

( )

ACC (IAA)

\*

( / / : / / : )

× cfu.ml<sup>-1</sup>  
( ) ( IAA) ACC  
( ) IAA+++ ACC  
IAA+++ ACC+++

(P<0.001)

Sm Rlv Rlp IAA IAA

IAA Bj Rlt  
-ACC+++ & IAA+++

( -ACC-- ) IAA+ +

(P<0.001)

( )

ACC (IAA) PGPR :

( )

( mg kg<sup>-1</sup>)  
. ( )

PGPR

( ) ( )  
. ( )

Auxein

IAA (PGPs)  
IAA ( )

( L-TRP, ) -

IAA ( )  
IAA L-TRP ( )

.... (IAA) /  
. ( )

IAA  
. ( ) L-TRP

(C2H4) )  
. ( ) / ( ) ( L-TRP  
(. ( ) (IAA)  
. ( )  
L-TRP ( / × mg kg<sup>-1</sup> )

.( )

---

1. L-Tryptophan

... ACC (IAA) :  
 PGPR (ACC) ACC  
 ( ) ( )  
 : (SAM) - -  
 SAM  
 - ACC  
 ACC ( ) ACC SAM  
 ACC  
 .( )  
 rhizobium  
 orhizobium Mesorhizobium  
 .( )  
 ACC (PGPR)  
 EMB ml  
 °C ACC ( )  
 ACC  
 .( ) IAA ( )  
 ) ( )  
 °C ( ) (PGPR)  
 ACC  
 ( OD )  
 (1100,USA; )  
 - CFU ) ( )  
 x cfu ml

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1. Aminocyclopropane-1-Carboxylic acid:ACC  
 2. Yang Cycle  
 3. Adenyl L-Methionine Synthetase: SAM

1. Aminocyclopropane-1-Carboxylic acid:ACC
  2. Yang Cycle
  3. Adenosyl-L-Methionine Synthetase: SAM-Synthetase

( ) (IAA)  
 (CD) (HD)<sup>3</sup>  
**IAA**  
 ( HD/CD )  
 ( ) IAA  
 IAA ( )  
 ( / / / / / )  
 / × /  
 ( )  


---

 IAA  


---

 L.B mM LB  
 ) IAA (LB )  
 ( IAA (LB)  
 ( ) IAA  


---

 mm LB  
 μl IAA  
 ( / )  
 mm  
 ( )  
 IAA  
 /  
 ( )  
 ACC IAA  
 /  
 ( )  
 RMM IAA  
 RMM  


---

- 3 . Hllo Diameter
4. Colony Diameter
- 5 ..Rhizobia Minimal Medium (RMM)

- 
1. Luria – Bertani
  2. Salkofski

... ACC	(IAA)	:	
) p0B0	(P1,P0)	( / M) ACC	$\mu$ l
(N.C)	(	.	
	) P1B0	$\mu$ l	RMM
	(		( / M) NH4Cl
	(P.C)		
	B3		(PC)
-ACC-	IAA+++		RMM
B6		(NC)	(NH4Cl) ACC
IAA-	_ACC++		

---

ACC	ACC	ACC	$^{\circ}$ C
	( )		
	(+) +		
	(++) +		
	(+++) +		RMM + ACC
	(++++) +	(RMM)	(RMM + NH4Cl)

ACC		
( )		
	( )	
		PGPR

SC647  
RCBD Contrast SAS V 6.12

( % )	( B6 B1)
	( B5 B1)

IAA

---

1 .Positive Control(P.C)  
2 .Negative Control(N.C)

IAA

(IAA)

LB

IAA

(% / )

PGPR

IAA++

( )IAA

IAA+ ++

IAA

(% )

LB-TRP

LB

1

NaCl

PGPR

-PGPR								
ACC	P-Solubilization	Sid	IAA	HCN	%S.E			
						( )		B0
+4		3			121	R125 Sm		B1
+2.7	403					R419 Mc		B2
-			2.7			R254 Rlp		B3
-				5		R340 Rlv		B4
+4	368	1.25	2.7			R320RIV	(Mixed)	B5
+2	302	1.25	1.33			R307Rlp		
-	315.6	1.33	1.08	5		R375 Mc		
-		3.8	1.25			R490 Rlv		
+3					121	R164 Sm		B6

