A Semi-Selective Agar Medium to Detect the Presence of *Xanthomonas axonopodis* pv. m*alvacearum* in Naturally Infected Cotton Seed

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ABSTRACT

A semi-selective agar medium was developed for detection of *Xanthomonas axonopodis* pv. *malvacearum* (Xam) in cotton (*Gossypium hirsutum*) seed. The basic medium was peptone-sucrose-agar (PSA). Criteria for the semi-selective medium were the typical colony characters of Xam and its pathogenicity on cotton. Several systemic fungicides and antibiotics in different concentrations were tested alone or in combination with others. The final composition of the semi-selective agar medium was established after several attempts in order to inhibit most of the fungal and bacterial saprophytes and favour the development of Xam. It contained PSA + cyclohexamide, cephalexin, pencycuron, triadimenol and tolylfluanid. The bacteria were recovered from naturally infected seeds by the direct plating of 2,000 surface disinfected seeds on the semi-selective medium. The recovery of the pathogen from naturally infected leaf tissues and in dilution plating, on semi-selective medium and on nutrient agar, were comparable. Among the three detection methods tested, the semi-selective medium was found to be the most reliable and quantifiable. Degree of severity of angular leaf spot in the field was not always correlated with the level of infection in the seed. This is the first report of a semi-selective agar medium to detect the presence of Xam in naturally infected cotton seed.

Additional keywords: *Gossypium hirsutum*, black arm of cotton, angular leaf spot.

RESUMO

Meio semi-selectivo para detectar a presença de *Xanthomonas axonopodis* pv. *malvacearum* em sementes de algodoeiro naturalmente infetadas

Um meio de cultura semi-seletivo foi desenvolvido para detectar *Xanthomonas axonopodis* pv. *malvacearum* (Xam) em sementes do algodoeiro (*Gossypium hirsutum*). O meio básico foi peptona-sacarose-ágar (PSA). Os critérios para o meio semi-seletivo foram; as características típicas das colônias de Xam e sua patogenicidade em algodoeiro. Vários fungicidas sistêmicos e antibióticos foram testados sozinhos ou em combinação com outros em diferentes concentrações. A composição final do meio semi-seletivo foi estabelecida após várias tentativas no sentido de inibir a maioria dos fungos e bactérias saprofíticas e favorecer o desenvolvimento de Xam. O meio contém: PSA + ciclohexamida, cefalexina, pencycuron, triadimenol e tolylfluanid. A bactéria foi recuperada de sementes naturalmente infetadas através do plaqueamento direto de 2.000 sementes externamente desinfestadas em meio semi-seletivo. A recuperação do patógeno de tecidos homogenizados da folha infetada e a recuperação em plaqueamento por diluição, em meio semi-seletivo e em agar nutriente, foram comparáveis. Entre os três métodos de detecção, o meio semi-seletivo foi o mais preciso e confiável. O nível da severidade de infecção em campo não era sempre correlacionado com o nível de infecção nas sementes. Este é o primeiro relato de meio semi-seletivo para detectar a presença de Xam em sementes do algodoeiro.

Palavras-chave adicionais: *Gossypium hirsutum*, mancha angular.

INTRODUCTION

Angular leaf spot caused by *Xanthomonas axonopodis* pv. *malvacearum* (Xam) [Sin. *X. campestris* pv. *malvacearum* (Smith) Dye)] is one of the most important diseases of tetraploid cotton (*Gossypium hirsutum* L. and *G. barbadense* L.) in many countries, including Brazil. The disease can cause heavy yield losses depending on the year and the cultivar. Cotton seed is considered to be an important transmission vehicle of Xam and a source of primary inoculum (Bain, 1939; Brinkerhoff & Hunter, 1963; Hunter & Brinkerhoff, 1964; Mohan, 1983a, 1983b). Tarr (1961) and Brinkerhoff & Hunter (1963) reported that internally infected seed between 0.017 and 2.0% resulted in severe losses in commercial cotton fields in Sudan and in the USA. So far, in Brazil, there is no regulation, either for field inspections for angular leaf spot or for laboratory testing, of seeds infected with the pathogen. As a result, the disease is on the increase in the States of Paraná, São Paulo, Goiás, and Mato Grosso, and severe epiphytotics of the disease

