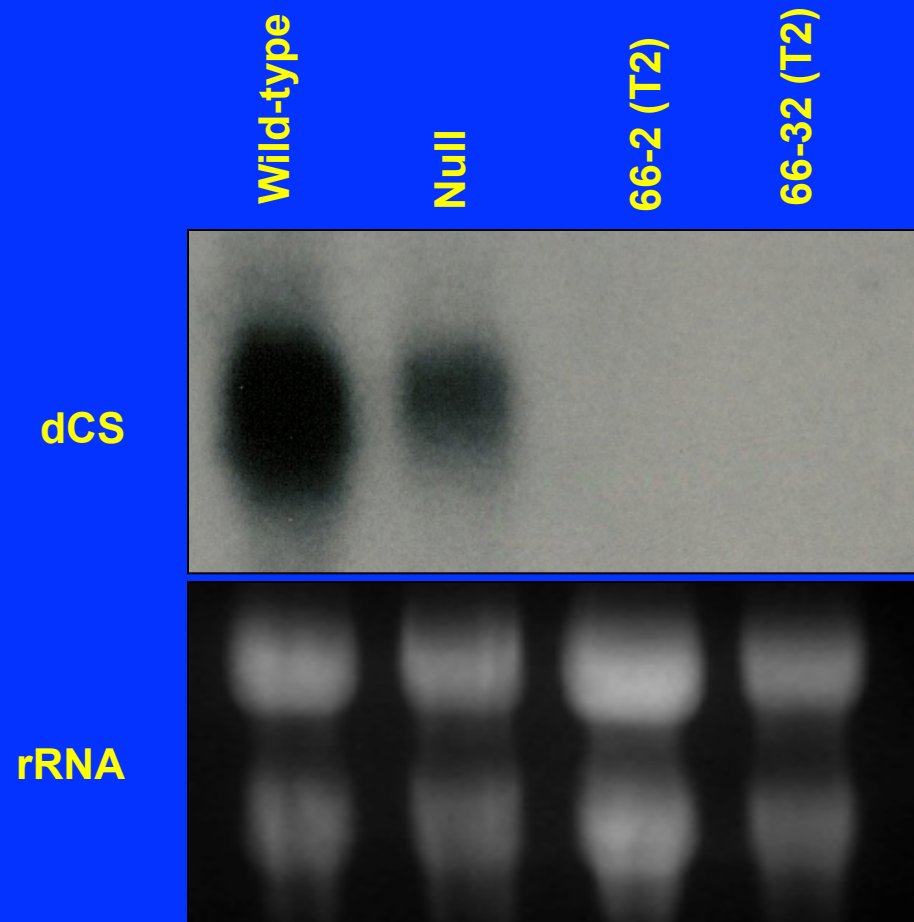
Levels of δ -cadinene Synthase Transcripts in Individual, Wild-type and T1 Transgenic Embryos at 35 dpa



Sunilkumar Rathore (2006)

PNAS 103:18054-18059

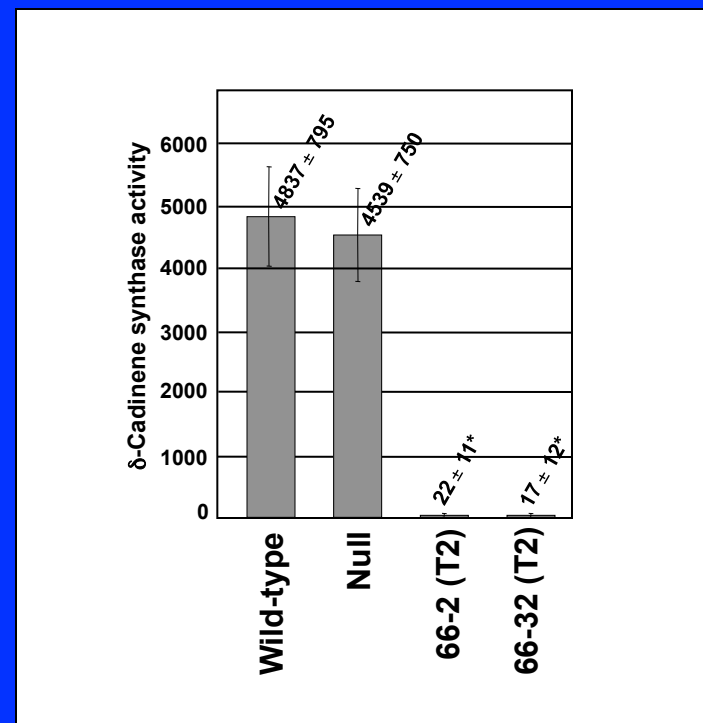
δ -Cadinene Synthase Transcripts in Embryos (35 dpa) from Wild-type, Null Segregant and Transgenic T1 Plants



Sunilkumar Rathore (2006)