HCN

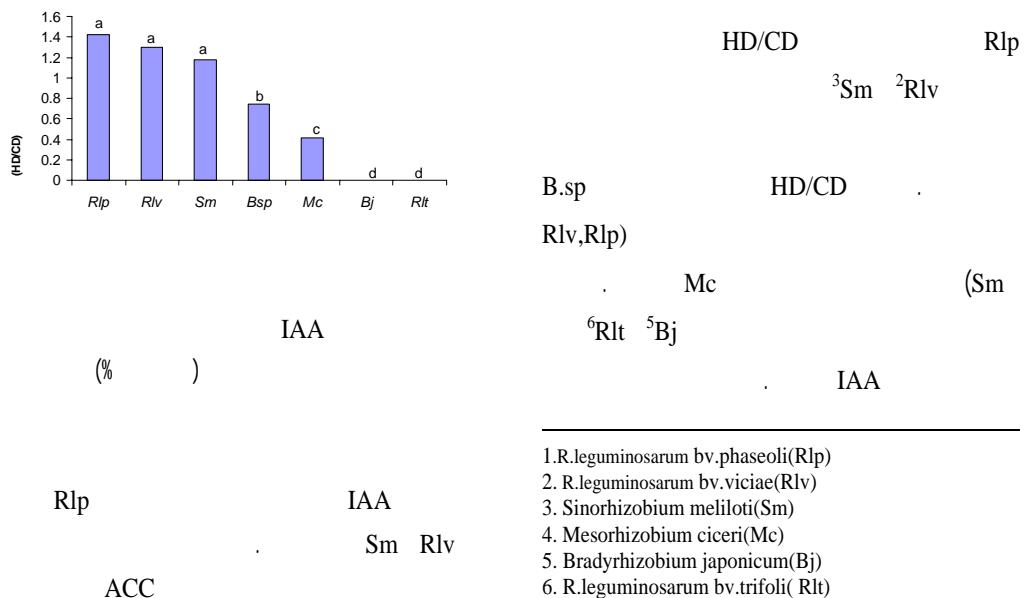
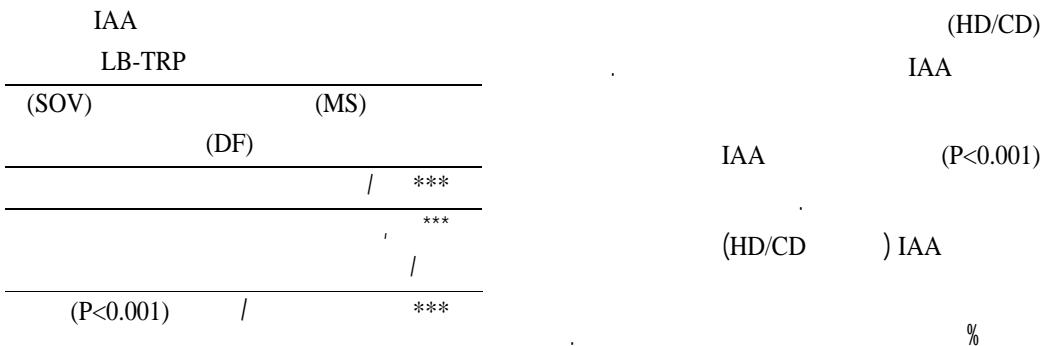
IAA

(mg l<sup>-1</sup>)

ACC

... ACC (IAA) :

LB-TRP		LB	IAA+			IAA			
( )				(%)		0	1	2	3
R.leguminosarum	bv.phaseoli			/	/				
	bv.viciae			/	/				
	bv.trifolii								
				/	/				
Sinorhizobium meliloti				/					
Mesorhizobium ciceri (& M.mediterraneum)				/	/				
Bradyrhizobium japonicum									
B. spp(groundnut)					/				
				/					



## PGPR

## ACC

		ACC +	
( )			(%)
R.leguminosarum	bv.phaseoli		/
	bv.viciae		/
	bv.trifolii		
			/
Sinorhizobium meliloti			
Mesorhizobium ciceri (& M.mediterraneum)			
Bradyrhizobium japonicum			
Bradyrhizobium. spp(groundnut)			
			/

## ACC

## PGPR

)  
ACC  
( )  
ACC  
ACC (%) / )

% % Mc Sm

ACC+++

Rlv Rlp

## ACC

ACC (P<0.001)

## SANDY LOAM

16.0	%Clay
20.2	% Silt
63.8	% Sand
7.88	pH
0.83	ECe (dS m <sup>-1</sup> )
23.5	SP
17.13	%F.C
0.49	%O.M
4.5	Ca2+ (mel -1)
0.6	Mg2+ (mel-1)
2.0	Na+(mel -1)
0.13	K+(mel -1)
8.47	% CCE
0.081	%N(total)
887.2	P-total (mg/kg)
2.52	P-ava.( mg/kg)
3.6	* Fe-ava.( mg/kg)
150	**PSM(Cells gr-1soil)
31	S.meliloti (Cells gr-1soil)

## ACC

## (PGPR)

(SOV) (DF) (MS)

/ \*\*\*

/ \*\*\*

(P<0.001)

/

\*\*\*

DTPA \* Phosphate Solubilizing Microorganisms \*\*



.( )

ACC

( PGPR )

.( )

PGPR

)  
( " "

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