

Characterization of Resistance to the Green Rice Leafhopper (*Nephotettix cincticeps* Uhler) in a Core Collection of Landraces in Rice (*Oryza sativa* L.)

Tan Van Mai^{1,2}, Atsushi Yoshimura¹, Hideshi Yasui^{1*}

¹Plant Breeding Laboratory, Faculty of Agriculture, Graduate School, Kyushu University, Fukuoka, Japan ²Crop Research and Development Institute, Vietnam National University of Agriculture, Hanoi, Vietnam Email: *hyasui@agr.kyushu-u.ac.jp

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Abstract

The green rice leafhopper (GRH; Nephotettix cincticeps Uhler) is one of the most devastating insect pests of cultivated rice (Oryza sativa L.) in temperate regions in Asia. Using the rice germplasms with biotic stress resistance is the most effective and environmentally-friendly way to control the insect pests in the paddy. Sixty accessions from a core set of worldwide collection of rice were characterized for resistance to the GRH by antibiosis test both at the seedling and at the booting stages. The positive correlations of average nymph mortality (ANM) were observed between at the seedling stage and at the booting stage on 3 days after infestation (DAI) (r = 0.684**), 5DAI (r = 0.680^{**}), and 7DAI (r = 0.652^{**}), respectively. This result will give us the opportunity to screen resistance to the GRH with the cost-efficient way using rice seedlings in a growth chamber. To classify the 60 accessions evaluated, the ANM of the GRH of each accession was compared to the respective ANM of resistant and susceptible controls with the least significant difference (LSD) value. Based on the statistical difference or similarity of the ANMs to the resistant and the susceptible controls, we proposed the four groups of resistance to the GRH, (I) high level of resistance, (II) considerable level of resistance, (III) moderate level of resistance, and (IV) susceptibility. At the seedling stage, a total of 26 accessions were highly resistant in addition to other 6 for considerable level of resistance and other 10 for moderate level of resistance. At the booting stage, on the other hand, a total of 18 accessions were highly resistant in addition to other 3 for considerable level of resistance and other 5 for moderate level of resistance. A total of 42 accessions with high to moderate level of resistance were distributed across 16 countries in Asia in addition to each one

for Madagascar and USA. The classification of landraces based on the present protocol for screening resistance to the insect provided fundamental information for genetics and breeding on resistance to the GRH in rice.

Keywords

Rice, Core Collection, Green Rice Leafhopper, Correlation, Germplasm

1. Introduction

The green rice leafhopper (GRH: *Nephotettix cincticeps* Uhler) is one of the most devastating insects of cultivated rice (*Oryza sativa* L.) and distributed in Asia [1]. The GRH not only causes damage by sucking sap from phloem and xy-lem of susceptible cultivars [2] [3], but also by transmitting viruses such as the rice dwarf and the yellow dwarf as a vector [1] [4]. Rice plants are most sensitive to the GRH at both seedling and booting stages. The sucking and oviposition by the GRH often make plants dwarfish. It causes feasible condition to the plants for the infection by either fungi or bacteria, or both of them. Heavy infestation of the GRH around the booting to the heading stages makes the rice plants ne-crotic and often leads severe yield losses in rice.

Among the various kinds of pest managements, deployment of resistant cultivars is one of the most effective ways to control the GRH in the paddy without any environmental burden. Several GRH resistance genes derived from landraces and wild relatives have been introduced to the elite genetic background of rice cultivars [5], such as Norin PL2 (*GRH1*) [6], Aichi 80 (*GRH3*(t)) [7], Norin-PL5 (GRH1 + GRH2) [8], and Norin-PL6 (GRH2 + GRH4) [9]. One of the constraints to the deployment of resistant cultivars is the dominance of virulent insects against the resistance genes soon after the release of the resistant cultivars. Screening of a large-scale germplasm for GRH resistance is required to ensure useful resources of resistance genes and to utilize them for future breeding programs in rice. Screening "core collections" of crops is recommended to explore resistance resources to abiotic and biotic stresses [10]. In the concept of the core collection, it consists of a limited number of accessions represented as much as possible of the genetic diversity of the whole germplasms [10] [11]. Core collections have been developed based on the genome-wide DNA markers as well as their agronomical traits in several crop species such as rice, maize, and sorghum [12] [13] [14] [15]. Evaluations of core collections were performed in the tolerance or resistance to abiotic and biotic stresses and favorable resources have been effectively screened [16] [17].