PNAS 103:18054-18059

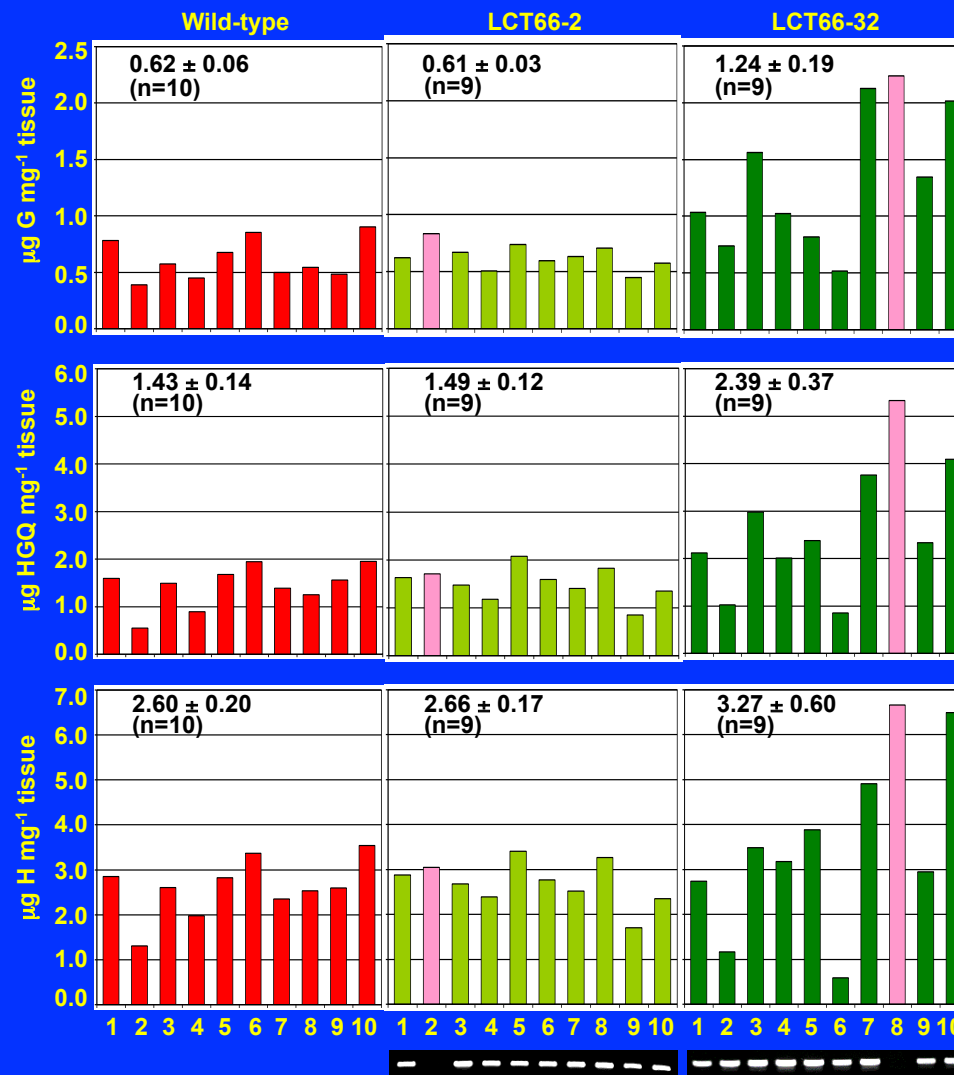
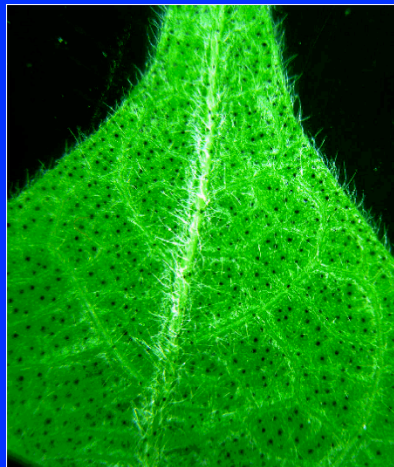
δ -Cadinene Synthase Activities in Embryos (35 dpa) from Wild-type, Null Segregant and Transgenic T1 Plants



Sunilkumar Rathore (2006)

PNAS 103:18054-18059

Levels of Gossypol, Hemigossypolone, and Heliocides in Leaf Tissues from Individual Wild-type and Transgenic T1 Progeny Plants

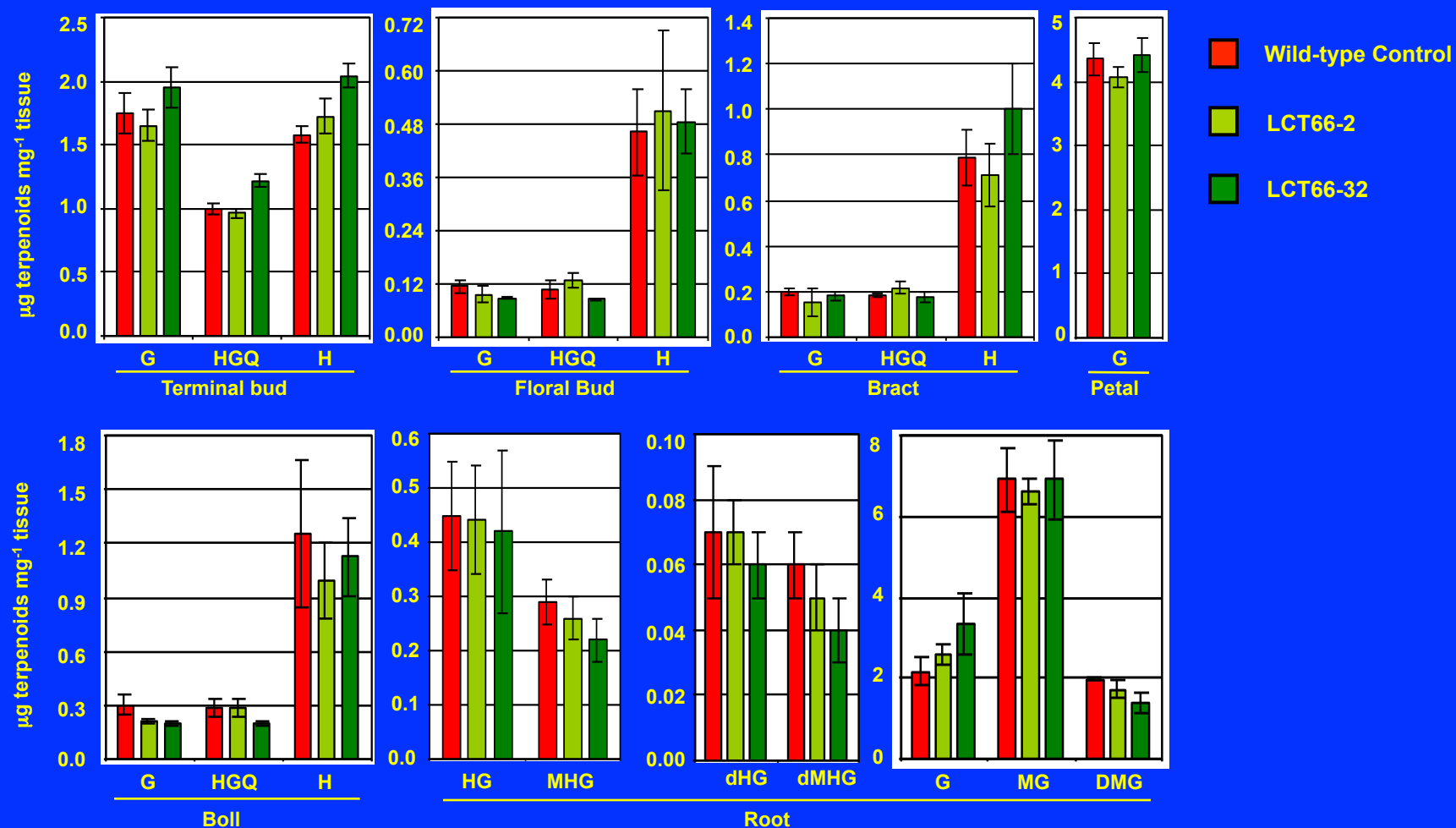


Sunilkumar Rathore (2006)

PNAS 103:18054-18059

U.S. Patent #7999148

Levels of Gossypol and Related Terpenoids in Terminal Bud, Floral Bud, Bract, Petal, Boll, and Root Tissues



