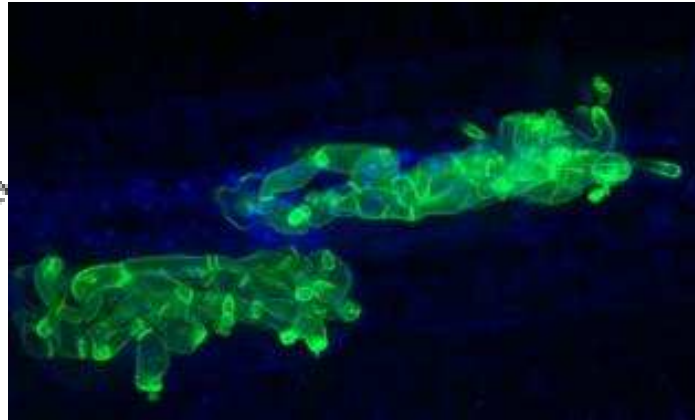


**Problématique:** comprendre les interactions à l'échelle moléculaire pour construire des résistances plus durables des céréales aux champignons phytopathogènes



*Magnaporthe oryzae*



Pyriculariose

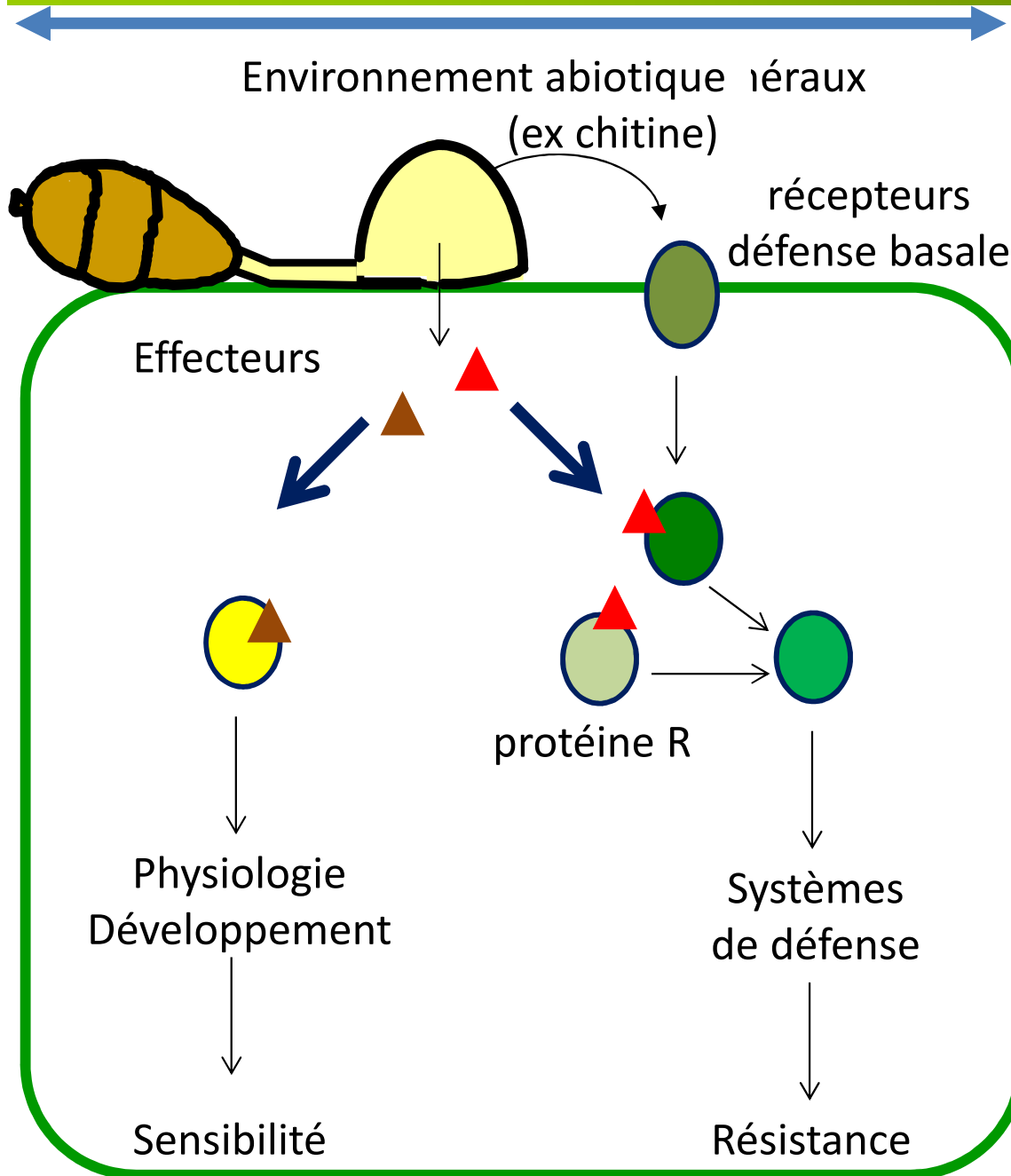


## La pyriculariose

Principale maladie du riz  
Emergent sur blé en Amérique du Sud

## Un couple modèle pour l'étude des interactions

apports à un système d'importance économique  
connaissances transférables à d'autres céréales



## Comment le pouvoir pathogène se construit-il?

### Axe 1: Effecteurs et cibles

- Quels effecteurs fongiques?
- Quelle contribution à l'agressivité?
- Quelles fonctions dans la plante?

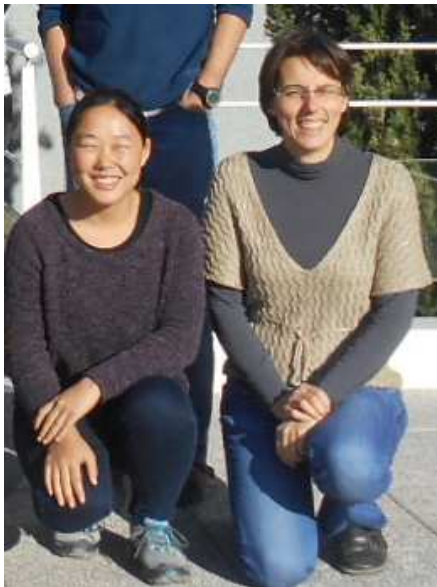
### Axe 2: Mécanismes de résistance

- Comment les effecteurs sont-ils reconnus?
- Quels régulateurs en aval?

### Axe 3: Effets de l'environnement abiotique

Comment l'environnement affecte-t-il le fonctionnement de ces systèmes?

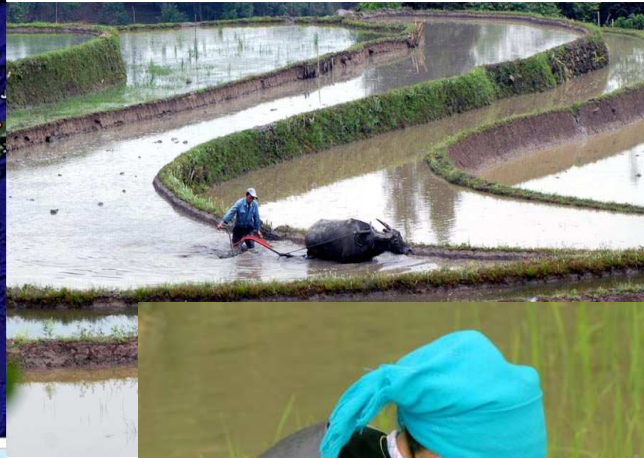
# Survey of the Durability of YuanyangTerrace



Jingjing Liao, Isabelle Meusnier, Huichuan Huang, Xiahong He, Youyong Zhu, Elisabeth Fournier, Didier Tharreau, Thomas Kroj, Jean-Benoit Morel



