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Original article

Rhizobium gallicum as an efficient symbiont for bean cultivation

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Abstract – Rhizobia are soil bacteria that fix atmospheric nitrogen in symbiosis with legumes in specialized organs called nodules. The legumes thus acquire the autonomy to grow in nitrogen-deficient soils. When nitrogen fixation by indigenous rhizobia is limited, field inoculation with efficient and competitive strains is an economically feasible way to increase production. When the inoculant is made from native strains of rhizobia the success of inoculation should be increased, since local strains are better adapted than commercial inoculants. Here, a *Rhizobium gallicum* strain, 8a3, previously selected for its competitiveness and symbiotic effectiveness with common bean under laboratory conditions, was tested in field trials in Tunisia. The experiments were conducted in six fields using three common bean cultivars. The majority of the fields showed a low density of the native rhizobia and inefficient nodulation by *Sinorhizobium meliloti*, a known symbiont of *Medicago*. Our results show that inoculation with *R. gallicum* strain 8a3 induced an increase in nodule numbers accompanied by a more than twofold increase in shoot dry yield. Monitoring of the nodulation occupancy through the fingerprinting of the repetitive extragenic palindromic sequences (REP-PCR) showed that strain 8a3 was competitive even in the soil showing a high population density of indigenous *R. gallicum*, and occupied more than 40% of the nodules. Moreover, in vitro antibiosis assays indicated that strain 8a3 produces antimicrobial activity on agar medium against indigenous common bean rhizobia, including the inefficient strains of *S. meliloti*. These results point out the benefits that could be achieved by selecting efficient and competitive strains among natural populations of rhizobia.

competitiveness / N2 fixation / Phaseolus vulgaris / Rhizobium gallicum

1. INTRODUCTION

Rhizobia are soil bacteria capable of inducing the formation of nitrogen-fixing nodules on the roots or stems of particular legume host plants (Spaink et al., 1998). The root nodule formation is the result of a complex molecule dialogue between rhizobia and legumes (Garg and Geetanjali, 2007). The symbiotic nitrogen fixation resulting from the rhizobialegume interaction can act as a renewable and environmentally sustainable source of nitrogen and can complement or replace fertilizer inputs (Peoples et al., 1995). Nitrogen offered by the bacteria is used directly by the plant, and so is less susceptible to volatilization, denitrification and leaching (Garg and Geetanjali, 2007).

Common bean (*Phaseolus vulgaris* L.) is an important food crop worldwide. However, it is classified among the least cultivated grain legumes in Tunisia. The low productivity of this crop has contributed to a progressive reduction in the area used for its cultivation, and currently national production does not cover more than 40% of consumption needs. A broad range of *Rhizobium* species are reported to nodulate this legume. However, bean is considered a poor nitrogen–fixing pulse in comparison with other grain legumes (Hardarson, 1993). Sparse nodulation and the lack of response to inoculation in field experiments has frequently been reported worldwide, raising questions about the benefits of inoculation (Graham, 1981;

Buttery et al., 1987). This fact is attributed to intrinsic characteristics of the host plant, particularly the nodulation promiscuity (Michielis et al., 1998), as well as the great sensitivity to other nodulation-limiting factors, such as the high rates of N fertilizer used in intensive agriculture, high temperatures and soil dryness (Graham, 1981).

In Tunisia we showed that P. vulgaris is nodulated by a diversity of species, including Rhizobium gallicum, R. leguminosarum bys. phaseoli and viciae, R. etli, R. giardinii, Sinorhizobium fredii, S. meliloti and S. medicae (Mhamdi et al., 2002). N₂-fixing nodulation was, however, found only in soils routinely cultivated with this legume. In the other soils, nodulation was very scarce or absent (Mhamdi et al., 1999). The nodules recovered from these regions were ineffective at N₂ fixation and caused by non-specific nodulation with S. meliloti, S. medicae and bv. viciae of R. leguminosarum (Mhamdi et al., 2002). The promiscuous nodulation of common beans with non-specific rhizobia is probably the result of low abundance or absence of natural populations of bean-compatible rhizobia in the Tunisian soils. If so, inoculation of common beans with efficient and competitive rhizobial strains may be the means to improve biological N₂ fixation and common bean productivity. However, inoculation trials conducted with the commercial strain R. tropici CIAT899 in the valley of Medjerda failed (Trabelsi, pers. com.).