occurred in 2001, 2002 and 2003 in these states. Currently, the Brazilian Ministry of Agriculture is considering the adoption of rules and regulations requiring field inspections for the presence of angular leaf spot disease. A zero tolerance for fields infested with the disease is being proposed. Any seed production field with the disease, irrespective of the level of infection, will be destined for commercial cotton and not for seed production. This stringent regulation may create seed shortages and other problems. The year 2003 was very favourable for angular leaf spot, and in the State of Mato Grosso, there was not a single field which was completely free from the disease. Thus, the success of a seed health certification program, as in other cases (Mohan & Schaad, 1987), would depend on the availability of a reliable method to detect Xam in naturally infected seeds.

Several semi-selective agar media were reported for phytopathogenic bacteria like *Corynebacterium* spp. (*Clavibacter*), *Erwinia* spp., *Pseudomonas* spp. and *Xanthomonas* spp. (Kado & Heskett, 1970), *Xanthomonas translucens* pv. *undulosa* (Schaad & Forster, 1985), *X. axonopodis* pv. *phaseoli* (Clafin *et al.*, 1987), and *Pseudomonas syringae* pv. *syringae* (Mohan & Schaad, 1987). In Israel, a method to detect the pathogen in the seed lot, but not quantitatively, was reported by Halfon-Meiri & Volcani (1977). A semi-selective medium to detect and to quantify the presence of Xam in cotton seed has not been reported. Because of the lack of an appropriate, sensitive, selective medium, research on control measures including methods to erradicate the bacteria from the seed is hampered. A preliminary report about semi-selective medium has been published (Mehta & Bolognini, 2003). This paper describes a semi-selective agar medium and compares other methods to detect the presence and longevity of Xam in naturally infected cotton seed.

MATERIALS AND METHODS

Development of semi-selective agar medium

Some of the semi-selective media reported for xanthomonads (Schaad & Forster, 1985; Claflin *et al*., 1987) were evaluated in preliminary tests. Some systemic fungicides were tested alone or in combination with others in different concentrations. Some of the antibiotics were also evaluated by paper disc method (Randhawa & Schaad, 1984). The basic medium was peptone-sucrose-agar (PSA), containing 0.35 g Ca (NO_3), 0.35 g $FeSO_4$, 1.4 g Na_2HPO_4, 3.5 g peptone, 14.0 g sucrose, 10.5 g bacto-agar, 700 ml distilled water, pH 6.8. Criteria for the semi-selective medium were the typical colony characters of Xam and its pathogenicity on cotton. The final composition of the semi-selective agar medium was established so as to inhibit most of the fungal and bacterial saprophytes and to favour the isolation of Xam. It contained: PSA + cycloheximide (Sigma) 100 mg (2 ml of a stock solution of 500 mg in 10 ml of 75% alcohol); cephalexin (Sigma) 10 mg (1 ml of a stock solution of 100 mg in 10 ml of 75% alcohol); pencycuron 0.05 g (Monceren PM, Bayer S.A.); triadimenol 1 ml (Baytan SC, Bayer S.A.); tolylfluanid (Euparen 500PM, Bayer S.A.) (1 ml of a stock solution of 0.125 g in 10 ml of sterilized water). The fungicides pencycuron, triadimenol and tolylfluanid were efficient in inhibiting the development of most of the seed-borne fungal saprophytes. Tolylfluanid as a seed dressing fungicide is also known to control *Rhizoctonia solani* Kühn in cotton seedlings (Goulart & Ferraz, 2001). Antibiotics and fungicides were added, after the medium was autoclaved, in the order presented above. In all the tests, young culture of Xam was streaked onto two others containing PSA and two plates containing semi-selective medium and maintained as controls to ensure that in none of the tests Xam was inhibited by the semi-selective medium.

Plating efficiency

Plating efficiency was determined in comparison with Difco nutrient agar (NA) as a standard medium and was evaluated by colony counts at 10 fold serial dilutions (10^{-4}, 10^{-5}, 10^{-6}, 10^{-7}), using young cultures of 13 strains of *Xanthomonas* sp. in three replicates. Initial cell suspension was adjusted to an absorbance of 0.1 at 640 nm using a spectrophotometer. One hundred microlite of each diluted suspension was distributed on each Petri dish containing either NA or semi-selective medium. Dilutions yielding over 300 colonies per plate were not considered for counting. The mean percentage of recovery of the colony forming units (cfu) were compared with NA. The percentage of plating efficiency was calculated in relation to the cfu developed on NA, three days after incubation at 24 °C. Recovery of Xam from homogenized diseased leaf tissues was also verified on both NA and semi-selective media.