# The yuanyang terraces, Yunnan



~ 6500 hectares

Acuce genotype for more than 110 years

Yield ~ 5-7 t/ha

Traditional varieties

No chemical fertilizer

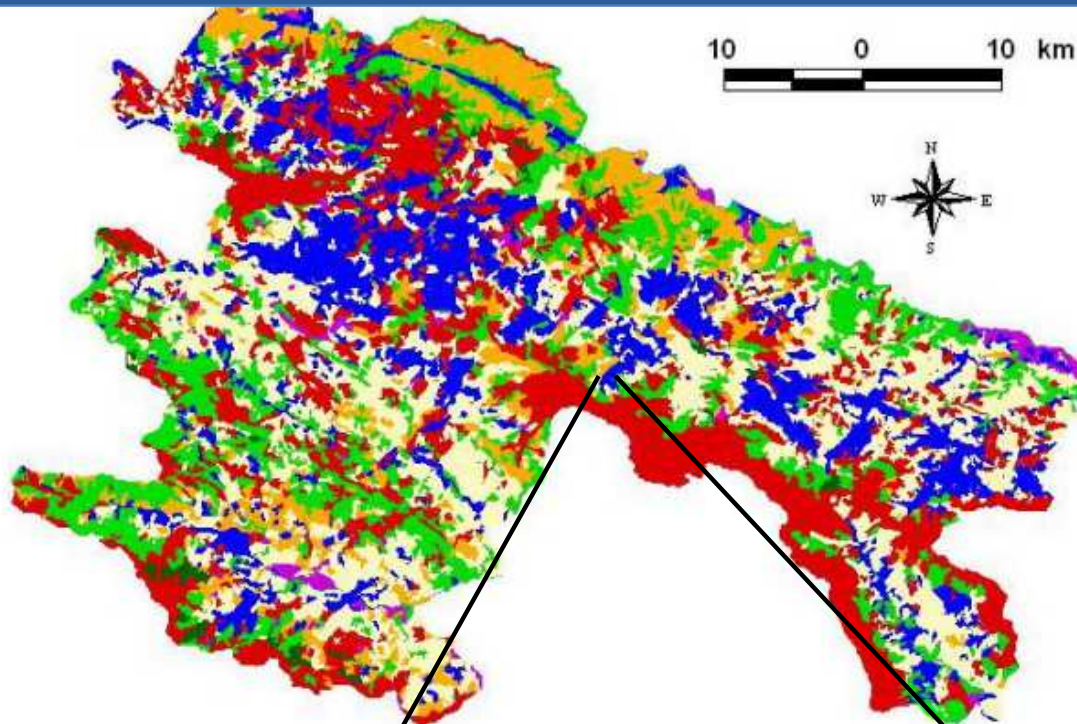
High microbial diversity in the soil

No fungicide

Area close to diversity center of *M. oryzae*



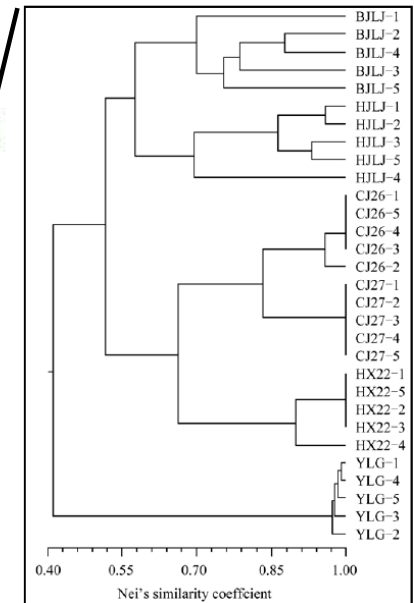
# The yuanyang terraces: diversity at all levels