Since native rhizobia are believed to be better adapted to local environmental conditions than commercial inoculants, we conducted a screening to select for the most efficient and

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competitive strains among indigenous rhizobia. The strain 8a3 of *R. gallicum* isolated from root nodules of common bean cultivated in Cap Bon was selected for its high symbiotic effectiveness (Mhamdi et al., 2002). The competitiveness of strain 8a3 on soil cores, monitored by using the reporter gene *gusA*, showed that this strain was highly competitive when inoculated into the cultivars Coco and Royalnel (Mrabet et al., 2005). The efficiency of utilization of this strain as an inoculant needs further confirmation under field conditions. On the other hand, the low productivity of common bean in Tunisia has led during recent years to the introduction of new cultivars. However, these cultivars were selected on the basis of simple agronomic characteristics and their symbiotic capabilities were generally neglected.

In this work, we aimed to assess the symbiotic effectiveness and competitiveness of strain 8a3 with these new cultivars in on-farm trials.

2. MATERIALS AND METHODS

2.1. Field experiments

On-farm trials were conducted in close partnership with farmers and extension services. They were replicated across farmers' fields but not within fields. Six farmers were selected according to their willingness to cooperate with the experiment (Tab. I).

The experimental design consisted of three treatments, not inoculated, inoculated and fertilized (50 kg of NH₄NO₃/ha). Each treatment was subdivided into three subplots cultivated with three common bean cultivars, Coco, BRB17 and Flamingo. The subplots were 13.5 m \times 6 m in size and contained 30 rows. The sowing density was 20 seeds/m² (10 cm/50 cm). A margin of 2 m separated the inoculated and the non-inoculated plots. The experiment was established in March 2005. The native strain 8a3 of R. gallicum by. gallicum (Mhamdi et al., 2002) was grown to the late exponential phase in yeast-extract-mannitol broth (Vincent, 1970) and diluted to 1/20 with well water (approximately 10⁸ bacteria mL⁻¹). This suspension was used to inoculate the emerging seedlings using a watering can. Each subplot received 10 liters of inoculum along the seedling rows (approximately 5 mL for each seedling). Thirty plants were randomly collected from each subplot and used for nodule counting and shoot dry weight analysis at the flowering stage. Results were submitted to analysis of variance and mean comparison by the HSD test (P < 0.05).

2.2. Characterization of indigenous rhizobia

The abundance of indigenous populations of rhizobia nodulating common bean in each soil was estimated by the most probable number (MPN) method (Vincent, 1970). Native rhizobia were isolated from root nodules of common bean according to standard procedures (Mhamdi et al., 1999). Polymerase chain reaction coupled to restriction fragment

length polymorphism (PCR-RFLP) of 16S rRNA genes was conducted on cells treated with proteinase K using primers fD1 and rD1 (Weisburg et al., 1991) according to the procedure described by Mhamdi et al. (2002). Digestion was performed by five endonucleases, *MspI*, *NdeII*, *CfoI*, *HaeIII* and *RsaI*. Species assignation was done according to 16S rDNA typing and comparison with the published database of mapped restriction sites in the 16S rRNA genes of rhizobia (Laguerre et al., 1997).

2.3. Competitiveness monitoring

Fingerprinting of the repetitive extragenic palindromic sequences amplified by polymerase chain reaction (REP-PCR) was used to monitor the nodulation occupancy by the introduced strain in the on-farm trials. The primers used were those described by Versalovic et al. (1991). Proteinase-treated cells were used as a DNA template for REP-PCR, as previously described (Zribi et al., 2004). The frequency of the REP-PCR pattern of strain 8a3 was estimated in the inoculated and non-inoculated plots for the three cultivars used.