Seed samples

Samples were obtained from six seed lots belonging to *G. hirsutum* (three of cv. ITA 90, one of cv. Fabrika and two of cv. Makina) from a 2001 harvest in Mato Grosso of fields producing seeds with angular leaf spot disease (Table 3). The level of disease incidence in these fields was not known. After delinting with concentrated sulfuric acid (Brinkerhoff & Hunter, 1963) the samples were gently washed three times with tap water, air-dried on filter paper and stored at 5 °C for further use.

Seed testing assays

To detect the presence of seed infection in six seed lots, the following methods were used. For all pathogenicity tests, first leaves of 30-35-day-old plants of cvs. Ita 90 and Saturno were inoculated (three plants per cultivar) using a tooth pick method (Cia *et al*., 1973). Inoculated plants were incubated in a dew chamber for 48 h and later placed on the greenhouse bench and examined periodically for symptom expression.

Seed washing. Seeds (200 g) were disinfested with 70% ethyl alcohol for 1 min, immersed 4 min in sodium hypoclorite (2.5% of NaClO), followed by three quick

washings with sterilized water. The disinfested seeds were finally washed in 300 ml of saline water (0.5% NaCl) for 24 h in a mechanical shaker in three replicates. The suspension was filtered through Whatman filter paper No. 1, centrifuged for 10 min at 10,000 rpm. The pellet was suspended in 5 ml of distilled water and used for pathogenicity tests in the greenhouse;

Direct seed plating. Two thousand seeds were surface disinfected as mentioned earlier for seed washing, but instead of washing again for 24 h, were placed in Petri plates (ten seeds per plate) each containing 15 ml of semi-selective medium. Plates were examined periodically seven days after incubation. Within seven-12 days after incubation, typical colonies of Xam growing around the seeds were observed and were tested directly for pathogenicity. The identification of the Xam colonies was based on morphological characteristics like size, colour and consistency which were also confirmed by streaking onto PSA plates. After 12 days no new colonies of Xam were observed.

Growing-on-test. Two thousand seeds of each seed lot were sown in plastic trays (20 x 35 cm) containing unsterilized soil in the greenhouse (100 seeds per tray), trays were irrigated once a day and the plants were examined for symptom development 15-35 days after sowing. The average greenhouse temperatures during the experimental period varied between 18 and 26 ^{o}C, and the relative humidity between 60 and 85%. Care was taken to avoid counting plants with secondary infections. For this purpose, plants were examined daily and the infected ones were marked within the first 25-30 days after sowing. Although secondary infection was rare, it was normally identified as a few water-soaked angular spots on the first or second leaves on plants with no other symptoms. Invariably such plants were surrounded by the initially infected plants.

Longevity

Longevity of Xam in two seed lots was studied by plating 2,000 surface desinfested seeds periodically onto the semi-selective medium using the same procedure as stated for direct seed plating. The coefficient of correlation between the period of seed storage and counting the number of bacterial colonies was evaluated by regression analysis using the "Cricket graph" program, version 1.01 of Macintosh.

Correlation between disease severity in the field and seed infection

The disease severity levels of angular leaf spot in the field (estimated as the percentage of leaf area infected) were correlated with the level of infection in the seed by plating upto a total of 4,800 seeds of some samples onto the semi-selective medium. For this purpose, the disease severity scale of 0-4 for field infection was used, where, 0= no disease, 1=trace - less than 5% of the leaf area infected (LAI); 2= low (<25% LAI); 3=medium (<50% LAI); 4=severe (>50% LAI).

