

Original Article

Production Bacteriosin by Rhizobia obtained from Root nodules of *Macrotyloma uniflorum* (Lam.) Verdc. (Horse Gram)

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Thirty two *Rhizobium* isolates obtained from the root nodules of Horse gram plants grown in 32 different soil samples collected from various regions in Andhra Pradesh, India. They were identified as rhizobia by nodulation test. These isolates were used to study the production of bacteriocins. All the isolates produced bacteriocins against the remaining isolates. The isolates HGR-4 and 9 showed maximum activity/inhibition zone among all the isolates, and also found to be inhibiting more number of isolates than that of other bacteriocin producing isolates. The bacteriocin protein substance produced by the isolates HGR-4 and 9 showed a molecular weight of ~ 50 kDa by SDS-AGE and bacteriocin plasmids molecular weight is > 22 kb and > 24 kb.

Key words: bacteriocin, rhizobia, competition.

Introduction

The *Rhizobium* legume symbiosis is the most promising plant bacterium association so far known. Inoculated *Rhizobium* strains often fail to compete with the indigenous soil rhizobia and do not increase nodulation^{1,2}. Thus the successful use of rhizobial inoculants requires the knowledge of factors affecting the effectiveness and competitive ability of the rhizobia. One of the major factors reported to be affecting competition among rhizobia are bacteriocins³⁻⁵. Bacteriocins are proteins or protein complexes with bacteriocidal activity against usually closely related to producer bacterium⁶. Bacteriocins are ribosomally encoded peptide antibiotics produced by both Gram negative and Gram positive bacteria. Bacteriocins produced by the Gram negative are well studied. *Rhizobium leguminosarum* has been shown to produce bacteriocins which have been characterized small, medium or large based on their assumed size characteristics^{7,8}. Small bacteriocins were found to be chloroform soluble and have molecular mass less than 2000 Da^{7,9}. Oresnik *et al.*⁵ found that the bacteriocins appear to play a major role in determining competitiveness for nodulation when assayed against some strains.

Genetic evidence suggests that some of the genes involved in *Rhizobium*-legume symbiosis may be located on plasmids¹⁰⁻¹². Many bacteriocins of enterobacteria are plasmid determined¹³ and this may be a feature common to other bacterial genera. Bacteriocin production by *Rhizobium* spp. has been

described^{4,14-16} but, there are no data to suggest that it may be plasmid determined.

So the successful preparation of mixed inoculum requires the knowledge of bacteriocin producing ability of the inocula strains as well as their effect on the related rhizobia. In the present study 32 rhizobia isolated from various regions in Andhra Pradesh, India has been investigated for their bacteriocin production.

Materials and Methods

Isolation of root nodulating bacteria from horse gram

Root nodules were collected from the horse gram plants. Rhizobia were isolated from fresh nodules by the standard method on Yeast Extract Mannitol Agar (YEMA) medium¹⁷. They were designated as HGR-1 (Horse Gram Rhizobia) to HGR-32. Root nodulating ability of these isolates was determined by nodulation test¹⁸. Morphological, cultural and biochemical activities of the isolates were studied using standard methods. Four strains such as HGR-4, 6, 13, and 25 were selected for 16S rRNA sequence and were submitted to the NCBI GenBank under the accession numbers GQ483457, GQ483458, GQ483459 and GQ483460.

Bacteriocin Activity Assay

The bacteriocin producing ability of the strains was bioassayed by simultaneous antagonism method⁶. Bacteriocin activity was examined by adding 1 ml of each diluted, sterile filtered sample on to the wells made on Tryptone Yeast extract (TY) medium (0.6%

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w/v agar) seeded with log phase indicator strains (0.5µL of the medium). Activity was quantified by two fold serial dilution¹⁹ and is expressed in arbitrary units mL⁻¹ (AU mL⁻¹).

Bacteriocin protein purification

Purification of proteins was carried out using the procedure of Yang *et al.*²⁰. Cell Free Supernatant (CFS) was used to carry out protein extractions. Twenty percent chloroform was added to the CFS in a separatory funnel. The aqueous phase formed was separated and used for precipitating out the proteins. Protein precipitation was carried out on ice or at 4°C by the addition of analytical grade ammonium sulphate. The aqueous phase was saturated with cold ammonium sulphate from 20 to 80% (w/v) saturations and gradually stirred with a glass stirrer for 10-15 min. The aqueous phase was kept overnight at 4°C.

