dons, however is not affected by chloramphenicol¹⁶ while in rice^{17,18} and maize seedlings¹⁹ it is induced by the antibiotic. Dixit et al.²⁰ reported the inhibition of induction of nitrate reductase by chloramphenical in maize leaves. Such conflicting results exist in the literature. The present study therefore suggests the need for investigating and understanding the effect of chloramphenical directly or indirectly on eukaryotic protein synthesis inhibition.

TABLE 3

Effect of chloramphenicol on cytosolic isocitrate dehydrogenase malate dehydrogenase and isocitrate lyase activities from Aspergillus niger

Conditions of the growth	Isocitrate dehydro- genase*	Malate dehydro- genase*	Isocitrate lyase*	
Control	13.9	556.2	34.8	
With chloram- phenicol (2 mg/ml)	6.6	328.4	18.4	
With chloram- phenicol (4 mg/ml)	5.6	206.7	18.5	

^{*(}units/mg protein)

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REGULATION OF NITROGENASE EXPRESSION IN RHIZOBIUM

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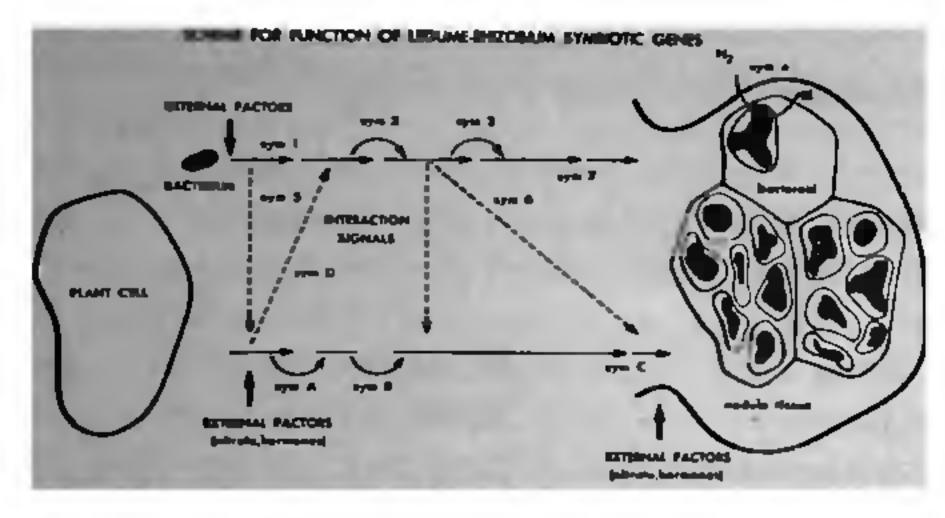
Introduction

The catalytic reduction of atmospheric dinitrogen to ammonia by nitrogenase and its subsidiary enzyme systems (nitrogen fixation), is of immense agricultural importance. Research on biological nitrogen fixation was pioneered by intensive studies of legume-Rhizobium symbioses, since legumes contribute more to the nitrogen economy of the world than any other system. Reductionist efforts to characterise these symbioses have been made with excised root cultures^{1,2}, detached leaves and pods³,

nodules⁴, nodule protoplasts⁵; and isolated bacteroids⁸. Although potentially valuable, the application of these isolated experimental systems is limited because of the enormous complexities involved in the symbiosis. Development of an effective symbiosis comprises a multi-step cascade of events, namely root colonisation, root hair adhesion, infection, nodule initiation and development, bacteroid development and ultimately nitrogen fixation⁷. This requires a tightly co-ordinated reciprocal communication between plant and bacterium^{7,8} as shown in figure 1. The range of non-fixing *Rhizobium* mutants isolated⁹

¹³ December 1982

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TWO DEVELOPMENTAL PATHWAYS OF DIFFERNIATION WITH DIGHTLY COORDINATED RECPROCAL ENTERACTION

Figure 1: A scheme for function of Legume-Rhizobium symbiotic genes. The success of symbiotic process requires the co-ordinated read-off of genetic signals between plant and the bacterium, both being subjected to control by environment. Plants cells undergo differentiation (Sym. A, B etc) for cortical via meristematic to bacteroid filled cells. Meanwhile the vegetative bacterial cells undergo a morphogenesis (Sym. 1, 2 etc) and develop into nitrogen fixing bacteroids.

indicates that each step of the symbiosis is governed by at least one prokaryotic gene. Further complications at different biochemical and physiological levels arise in the study of the nodule, an organ sui generis often plagued by changes in macro and microenvironmental conditions¹⁰. Thus, isolated plant organs have a restricted value for the study of the central features of the symbiosis such as the expression of nitrogenase.