Cao et al, 2012



Gao et al, 2012



Trad.  
modern  
Acuce

> 30 varieties

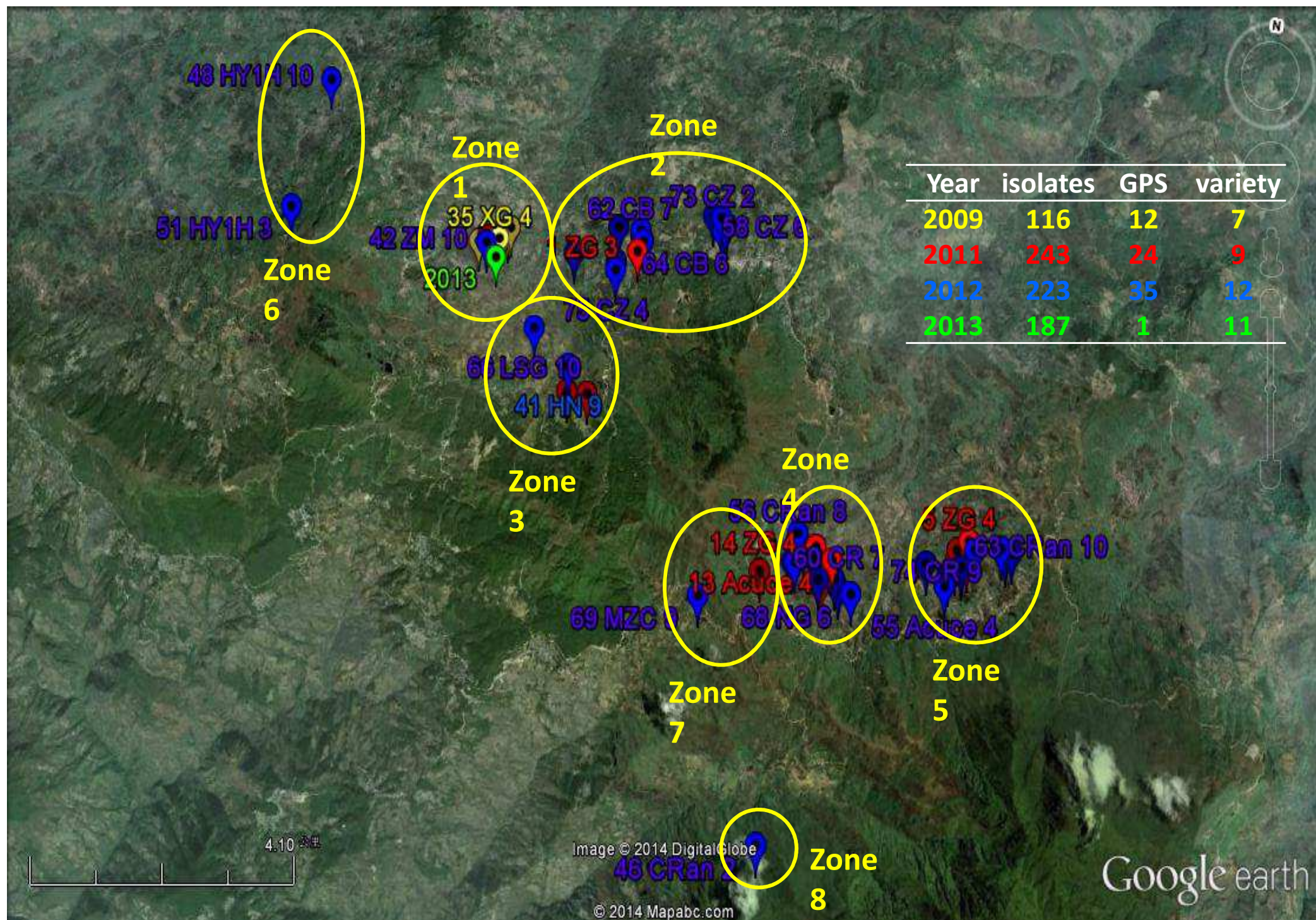
Mixtures

Glutinous and non-glutinous



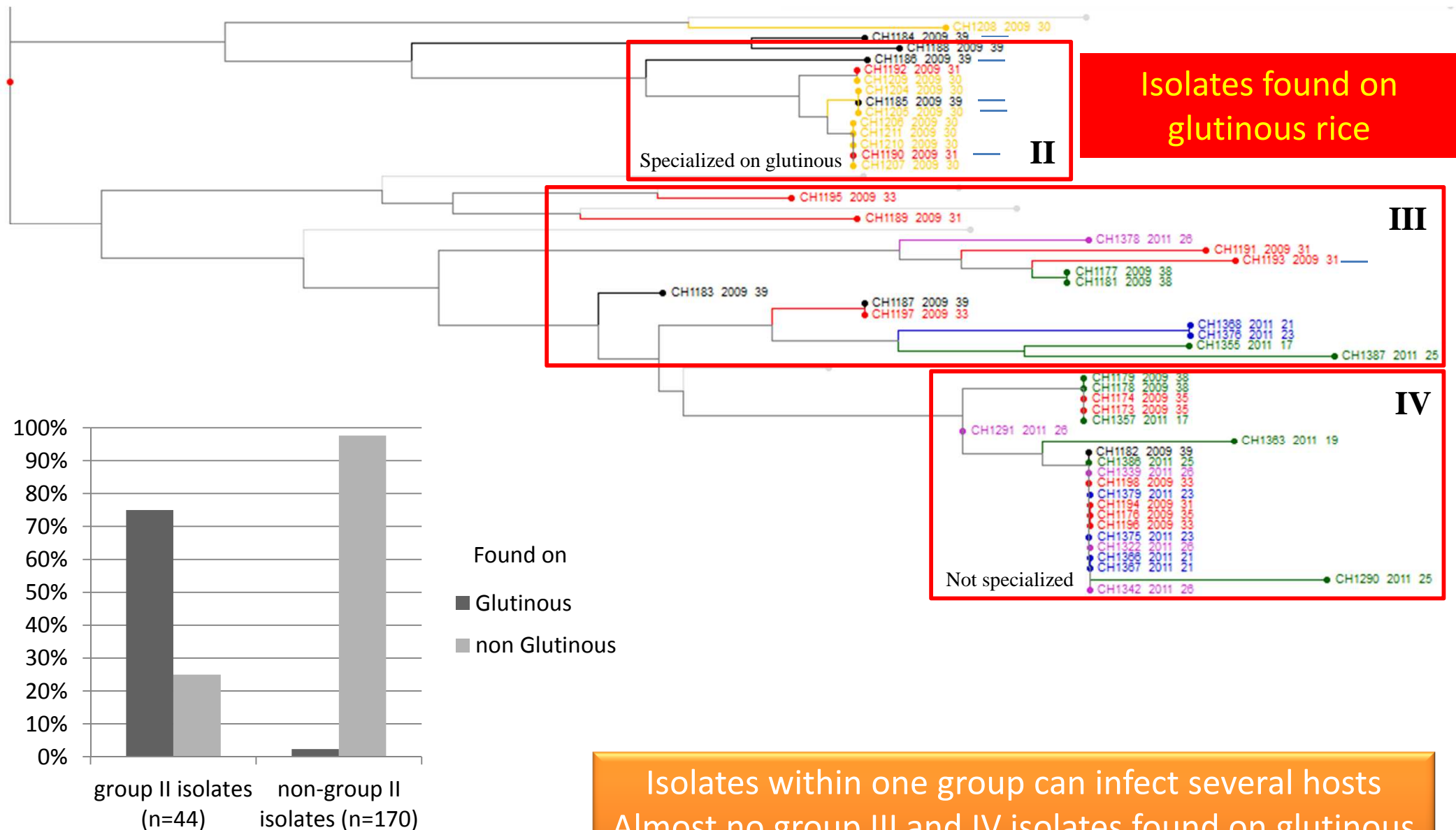


# Sampling *Magnaporthe* in Yuanyang terraces



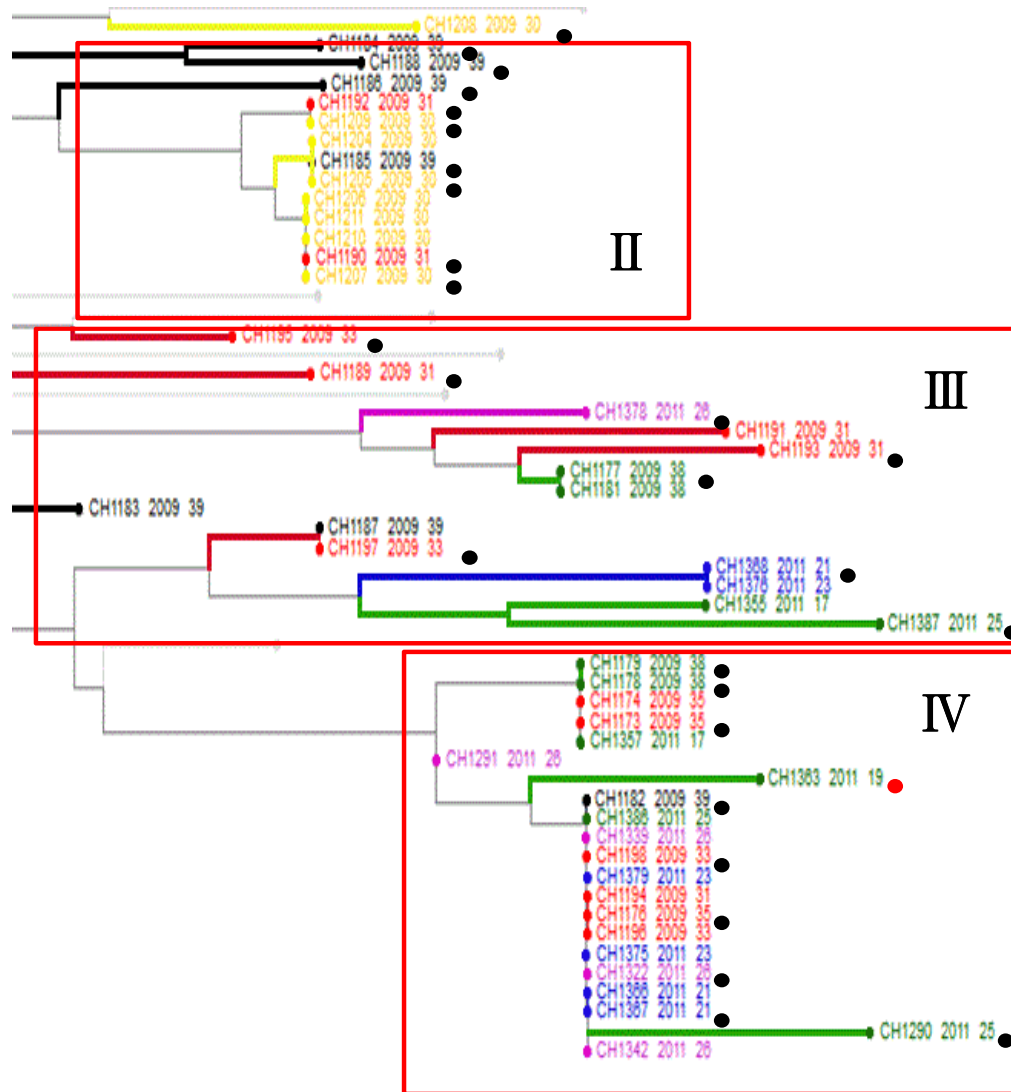


# Microsatellite analysis of 54 YYT isolates and 10 reference isolates



Isolates within one group can infect several hosts  
 Almost no group III and IV isolates found on glutinous  
 Most isolates of group II are found on glutinous