RESULTS

Plating efficiency

In the dilution plating, the percentage of Xam recovery on semi-selective agar medium compared well with NA. Colonies were recovered in both media even at a very high dilution indicating the high sensitivity of the semi-selective medium to detect the pathogen. (Table 1, Figure 1). The colonies of Xam grew faster and were larger on semi-selective medium than on NA. Plating efficiencies (% mean recovery) on semi-selective medium varied between zero and 187%. Whereas the semi-selective medium favoured good recovery for all the strains of *Xanthomonas* sp. from cotton, the recovery of one strain of *X. axonopodis* pv. *phaseoli* (Smith) Dye was zero. The semi-selective medium inhibited the growth of *X. axonopodis* pv. *phaseoli* during the first three days after incubation (Table 1). Few and very small colonies of this strain developed five days after incubation and hence were not considered for analysis. Amongst the xanthomonads the percentage recovery of *X. axonopodis* pv. *citri* was one of the highest on semi-selective medium.

Recovery of Xam from homogenised diseased leaf tissues was similar on both NA and semi-selective medium indicating the non-toxicity of the semi-selective medium (Figure 2).

Seed washing

When suspensions of seed washings were inoculated on susceptible cotton plants, typical disease symptoms were reproduced 12 days after inoculation in seed sample Nos. 5 and 6 in all the three replications (Table 2).

Direct seed plating

Growth of most of the saprophytic bacteria and fungi was inhibited by the semi-selective agar medium and by the disinfestation of the seeds by alcohol and by sodium hypoclorite. Since the pathogen is internally seed-borne, detection of Xam was not affected by disinfestation as evidenced by some preliminary tests. The addition of gentamycin inhibited the development of most of the fungi and bacteria (Schaad & Foster, 1985), but it was toxic to Xam. Susbstitution of gentamycin by pencycuron, triadimenol and tolylfluanid, inhibited the development of most of the fungi but at the same time did not affect the development of Xam. Addition of kasugamycin did not improve the medium. It was non toxic to Xam as well as to other saprophytic bacteria. Laboratory tests indicated that some of the saprophytic bacteria associated with the cotton seed were not antagonistic to Xam. The typical colonies of Xam growing around the seed were slightly yellowish, smooth, convex and glistening (Figure 3). The identity of such colonies was always confirmed by streaking onto the PSA medium and by the pathogenicity test in the greenhouse. Out of six seed lots two were free from the bacteria. The semi-selective medium could detect a very small percentage

TABLE 1 - Number of strains of *Xanthomonas* spp. on semi-selective agar medium compared to growth on Difco nutrient Agar

Strain/Pathovar*	Host/State	Mean recovery (%)**
Xan. 13403/*malvacearum*	Cotton/Mato Grosso	66 ed
Xan. 13355/*malvacearum*	Cotton/Mato Grosso	86 ced
Xan. 13360/*malvacearum*	Cotton/Mato Grosso	100 cbd
Xan. 13362/*malvacearum*	Cotton/Mato Grosso	87 ced
Xan. 13601/*malvacearum*	Cotton/Mato Grosso	104 cbd
Xan. 13838/*malvacearum*	Cotton/Mato Grosso	114 cbd
Xan. 13856/*malvacearum*	Cotton/Mato Grosso	65 ed
Xan. 13805/*malvacearum*	Cotton/Mato Grosso	158 cb
Xan. 15219/*malvacearum*	Cotton/Mato Grosso	103 cbd
Xan. 6667/*undulosa*	Wheat (*Triticum aestivum* L.)/Paraná	57 ed
Xan. 12917/*citri*	Citrus (*Citrus* spp.)/Paraná	187 a
Xan. 3701/*phaseoli*	Beans (*Phaseolus* spp.)/Paraná	0 e
Xan. 11370/*passiflora*	Passionfruit (*Passiflora edulis* Sims)/Paraná	56 ed

*Xan=*Xanthomonas*. All strains belong to the culture collection of IAPAR, Londrina;

**(Number of colonies recovered on semi-selective agar medium x 100)/number of colonies on Difco nutrient agar). Calculated from the mean number of colonies per plate in three replications per strain. Columns with the same letter do not differ from each other according to Tukey at 5% (p=0.05). *Xanthomonas* strains from passion fruit, citrus, and beans, were collected by Rui Perreira Leite, IAPAR, Londrina, PR.

of infection (0.05%), which otherwise was not detected by any other method (Table 2).