The last decade has seen considerable progress towards an understanding of the more subtle aspects of the nitrogenase regulation. Attempts have been made to mimick the plant-Rhizobium symbiosis in explanta co-cultures of plant cells and Rhizobium. Additionally, under appropriate conditions nitrogenase activity (the final operative phenotype in the root nodule symbiosis previously thought to be expressed only in bacteroids) has been demonstrated in axenic cultures of Rhizobium. Although it is not known yet, whether or not such asymbiotic activity by Rhizobium occurs in the rhizosphere, this discovery was signicant as it permitted the development of the genetics of Rhizobium nitrogenase and its regulation. This article discusses these developments.

IN VITRO SYMBIOSIS

Recent advances in cell and tissue culture opened up new avenues for studying symbiotic nitrogen fixation at the cellular level. Co-cultures of plant callus or cell suspension with rhizobia were initially used as model systems. Progress in this area has been recently reviewed 11,12 and is summarised in table 1.

(a) Infection and nodulation:

Original interest in in vitro co-culture was directed towards its potential as a system to study the infection process. Although many workers demonstrated Rhizobium infection of plant cell cultures to be similar to the in planta situation (table 1), others 20-22 have referred to the phenomena as a superficial, morphological artifact rather than a genetically-controlled developmental process. Furthermore, the behaviour of cultured plant cells does not always correspond to the genetic constitution of the plant from which they were derived. Thus, the conclusions drawn from infection studies made in vitro were anomalous. Ability of Rhizobium strains to interact with legume stems 22 and non-legume derived callus culture⁸⁸ conclusively demonstrated that genetic barriers to in planta nitrogen fixation (infection, nodule initiation, etc.) either were not operable or were bypassed in vitro. The recent discovery of promiscuous Rhizobium strains (belonging to the cowpea miscellany) capable of nodulating legume stems³³ and various non-leguminous tree species (belonging to the *Parasponia* genus)⁷⁷ shows a relaxed stringency of plant—Rhizobium association similar to in vitro studies. These findings have thus helped to dispel the existing dogma that a Rhizoblum strain is capable of nitrogen fixation only when associated with the root of particular legumes defined by its cross inoculation specificity.

(b) Nitrogen fixation in vitro

The capability of Rhizobium strain 32H1 to derepress nitrogenase activity (as measured by acetylene reduction or ¹⁵N₂ incorporation) in the presence of both legume and non-legumes callus and later in the absence of any plant cells elicited two important features: (a) the genetic information for nitrogenase is encoded in the bacterium and (b) that the diffusable but yet nutritionally substitutable substances produced by the plant cells were probably responsible for stimulating Rhizobium nitrogenase activity. Interestingly some of the nutritional and environmental factors controlling success of the in planta symbiosis also regulate expression of nitrogenase in vitro. For example, nitrate, ammonia and glutamine as well as oxygen have repressing effects. Succinate, a carbon metabolite passing from the plant cytoplasm to the bacteroid, was a key compound to supporting high levels of in vitro nitrogenase activity. Further investigations using a trans-filter apparatus (i.e. Rhizobium-plant cell suspension co-cultures)28 extended these initial observations made on separated agar cultures.

Nitrogenase activity was derepressed and/or stimulated in normally non-derepressable or moderately

TABLE 1
Summary of nitrogen fixation studies in cell culture-Rhizobium associations

