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	df	Lint percentage	100 seed weight	Boll weight	Yield
1988	2	3.4850	0.1052	0.4007	1425813.284
1989	2	19.1771	0.4847	1.4670	2506028.659
1990	4	8.7835	0.9112	0.7788	549043.712
1991	9	22.0957	2.3763	1.3129	986179.092
1992	18	3.9470	1.1352	0.3033	353806.199
1993	45	5.5265	0.7716	0.4369	462285.610
1988-1992					
1988	2	24.7805	1.3981	1.3909	956400.719
1989	9	29.8224	2.6713	0.5850	321605.174
1990	17	24.9812	1.2833	0.4997	244179.410
1988-1992					
1988	2	9.7359	0.2652	0.1441	198046.516
1989	9	10.9025	2.1128	0.8385	340411.249
1990	13	15.1750	0.4328	0.1808	381871.797
1988-1992					
1988	2	3.2203	0.2019	0.2365	1494034.057
1989	9	7.8772	0.9269	0.5466	1009602.560



GENETIC ASSOCIATION AMONG YIELD AND FIBER TRAITS IN F₂ HYBRID COTTON
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Abstract

Commercial cultivars continue to occupy a portion of the cotton (*Gossypium hirsutum*) production areas in the United States and other countries. This study was conducted to determine the genetic relationship between traits and heritabilities characteristics in F₂ hybrid population. Sixty-four F₂ hybrids resulted from crosses of 4 commercial cultivars and 16 pest-resistant germplasm lines were evaluated for 5 fiber and 4 yielding traits in four environments at Mississippi State, MS. An additive-dominance model was employed for these traits with different genetic background in different environments. MINQUE(1) method was used for estimating genetic variance and covariance components and predicting the genetic correlation.

Results indicated that dominant variance was the major portion of the phenotypic variances for lint yield, lint percentage and boll size. A smaller proportion of additive variance for fiber traits and the significance on additive x environment variance components suggested that lack of substantial genetic variability for fiber traits within these F₂ hybrids. However, relatively high values of narrow-sense heritabilities for lint yield and yield component indicated that a sufficient proportion of additive genetic variance might be available in F₂ hybrids for effective selection. The estimates of the genetic and phenotypic coefficients of correlation between the pairs of characters, for the most part, seem to be of comparable magnitude. Fiber strength was positive-additively correlated with boll weight, and appeared dominant correlations between these traits with elongation and 2.5% boll length were also significant positive. Based on the analysis of heritability and genetic correlation, the indirect selection for yield improvement only by lint percentage were expected to be as effective as by lint yield itself for these F₂ populations; however, indirect selection for yield improvement by other traits, such as boll number, fiber elongation and micronaire, would be 66%, 79% and 65% less effective than selection for lint yield itself, respectively.



INTROGRESSION OF GLANDED-PLANT AND GLANDLESS-SEED TRAIT FROM *G. STURTIANUM* WILLIS INTO TETRAPLOID COTTON PLANTS.

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Abstract

Two trispecific synthetic allotetraploids have been obtained using the Australian diploid species *G. sturtianum*, the main cultivated amphidiploid, *G. hirsutum*, and two American wild diploid species: *G. thurberi* and *G. raimondii*. Observation in the progeny of these trispecific hybrids revealed the expression of the gossypol glands morphogenesis repressive mechanism of *G. sturtianum* in a rather high proportion of the BC₁ seeds (6 on 41). In these materials, the glandless-seed and glanded-plant trait seems to be linked to a lethal factor. Only one of the six totally glandless BC₁ seeds gave an adult plant. *In vitro* culture of the seed and grafting of the plantlet at a early stage on *G. hirsutum* were necessary to obtain a normal development of this genotype. All the other glandless materials died just after germination or never germinated. The survival plant will be used in a backcrossing programme to obtain the introgression of the glanded-plant and glandless-seed character into *G. hirsutum*.