In the present study, 60 out of 69 accessions of a core set of rice landraces were characterized for the response to the GRH [13]. The objectives of this study were to develop the efficient technique for the characterization of the response to the GRH in rice and to identify the resources with resistance to the GRH in a core set of rice landraces.

2. Materials and Methods

2.1. Plant Materials

A core set of rice cultivars consists of 69 accessions, which was provided by National Institute of Agrobiological Sciences, Japan, was used in the present study [13]. Among them, sixty-three are landraces and remaining 6 are modern cultivars. This core set of rice cultivars retained 91% of the alleles detected in the original 332 accessions and covered the variation of the initial set of accessions in terms of several agro-morphological traits. This core collection distributed across 16 counties in Asia and one accession in each USA, Brazil and Madagascar. Due to the limitation of the seed stocks, nine accessions were excluded for the evaluation of the resistance to the GRH. Those were WRC54, WRC60, WRC62, WRC64, WRC67, WRC97, WRC98, WRC99 and WRC100. The cultivar "DV85" (Oryza sativa L. ssp. indica) carrying GRH2 and GRH4 [18] [19] was used as a resistant control and "Taichung 65" (T65; Oryza sativa L. ssp. japonica) with none of the resistance gene to the GRH was used as a susceptible control.

2.2. The GRH Strain

The GRH strain was collected in Fukuoka (N33°38' E130°28') Japan in 1991. The GRH strain has been maintained by continuous rearing more than 200 generations on seedlings of the susceptible japonica cultivar Nipponbare under the condition of $25^{\circ}C \pm 1^{\circ}C$ and 16 h light: 8 h dark.

2.3. Evaluation of GRH Resistance

Antibiosis test for GRH resistance reported by Kishino and Ando [20] was modified and used in the present study. Seven-day-old rice seedlings of each accession were infested with 10 first-in star nymphs in a test tube with five replications for the seedling test. The average nymph mortality (ANM) of each accession was then calculated as the mean of the five replications ((ANM = (Rep.1 +Rep.2 + Rep.3 + Rep.4 + Rep.5)/5) at 3 days after infestation (DAI), 5DAI, and 7DAI, respectively. The flag leaves of each accession were similarly infested with 10 first-instar nymphs in a test tube with three replications for the booting test. The ANM of each accession was calculated as the mean of the three replications ((ANM = (Rep.1 + Rep.2 + Rep.3)/3) at 3DAI, 5DAI, and 7DAI, respectively.

2.4. Data Analysis

Analysis of variance (ANOVA) was performed to test the difference in the response to the GRH between the accessions and controls, T65 and DV85. Least significant difference (LSD) was computed and used as a basis for grouping the accessions into different kinds of resistance categories. To classify 60 accessions evaluated in the present study, the ANM of the GRH of each accession was compared to the respective means of resistant and susceptible controls with the LSD value. Finally, the classification was conducted based on the statistical difference to the ANM of DV85 as the resistant control, and the difference to the



ANM of T65 as the susceptible control. Correlation and regression analysis was conducted between the ANMs of the GRH at the seedling stage and at the booting stage.

3. Results

3.1. Response of Rice Landraces to the GRH at the Seedling Stage

The resistant control DV85 had a high ANM with 91.0% at 3DAI that reached to 100% at 7DAI (**Table 1**). On the other hand, the susceptible control T65 had an extremely low ANM with 2.5% at 3DAI that was still low (5.0%) at 7DAI (**Table 1**). Broad variation of the ANMs ranged from 0% to 100% was observed in the core collection of rice landraces in the response to the GRH at the three evaluation times (**Figures 1(a)–(c)**). The frequency distribution of the ANMs of the GRH on rice landraces was discrete at 3DAI where 26 accessions showed high ANMs and the remaining 34 accessions showed low to medium ANMs (**Figure 1(a), Table 1**). The frequency distribution became continuous and it was skewed to the high ANMs at 5DAI and 7DAI (**Figure 1(b), Figure 1(c)**). The accessions were grouped as (I) highly resistant (26 accessions), (II) considerably resistant (6 accessions) based on the comparison to the respective susceptible and resistant controls (**Table 1**, Supplementary **Table S3**). The each ANM of highly resistant accessions was distinctly higher than that of T65 throughout the period evaluated



Figure 1. Frequency distribution of the ANMs of the GRH on 60 rice accessions at the seedling stage at 3 days after infestation (DAI) (a), 5DAI (b) and 7DAI (c), and at the booting stage at 3DAI (d), 5DAI (e) and 7DAI (f).