Associations involved	Aspects of symbiosis studied	References	
1 Rhizobium-legume associations	<u></u>		
(1) R. japonicum-Glycine max	(a) Cellular differentiation	13, 14, 15, 16, 17	
	(b) Intracellular symbiosis	18, 19, 20	
	(c) Nutritional and cultural parameters:	13, 14, 15, 16, 17, 18, 22	
	Hormonal inorganic combined N ₂	15, 16, 17, 18, 23, 24, 25, 26	
	Carbohydrates	17, 27	
	(d) Host symbiont genetic effect	15	
	(e) Stimulatory/inhibitory effect of		
	diffusable factors	17, 25, 28, 29, 44, 45, 46	
	(f) Non-specificity	29. 30	
(2) R. vigna-Arachis hypogaea (cowpea miscellany)	Intracellular symbiosis	31, 32	
Stylosanthes gracilis non-specificity (stem callus used)		33	
Glycine max non-specificity		27, 34	
Pisum sativum	(differential cross inoculation group		
— Trifolium Spp	combinations)		
Vicia hajastana	Calmania and affine a control of the	34	
•	Cowpea Stimulatory effects of diffusable factors		
(3) R. trifolii - Trifolium Organogenesis and cellular different		33	
- T. repens	Stimulatory effects of diffusable factors	36	
(4) R. lupini - Lupinous			
polyphyllus	Stimulatory factors	37	
2. Rhizobium (strain 32H1) - Non-	•		
legume associations			
— Brassica napus	deviation of		
Bromus inermis	Rhizobium - legume	34	
Tritieum monococcum	symbiosis in vitro and stimulatory		
	effects of diffusable factors		
— Nicotiana tabacum		35, 39, 40	
— Nemesia strummusa		35, 39, 40	
Portulaca grandistora		38, 41	
Petunia hybrida		33	
Daucus carota		48	
Triticum aestivum Sorghum nigricans		48	
			

derepressed Rhizobium strains during co-culture with plant cells. This effect was also obtained when vegetative Rhizobium cells were exposed to co-culture supernatant. It was postulated that this phenomenon involved an exchange of molecular signals between plant and bacterial cells. Recently legume cell cultures of soybean, pea, and white clover were shown to respond to certain bacterial substances by synthesiz-

ing dialysable plant factors (also called nitrogenase factors) which were claimed to accumulate in the conditioned plant cell medium (PCM)^{29, 30}, subsequently causing derepression of nitrogenase.

Fractionation of plant cell conditioned medium indicated that only certain fractions were able to stimulate expression of nitrogenase activity in pure cultures of *Rhizobium*. PCM fractionation led to the

postulate by Reporter that *cupper*-containing metallothioneins and peptidoglucans were involved in the in vitro interaction 42. PCM was reported to increase oxidative phosphorylation⁴³. In parallel, the derepression of nitrogenase was apparently associated with the loss of exopolysaccharide (Reporter, personal communication). However, there are many gaps, inconsistencies and experimental weaknesses that cloud the reliability and interpretation of the above mentioned studies and the numerous related investigations by other workers. It remains obscure in the absence of reproducible results, whether PCM components represent complex molecular signals. Whether such 'signals' exist, or if they are a collection of 'single' organic molecules involved in normal cellular metabolism still remains unclear, although the in vitro culture on completely defined media of some strains (e.g. CB756, ANU289 or 32H1) indicated that at least for some strains there was no need for signal complexity.

NITROGENASE IN FREE-LIVING RHIZOBIUM

Although attempts to show asymbiotic nitrogen fixation by rhizobia were first documented in 1945⁴⁹, the conclusive evidence for such activity was not obtained until 30 years later. The reports on in vitro symbiosis between plant cells and rhizobia (table 1) provided the primary impetus to efforts and subsquently discovery of nitrogen fixation in free-living rhizobia 50-52. The demonstration of nitrogen fixation by ¹⁵N₂ incorporation^{51,52} and nitrogenase activity (acetylene reduction) inhibition by specific inhibitors⁵³ has confirmed the early observations. Since then either agar, stationary/shaken liquid or chemostat culture of Rhizobium in defined media has been used to produce nitrogenase activity. To date more than 50 strains (table 2) show the nitrogenase positive phenotype under in vitro conditions.

However, most of these strains belong to slow growing Rhizobium species such as R. japonicum and 'cowpea strains'. Several strains belonging to these species cannot be derepressed for nitrogenase under similar conditions as used for derepressable strains. Since the strains differ with regard to the requirements for expression of nitrogenase activity⁵⁴⁻⁵⁶, it may be that yet optimal conditions necessary for derepression of nitrogenase in these strains have not been found. Alternately, derepression of nitrogenase in the laboratory cultures may be under genetic control. The results of some relatively recent genetic exchange experiments⁵⁷ and DNA: DNA hybridization studies⁵⁸ indicated that strains labelled 'cowpea rhizobia' and R. japonicum may represent at least two and three different sub-species, respectively. Since many strains belonging to one sub-species of R. japonicum were

TABLE 2

Number of strains tested for the ability to reduce acetylene in free living state in various Rhizobium species.