ULGCS trait is completely tissue specific

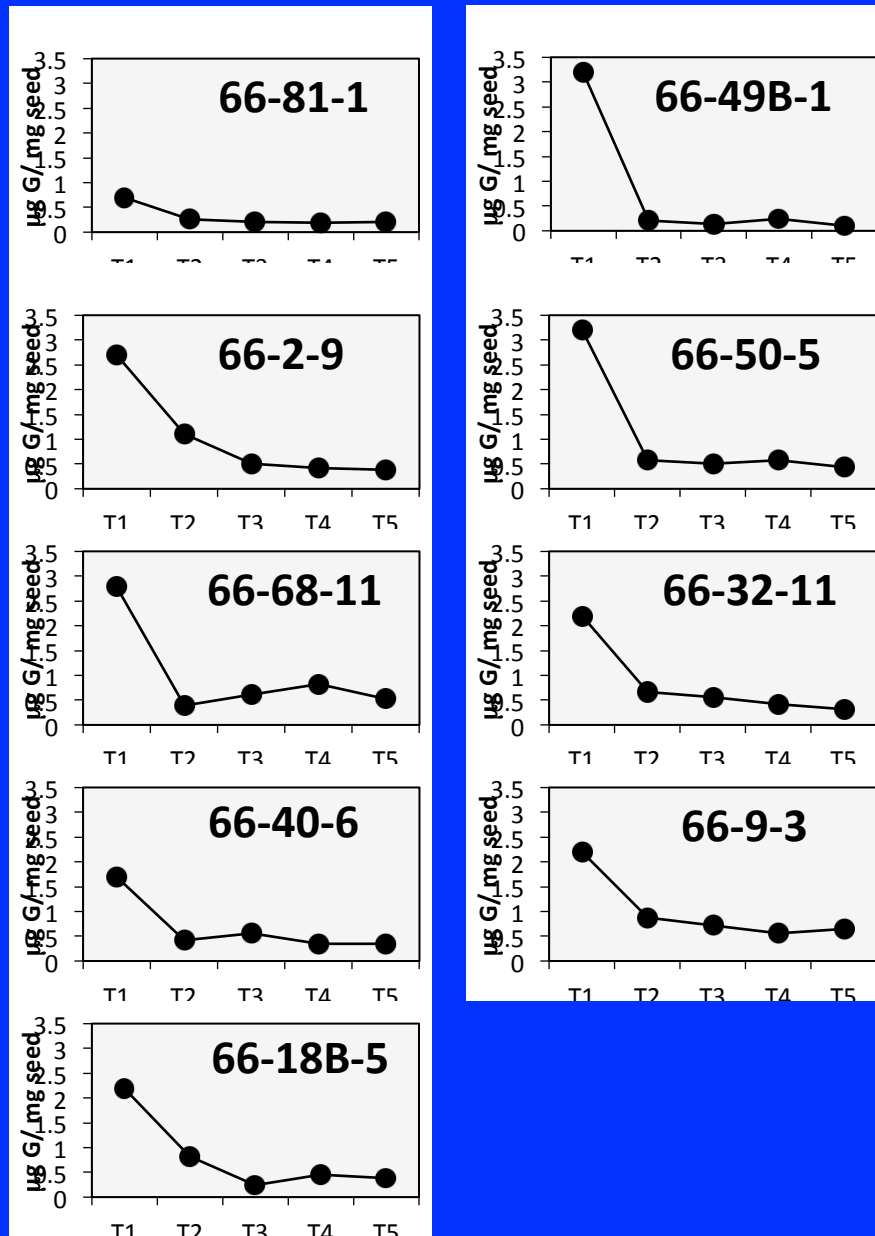
Sunilkumar Rathore (2006)

PNAS 103:18054-18059

U.S. Patent #7999148

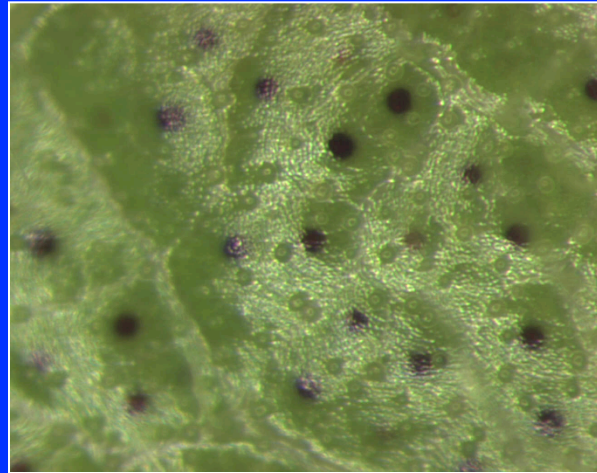
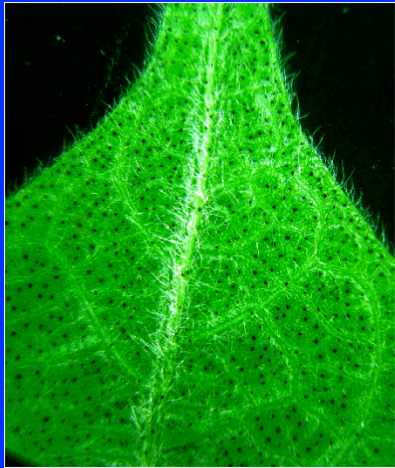
- Is the RNAi-mediated ULGCS trait stable?

Levels of gossypol (G) in T1, T2, T3, T4, and T5 seeds obtained from greenhouse-grown cotton plants



RNAi-mediated, ULGCS Trait is Stable and Heritable

Gossypol and related terpenoids are also induced in response to pathogens and serve as phytoalexins



- Are the very young seedlings from RNAi lines more susceptible to cotton diseases?
- Are the very young seedlings from RNAi lines capable of launching terpenoid-mediated defense response when challenged with a pathogen?