Growing-on-test

The pathogen transmission by growing-on-test in one seed lot was demonstrated under greenhouse conditions (Table 2). Twenty five to 30 days after sowing, typical water soaked translucent angular leaf spot symptoms developed on the first leaves of cv. Makina which in severe cases caused death of the seedling. The identity of the pathogen was confirmed by isolation and by pathogenicity tests in the glasshouse. As mentioned earlier, failure to demonstrate the transmission of the bacteria in other seed lots by growing-on-test was possibly due to a relatively low level of seed infection. Besides, an inapropriate temperature and humidity regime during the experimental period may inhibit the development of typical angular leaf spot symptoms, and consequently, a particular seed lot may be wrongly identified as free from Xam. The exact temperature and humidity regime for seed transmission is not well understood.

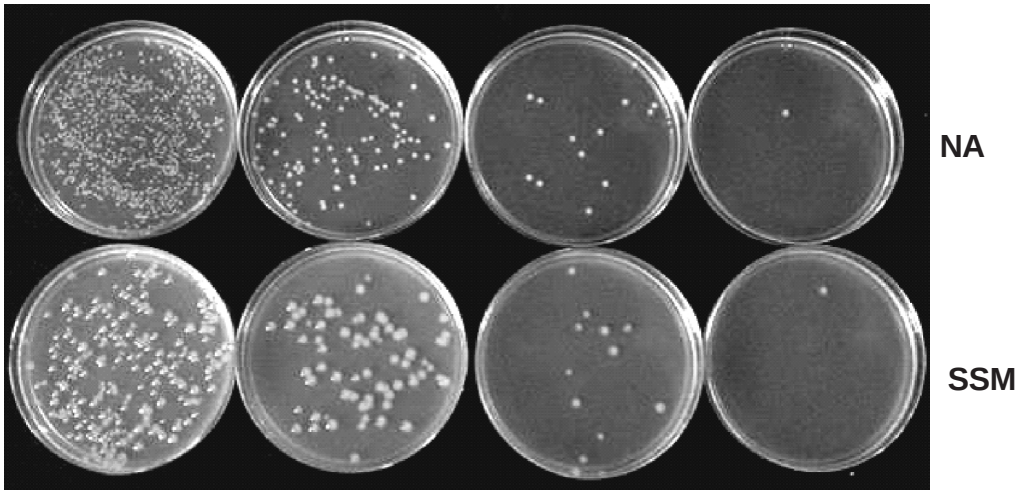

FIG. 1 - Comparison of colony forming units of *Xanthomonas axonopodis* pv. *malvacearum* in dilution plating (from left to right - 10^{-4},10^{-5},10^{-6},10^{-7}) between Difco nutrient agar (upper plates) and on semi-selective agar medium (lower plates), three days after incubation.

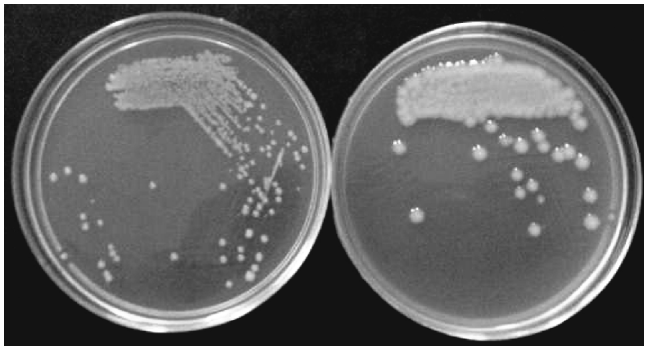

FIG. 2 - Recovery of *Xanthomonas axonopodis* pv. *malvacearum* from homogenised diseased cotton (*Gossypium hirsutum*) leaf tissues on Difco nutrient agar (left) and on semi-selective agar medium (right), three days after isolation.

FIG. 3 - Typical colony growth of *Xanthomonas axonopodis* pv. *malvacearum* indicated by an arrow, recovered on semi-selective agar medium from naturally infected cotton (*Gossypium hirsutum*) seed, eight days after incubation at 24° C.