Rhizobium species	Nitrogenase positive		Nitrogenase negative	
	sively	hquid as well as agar	agar	liquid
'Cowpea' strains	12	4	9	_
R. japonicum	8	26	5	13
R. Iupini			4	
R. meliloti		2(?)	4	2
R. trifolii		2(?)		16
R. leguminosarum Parasponia	1(?)		6	
Rhizobium	1	1	14(?)	_

^{-,} not known

nitrogenase positive in vitro⁶⁵ it was thought that nitrogenase derepression in culture may be confined to one sub-species. The organisation of nitrogen fixation genes was reported to be very different in R. japonicum strains USDA110 and 61A76⁷⁵, belonging to the same sub-species (on the basis of DNA homology classification), of which only strain 61A76 is capable of nitrogenase expression in vitro. Thus it appears that the genetic basis for in vitro nitrogenase expression may be strain specific. The precise molecular architecture for such differences between strains remains to be worked out.

In contrast, as shown in table 2 there are only a few unsubstantiated reports in the literature of nitrogenase activity with fast growing Rhizobium species. Demonstration of nitrogenase activity in R. trifolii strain T1, the spectinomycin resistant derivative T1 spec⁶¹ or by exposure to plant cell conditioned medium³⁶ was not repeatable in our laboratory and in that of others (Beringer pers. comm.). Furthermore, other reports of nitrogenase activity in fast growing Rhizobium strains 59,60 are either characterised by a lack of repeatability or proper contamination tests. It is thus our opinion that as yet a repeatable derepression of nitrogenase in fast growing Rhizabium strains such as R. melilotii, R. trifolii and R. leguminosarum under defined or in associated in vitro culture is not demonstratable.

^{(?),} not confirmed results

Several factors influence derepression of nitrogenase in *Rhizobium in vitro*. Some recent reviews 55, 62, 68 have discussed these factors. Here, we discuss some new developments which have not been reviewed previously.

(a) Regulation by Oxygen

Nitrogenase activity on agar cultures was measured under atmospheric oxygen tension (0.2 atm), which was later found to be the optimal concentration⁵². Attempts⁵² to derepress nitrogenase in liquid culture under air were unsuccessful as a consistently low O₂ tension was obligatory for derepression. Oxygen concentrations in the range of 0.06 to 0.36% were effective^{55,64}. Studies in chemostat cultures of strain 32H1 indicated that cultures growing with about 1 micromole dissolved oxygen gave high specific rates. The optimal level of oxygen in the gasphase differed with carbon and nitrogen sources used in the medium Different strains showed variable O₂ optima, all being within a small range generally classified as microaerobic⁵⁶.

It was not known then, whether the lack of activity under aerobic conditions was due to O2 repression of nitrogenase synthesis, O2 inactivation of nitrogenase or auto-oxidation of reduced electron donors. Measurement of antigenically cross-reacting material⁶⁶ in aerated continuous cultures of strain 32H1 showed oxygen repression of nitrogenase synthesis. Exposure to moderate oxygen concentrations (20-30 µM dissolved oxygen tension) of the culture with established nitrogenase activity also resulted in inactivation of nitrogenase. Similar effects of O2 on nitrogenase synthesis were observed by pulse labelling derepressed cells with a 14C labelled amino acid in free-living R. japonicum⁶⁷. In Klebsiella pneumoniae oxygen is thought to interact with the nif LA regulatory protein complex which in turn controls transcription of the nif HDK operon responsible for the synthesis of the Fe and Mo/Fe component protein of nitrogenase. Whether O₂ regulates nitrogenase synthesis in Rhizobium in a manner similar to that observed in Klebsiella remains to be answered.

(b) Regulation by Carbon sources

Rhizobium strains differ in their preference for a carbon source for growth as well as derepression of nitrogenase. Initially a combination of two carbon sources (a pentose sugar, e.g. arabinose and a TCA cycle intermediate, e.g. succinate) was advocated of the derepress nitrogenase activity in strain 32H1. From our work with Parasponia-Rhizobium

strain ANU289, it appears that of the two, succinate seems to play a dominating role in derepression. Comparable results were also obtained using a single carbon source (gluconate or succinate) in strain 32H1 and several R. japonicum strains. However, it must be noted that the later reports utilised the liquid derepression rather than agar culture system as well as different oxygen levels.