# Group II isolates show reduced virulence on Non-Glutinous rice



	Glutinous		Non-Glutinous			
	HPN	NG	XG	LJG	AZG	Acuce
CH1184	S	S	R	R	R	R
CH1188	S	S	R	R	R	S
CH1186	S	S	R	R	R	R
CH1192	S	S	R	R	R	R
CH1209	S	S	S	S	R	S
CH1185	S	S	R	R	S	R
CH1205	S	S	R	R	R	R
CH1190	S	S	R	R	R	R
CH1207	S	S	R	R	R	R
CH1195	S	S	S	S	R	R
CH1189	S	S	S	S	S	S
CH1378	S	R	S	S	S	S
CH1193	S	S	S	S	S	S
CH1177	S	S	S	S	S	S
CH1197	S	S	S	S	S	S
CH1368	S	S	S	S	S	S
CH1387	S	S	S	S	S	S
CH1179	S	R	S	S	S	S
CH1178	R	R	S	S	S	S
CH1173	S	S	S	S	S	S
CH1383	S	R	S	S	S	S
CH1182	S	S	S	S	S	S
CH1198	S	S	S	S	S	S
CH1176	S	S	S	S	S	S
CH1322	S	R	S	S	S	S
CH1367	S	R	S	S	S	S
CH1290	S	R	S	S	S	S

Group II isolates resemble specialists  
Group III and IV isolates seem generalists  
Glutinous rice is a universal host

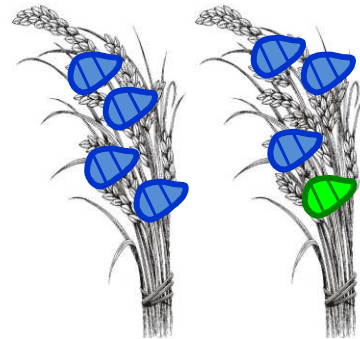


# Unbalanced situation between glutinous and non-glutinous rice

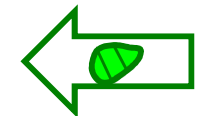
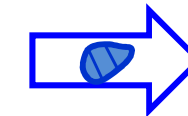
	HPN	ND	XD	LJB	A29	Acute
CH1184	S	S	R	R	R	S
CH1185	S	S	R	R	R	S
CH1186	S	S	R	R	R	R
CH1187	S	S	R	R	R	R
CH1188	S	S	R	R	R	R
CH1189	S	S	R	R	R	R
CH1190	S	S	R	R	R	R
CH1191	S	S	R	R	R	R
CH1192	S	S	R	R	R	R
CH1193	S	S	R	R	R	R
CH1194	S	S	R	R	R	R
CH1195	S	S	R	R	R	R
CH1196	S	S	R	R	R	R
CH1197	S	S	R	R	R	R
CH1198	S	S	R	R	R	R
CH1199	S	S	R	R	R	R
CH1200	S	S	R	R	R	R
CH1201	S	S	R	R	R	R
CH1202	S	S	R	R	R	R
CH1203	S	S	R	R	R	R
CH1204	S	S	R	R	R	R
CH1205	S	S	R	R	R	R
CH1206	S	S	R	R	R	R
CH1207	S	S	R	R	R	R
CH1208	S	S	R	R	R	R
CH1209	S	S	R	R	R	R
CH1210	S	S	R	R	R	R
CH1211	S	S	R	R	R	R
CH1212	S	S	R	R	R	R
CH1213	S	S	R	R	R	R
CH1214	S	S	R	R	R	R
CH1215	S	S	R	R	R	R
CH1216	S	S	R	R	R	R
CH1217	S	S	R	R	R	R
CH1218	S	S	R	R	R	R
CH1219	S	S	R	R	R	R
CH1220	S	S	R	R	R	R
CH1221	S	S	R	R	R	R
CH1222	S	S	R	R	R	R
CH1223	S	S	R	R	R	R
CH1224	S	S	R	R	R	R
CH1225	S	S	R	R	R	R
CH1226	S	S	R	R	R	R
CH1227	S	S	R	R	R	R
CH1228	S	S	R	R	R	R
CH1229	S	S	R	R	R	R
CH1230	S	S	R	R	R	R

why do group II isolates are not virulent on non-glutinous rice?

Glutinous rice

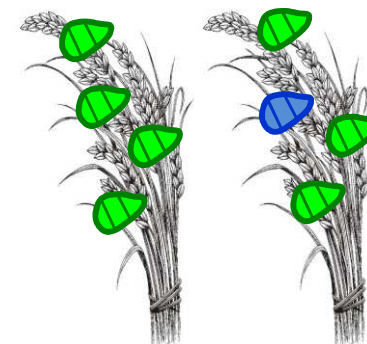


difficult

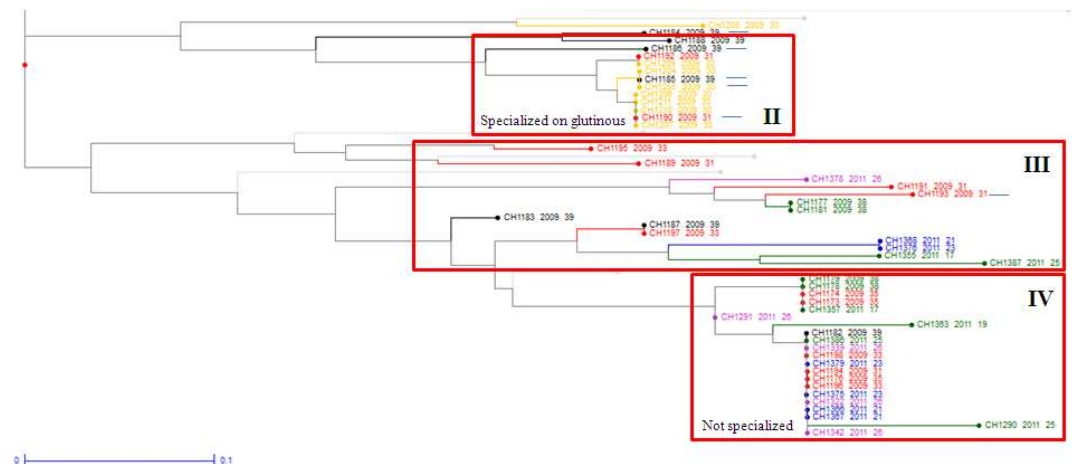


possible but...

Non-Glutinous rice



why so few non-group II isolates on glutinous rice?