Cotton seedlings after five days of growth on *Rhizoctonia solani*-infested soil



Coker 312



(Glandless) GL



49B-1-1-1



49B-1-2-1



81-5-5-6



81-5-2-10

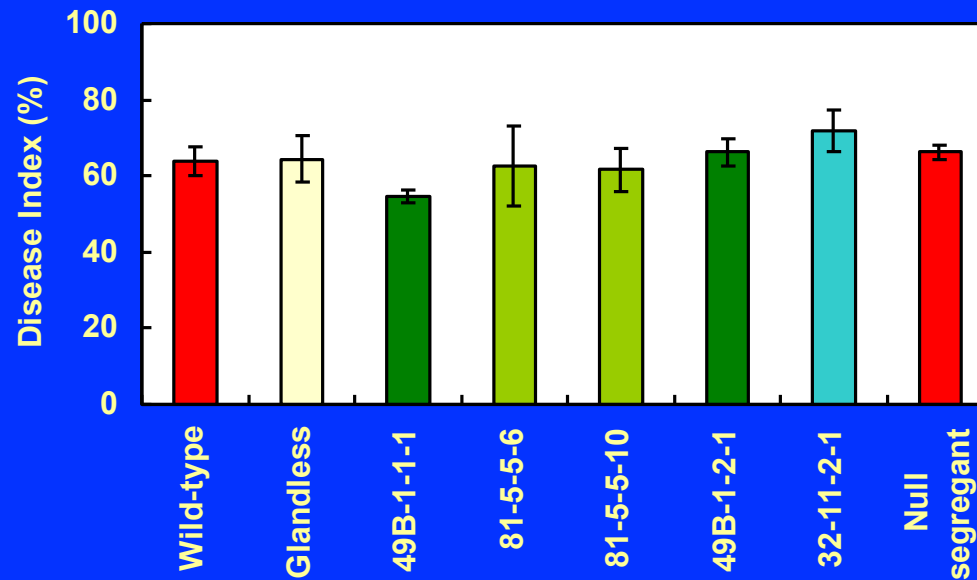


(Null segregant) 32/14

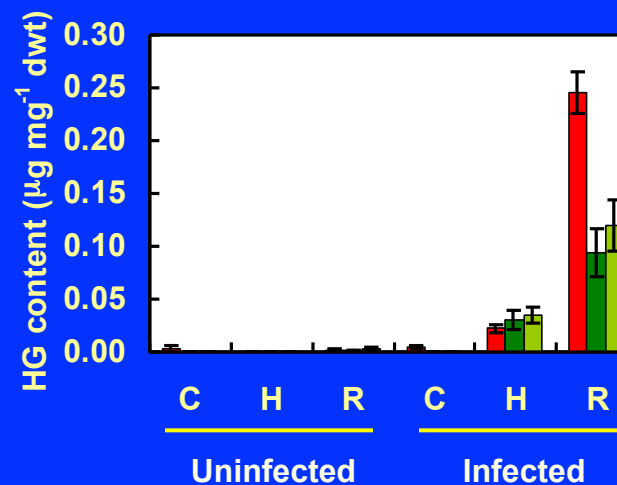
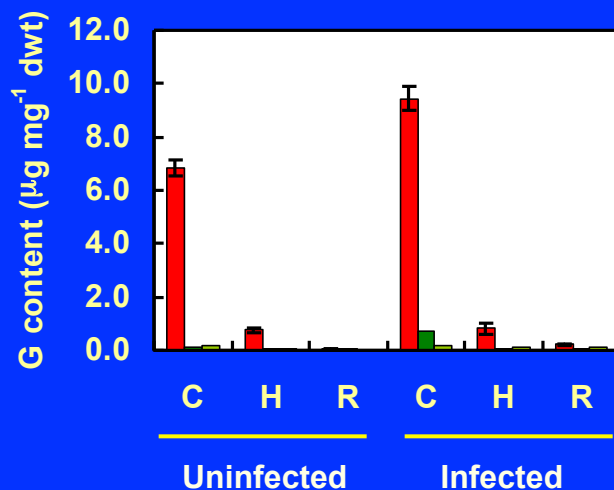
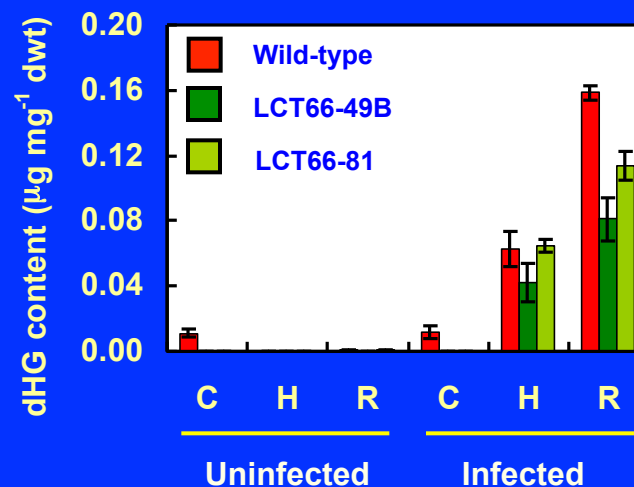
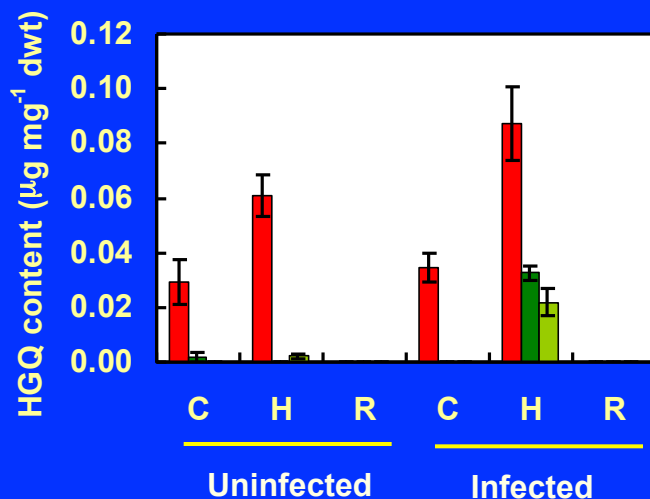


32-11-2-1

Susceptibility of the RNAi lines to *Rhizoctonia solani* is not greater than that of the wild-type plants



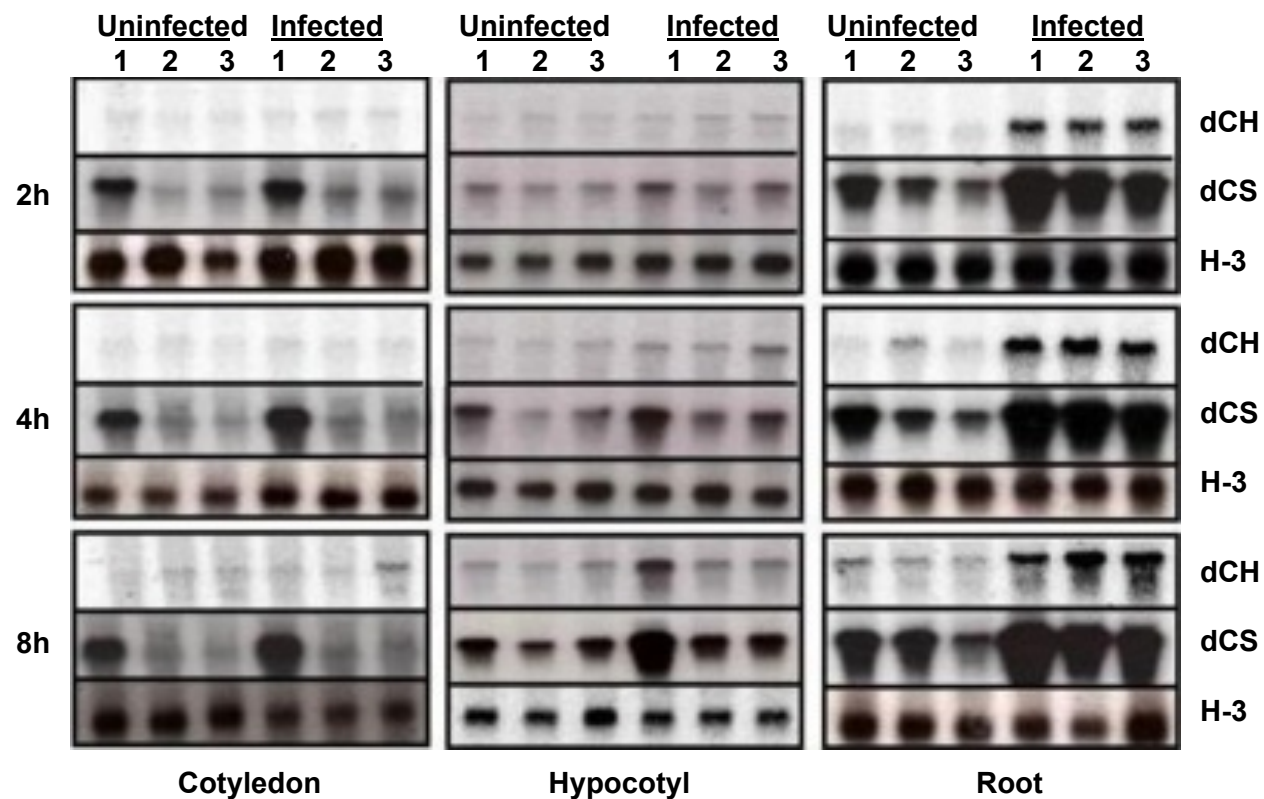
RNAi lines are Capable of Launching Terpenoid-based Defense in Response to *Rhizoctonia solani* infection of Roots