The exact role of a carbon source in derepression of nitrogenase is yet not understood. Using an experimental approach that allows the quantitative determination of the *de-novo* biosynthesis of the constituent polypeptides of nitrogenase, the amount of synthesis was shown to be dependent upon the carbon source used in free living R. japonicum⁶⁷. Cells grown on mannitol or glycerol produced only trace amounts of the nitrogenase polypeptides whereas gluconate supported maximal synthesis and whole cell nitrogenase activity.

Since catalytic activity of nitrogenase depends on the supply of reducing equivalents and energy, the carbon source may exert its effect by modulating general metabolism of the cell⁵⁵. A negative correlation between exopolysaccharide production and nitrogenase activity was found in liquid cultures in a survey of 20 Rhizobium strains 65. It was envisaged that nitrogenase and exopolysaccharide (EPS) synthesis compete for energy (in energy limiting microaerobic conditions). Thus in strains, which produce large amounts of EPS, most available energy was utilized in the production of EPS rather than synthesis and maintenance of nitrogenase. EPS synthesis and nitrogenase depression in Parasponia-Rhizobium strain ANU289 (nonmucoid on mannitol containing medium) and its isogenic mucoid derivative strain ANU288^{56,77} supported the above hypothesis⁶⁵. Further studies with inhibitors specific for exopolysaccharides synthesis or further isolation of specific mutants such as strain ANU288 and ANU289 lacking any one of the enzymes necessary for exopolysaccharide synthesis may be helpful in understanding the correlation between EPS synthesis and nitrogenase activity.

(c) Regulation by ammonium

Rhizobium strains with the exception of Sesbania (Dreyfus, pers. comm.) strains in contrast to other free-living nitrogen fixing organisms exhibit the nitrogen fixing phenotype only in a developmental state which is different from their normal vegetative growth state. Attempts to grow Rhizobium on its own fixed nitrogen have been difficult because derepression of nitrogenase occurred only in the presence of a utilizable nitrogen source, such as glutamate, glutamine, ammonium chloride, potassium nitrate, aspartate,

aspargine and casaminoacids. The type and amount of the particular nitrogen sources varies between strains.

For example, inhibition of nitrogenase activity by 10 mM ammonium was observed in agar culture 40,60 at 20% oxygen in the gas phase. In contrast, similar concentration of ammonium had no inhibitory effect on nitrogenase in strains 32H1 and 31-1b-83 in dilute shaken culture under low oxygen tension⁶⁴. Thus a close interaction between oxygen and ammonium concentrations and culture method was thought to be involved. Furthermore, studies with R. japonicum strain 31-1b-83 revealed that the degree of ammonium inhibition was pH dependent and the maximum inhibition was found at the optimum pH for nitrogenase activity⁵⁵. Variable effects of ammonium depending upon the carbon source used in the medium observed in our laboratory indicated that the inhibition may be mediated by a possible metabolic effect. Thus the oxygen and carbon effects can be explained by differential growth and thus depletion. As ammonium effects on nitrogenase activity are strain specific and affected by carbon sources, oxygen levels, culture regime and the presence of other nitrogenous compounds, it is at present difficult to develop a generalised mechanism of ammonium regulation in Rhizobium. In the absence of evidence for lack of polypeptide synthesis, it is not known whether inhibition is due to repression of nitrogenase synthesis or simply a modulation of activity. More recently, some new findings on ammonia assimilation in Rhizobium have been reported. Glutamine (Gln) auxotrophs of Rhizobium strain 32H1 failed to derepress synthesis of nitrogenase both in culture and in planta⁶⁹. The nitrogen fixation (nif) defects in these strains were shown to be the direct result of glutamine auxotrophy as reversion to prototrophy simultaneously recovered nitrogenase derepression ability.