Groups	No. of		Nymph mortalityª	<i>P</i> val Differen	ues ce from	Response to the	
	accessions		(%)	T65	DV85	GRH	
		3DAI	76.6 ± 5.4	< 0.001	ns		
Ι	26	5DAI	90.2 ± 2.6	< 0.001	ns	HR	
		7DAI	92.8 ± 2.1	< 0.001	ns		
		3DAI	12.0 ± 4.5	ns	< 0.001		
II	6	5DAI	39.4 ± 3.6	< 0.01	< 0.001	CR	
		7DAI	62.8 ± 1.6	< 0.001	< 0.05		
		3DAI	7.1 ± 2.0	ns	< 0.001		
III	10	5DAI	22.6 ± 3.6	< 0.05	< 0.001	MR	
		7DAI	47.1 ± 2.2	< 0.05	< 0.01		
		3DAI	3.6 ± 0.8	ns	< 0.001		
IV	18	5DAI	9.0 ± 1.5	ns	< 0.001	S	
		7DAI	17.4 ± 2.1	ns	< 0.001		
Total	60						
		3DAI	91.0 ± 1.5	< 0.001	-		
DV85 (Resistant control)	-	5DAI	97.1 ± 1.3	< 0.001	-	HR	
``````````````````````````````````````		7DAI	$100.0\pm0.0$	< 0.001	-		
		3DAI	2.5 ± 1.2	-	< 0.001		
T65 (Susceptible control)	ntrol)	5DAI	$5.0 \pm 1.5$	-	< 0.001	S	
-		7DAI	$5.0 \pm 0.5$	-	< 0.001		

Table 1. Classification of the rice landraces based on the response to the GRH by the seedling test

ns, not significant different; HR, high resistance; CR, considerable resistance; MR, moderate resistance; S, susceptible. ^aNymph mortality of total accessions in the group. CR: distinctly higher than that of T65 and slightly lower than that of DV85. MR: higher than that of T65 but distinctly lower than that of DV85.

and was the same as DV85 at 7DAI. The ANMs of considerably resistant accessions were the same as that of T65 at 3DAI, but became distinctly higher than that of T65 at 7DAI. These ANMs were slightly lower than that of DV85 one landrace from Bangladesh (WRC55) which showed an equivalent level of resistance on DV85 at 7DAI. Similarly, the ANMs of moderately resistant accessions were the same as that of T65 at 3DAI, but became significantly higher than that of T65 at 7DAI. These ANMs were distinctly lower than that of DV85 throughout the period evaluated. The ANMs of susceptible accessions were the same as that of T65, and distinctly lower than that of DV85 throughout the period evaluated. Among the 26 highly resistant accessions, 9 were from India, each 3 were from China, Cambodia, and Philippines, 2 were from Nepal and each one was from Korea, Thailand, Myanmar, Malaysia, Indonesia, and Madagascar, respectively (Supplementary Table S1).



#### 3.2. Response of Rice Landraces to the GRH at the Booting Stage

The resistant control DV85 had a considerably high ANM of 66.7% at 3DAI that reached to high ANM of 80.0% at 7DAI. On the other hand, the susceptible control T65 had an ANM of 0.0% throughout the period evaluated (**Table 2**). Broad variation of the ANMs ranged from 0% to 96.7% was observed in sixty accessions of rice landraces in the response to the GRH at the three evaluation times as similarly as at the seedling stage (**Figures 1(d)-(f)**). The frequency distribution of the ANMs of the GRH on rice landraces was continuous and skewed toward the susceptible throughout the period evaluated (**Figures 1(d)-(f)**). The accessions were grouped as (I) highly resistant (18 accessions), (II) considerably resistant (3 accessions), (III) moderately resistant (5 accessions) and (IV) susceptible (35 accessions) (**Table 2**, Supplementary **Table S4**). Among the 18 highly resistant accessions, 5 were from India, each 3 were from China and Cambodia, each 1 was from Korea, Myanmar, Philippines, Malaysia, Bhutan, and Madagascar, respectively (Supplementary **Table S2**).