Introduction

The introgression of the glandless-seed and glanded-plant character from the Australian wild diploid species into upland cotton would transform it in a true food crop whilst preserving one of its natural defence mechanism. Either bispecific or trispecific hybrids can be used to reach this goal (Mergeai, 1993). To date, the main efforts have concerned the exploitation of bispecific hybrids issued from direct crosses between *G. hirsutum* and different Australian species. Most of the hexaploids and pentaploids obtained from those initial crosses are characterised by lower seed gossypol contents than the cultivated parent and a normal gossypol concentration in the other parts of the plant (Muramoto, 1969; Dilday, 1986; Koto, 1989). The addition families produced by backcrossing the pentaploids with *G. hirsutum* show varying gossypol glands density in the seeds and leaves. The only glanded plants issued from glandless seeds produced by this way are multiple addition materials of *G. sturtianum* or *G. australe* on *G. hirsutum* (Altman *et al.*, 1987; Koto, 1989). No monosomic addition line express the glandless seed and glanded-plant trait (Koto, 1989; Rooney *et al.* 1991). Another way to achieve the introgression is to create trispecific hybrids combining *G. hirsutum* with an Australian species and either a D or a A diploid species. Shujing and Biling (1993) followed this last scheme. They crossed *G. bickii* with *G. arboreum* to obtain a sterile diploid hybrid whose fertility was partially restored by chromosome doubling. The resulting allotetraploid was then crossed with *G. hirsutum* to produce a fertile trispecific hybrid. The bispecific and trispecific allotetraploids showed differences in the expression of the glandless seeds-glanded plant trait. All plants of each hybrid showed glands in their aerial parts but only the kernels of the bispecific allotetraploid seeds were totally glandless. A similar observation was made by Mergeai (1992) on an (*G. arboreum* x *G. australe*) allotetraploid. All the seeds of the *bickii*-*arboreum*-*hirsutum* triple hybrid showed a few small and light coloured glands on the edges of the cotyledons. No details are available on the glands distribution of the seeds produced from the backcross of the *arboreum*-*bickii*-*hirsutum* trispecific hybrid. From the literature (Shujing and Biling, 1993) it appears that no plant expressing the glandless seed-glanded plant trait has been produced yet through this method. Cytogenetics observations on the trispecific hybrid showed an extremely low number of bivalents in Metaphase I. This result questions the actual possibilities to exploit AADC or AADG trispecific hybrids in an hybridization programme aiming at the introgression of the glanded-plant and glandless-seed trait into *G. hirsutum*. The phyletic distance between D and G genomes seems to be too large to allow the introgression of the concerned character in the D subgenome of *G. hirsutum*. Given the closer phyletic relations existing between A and C species, it is expected to obtain better results using ACDD trispecific hybrids. We decided therefore to develop this kind of material and to check its potentialities for the introgression of this character.

Material and methods

Two trispecific allotetraploid hybrids including an Australian diploid species have been created within the cotton collection maintained at the Tropical Crop Husbandry Department of the Faculty of Agricultural Sciences of Gembloux (Marchal, 1983). The first trispecific allotetraploid was obtained according to the method developed by Beasley (1940). In a first step, a fertile bispecific allotetraploid was created by doubling the chromosome number of the allo-diploid

Improvement

hybrid resulting from the cross: *G. thurberi* x *G. sturtianum*. This synthetic allotetraploid was then backcrossed by *G. hirsutum* to give the trispecific hybrid *G. thurberi* - *G. sturtianum* - *G. hirsutum* (CID1AhDh) (designated by the initials TSH). The second trispecific allotetraploid was obtained following the scheme developed by Deodikar (1949). The first step was to create an allohexaploid which resulted from colchidiodiploidization of the triploid hybrid issued from the cross *G. hirsutum* x *G. raimondii*. In a second step, this allohexaploid was crossed by *G. sturtianum* to develop the trispecific hybrid *G. hirsutum* - *G. raimondii* - *G. sturtianum* (AhDhDSC1) (designated by the initials HRS).