R: Root; H: Hypocotyl; C: Cotyledon

HGQ: Hemigossypolone; G: Gossypol; DHG: Desoxyhemigossypol; HG: Hemigossypol

δ -Cadinene Synthase (dCS) Transcripts are Induced in the Roots and Hypocotyl of RNAi Lines in Response to a Challenge from *Rhizoctonia solani*



1 = Wild type 2 = 66-81 3 = 66-49B

dCH: δ -cadinene hydroxylase

dCS: δ -cadinene synthase

H-3: Histone

- Young ULGCS seedlings are capable of launching terpenoid-based defense response
- δ -cadinene synthase gene is induced in ULGCS seedlings in response to a pathogen challenge
- The basic defense mechanism remains functional in the ULGCS seedlings

**Do RNAi lines maintain the ULGCS
trait under field conditions?**

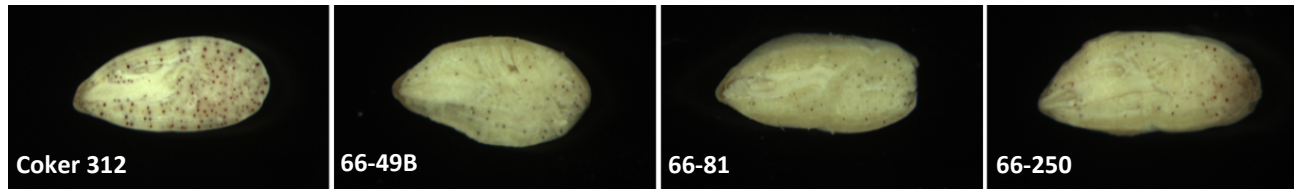
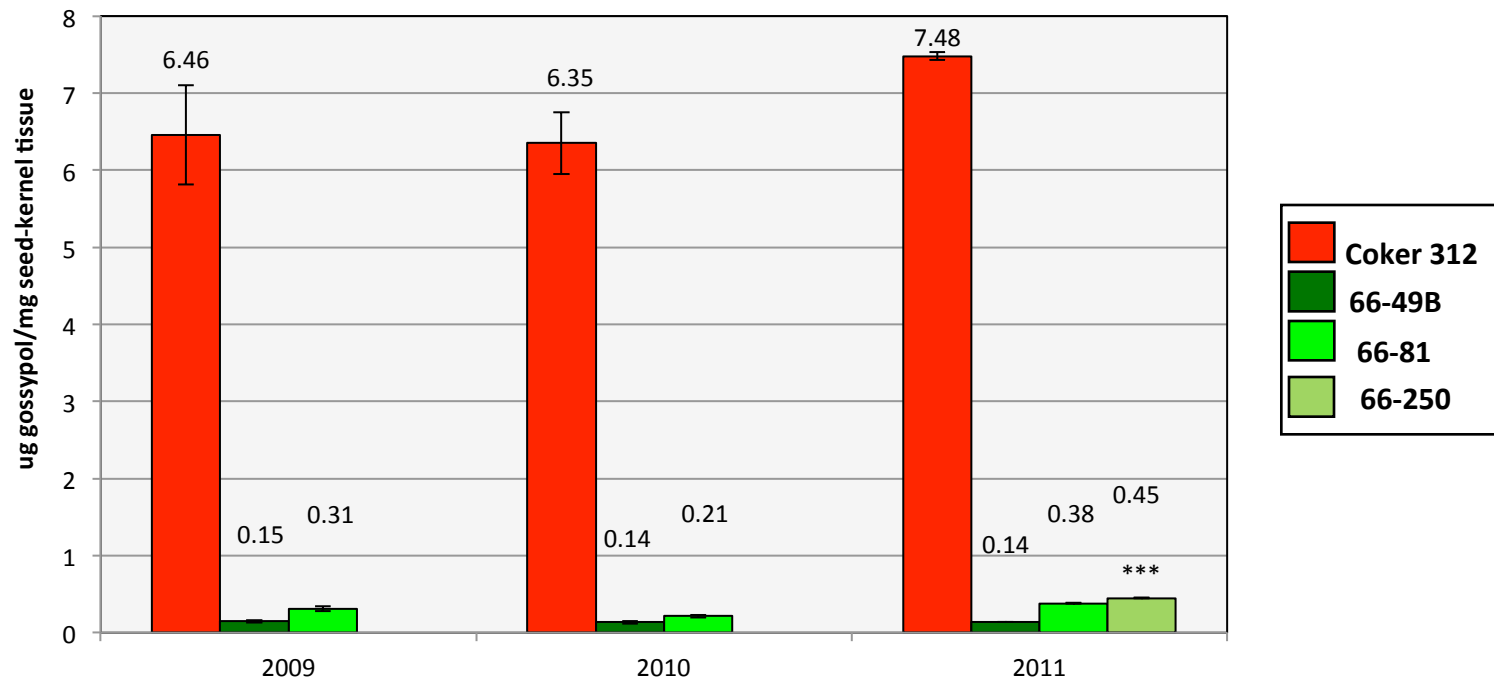
**Do RNAi lines maintain normal levels of
defensive terpenoids in the non-seed parts of the plant
under field conditions?**