 Table 2. Classification of the rice landraces based on the response to the GRH by the booting test.

Groups	No. of		Nymph mortality ^a	P va Differei	alues nce from	Response to the GRH
			(%)	T65	DV85	
		3DAI	71.1 ± 5.0	< 0.001	ns	
Ι	18	5DAI	$80.4\pm4.5$	< 0.001	ns	HR
		7DAI	81.6 ± 4.2	< 0.001	ns	
		3DAI	36.6 ± 8.3	< 0.01	< 0.05	
II	3	5DAI	$43.3\pm3.3$	< 0.001	< 0.05	CR
		7DAI	50.3 ± 1.9	< 0.001	< 0.05	
		3DAI	19.6 ± 5.1	ns	< 0.001	
III	5	5DAI	$25.5 \pm 5.3$	< 0.05	< 0.001	MR
		7DAI	$32.2\pm3.1$	< 0.05	< 0.001	
		3DAI	$2.6\pm0.3$	ns	< 0.001	
IV	34	5DAI	$3.9\pm0.7$	ns	< 0.001	S
		7DAI	$5.7 \pm 0.8$	ns	< 0.001	
Total	60					
		3DAI	$66.7\pm4.3$	< 0.001	-	
DV85 (Resistant control)	-	5DAI	73.3 ± 5.4	< 0.001	-	HR
(Resistant control)		7DAI	$80.0\pm3.7$	< 0.001	-	
		3DAI	$0.0 \pm 0.0$	-	< 0.001	
T65 (Susceptible control)	-	5DAI	$0.0 \pm 0.1$	-	< 0.001	S
		7DAI	$0.0 \pm 0.1$	-	< 0.001	

ns, not significant different; HR, high resistance; CR, considerable resistance; MR, moderate resistance; S, susceptible. ^aNymph mortality of total accessions in the group. CR: distinctly higher than that of T65 and slightly lower than that of DV85. MR: higher than that of T65 but distinctly lower than that of DV85.

#### 3.3. Correlation between ANMs of the GRH at the Seedling and at the Booting Stages

The relationship between ANMs of the GRH at the seedling and at the booting stages was demonstrated (Figures 2(a)-(c)). Significant positive correlations were found between the ANMs of the GRH at seedling and booting stages at significant level of 1%. The correlation coefficients between ANMs of the seedling and the booting stages were 0.684 at 3DAI, 0.680 at 5DAI and 0.652 at 7DAI.

The growth stage-specific resistance was observed at the seedling stage but not at the booting stage (Supplementary **Table S3**, Supplementary **Table S4**). The four accessions (WRC25, WRC34, and WRC40) were highly resistant and one (WRC55) was considerably resistant at the seedling stage but obviously susceptible at the booting stage. There was nothing to be resistant only at the booting stage.

#### 4. Discussion

#### 4.1. Phenotypic Coverage of the Rice Core Collection

Core collection of several crops has been developed based on phenotypic and passport data [12]. In the present study, we used a core collection that was developed by using DNA markers. This core set of rice landraces covered 80% -100% of agronomical traits [13]. Sixty accessions originated across 19 countries including 16 countries in Asia, 1 country in Africa and 2 countries in North and South America. Sixty accessions covered a whole range of the resistance to the GRH, from susceptible to