The current model which states that unadenylated GS mediates derepression of nitrogenase is consistent if not an extrapolation by hyperbole, with the regulation of GS by the classical adenylylation cascade established by Stadtman and his colleagues for $E. coli^{n}$, and more recently in Klebsiella⁷⁶. Regulation of other nif operons in K. pneumoniae by the nif LA operon which is subjected to control by products of gln operons is shown in figure 2. Whether the regulatory patterns as shown for Klebsiella also hold true for Rhizobium in general is still unknown. Perhaps arguing against a general concept is the fact that (a) Rhizobium has two GS (GSI+GSII) enzymes, (b) the sequence homology of the regulatory region of K. pneumoniae nif HDK is minimal, compared to that of R. meliloti and R. trifolii and (c) in some Rhizobium

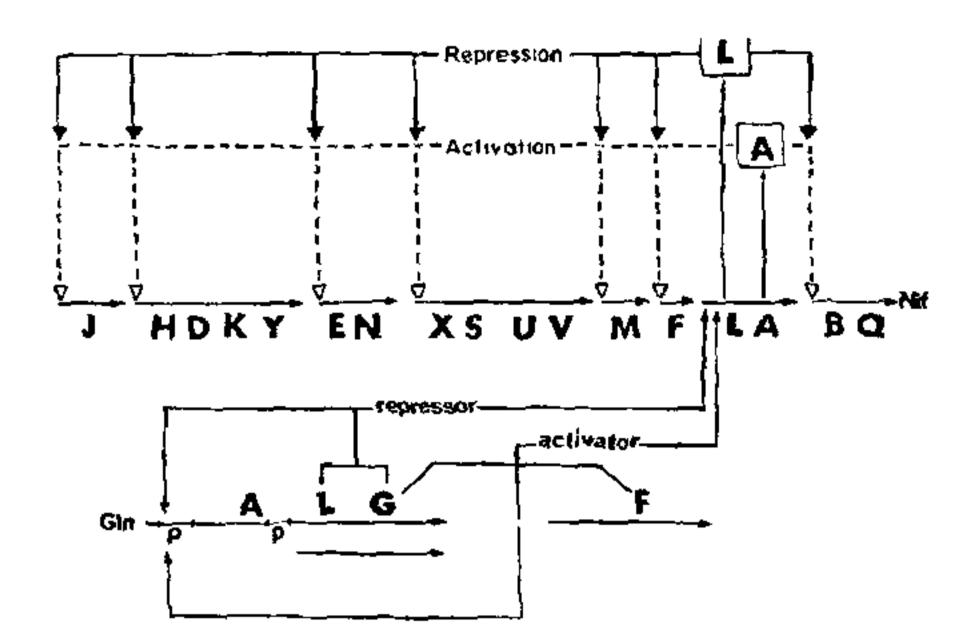


Figure 2: The current model of nif regulation in Klebsiella pneumoniae. General nitrogen control, genes such as gln G(ntr C), gln F(ntr A) and gln L(ntr B) are involved in regulation of nif. gln F product activates gln G, which in turn activates transcription from nif LA promoter. gln L product either by itself or in combination of gln G repress nif LA transcription. In nif cluster, nif L and nif A gene act respectively to repress (solid line) or activate (dotted line) all other nif operons.

ANU289) the nifH (Fe-protein) and nifD (component of the Mo-Fe protein) are not in the same transcriptional unit as they are either in K. pneumoniae or R. trifolii and R. meliloti. (Scott and Shine, Personael Communication).

Before one builds models for Rhizobium nitrogenase regulation based on Klebsiella data, it is essential that more fundamental Rhizobium work is carried out.

CONCLUDING REMARKS

In essence, the so-called reductionist approach using in vitro co-cultures of plant callus-cell suspension with rhizobia has not provided such a similified experimental system compared to the intact nodule as was formerly anticipated. The approaches outlined however permitted an elaboration of factors controlling nitrogenase activity in vitro and ultimately led to the discovery that the genes for nitrogenase are encoded by the Rhizobium genome. Additionally, the derepression of nitrogenase activity in free-living Rhizobium has undoubtedly initiated investigations aimed towards the understanding of several aspects of regulation of nitrogenase and related assimilatory enzymes.

Understanding the expression the Rhizohium nitrogenase, has recently acquired a commercial character stimulated by the need to increase protein

production and to reduce the use of expensive inorganic fertilizers on a world-wide basis. The advent of new recombinant DNA techniques has advanced our knowledge of *Rhizobium* participation in symbiosis. For example some of the genes responsible for both nodulation and nitrogen fixation processes have been identified and cloned ⁷²⁻⁷⁴. The physical mapping of nif genes in *Rhizobium* is currently undertaken in several laboratories. Whether our increased understanding of the molecular biology, genetics and biochemistry of nitrogenase and symbiotic nitrogen fixation actually will result in increased crop production or just constitute a further indepth analysis of a complex developmental process is beyond the scope of this paper.

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