Ultra-low Gossypol Cottonseed (ULGCS) – Field Trials **2009, 2010 & 2011**



- **ULGCS Stability**
- **ULGCS Specificity**
- **Fiber/seed Yield & Quality**

Seed gossypol content of field-grown plants



ULGCS Trait is stable under field conditions

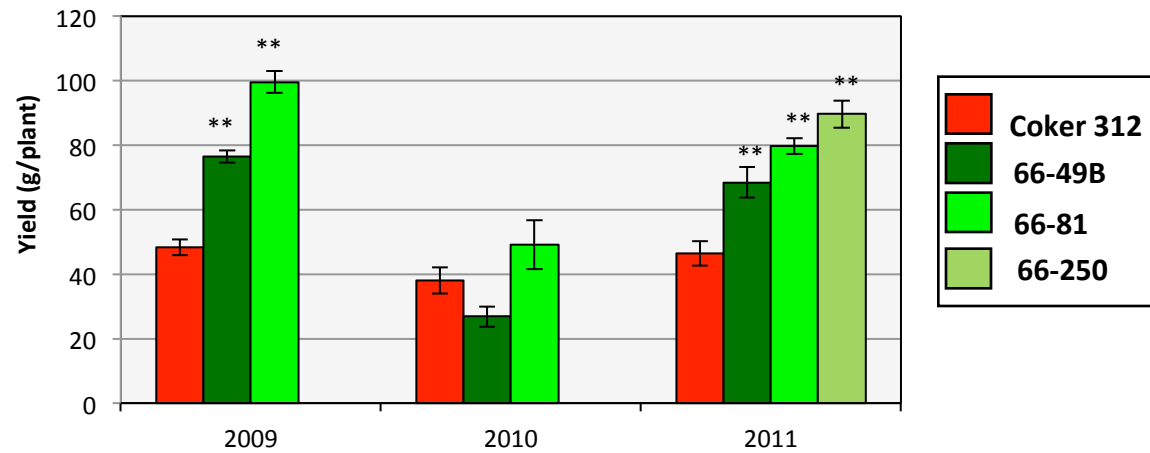
Levels ($\mu\text{g}/\text{mg}$ dry weight) of gossypol and related terpenoids

ULGCS Foliage & Floral Tissues Retain Normal Levels of Terpenoids

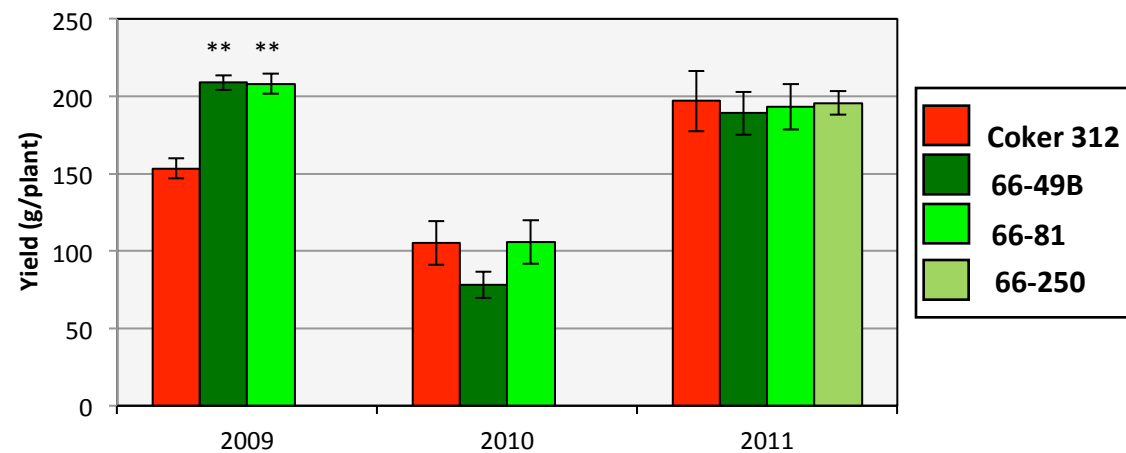
		2009			2010			2011			
		Coker 312	66-49B	66-81	Coker 312	66-49B	66-81	Coker 312	66-49B	66-81	66-250
Bracts	G	0.08 (± 0.008)	0.06 (± 0.003)	0.11 (± 0.011)	0.11 (± 0.01)	0.10 (± 0.03)	0.12 (± 0.02)	0.08 (± 0.00)	0.06 (± 0.00)	0.16 (± 0.02)	0.11 (± 0.00)
	HGQ	0.15 (± 0.02)	0.10 (± 0.012)	0.10 (± 0.014)	0.17 (± 0.04)	0.12 (± 0.04)	0.11 (± 0.03)	0.25 (± 0.04)	0.24 (± 0.02)	0.32 (± 0.08)	0.38 (± 0.02)
	H1-H4	1.25 (± 0.05)	0.89 (± 0.06)	1.05 (± 0.10)	1.78 (± 0.32)	1.59 (± 0.46)	1.44 (± 0.32)	1.51 (± 0.25)	1.07 (± 0.59)	1.75 (± 0.47)	1.82 (± 0.09)
	G	5.65 (± 0.42)	5.04 (± 0.39)	6.41 (± 0.91)	3.15 (± 0.28)	3.14 (± 0.43)	3.47 (± 0.19)	3.09 (± 0.10)	3.17 (± 0.05)	4.01 (± 0.08)	3.93 (± 0.07)
Floral buds	HGQ	1.11 (± 0.14)	0.96 (± 0.11)	0.88 (± 0.23)	0.45 (± 0.09)	0.36 (± 0.06)	0.40 (± 0.07)	0.51 (± 0.06)	0.61 (± 0.09)	0.66 (± 0.02)	0.87 (± 0.04)
	H1-H4	2.39 (± 0.09)	1.99 (± 0.22)	1.90 (± 0.40)	1.47 (± 0.32)	1.46 (± 0.19)	0.90 (± 0.03)	1.89 (± 0.13)	2.07 (± 0.19)	2.30 (± 0.13)	2.40 (± 0.12)
	G	1.04 (± 0.30)	0.99 (± 0.26)	0.82 (± 0.03)	1.02 (± 0.03)	0.98 (± 0.24)	1.15 (± 0.12)	0.92 (± 0.14)	1.15 (± 0.08)	1.32 (± 0.31)	1.17 (± 0.19)
Terminal part of axillary branch	HGQ	1.62 (± 0.16)	1.22 (± 0.19)	1.15 (± 0.17)	1.24 (± 0.06)	0.98 (± 0.05)	1.12 (± 0.09)	2.22 (± 0.21)	1.89 (± 0.19)	1.96 (± 0.27)	2.27 (± 0.12)
	H1-H4	2.84 (± 0.27)	1.77 (± 0.23)	2.43 (± 0.40)	1.64 (± 0.14)	1.74 (± 0.25)	1.94 (± 0.17)	3.19 (± 0.43)	2.70 (± 0.11)	2.91 (± 0.21)	3.67 (± 0.16)
	G	0.34 (± 0.08)	0.19 (± 0.02)	0.48 (± 0.13)	0.80 (± 0.01)	0.43 (± 0.04)	0.97 (± 0.04)	0.65 (± 0.09)	0.62 (± 0.04)	1.12 (± 0.12)	0.63 (± 0.01)
Leaves	HGQ	1.66 (± 0.49)	0.78 (± 0.06)	1.37 (± 0.37)	2.25 (± 0.23)	1.25 (± 0.09)	2.09 (± 0.08)	4.04 (± 0.40)	3.73 (± 0.30)	4.38 (± 0.18)	3.57 (± 0.15)
	H1-H4	2.68 (± 0.36)	1.77 (± 0.23)	2.46 (± 0.76)	1.81 (± 0.38)	0.91 (± 0.07)	1.14 (± 0.09)	2.94 (± 0.25)	2.77 (± 0.59)	3.51 (± 0.47)	2.57 (± 0.11)
	G	1.69 (± 0.09)	1.37 (± 0.08)	1.88 (± 0.07)	2.02 (± 0.24)	1.22 (± 0.05)	1.47 (± 0.06)	1.06 (± 0.06)	0.86 (± 0.02)	1.04 (± 0.06)	0.89 (± 0.01)
2 day bolls	HGQ	7.29 (± 0.05)	6.43 (± 0.23)	6.58 (± 0.41)	4.29 (± 0.28)	3.69 (± 0.24)	3.05 (± 0.05)	4.95 (± 0.41)	4.57 (± 0.26)	3.98 (± 0.25)	4.32 (± 0.03)
	H1-H4	11.03 (± 0.14)	9.05 (± 0.54)	7.64 (± 0.39)	11.27 (± 3.58)	9.96 (± 1.88)	5.81 (± 1.31)	8.66 (± 0.73)	8.74 (± 0.54)	5.51 (± 0.60)	7.62 (± 0.38)
Petals	G	5.70 (± 0.15)	6.28 (± 0.22)	6.25 (± 0.25)	4.77 (± 0.26)	5.92 (± 0.13)	5.97 (± 0.05)	3.60 (± 0.19)	3.78 (± 0.30)	4.15 (± 0.18)	4.22 (± 0.14)