highly resistant. Total of 26 out of 60 accessions were highly resistant at the seedling stage. Among the eight loci conferring resistance to GRH, four loci GRH1, GRH2, GRH3 and GRH4 were found in rice landraces [9] [21] [22] [23]. GRH6 was found in both rice landraces and wild relative of rice, Oryza sativa and Oryza nivara [24] [25]. Rice landraces are sources of useful genetic resources for GRH resistance. The three resistance genes GRH1, GRH2 and *GRH3* were lost its effectiveness by the GRH strains known as biotype 1, biotype 2 and biotype 3 in the laboratory condition [26]. Therefore, it is necessary to screen large-scale germplasm in order to identify new genes/alleles that contribute to resistance to GRH in rice.



**Figure 2.** Correlation and regression between the ANMs of the GRH at the seedling stage and the ANMs at the booting stage in a rice core collection at 3DAI (a), 5DAI (b) and 7DAI (c). ANM; average nymph mortality.



## 4.2. Positive Correlation of GRH Resistance between the Rice Growth Stages

We found significant positive correlations of GRH resistance between at the seedling and at the booting stages (Figures 2(a)-(c)). This indicated that genetic components existed in landraces are commonly stable from the seedling to the booting stages. In addition, the correlation of GRH nymph mortality between at the seedling stage at 3DAI and at the booting stage at 7DAI was observed (r = 0.683, data was not shown). The evaluation at the seedling stage, therefore, promises us to identify the genetic resources with GRH-resistance accurately.

#### 4.3. Seedling Specific Resistance Found in Rice Landraces

In rice, growth stage plays an important role against the insect's attack. It seems to be that damages to the rice plant caused by the GRH are most serious at the seedling and the booting stages. In the present study, we evaluated the resistance to the GRH of the core collection at the seedling and the booting stages. Four highly resistant accessions (WRC25, WRC34, WRC37, and WRC40), four considerable resistant accessions (WRC6, WRC7, WRC48, and WRC55) and eight moderate resistant accessions (WRC4, WRC9, WRC13, WRC15, WRC44, WRC45, WRC46, and WRC50) at the seedling stage. But all of the above-mentioned accessions were obviously susceptible at the booting stage. These accessions are useful for exploitation of new genes/alleles confer seedling-specific resistance to the GRH at the seedling. On the other hand, there were no accessions with booting-specific resistance among the 60 accessions.

#### **5.