2.4. In vitro antibiosis assay

The overlay plate technique (Oresnik et al., 1999) was used to check the in vitro inhibition of the indigenous rhizobia by the introduced *R. gallicum* strain 8a3. Ten representatives from each indigenous species were tested. A saturated culture (48 h old, OD_{630 nm} > 1) of the indicator strain (indigenous strain) grown in liquid tryptone-yeast extract medium (TY) was diluted to 10^{-3} in 5 mL of soft TY medium (1% agar). The diluted suspension was poured on a solid TY (2% agar) layer in Petri dishes. Two hours later, 3 μ L of a saturated culture of the producing strain (strain 8a3) were inoculated onto the soft agar surface. Inhibition zones were measured after four days of incubation at 28 °C. Each antibiosis test consisted of three replicates.

3. RESULTS AND DISCUSSION

3.1. Diversity of indigenous rhizobia

The density and the species diversity of the indigenous rhizobia nodulating common bean in the non-inoculated fields are given in Table II.

The density of the native rhizobia nodulating common bean was very low or null in the different fields, except in the Ben Mustapha soil. The 16S rDNA typing of field isolates showed that the native rhizobia isolated from the Ben Mustapha field were all assigned to *R. gallicum*. However, in the other soils, where nodulation was poor, a significant proportion of nodules were induced by non-specific nodulation by *S. meliloti* and *S. fredii*. Some *Agrobacterium* strains were also recovered from the Tej soil; however, they failed to re-nodulate their original host when re-examined for nodulation. This phenomenon

Table I. Fields used for inoculation trials in this study.

Field	Climate	Irrigation practice	Cultivation history
Ben Mustapha	Sub-humid	Irrigated	Potato, Common bean, Faba bean, Cereals
Tej	Sub-humid	Rain-fed	Faba bean, Chickpea, Cereals
Nassereddine	Sub-humid	Rain-fed	Cereals
Lazhar	Sub-humid	Rain-fed	Chickpea, Cereals
Mghraoui	Sub-humid	Irrigated	Potato
Mograne	Semi-arid	Irrigated	Experimental field (horticulture)

Table II. Density and species diversity of the indigenous rhizobia nodulating common bean in the different soils before inoculation. The numbers between parentheses indicate the number of samples in which each species occurred.

	Indigenous rhizobia nodulating common bean				
Site	MPN*	Species diversity			
Ben Mustapha	High	R. gallicum (19/19)			
Tej	Low	R. etli (3/19); S. fredii (14/19); Agrobacterium (2/19)			
Nassereddine	Low	S. meliloti (20/20)			
Lazhar	Null	None			
Maghraoui	Low	R. gallicum (11/20); S. meliloti (9/20)			
Mograne	Low	R. etli (2/5); R. gallicum (1/5); S. meliloti (2/5)			

^{*} MPN, Most Probable Number (Vincent, 1970): high, = 10⁶ rhizobia g⁻¹ soil; low, < 10³ rhizobia g⁻¹ soil; null, below the detectable limit.

has been described by different authors (De Lajudie et al., 1999; Mhamdi et al., 2002; Hameed et al., 2004). Mhamdi et al. (2005) showed that these non-nodulating agrobacteria were able to colonize root nodules of common bean. Both rhizobia and agrobacteria co-existed in the infected nodules. The mechanism by which these isolates integrated nodules is still not known. More recently, Mrabet et al. (2006) found that *Agrobacterium* strains isolated from root nodules of common bean specifically reduce nodulation by *R. gallicum* in non-sterile soil samples and in vitro antibiosis assays. The introduced strain 8a3 was tested in in vitro antibiosis assays towards the agrobacteria recovered from Tej soil and was found to be highly sensitive. However, the extent of this impact on the introduced strain 8a3 needs to be evaluated further in field experiments.