Yield data for field-grown plants

Fiber

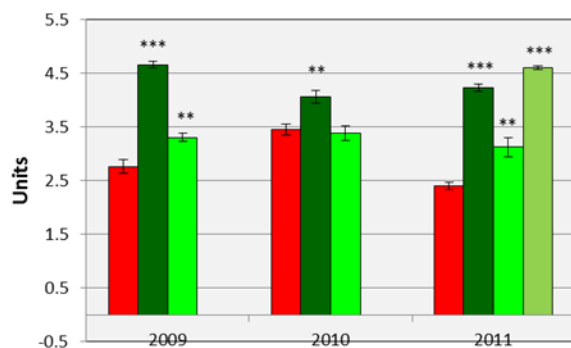


Seed

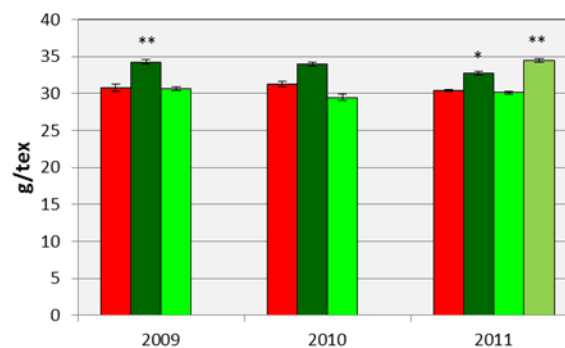


Fiber Quality under Field Conditions

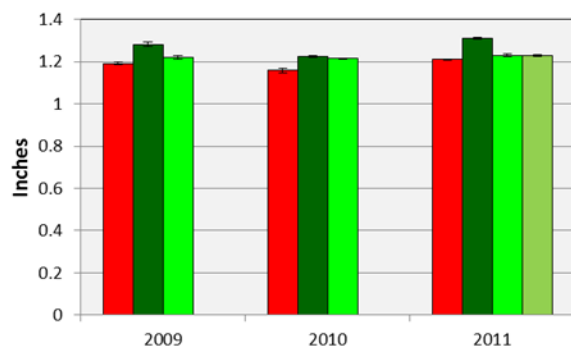
MIC



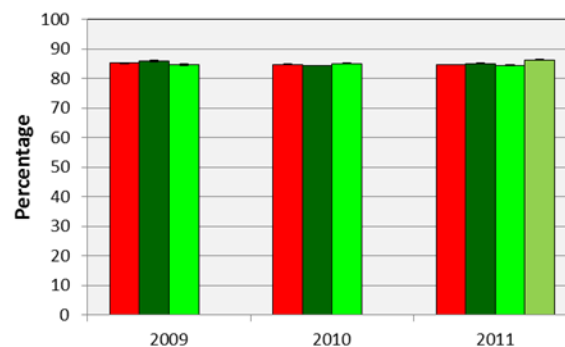
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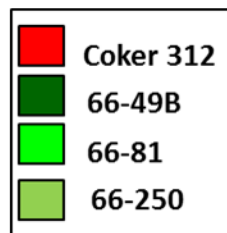
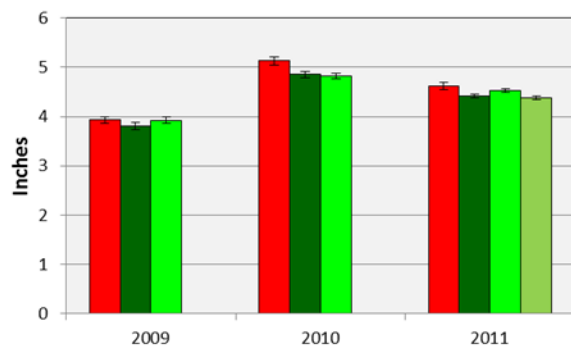
UHM