** Conclusion

In this study, we identified a large number of accessions that were a wide range of resistance to the GRH at both seedling and booting stages. Based on positive correlations of the GRH resistance between seedling and booting stages, we suggested that seedling stage is an effective stage to screen resistance to the GRH with the cost-efficient way using rice seedlings in a growth chamber. The four groups of resistance to the GRH were identified as (I) high level of resistance, (II) considerable high level of resistance, (III) moderate level of resistance, and (IV) susceptibility. In addition, we identified the unique phenomenon in some accessions, which showed resistance at the seedling stage but susceptible at booting stage. The exploitation of resistant landraces based on the present protocol will provide fundamental information for genetics and breeding on resistance to GRH in rice.

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### **Supplements**

Table S1. Geographical origin and classification of resistance to the green rice leafhopper (Nephotettix cincticeps Uhler) of a core collection at the seedling stage.

	Country of -		3D	AI		5DAI				7DAI				
	Country of	No.	of ac	cessio	ns ^a	No.	of ac	cessic	ons ^a	No. of accessions ^a				Total
	origin	HR	CR	MR	S	HR	CR	MR	S	HR	CR	MR	S	
	Korea	1	0	0	0	1	0	0	0	1	0	0	0	1
East Asia	Japan	0	0	0	2	0	0	0	2	0	0	0	2	2
	China	3	0	0	4	3	0	1	3	3	0	2	3	7
	Vietnam	0	0	1	1	0	0	1	1	0	1	0	1	2
	Laos	0	0	0	2	0	0	0	2	0	0	1	1	2
South East Asia	Cambodia	2	0	1	0	3	0	0	0	3	0	0	0	3
	Thailand	0	0	1	0	1	0	0	0	1	0	0	0	1
	Myanmar	1	0	0	2	1	0	1	1	1	0	1	1	3
	Philippines	3	0	0	4	3	0	2	2	3	1	2	1	7
Pacific Asia	Malaysia	0	0	1	0	0	1	0	0	1	0	0	0	1
	Indonesia	1	0	0	2	1	0	1	1	1	1	0	1	3
	Bangladesh	0	0	0	3	0	0	0	3	0	1	0	2	3
South Asia	India	5	0	4	4	8	1	1	3	9	1	1	2	13
	Sri Lanka	0	0	0	1	0	0	0	1	0	0	0	1	1
TT:	Bhutan	0	0	0	3	0	0	1	2	0	1	1	1	3
Himalaya	Nepal	2	0	0	3	2	0	0	3	2	0	1	2	5
Africa	Madagascar	1	0	0	0	1	0	0	0	1	0	0	0	1
North America	USA	0	0	0	1	0	0	0	1	0	0	1	0	1
South America	Brazil 0 0		0	0	1	0	0	0	1	0	0	0	1	1
	Total	19	0	8	33	24	2	8	26	26	6	10	18	60

^aClassification of accessions based on their nymph mortality of the GRH at the seedling stage by comparing their mean and LSD with resistant cultivar (DV85) and susceptible cultivar (T65). HR: highly resistant, R: resistant, MR: moderate resistant, S: susceptible.

			3DAI			5DAI				7DAI				
	Country of origin	No.	ofac	cessio	ns ^a	No.	No. of accessions ^a			No. of accessions ^a			Total	
	8	HR	CR	MR	S	HR	CR	MR	S	HR	CR	MR	S	-
	Korea	1	0	0	0	1	0	0	0	1	0	0	0	1
East Asia	Japan	0	0	0	2	0	0	0	2	0	0	0	2	2
	China	2	0	1	4	2	1	0	4	3	0	1	3	7
	Vietnam	0	0	0	2	0	0	0	2	0	0	0	2	2
	Laos	0	0	0	2	0	0	0	2	0	0	0	2	2
South Fast Asia	Cambodia	2	1	0	0	3	0	0	0	3	0	0	0	3
East Asia	Thailand	0	0	0	1	0	0	1	0	0	1	0	0	1
	Myanmar	1	0	0	2	1	0	0	2	1	0	0	2	3

Table S2. Classification of resistance to the green rice leafhopper (Nephotettix cincticeps Uhler) of a core collection at the booting stage.



Continued														
	The Philippines	0	1	3	3	1	1	2	3	1	1	2	3	7
Pacific Asia	Malaysia	0	0	1	0	1	0	0	0	1	0	0	0	1
	Indonesia	0	0	0	3	0	0	0	3	0	0	0	3	3
	Bangladesh	0	0	0	3	0	0	0	3	0	0	0	3	3
South Asia	India	0	6	0	7	5	1	1	6	5	1	1	6	13
	Sri Lanka	0	0	0	1	0	0	0	1	0	0	0	1	1
Uimalawa	Bhutan	0	0	1	2	1	0	0	2	1	0	0	2	3
Tiiilalaya	Nepal	0	1	1	3	1	0	0	4	1	0	1	3	5
Africa	Madagascar	0	1	0	0	1	0	0	0	1	0	0	0	1
North America	USA	0	0	0	1	0	0	0	1	0	0	0	1	1