3.2. Inoculation yield

The nodule numbers and the shoot dry yields of the three common bean cultivars in the six fields analyzed are given in Table III. Inoculation with *R. gallicum* 8a3 induced a significant increase in nodule number in all the fields and with the three cultivars. This increase was significant even in the Ben Mustapha soil where the density of the native rhizobia was high (10⁶ rhizobia g⁻¹ soil). The nodulation increase was accompanied by a significant enhancement of the shoot dry yield in the different fields and with the different cultivars. More than a twofold increase was obtained in Ben Mustapha, Nassereddine, Lazhar and Mograne with the three cultivars. The shoot dry yields obtained in the irrigated fields, Ben Mustapha, Maghraoui and Mograne, were higher than those

obtained in the non-irrigated fields. The introduced strain 8a3 induced an increase in nodule number even in Tej soil, which contained native populations of agrobacteria colonizing nodules. It seems that the density of these agrobacteria is not high enough to exercise a perceptible inhibition on the nodulation by the introduced strain.

3.3. Competitiveness monitoring

Analysis of nodule occupancy is a useful method for investigating the establishment, competitiveness and persistence of the introduced strains. Many technical approaches have been applied for this purpose. Inoculated strains have been monitored through antibiotic resistance (Ramirez et al., 1998), serological typing (Evans et al., 1996) or melanin production (Castro et al., 2000). Nevertheless, it has been proven that these techniques suffer from inaccuracy caused by instability or cross-reaction with indigenous rhizobia. Mrabet et al. (2005) used gus-labeled transconjugants to assess the competitiveness of the strain 8a3 under laboratory conditions. However, we did not use this gene labeling approach under field conditions because of environmental considerations. We adopted an approach based on the amplification by polymerase chain reaction of the repetitive extragenic palindromic sequences (REP-PCR). This technique has been shown to be highly discriminatory and suitable for evaluating intraspecific diversity of rhizobial strains (De Bruijn, 1992). However, this technique is laborious and expensive and could not be applied to large samples. This analysis was therefore only conducted on three fields, Ben Mustapha, Maghraoui and Mograne. These soils were selected because they contain

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Table III. Effect of inoculation with *R. gallicum* 8a3 in the different fields.

Site	Cultivar	Nodule number		Shoot dry weight (g/plant)			
		T_0	T_{8a3}	T_{N}	T_0	T _{8a3}	T_N
Ben Mustapha ^I	BRB17	21 ^b	35 ^a	4 ^c	2.0 ^b	4.4 ^a	3.5 ^a
	Coco	24 ^b	35^{a}	7 ^c	3.0^{c}	9.7^{a}	7.5^{b}
	Flamingo	21 ^b	58 ^a	5°	$4.0^{\rm c}$	9.7ª	7.5 ^b
Tej	BRB17	3^{b}	11 ^a	0_{p}	2.0^{b}	3.8^{a}	3.2^{a}
	Coco	0_{p}	11 ^a	$0_{\rm p}$	1.9 ^c	3.7^{a}	2.4 ^b
	Flamingo	0_{p}	20 ^a	0_{p}	$2.0^{\rm c}$	5.1a	3.8 ^b
Nassereddine	BRB17	0_{p}	10 ^a	$0_{\rm p}$	1.1 ^b	2.5 ^a	2.5^{a}
	Coco	1 ^b	18 ^a	$0_{\rm p}$	1.5 ^b	3.0^{a}	2.7 ^a
	Flamingo	2^{b}	21ª	0_{p}	$2.0^{\rm c}$	4.7 ^a	3.2^{b}
Lazhar	BRB17	0_{p}	12 ^a	$0_{\rm p}$	0.9^{b}	2.0^{a}	2.1a
	Coco	0_{p}	12 ^a	$0_{\rm p}$	1.1 ^b	4.3a	3.4^{a}
	Flamingo	O_p	17 ^a	$0_{\rm p}$	1.9 ^b	4.7 ^a	4.3^{a}
$Maghraoui^{I}$	BRB17	nd	nd	nd	nd	nd	nd
	Coco	3 ^b	15 ^a	$0_{\rm p}$	11.0^{c}	16.0^{a}	13.0 ^b
	Flamingo	1 ^b	12 ^a	0_{p}	12.0°	23.0^{a}	18.0 ^b
Mograne ^I	BRB17	nd	nd	nd	nd	nd	nd
	Coco	nd	nd	nd	nd	nd	nd
	Flamingo	0.5^{b}	27 ^a	$0_{\rm p}$	4.9 ^b	11.0 ^a	8.9 ^a

¹ This soil received a complement of irrigation according to plant need; T_0 , non-inoculated treatment; T_{8a3} , inoculated with *R. gallicum* strain 8a3; T_N , fertilized treatment. For each field and each cultivar, means with different letters are significantly different (HSD test, P < 0.05).