Longevity

By using the semi-selective medium it was possible to study the longevity of Xam in two naturally infected cotton seed lots. The viability of bacteria started declining gradually 12 months after harvest and reached the lowest level after 23-26 months of storage (Table 3). The reduction in the level of seed infection was inversely proportional to the increase in storage period. For the seed lots No. 5 and 6, the coefficient of correlation (r) was 0.890 and 0.975, respectively. Unfortunately, there is no information regarding whether there was a decline or not in the level of seed infection during the first 12 months after harvest, since the semi-selective medium was developed after this period. The results showed that the bacteria can survive in the infected seed for a period of at least 28 months, especially when the seed is stored at 5 °C. The longevity of Xam in dried leaves preserved under laboratory conditions for seven years and in the seed for 56 months was demonstrated by Schnathorst (1964) and Hunter & Brinkerhoff (1964), respectively. It is also known that *X. campestris* pv. *undulosa* Hagb of wheat, for example, can survive in the infected seed for a period of over four years (Mehta, 1990; Mehta & Bassoi, 1993; Bragaard *et al*., 1993). Further research is needed to verify the longevity of Xam in the infected seed at the normal storage conditions (about 24 °C) commercially practiced in Brazil.

Correlation between disease severity in the field and seed infection

Results indicate that the disease severity levels of angular leaf spot in the field did not always correlated with the level of infection in the seed (Table 4).

DISCUSSION

In general, the semi-selective medium showed good plating efficiency for the strains of *X. axonopodis* pv.

TABLE 2 - Detection of *Xanthomonas axonopodis* pv. *malvacearum* (Xam) in six naturally infected commercial cotton (*Gossypium hirsutum*) seed lots by three different methods

Seed lot and cotton cultivar	Origin Location/State/Year	Seed infected with Xam		
		Method of detection*		
		Seed washing	Seed plating on semi-selective medium**	Growing on-Test
Fabrika	Rondonó polis,MT/2001	Negative	4 (0.20%)	0
Ita 90 A	Serra de Petrolina, MT/2001	Negative	0	0
Makina	Rondonó polis, MT/2001	Negative	1 (0.05%)	0
Ita 90 B (treated with unknown fungicide)	Rondonó polis (Salles), MT/2001	Negative	0	0
ITA 90 C	Itiquira, MT/2001	Positive***	21 (1.05%)	No data due to short of seed
Makina	Costa Rica, MT/2001	Positive***	20 (1.00%)	31 (1.55%)

*See text for details of the three methods of detection.
**Number (percentage of infected seed with Xam).
***After seed washing the pellet containing Xam produced typical symptoms of the disease 12 days after inoculation. Delinting was done using concentrated sulphuric acid (Brinkerhoff & Hunter, 1963).

TABLE 3 - Longevity of *Xanthomonas axonopodis* pv. *malvacearum* (Xam) in naturally infected cotton (*Gossypium hirsutum*) seeds stored at 5° C soon after harvest

Date of seed analysis/ recovered on months after harvest	Number of seeds which anabled the recovery of Xam on semi - selective medium from two seed lots*	
	Seed lot - cv. Ita 90	Seed lot - cv. Makina
August 5, 2002/12	21	20
October, 7, 2002/14	-	13
February 24, 2002/18	-	11
March 19, 2003/19	-	6
April 2, 2003/20	5	-
May 12, 2003/21	2	-
Ju ne 5, 2003/22	-	1
July 5, 2003/23	2	0
October 1, 2003/26	4	- (Short of seed)
December 5, 2003/28	4	- (Short of seed)

*Seed samples originated from fields with angular leaf spot from the State of Mato Grosso, and soon after harvest in July, 2001, stored at 5 °C. For each analysis 2,000 surface sterilised seeds were placed on Petri dishes containing semi-selective medium. Typical Xam colonies were observed eight-ten days after incubation at 24 °C in dark. (-) indicates lack of data. The coefficient of correlation (r) between the increase in storage period and the number of colonies of Xam, for cv. Ita 90 and for cv. Makina was 0.890 and 0.975 respectively.