South America	Brazil	0	0	0	1	0	0	0	1	0	0	0	1	1
Total		6	10	7	37	17	3	4	36	18	3	5	34	60

^aClassification of accessions based on their nymph mortality of the GRH at seedling stage by comparing their mean and LSD with resistant and susceptible controls. HR: highly resistant, R: resistant, MR: moderate resistant, S: susceptible

Table S3	The nymph	mortality of t	he green r	rice leafhopper	on the rice l	andrace acce	ssions at s	eedling stage

Name	Cultivar name	Country of origin	Туре		ANM (%)	Different from T65	Different from DV85	Classification
WRC2	Kasalath	India	Landrace	3DAI	89.8	***	ns	
				5DAI	94.3	***	ns	HR
				7DAI	97.1	***	ns	
WRC3	Bei Khe	Cambodia	Landrace	3DAI	94.6	***	ns	
				5DAI	97.1	***	ns	HR
				7DAI	100.0	***	ns	
WRC5	Naba	India	Landrace	3DAI	31.9	**	***	
				5DAI	71.9	***	*	HR
				7DAI	81.9	***	ns	
WRC10	Shuusoushu	China	Landrace	3DAI	92.1	***	ns	
				5DAI	100.0	***	ns	HR
				7DAI	100.0	***	ns	
WRC16	Vary Futsi	Madagascar	Landrace	3DAI	96.7	***	ns	
				5DAI	100.0	***	ns	HR
				7DAI	100.0	***	ns	
WRC17	Keiboba	China	Landrace	3DAI	97.5	***	ns	
				5DAI	100.0	***	ns	HR
				7DAI	100.0	***	ns	
WRC18	Qingyu (Seiyu)	China	Landrace	3DAI	95.0	***	ns	
				5DAI	100.0	***	ns	HR
				7DAI	100.0	***	ns	

#### Continued

WRC20	Tadukan	Philippines	Landrace	3DAI	84.8	***	ns	
				5DAI	97.1	***	ns	HR
				7DAI	97.1	***	ns	
WRC22	Calotoc	Philippines	Landrace	3DAI	88.5	***	ns	
				5DAI	100.0	***	ns	HR
				7DAI	100.0	***	ns	
WRC24	Pinulupot 1	Philippines	Landrace	3DAI	81.8	***	ns	
				5DAI	91.0	***	ns	HR
				7DAI	91.0	***	ns	
WRC25	Muha	Indonesia	Landrace	3DAI	82.9	***	ns	
				5DAI	85.7	***	ns	HR
				7DAI	88.6	***	ns	
WRC26	Ihona 2	India	Landrace	3DAI	48.9	***	***	
	,			5DAI	73.6	***	ns	HR
				7DAI	73.6	***	ns	
WRC30	Aniana Dhan	Nepal	Landrace	3DAI	73.0	***	ne	
WIKC50		ivepai	Landrace	5DAI	96.7	***	113	ЦD
					96.7	***	115	ш
WDC24	ADC 7201	T., J.,	T J		90.7	***	115	
WRC34	ARC 7291	India	Landrace	3DAI	91.4	***	ns	IID
				5DAI	97.1	***	ns	ΠК
WDC25	ADC FOFF	India	Landraca	7DAI 2DAI	100.0	***	ns	
WRC55	AKC 5955	India	Landrace	5DAI	100.0	***	ns	НR
				7DAI	100.0	***	ns	Ш
WRC36	Ratul	India	Landrace	3DAI	97.1	***	ns	
				5DAI	100.0	***	ns	HR
				7DAI	100.0	***	ns	
WRC37	ARC 7047	India	Landrace	3DAI	40.0	***	***	
				5DAI	74.3	***	ns	HR
				7DAI	74.3	***	ns	
WRC39	Badari Dhan	Nepal	Landrace	3DAI	94.3	***	ns	
		1		5DAI	100.0	***	ns	HR
				7DAI	100.0	***	ns	
WRC40	Nepal 555	India	Landrace	3DAI	82.9	***	ns	
				5DAI	94.3	***	ns	HR
				7DAI	94.3	***	ns	
WRC42	Local Basmati	India	Landrace	3DAI	35.4	**	***	
				5DAI	78.9	***	ns	HR
				7DAI	94.6	***	ns	

Continued								
WRC57	Milyang 23	Korea	Modern	3DAI	86.6	***	ns	
				5DAI	92.3	***	ns	HR
				7DAI	95.1	***	ns	
WRC58	Neang Menh	Cambodia	Landrace	3DAI	81.3	***	ns	
				5DAI	89.2	***	ns	HR
				7DAI	94.3	***	ns	
WRC59	Neang Phtong	Cambodia	Landrace	3DAI	50.4	***	***	
				5DAI	75.4	***	ns	HR
				7DAI	81.1	***	ns	
WRC61	Radin Goi Sesat	Malaysia	Landrace	3DAI	40.0	***	***	
				5DAI	65.7	***	*	HR
				7DAI	77.1	***	ns	