# Group II isolates possess many *Avr*-genes

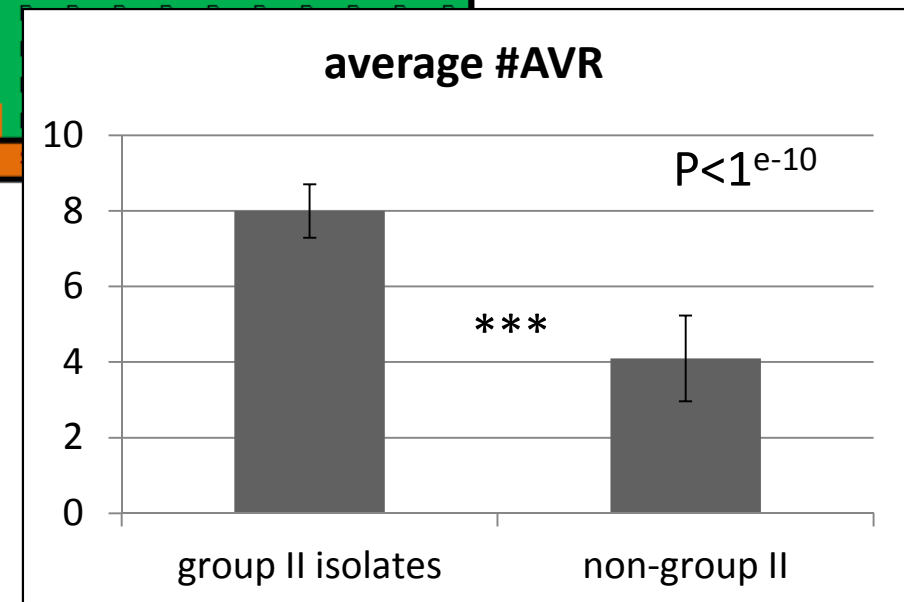
	Group II isolates									Non-Group II isolates																					
	CH1184	CH1188	CH1186	CH1192	CH1209	CH1185	CH1205	CH1190	CH1207	CH1208	CH1180	CH1368	CH1195	CH1189	CH1378	CH1193	CH1177	CH1197	CH1368	CH1387	CH1179	CH1178	CH1173	CH1383	CH1182	CH1198	CH1176	CH1322	CH1367	CH1290	
a+19	R	R	R	R	S	R	R	R	R	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
R-CO39	R	R	R	R	S	R	R	R	R	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
ta	R	R	S	S	R	R	R	R	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
1	R	R	R	R	R	R	R	R	R	S	S	S	R	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
3	S	R	R	R	S	R	R	R	R	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
i+ks	R	R	R	R	R	R	R	R	R	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
ks	R	R	R	R	R	R	R	R	R	S	S	S	S	S	R	S	S	S	S	S	R	R	S	S	S	S	R	R	R	R	S
kp	R	S	R	R		R			R	S	S	S	R	S	R	S	S	S	S	S	S	S	S	S	S	R	S	S	R	S	S
km	R	R	R	R	R	R	R	R	R	S	S	R	R	R	S	S	S	S	R	S	S	S	S	S	S	R	S	S	S	S	S
kh	R	R	R	R	R	R	R	R	R	S	S	R	R	S	S	S	S	R	R	R	R	S	S	R	S	S	S	R	R	S	S
k	R	R	R	R	R	R	R	R	R	S	S	S	R	S	S	S	S	S	S	S	R	R	S	S	R	R	R	R	R	R	S
z+sh	S	S	S	S	S	R	S	S	S	R	R	R	S	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R
z	R	S	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R
zt	R	R	R	R	R	R	R	R	R	R	R	R	S	S	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R
b	R	R	R	R	R	R	R	R	R	R	R	R	S	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R
9	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R
33	R	R	S	R	R	S	R	S	R	S	S	R	S	S	S	S	S	R	R	R	R	R	R	R	R	R	R	R	R	R	R
2=z5	R	R	R	R	R	R	R	R	R	R	R	R	R	S	R	R	R	S	S	R	R	R	R	R	R	R	R	R	R	R	R
ta <sup>2</sup>	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	S	R	R	R	R	R	R	R	R	R	R	R	R	R
t	R	R	R	R	R	R	R	R	R	S	R	R	S	S	R	R	R	R	R	S	R	R	R	R	R	R	R	R	R	R	R
f	S	S	S	S	S	S	S	S	S	S	S	S	S	S	R	S	R	S	S	S	S	S	S	S	S	S	S	S	S	S	S

10

averag

14 R-genes tested

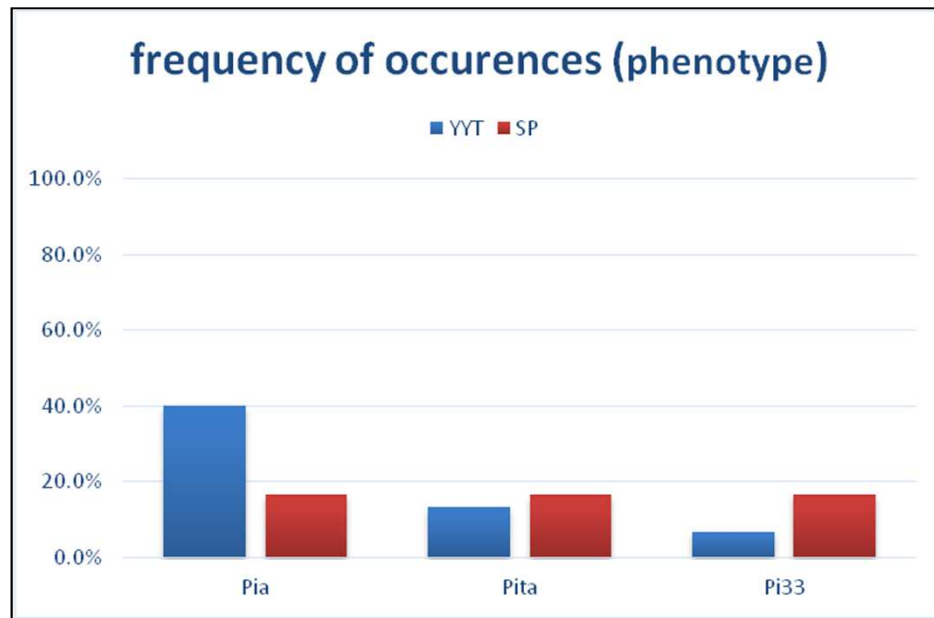
10 R-genes different between group II and other





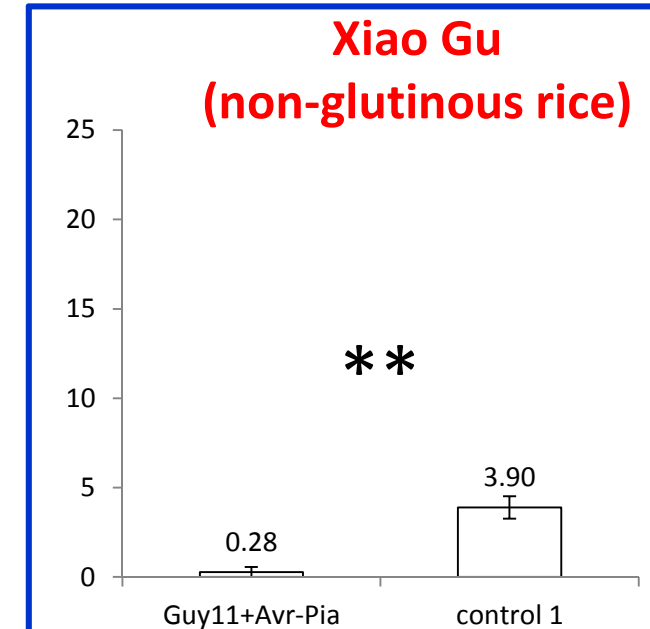
# R-genes deployed in Yuanyang terraces

- 22 R-genes found in more than 50 varieties
- Pik, Pi-ta, Pib and Pi-a found with high frequencies