The precipitate was collected by centrifugation at 15000xg for 20 min. The solid pellet dissolved in distilled water and dialyzed against distilled water at room temperature for 24 h. The suspension obtained was designated as proteinaceous fraction or crude bacteriocin fraction. All the different dialysates obtained were lyophilized. Lyophilized material of 0.01 g was added in 100ml Tris-HCl (pH 6.5) buffer and tested for inhibitory activity. The quantification of protein concentration was done by standard Bradford method²¹. Bovine serum albumin (BSA) was used to construct the standard curve.

Bacteriocin protein purification

Analysis of total cellular proteins of rhizobia was carried out by SDS-PAGE described by Laemmli²² with slight modifications. 3 ml of YEM broth was inoculated and incubated for 12 h at room temperature. It was centrifuged at 10000g for 4 min at 4°C. To the pellet sample buffer was added at 1:1 ratio. The samples were lysed by heating at 95°C for 2 min and immediately transferred to the cold water. Lysed samples and marker protein samples were loaded into the different gel slots. Electrophoresis was carried out at 100V for 4 h at room temperature. The gel was stained with coomassie blue (Sigma). The image was photographed using Gel documentation unit (Alpha Innotech, USA).

Plasmid analysis

An overnight culture of rhizobia in YEM broth was centrifuged at 14000xg, for 4 min at 4°C and the plasmid isolation and the analysis was carried out according to Sambrook and Russell²³.

Results and Discussion

The cells are Gram negative rods, colonies are small, white on YEMA medium at room temperature. The optimum pH was in the range of 7-7.5. These strains grow at a temperature between 10 to 40°C. Lower concentrations of NaCl favoured growth of these rhizobia. All the isolates were resistant to ampicillin and rifampicin. The isolates HGR-11, 22 and 23 showed resistance to most of the antibiotics tested, where as the isolate HGR-4 showed susceptibility towards most of the antibiotics.

Plasmid profile analysis is a useful tool for differentiating among strains of *Rhizobium* within the same species. The objective of this study is to determine the diversity in plasmid profiles and relationships between the isolates collected from the horse gram plants grown in 32 soil samples collected from various regions in Andhra Pradesh, India.

For comparative taxonomical identification the sequences of the four strains were submitted to the Ribosomal Database Project (RDP). According to NCBI and RDP first isolate (HGR-4) belongs to the genus *Rhizobium*, second one (HGR-6) is an unclassified member of *Rhizobiaceae*, third isolate (HGR-13) belongs to the genus *Rhizobium* and fourth isolate (HGR-25) belongs to the genus *Caulobacter* of the family *Caulobacteriaceae* in the order *Caulobacterales* of Alphaproteobacteria. It clearly shows that the horse gram rhizobia are phylogenetically distinct.

The production of bacteriocins is an advantageous character for strain competition²⁴. Bacteriocin production was indicated by the formation of clear zones around the colonies. All the thirty two isolates produced clear zones on YEMA medium (0.6% w/v) inoculated with the isolates from the same host. No test strains inhibited its own growth, which is characteristic of bacteriocin producers^{13,25}.

The bacteriocin production started after 48 h and reached maximum after 72 h of incubation. Further incubation does not affect the zone size, therefore, 72 h of growth of the producer colonies at 30°C was considered as optimum conditions for bacteriocin production in this study. That the production of bacteriocin is closely related with bacterial growth of producing organism and bacteriocin activity decreases more or less sharply at the end of the growth phase as a result of degradation by proteases was reported earlier in *Micrococcus* sp.²⁶. All the strains of *Rhizobium* were found to produce antimicrobial activity, which inhibited the growth of the related strains on the agar medium.

In the present study maximum activity/inhibition zone was observed with the isolate HGR-9 and the isolate HGR-4 was also found to be inhibiting more number of isolates than that of other bacteriocin producing isolates, further studies were carried out with these two strains (Table1). Thus the activity spectrum varied from strain to strain was reported earlier in *Rhizobium leguminosarum* bv. *viciae*. This is consistent with the earlier report that the strains of *Rhizobium leguminosarum* bv. *viciae* differ in their activity spectrum²⁷.