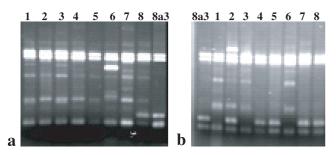


Figure 1. REP-PCR Fingerprinting of nodule isolates from the non-inoculated (a) and the inoculated (b) plots in the soil of Ben Mustapha. Lanes 1–8, different isolates from each plot; 8a3, REP-PCR pattern given by *R. gallicum* strain 8a3. REP-PCR: repetitive extragenic palindromic sequences amplified by polymerase chain reaction.

native populations of rhizobia nodulating common bean of the same species as the inoculant. This situation may inhibit the establishment of the introduced strain and the success of inoculation (Kamicher and Brill, 1986). Figure 1 shows examples of REP-PCR printing of nodule isolates from the inoculated and the non-inoculated plots. The nodule occupancy levels as determined according to the frequency of the REP-PCR pattern of the 8a3 strain are given in Table IV.

The REP-PCR typing of nodules recovered from the non-inoculated plot in Ben Mustapha showed that the REP-PCR 8a3 type was identified in 3/99 of the nodules analyzed. However, in the inoculated treatment, the REP-PCR 8a3 type represented 45/99 of the nodules analyzed, thus giving nodule occupancy of 42%. In the Maghraoui field, where the few nodules observed in the non-inoculated plot were induced by *R. gallicum* and *S. meliloti*, the REP-PCR 8a3 type was found in 2/30 of the nodules. In the inoculated plot, all the nodules tested showed the same REP-PCR 8a3 type. The nodule occupancy by 8a3 was thus higher than 90%. In Mograne soil, very few nodules were obtained with the non-inoculated plants. The REP-PCR 8a3 type was detected only in the inoculated treatment, where it represented 100%.

In vitro antibiosis assays by the overlay plate technique indicated that strain 8a3 exercises an antagonistic activity against all the tested strains of indigenous rhizobia from the species *R. gallicum*, *R. etli*, *S. meliloti* and *S. fredii*.

These results point out the benefits that could be achieved by selecting efficient and competitive strains among natural populations of rhizobia. However, these inoculation trials need to be replicated in separate experiments. The establishment and the persistence of the introduced strain need to be monitored for several years. The antimicrobial activity of this strain may play an important role in interspecific and intraspecific

Site	Cultivar	Frequency of the	REP-pattern	Nodule occupancy by the	
		of R. gallicum	strain 8a3	inoculated strain (%)	
		Non-inoculated	Inoculated	•	
Ben Mustapha	BRB17	2/33	16/33	42	
	Coco	1/33	11/33	30	
	Flamingo	0/33	18/33	54	
	All cvs.	3/99	45/99	42	
Maghraoui	BRB17	nd	nd	nd	
	Coco	0/15	15/15	100	
	Flamingo	2/15	15/15	87	
	All cvs.	2/30	30/30	93	
Mograne	BRB17	nd	nd	nd	
	Coco	nd	nd	nd	
	Flamingo	0/5	30/30	100	
	All cvs.	0/5	30/30	100	

Table IV. Nodule occupancy by the inoculated strain 8a3 of R. gallicum as estimated by REP-PCR profiling under field conditions.

nd, not determined.

competition, and their importance in soil microbial diversity needs to be investigated further.

4. CONCLUSION

Optimal performance of N-fixing symbiosis depends upon pre-selection of both symbiotic partners for adaptation to the target environment, which may in some form present a challenge to rhizobial survival or nodulation. When the inoculant is made from native strains of rhizobia the success of inoculation should be increased, since local strains are better adapted than commercial inoculants. The utilization of *R. gallicum* strain 8a3 as inoculant in different soils demonstrated the ability of this strain to outcompete indigenous populations of rhizobia of the same species and to occupy a high proportion of nodules.

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