TABLE 4 - Relationship between angular leaf spot severity in the field and the percentage of infected cotton (*Gossypium hirsutum*) seeds with *Xanthomonas axonopodis* pv. *malvacearum* (Xam) determined by using the semi-selective medium

Cotton Cultivar*	Disease severity in the field**	No. of seeds analysed using semi - selective medium***	Average No. of infected seeds	% infected seed
Breder's line	3	4,800	0	0
Saturno	3	4,800	9	0.38
ITA 90 - D	2	4,800	0	0
ITA 90 - E	3	4,800	1	0.04
ITA 90 - F	4	4,800	5	0.21
Alcapi	0	800	0	0
Alcapi	3	800	4	0.5
Delta Opal	0	800	0	0
STO 474	1	800	0	0
Saturno	2	800	0	0

*Seed samples originated from different fields or without angular leaf spot from the State of Mato Grosso, with the exception of the Breeder´s line which was from the State of Paraná. Soon after harvest in June, 2003, the seeds were disinfested and were analised using semi-selective medium. Typical Xam colonies growing around the infected seeds were observed 8-12 days after incubation at 24 °C in dark.

**Disease severity scale of 0-4 for field infection was used, where, 0= no disease, 1=trace - less than 5% of the leaf area infected (LAI); 2= low (<25% LAI); 3=medium (<50% LAI); 4=severe (>50% LAI).

***For the first five seed lots 4,800 seeds were analysed in two replications of 2,400 seeds each. Due to the limited seed of the rest of the seed lots, only 800 seeds were analysed.

malvacearum as compared to other xanthomonads. Among the three detection methods tested, the semi-selective medium was found to be the most reliable for quantitative assays. The identity of Xam colonies appearing around the infected seeds on the semi-selective agar medium was confirmed by pathogenicity tests, which are considered desirable since some of the colonies of Xam may loose their pathogenicity, or else they may still be confused with some of the yellow pigmented saprophytic bacteria. Alternatively, identification of Xam colonies may also be confirmed by morphological characteristics by streaking onto the PSA medium and comparing with the type culture colonies of Xam. Our experience shows that the colonies, once confirmed by streaking onto the PSA medium, were always pathogenic to cotton. Pathogenicity tests may be dispensable if identification is coupled with molecular analysis using a highly specific DNA probe. Nonetheless, the semi-selective agar medium presented here is practical and can be used for screening large number of seed samples. The medium can also be used for quarantine purposes.

Normally, the percentage of seed infection by Xam is very low. A relatively low percentage of seed infection, however, would be sufficient to create a severe epidemic of the disease under favourable field conditions (Brinkerhoff & Hunter, 1963). Considering the low percentage of seed infection and the possible establishment of a zero tolerance level for seed infection by the Ministry of Agriculture of Brazil, we suggest that 2,000 seeds per seed lot should be examined.

Although the pathogen can survive in seed and in volunteer cotton plants, no secondary or collateral hosts are known. The bacterium like most xanthomonads, is sensitive to drying and excessive heat, and is rapidly destroyed when

crop residue is buried soon after the cotton harvest (Arnold & Arnold, 1961; Brinkerhoff & Fink, 1964; Mehta, 1993). Such conditions prevail especially in the State of Mato Grosso, and thus reduce the possibility of survival of the pathogen in infected cotton plant residue. Although longevity and the role of inoculum present in crop residue under Brazilian conditions are not well studied, our results show that the seed could be an important source of primary infection of angular leaf spot in Brazil.

The disease severity levels of angular leaf spot in the field were not always correlated with the level of infection in the seed, even when the test was repeated and a total of 4,800 seeds of some samples were plated onto the semi-selective medium (Table 4). Similar results were obtained by Mohan & Schaad (1987) while working with bean blight bacteria. Because of the limited sample size, only 800 seeds were tested of some seed lots. Correlation between the level of field infestation and the level of seed infection would depend on the climatic conditions between the boll formation and boll opening. Future research using a large number of seed samples is considered essential to improve our understanding of the correlation between field infection and seed infection. Nonetheless, results indicate that the seed production fields infested with angular leaf spot disease can not be simply condemned. To the contrary, it would be worthwhile to adopt a zero tolerance level for seed infection but not for field infestation, especially now when the semi-selective medium has been made available. Besides, field inspections are not always very reliable and nor foolproof (Mohan & Schaad, 1987).

This is the first report of a semi-selective agar medium to detect the presence of Xam in commercial cotton seed. The development of such a medium would certainly provoke furture investigations into the establishment of appropriate methods designed to erradicate the pathogen from the cotton seed. The use of semi-selective medium would permit planting of only the healthy seeds for commercial cultivation. Along with healthy seeds the use of resistant cultivars would effectively control the angular spot disease of cotton.

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