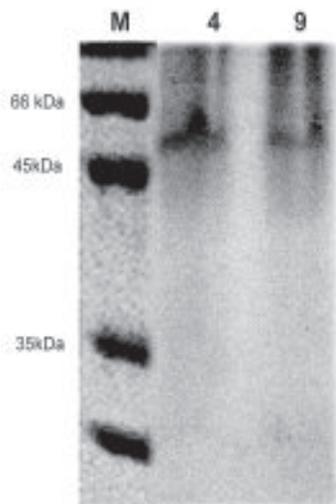
Present results showed that when the sample is successfully diluted, inhibition zone decreased until critical dilution was achieved, where no inhibition of the sensitive organism was observed. When the purified bacteriocin was tested against indicator strains, it showed highest activity at 50% ammonium sulphate saturated pellet. The activity does not depend on the quantity of the protein produced.

SDS-PAGE analysis of protein isolated from the strain HGR-4 and 9 showed the presence of ~ 50 kDa protein band (Fig 1). Plasmid analysis of this strains showed that it contains two plasmids similar to > 22 kb and 24 kb (Fig 2).

Table.1. Purification of isolated bacteriocin protein from HGR-4 and HGR-9

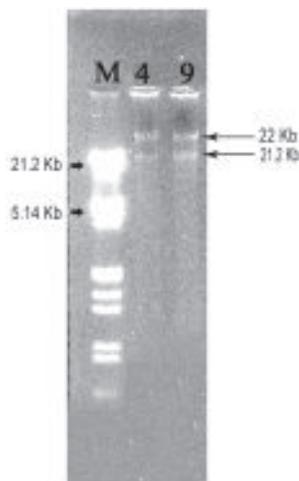
Sample material	Potency		Diameter of inhibition zone (mm)		Arbitrary Units (AU mL ⁻¹)		Protein concentration (mg mL ⁻¹)	
	HGR-4	HGR-9	HGR-4	HGR-9	HGR-4	HGR-9	HGR-4	HGR-9
Chloroform extract	+++	+++	22.0	15.0	190	160	6.0	7.25
Ammonium sulphate concentration (%)								
20	++	++	18.0	15.2	100	98	7.60	8.10
30	++	++	22.6	18.6	104	106	8.85	8.00
40	++	++	24.0	20.2	110	109	9.00	8.65
50	+++	+++	33.0	32.0	240	220	10.25	9.25
60	++	++	28.0	25.2	114	112	8.50	7.20
70	++	++	28.0	14.0	86	82	7.00	5.00
80	-	-	-	-	-	-	-	-

-: Ineffective, +: less effective, ++: Moderately effective, +++: Highly effective. All the results are means of triplicates.



(M) Marker lane (4) Isolate HGR-4 (9) Isolate HGR-9

Fig 1. SDS-PAGE of purified bacteriocin of rhizobia from horse gram.



(M) Marker lane (4) Isolate HGR-4 (9) Isolate HGR-9

Fig 2. Bacteriocin plasmids of horse gram rhizobia

A characteristic large plasmid (~4,000 bp) was present in thirteen isolates (HGR-2,3,4,5,6,10,11,12,14,20,22,24 and 26). Plasmid profile analysis of these isolates showed that these horse gram rhizobia showed similarity in their plasmid profiles although the rhizobia were isolated from different regions in Andhra Pradesh. Among the thirty two rhizobia, 13 isolates contained only one single plasmid i.e. ~4000 bp, the remaining isolates had no plasmid. According to the size and number of plasmids (plasmid profile) it was possible to classify these 32 horse gram rhizobia into two groups and suggests that the isolates are genetically different (Fig 3).

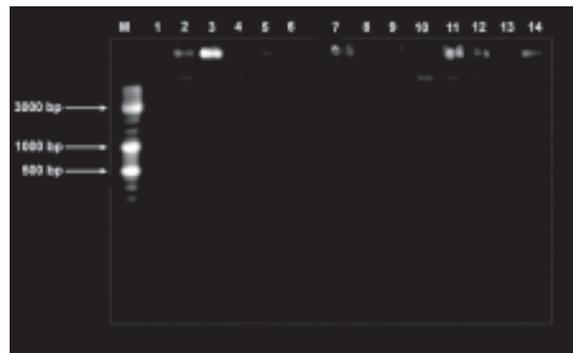


Figure 3. M- Marker Lane: Gene Ruler DNA Ladder Mix (Fermentas, USA)

Thus, these studies indicate that bacteriocin production is strain specific character. It is clear that bacteriocin production may play an important role in interspecific competition. These strains may help in the improvement of legume inoculants.

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