Both trispecific hybrids have been backcrossed by C2 and NC8 *G. hirsutum* varieties originating from Zaïre. To increase the rate of success in the backcrosses, we applied a growth regulator solution (50 mg/l NOA - 100 mg/l GA) on the ovaries according to the formula proposed by Altman (1988). To evaluate glands density of all the hybrids obtained, the seeds were soaked for one hour in water after removal of their integuments. The glands density of the hybrid seeds was assessed under a binocular microscope according to a visual scale ranging from 0, for totally glandless, to 10, for totally glanded.

After assessment of their glands density the seeds were either planted directly in jiffy-pots containing a mixture of sand, peat and compost in equal proportion, or transferred on the *in vitro* culture media developed by Stewart & Hsu (1977) for cotton germinated embryos. Adult plants were cultivated in greenhouses at Gembloux.

Results and discussion

Among the material created in Gembloux, the expression of the glanded-plant-glandless seed trait has been assessed in all hybrids including the Australian species *G. sturtianum* i.e. the bispecific allotetraploid *G. sturtianum* x *G. thurberi*, the trispecific allotetraploid hybrids TSH and HRS and their BC1 progenies. A similar expression of the character is observed in the bispecific allotetraploid and the two trispecific hybrids: the seeds have a reduced number of glands that are mainly located on the edges of the cotyledons. After germination, the number of glands increases to reach a normal density on the aerial parts of the plant. The trispecific hybrid seeds produced by Shuijing & Biling (1993) from the backcross of the bispecific allotetraploid *G. arboreum* x *G. bickii* with *G. hirsutum* showed a similar glands density. These results confirm data obtained by Mergeai (1992): the later pointed out a better expression of the repressive mechanism if an Asian diploid species (genome A) was associated with an Australian cotton species in a synthetic bispecific allotetraploid.

Thanks to the application of growth regulators, 41 viable seeds were obtained from the two trispecific allotetraploids (Table 1). For each hybrid, the backcross success rate is similar and very low. In our conditions, about 17 crosses are necessary to obtain one seed. Without application of growth regulators, we could not obtain BC1 seeds from the TSH trispecific hybrid while very few seeds were collected from the HRS hybrid (1 seed for more than 100 crosses).

The glands distribution of the BC1 seeds is very variable (Table 2). It ranges from "0" (totally glandless) to 9 (almost similar to *G. hirsutum* glanding pattern). The proportion of totally glandless seeds was relatively high (6 on 41). The distribution of the glands density frequencies of the rest of the seeds is characterised by an asymmetric bell shape.

In the BC1 population, the glandless trait appears to be linked with a lethal factor. Only one of the six glandless BC1 seeds gave rise to a viable plant. In order to facilitate the development of this plant, it was necessary to cultivate the seed *in vitro* for three weeks and then to graft the plantlet on a vigorous *G. hirsutum* seedling. All the other materials issued from glandless BC1 seeds ceased to grow before the spreading of the cotyledons. The only surviving plant from the six BC1 glandless seeds flowers abundantly and shows a normal density of gossypol glands on its aerial parts. This plant will be used in a backcrossing programme to obtain the introgression of the glanded-plant and glandless-seed character of *G. sturtianum* into *G. hirsutum*.

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Table 1. Production of BC1 seeds by HRS and STH trispecific hybrids.

Hybrids	Number of seeds produced	Number of crosses	Average numbers of crosses for one seed
HRS x NC8	5	175	35
HRS x C2	12	157	13
TSH x NC8	12	205	17
TSH x C2	12	150	13
Total	41	687	17

Table 2. Assessment of the gossypol glands density in trispecific BC1 seeds.

Glanding classes	Hybrids				Cumulated frequencies (%)
	HRS x C2	HRS x NC8	TSH x C2	TSH x NC8	
0	2	0	2	2	6(15)
1	0	0	1	0	1(2)
2	2	0	0	0	2(5)
3	1	1	0	2	4(10)
4	1	1	3	0	5(12)
5	0	1	0	3	4(10)
6	1	2	2	3	8(19)
7	2	0	2	2	6(15)
8	1	0	2	0	3(7)
9	2	0	0	0	2(5)
10	0	0	0	0	0(0)
Total	12	5	12	12	41(100)