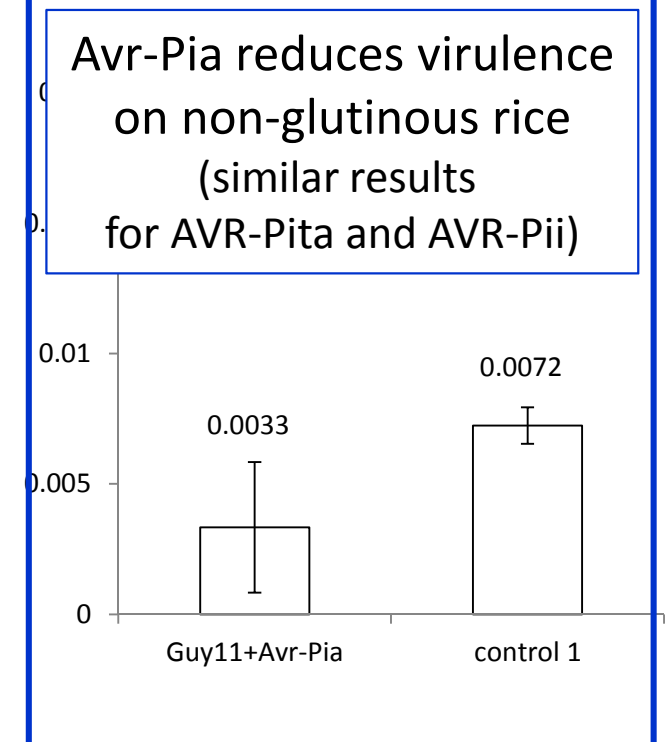


Transgenic isolates

Number of  
Susceptible lesions

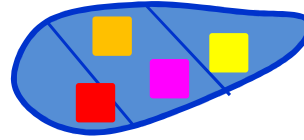


Surface per Susceptible  
lesions (cm<sup>2</sup>)



# Unbalanced situation between glutinous and non-glutinous rice

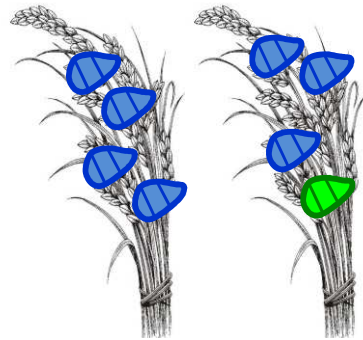
AVR genes



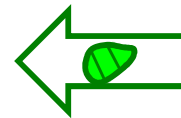
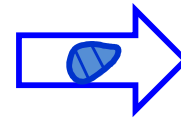
Probably because of R-AVR interactions  
since group II isolates have a large AVR complement

why do group II isolates are not aggressive on non-glutinous rice?

Glutinous rice

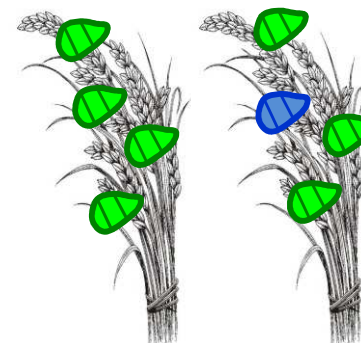


difficult



possible but...

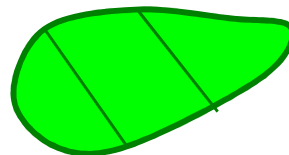
Non-Glutinous rice



R genes

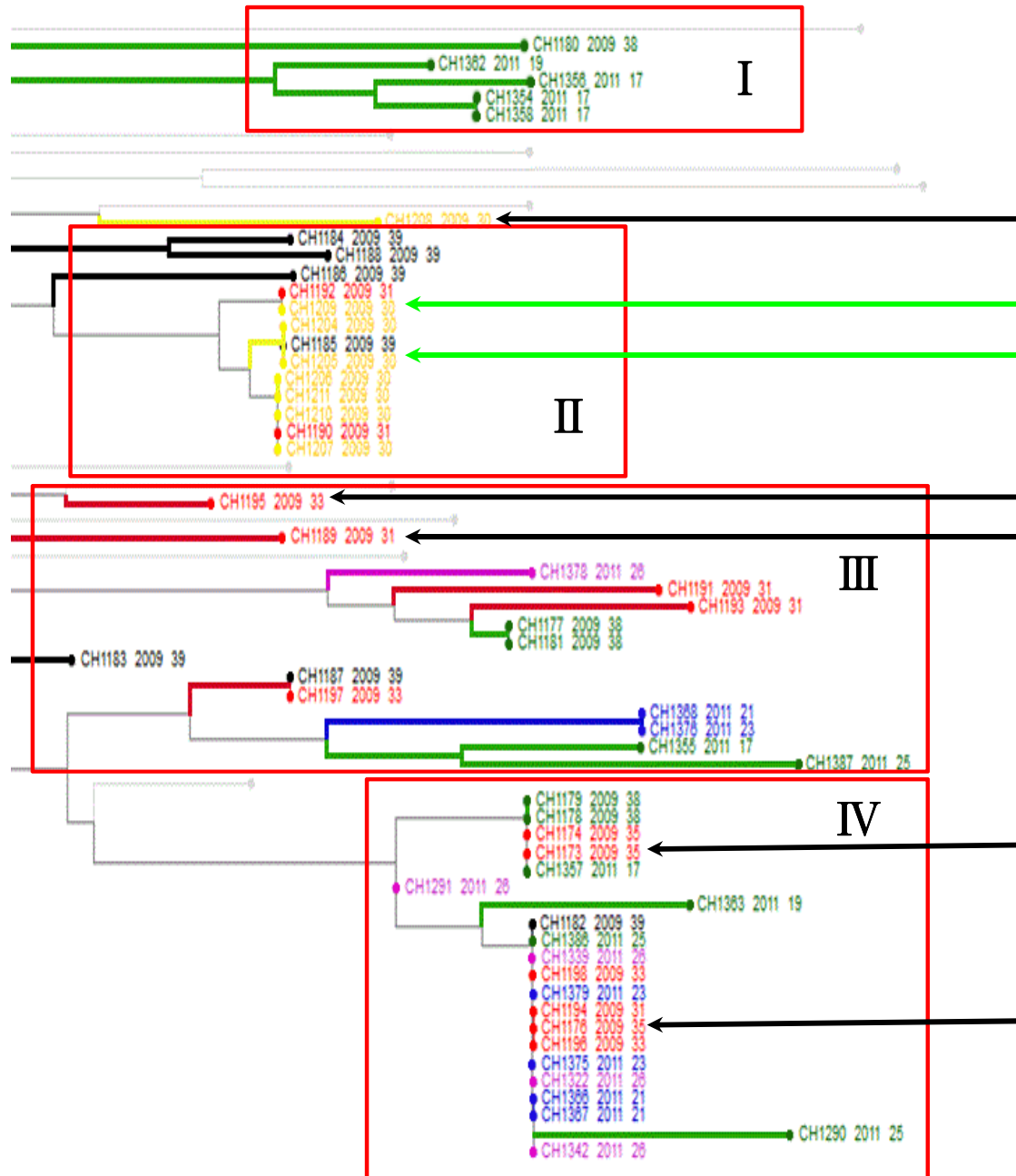
why so few non-group II isolates on glutinous rice?