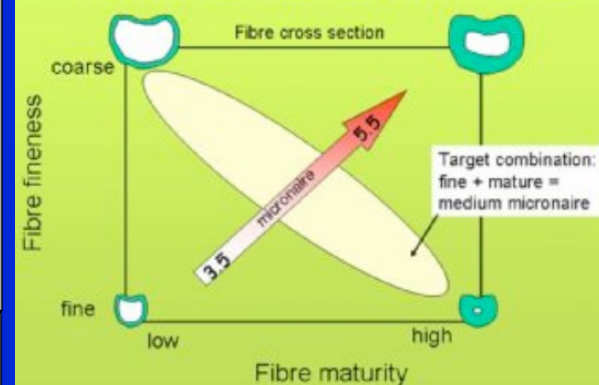
UI



ELO

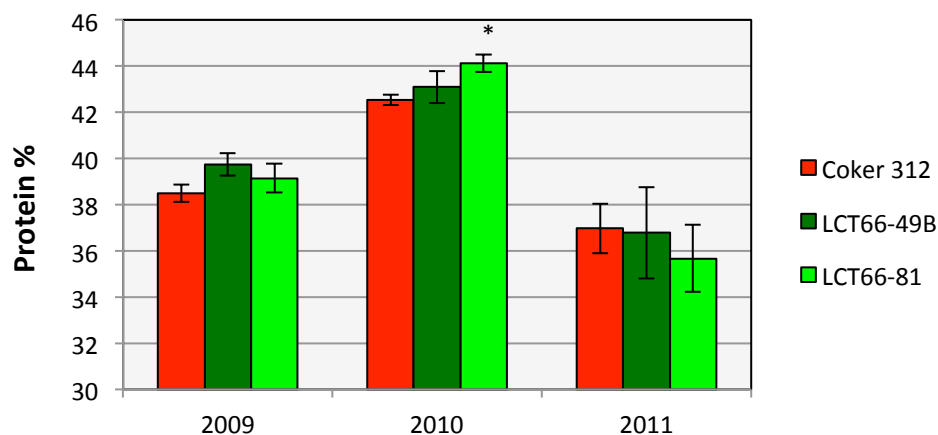


Micronaire = maturity x fineness

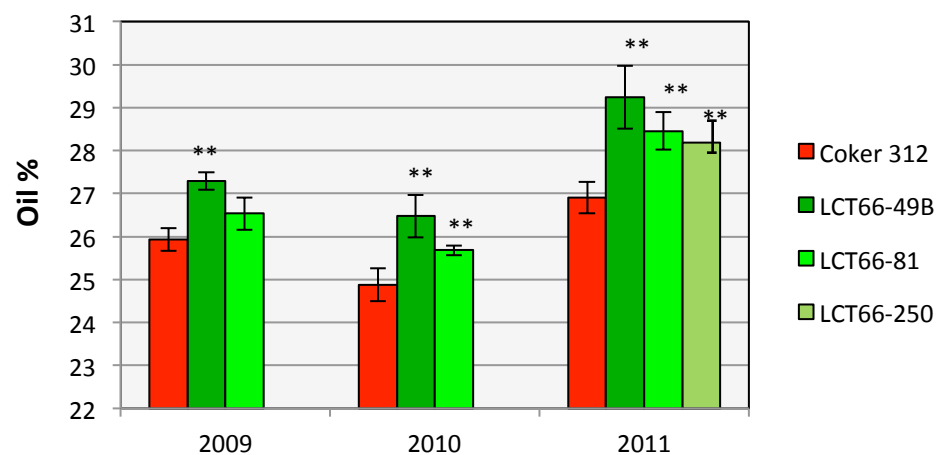


Protein and Oil levels in seeds of plants grown under field conditions

Seed Protein Content



Seed Oil Content



Note that the protein values were measured in seed kernels, while oil content was determined in whole seeds.

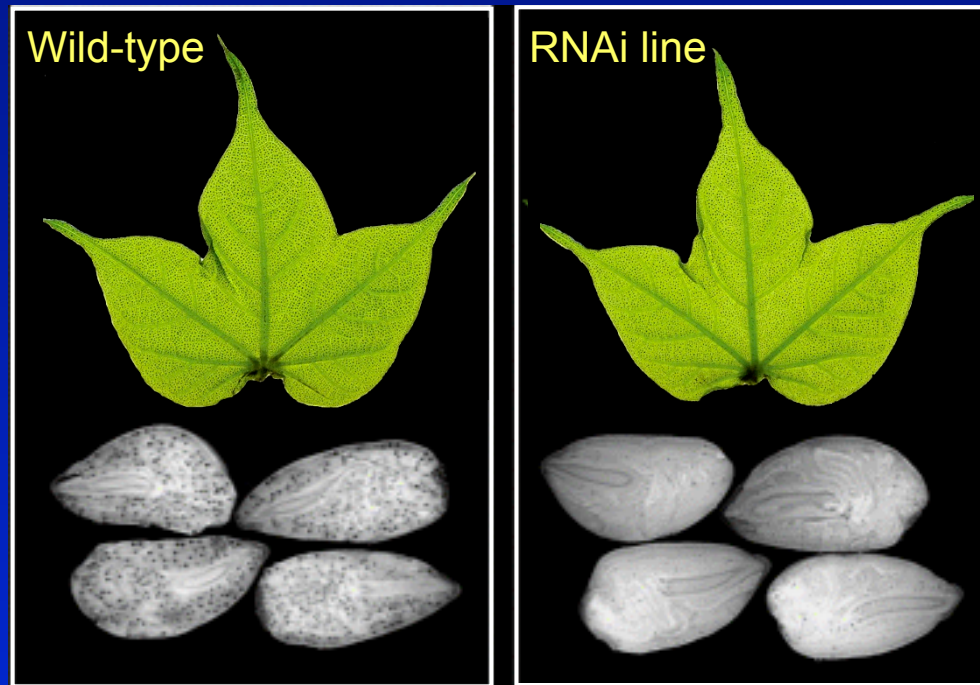
*Data represent mean $\pm$ SE, * $P < 0.05$; ** $P < 0.01$, $n = 3$.*

Compositional Analysis of Acid-delinted Cottonseed from Field (Dry Weight Basis)

Component (%)	LCT66-49B	LCT66-81	Coker 312	Acala GLS	DP 50
Moisture (fresh weight basis)	8.40	9.64	9.05	8.18	5.42
Protein	29.6	29.8	29.5	27.4	21.4
Total Fat	20.5	22.0	20.1	25.3	23.0
Ash	4.05	4.49	4.72	4.35	4.60
Carbohydrates	45.9	43.7	45.7	42.9	51.0
Phytic acid	1.80	1.75	2.08	2.04	1.85
Free gossypol (%)	0.0388	0.0284	0.671	0.00942	0.830
Total gossypol (%)	0.0479	0.0400	0.750	0.0150	0.996

Oil content (NMR)	24.69 ± 0.10	23.45 ± 0.17	22.89 ± 0.16
Total gossypol (% , HPLC, kernels)	0.0147 ± 0.0064	0.0312 ± 0.0026	0.6462 ± 0.0337

- **ULGCS trait is stable under field conditions**
- **Fiber- and seed-yields are not negatively impacted in the ULGCS lines**
- **Fiber-quality is not negatively impacted in the ULGCS lines**
- **With the exception of ultra-low levels of gossypol, no other major differences were observed in terms of seed composition**



- ❖ Some of the RNAi lines have seed-gossypol levels less than 200 ppm (FDA: < 450 ppm and FAO/WHO: < 600 ppm).
- ❖ Non-seed tissues DO NOT show reduction in the levels of gossypol and other defensive terpenoids. Upon germination, newly formed tissues of the plantlet produce normal levels of terpenoids.
- ❖ Trait is heritable (T1 –T8 generations) and stable under field conditions.
- ❖ Pathogen-induced terpenoid biosynthesis pathway remains functional and the RNAi lines do not show higher susceptibility to pathogens or insects.

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