HYP: A large AVR/Effector complement is required on glutinous rice





# Selection of isolate showing increasing number of AVR genes



CH1208: Avr-gene: 2

CH1209: Avr-gene: 7

CH1205: Avr-gene: 9

CH1195: Avr-gene: 3

CH1189: Avr-gene: 1

CH1173: Avr-gene: 4

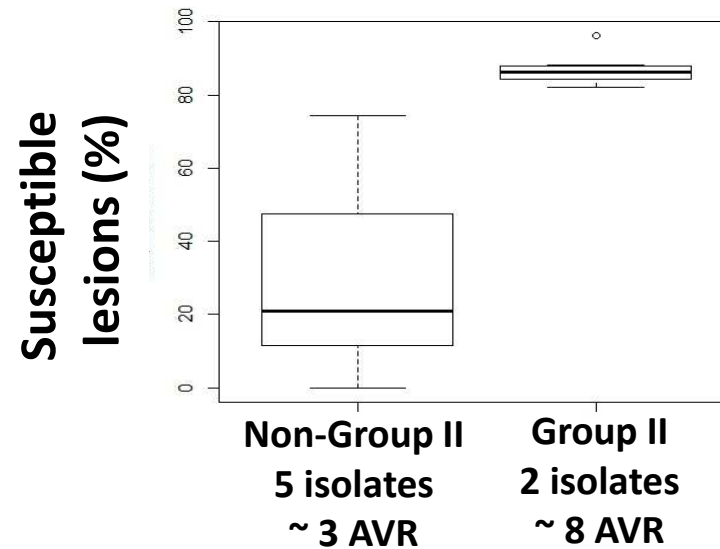
CH1176: Avr-gene: 5

**Glutinous rice:**  
Huangpinuo  
Nuogu

**Non-Glutinous rice:**  
Maratelli  
Nipponbare

# AVR number and disease severity on glutinous rice

## Nuo Gu



Non- Group II isolates



1 1 1 1 1 1 2 2 2 2 2 2 2 2 2 3  
4 5 6 7 8 9 0 1 2 3 4 5 6 7 8 9 0

Group II isolates

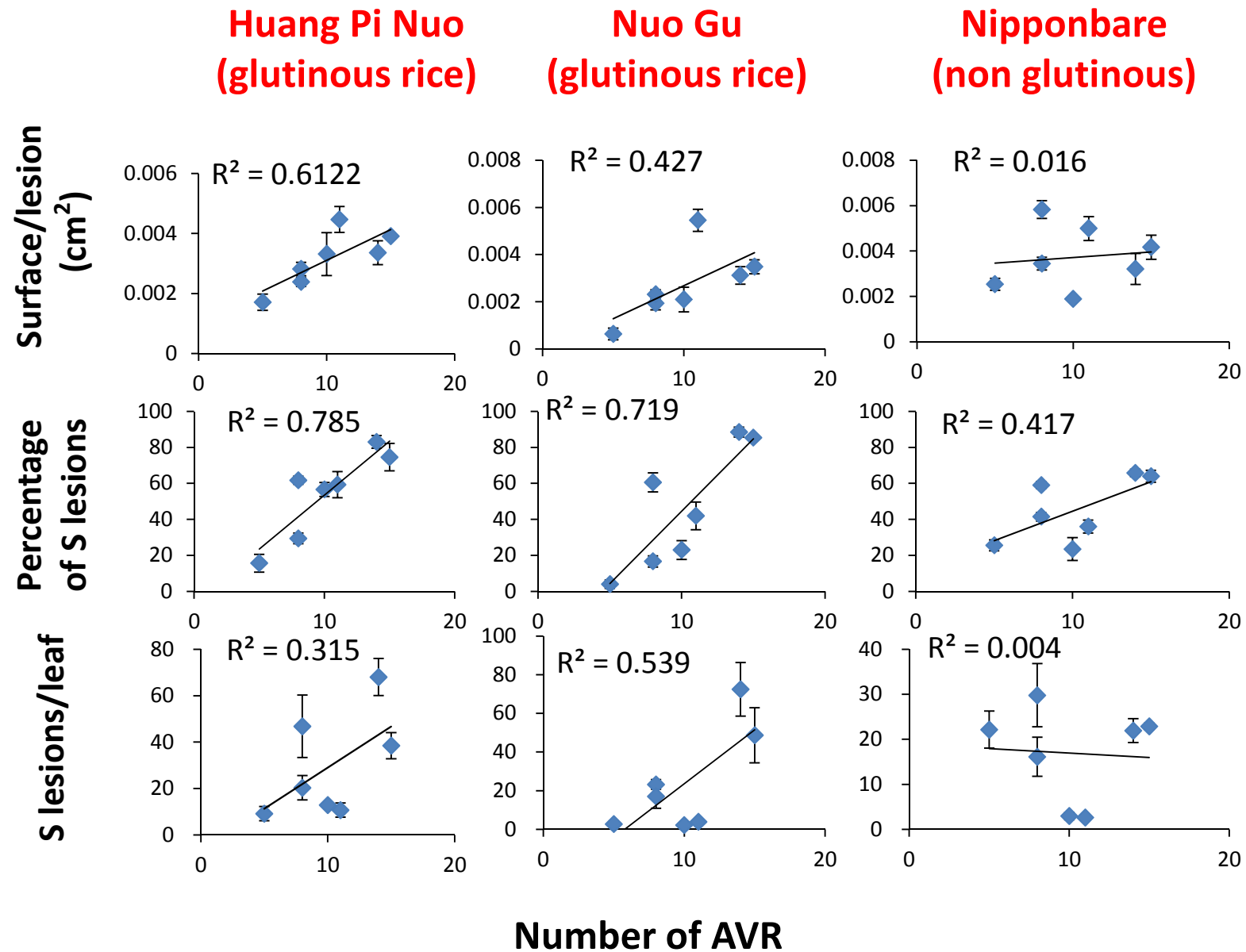


2 3 4 5 6 7 8 9 1 1 1 1  
0 1 2 3

Same results with Huangpinuo



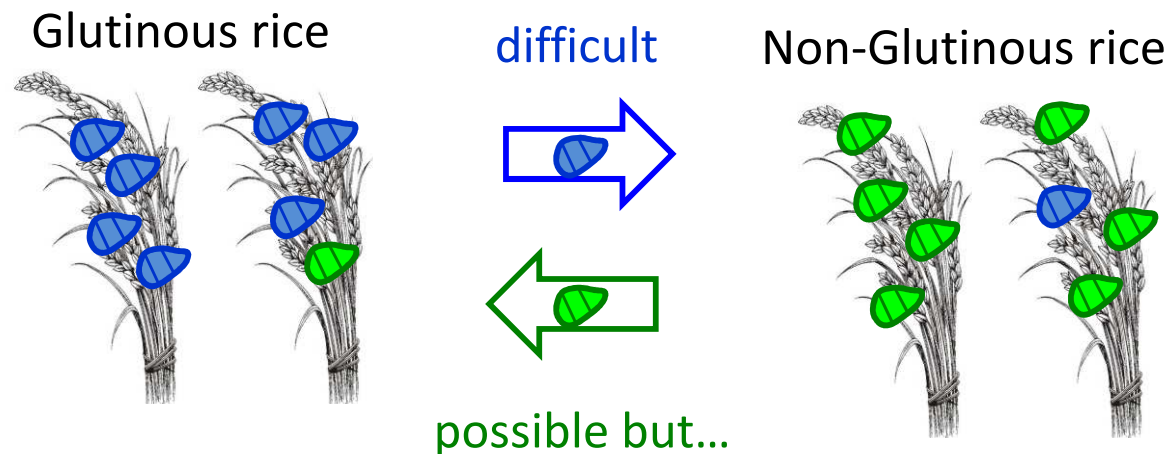
# AVR complement and fitness



# Unbalanced situation between glutinous and non-glutinous rice

Probably because of R-AVR interactions  
since group II isolates have a large AVR complement

why do group II isolates are not aggressive on non-glutinous rice?



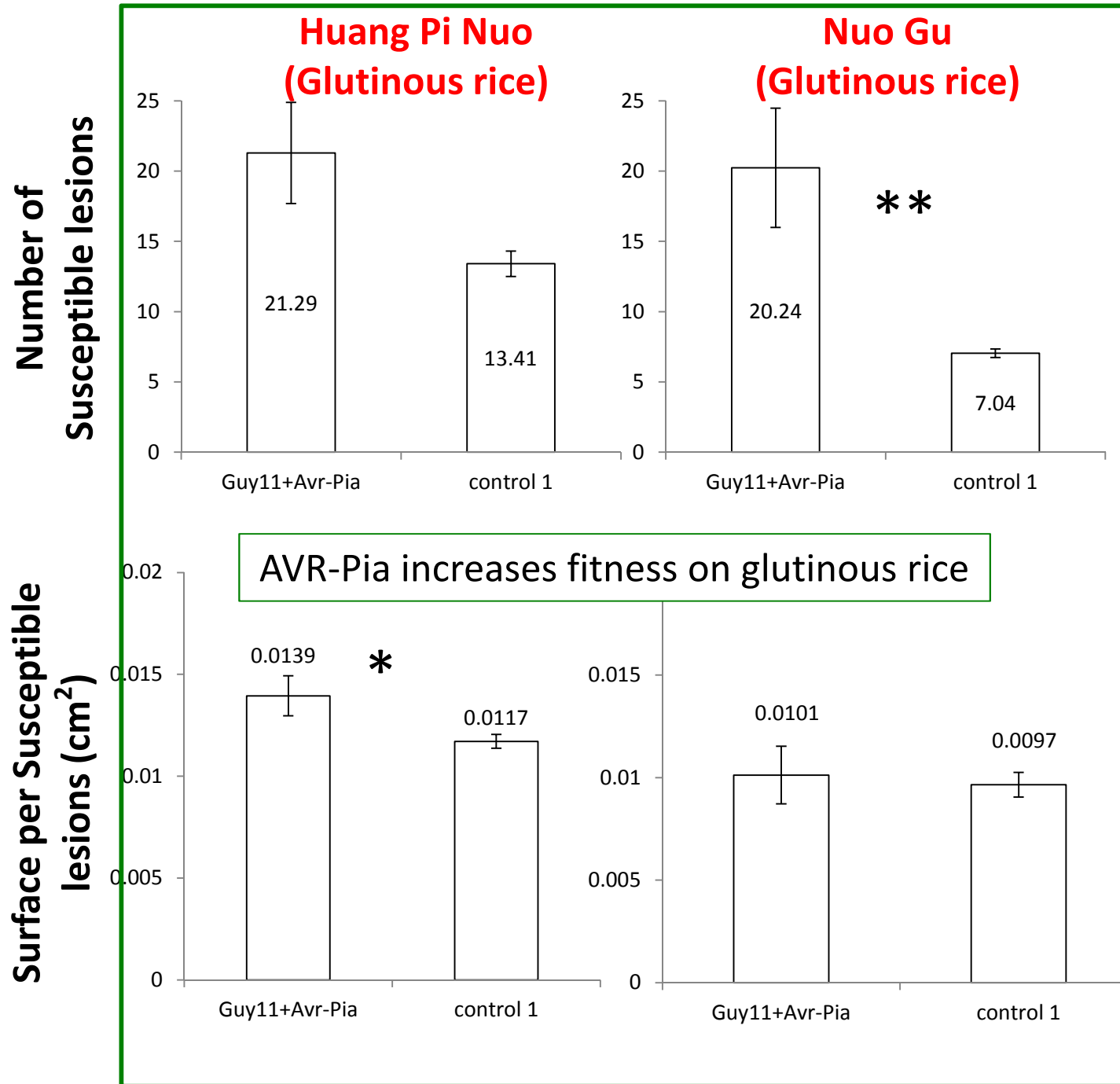
why so few non-group II isolates on glutinous rice?

Probably because a large AVR/effector complement is required on glutinous rice

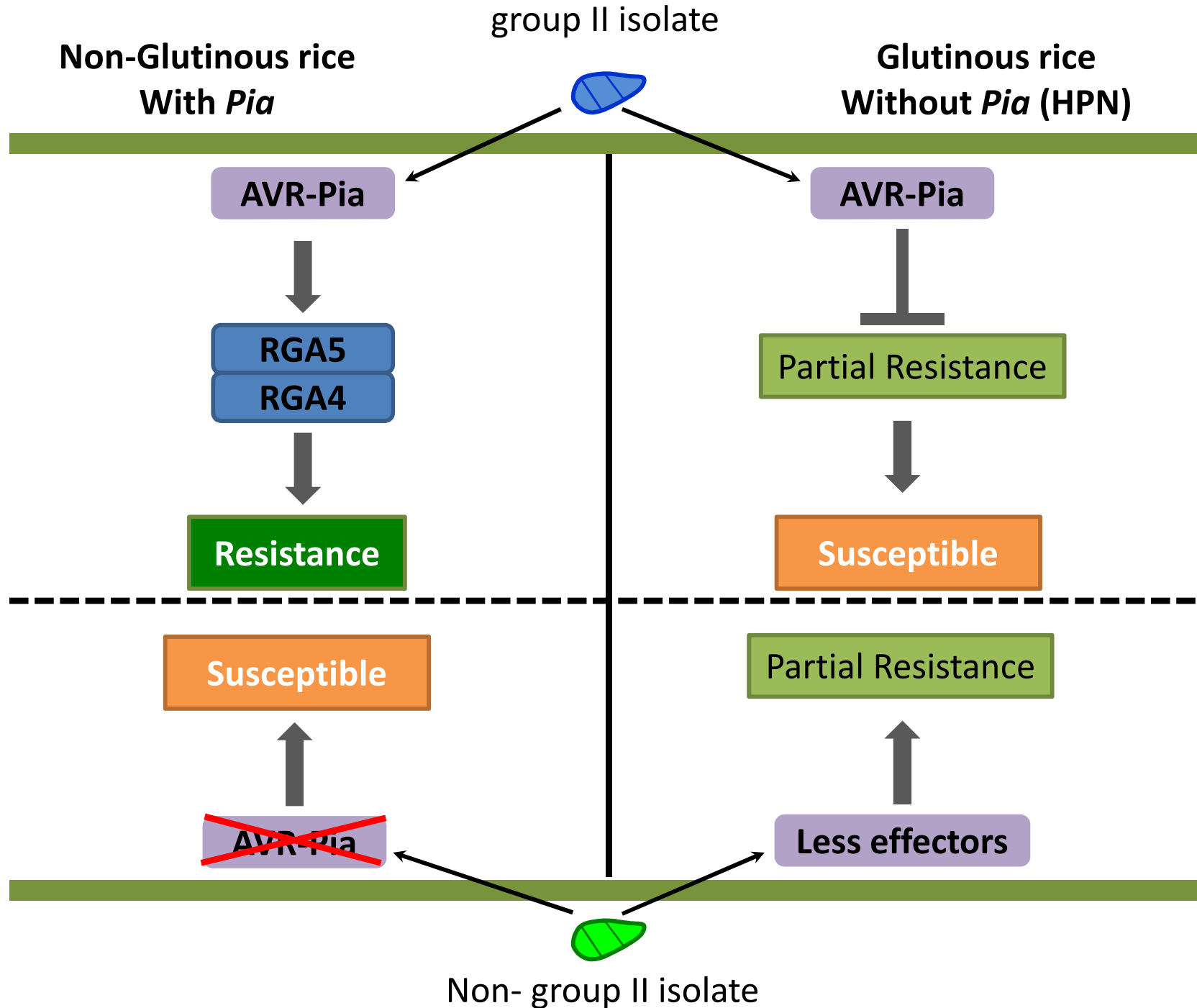




# Effect of AVR-Pia on fitness



# How did *AVR-Pia* make Group II isolates more virulent on Glutinous rice?





## Conclusions and prospects

- Effector complement shapes adaptation to host
- Hypothesis: glutinous rice has a type of resistance that forces adapted isolates to keep/acquire Effectors

An aerial photograph of terraced rice fields, showing a series of curved, stepped terraces carved into a hillside. The terraces are filled with water, reflecting the sky and surrounding landscape. The colors of the water vary from deep blue to light green, depending on the depth and the surrounding vegetation. The terraces are separated by narrow paths and small structures, likely for irrigation or access. The overall scene is a beautiful example of traditional agricultural engineering and landscape design.

Thank you