WRC63	Bleiyo	Thailand	Landrace	3DAI	43.3	***	***	
				5DAI	76.2	***	ns	HR
				7DAI	82.4	***	ns	
WRC66	Bingala	Myanmar	Landrace	3DAI	91.4	***	ns	
				5DAI	94.3	***	ns	HR
				7DAI	94.3	***	ns	
WRC6	Puluik Arang	Indonesia	Landrace	3DAI	10.0	ns	***	
				5DAI	39.4	**	***	CR
				7DAI	68.5	***	*	
WRC7	Davao 1	Philippine	Landrace	3DAI	3.9	ns	***	
				5DAI	37.8	**	***	CR
				7DAI	65.1	***	*	
WRC28	Jarjan	Bhutan	Landrace	3DAI	6.2	ns	***	
				5DAI	31.5	*	***	CR
				7DAI	59.4	***	*	
WRC48	Khau Mac Kho	Vietnam	Landrace	3DAI	31.2	**	***	
				5DAI	53.9	***	***	CR
				7DAI	60.1	***	*	
WRC55	Tupa729	Bangladesh	Landrace	3DAI	5.7	ns	***	
				5DAI	22.9	ns	***	CR
				7DAI	74.3	***	ns	
WRC65	Rambhog	India	Landrace	3DAI	8.6	ns	***	
				5DAI	34.3	*	***	CR
				7DAI	61.0	***	*	
WRC4	Jena 035	Nepal	Landrace	3DAI	7.5	ns	***	
				5DAI	25.5	ns **	***	MR
WECO	Rvou Suican Koumoi	China	Landrace		40.3 2 Q	ne	***	
¥¥ KC9	Kyou Suisan Kouniai	Ciillia	Landiace	5DAI	2.7	115	***	MD
					20.0	*	***	IVIK
				/DAI	43.2			

Continued								
WRC13	Asu	Bhutan	Landrace	3DAI	0.0	ns	***	
				5DAI	11.1	ns	***	MR
				7DAI	41.8	*	***	
WRC14	IR 58	Philippines	Landrace	3DAI	5.7	ns	***	
				5DAI	23.3	ns	***	MR
				7DAI	38.1	*	***	
WRC15	Co 13	India	Landrace	3DAI	0.0	ns	***	
				5DAI	12.5	ns	***	MR
				7DAI	43.9	*	***	
WRC19	Deng Pao Zhai	China	Modern	3DAI	11.4	ns	***	
				5DAI	32.4	*	***	MR
				7DAI	35.2	*	***	
WRC44	Basilanon	Philippines	Landrace	3DAI	19.9	ns	***	
				5DAI	44.2	**	***	MR
				7DAI	53.7	**	**	
WRC45	Ma sho	Myanmar	Landrace	3DAI	14.4	ns	***	
				5DAI	33.0	*	***	MR
				7DAI	57.6	***	**	
WRC46	Khao Nok	Laos	Landrace	3DAI	5.7	ns	***	
				5DAI	12.4	ns	***	MR
				7DAI	38.6	*	***	
WRC50	Rexmont	USA	Modern	3DAI	5.4	ns	***	
				5DAI	10.4	ns	***	MR
				7DAI	40.8	*	***	
WRC1	Nipponbare	Japan	Modern	3DAI	7.3	ns	***	
				5DAI	16.3	ns	***	S
				7DAI	16.3	ns	***	
WRC11	Jinguoyin	China	Landrace	3DAI	2.9	ns	***	
				5DAI	20.0	ns	***	S
				7DAI	25.7	ns	***	
WRC21	Shwe Nang Gyi	Myanmar	Landrace	3DAI	0.0	ns	***	
				5DAI	8.2	ns	***	S
				7DAI	16.9	ns	***	
WRC23	Lebed	Philippines	Landrace	3DAI	2.5	ns	***	
				5DAI	2.5	ns	***	S
				7DAI	28.2	ns	***	
WRC27	Nepal 8	Nepal	Landrace	3DAI	0.0	ns	***	
				5DAI	0.0	ns	***	S
				7DAI	15.7	ns	***	
WRC29	Kalo Dhan	Nepal	Landrace	3DAI	0.0	ns	***	
				5DAI	0.0	ns	***	S
				7DAI	22.9	ns	***	

Continued								
WRC31	Shoni	Bangladesh	Landrace	3DAI	0.0	ns	***	
				5DAI	6.2	ns	***	S
				7DAI	6.2	ns	***	
WRC32	Tupa 121-3	Bangladesh	Landrace	3DAI	0.0	ns	***	
				5DAI	2.9	ns	***	S
				7DAI	30.8	ns	***	
WRC33	Surjamukhi	India	Modern	3DAI	2.9	ns	***	
				5DAI	5.7	ns	***	S
				7DAI	5.7	ns	***	
WRC38	ARC 11094	India	Landrace	3DAI	0.0	ns	***	
				5DAI	20.0	ns	***	S
				7DAI	32.4	ns	***	
WRC41	Kaluheenati	Sri Lanka	Landrace	3DAI	6.2	ns	***	
				5DAI	12.4	ns	***	S
				7DAI	15.2	ns	***	
WRC43	Dianyu 1	China	Modern	3DAI	5.4	ns	***	
				5DAI	8.7	ns	***	S
				7DAI	11.5	ns	***	
WRC47	Jaguary	Brazil	Landrace	3DAI	8.6	ns	***	
				5DAI	14.3	ns	***	S
				7DAI	14.3	ns	***	
WRC49	Padi Perak	Indonesia	Landrace	3DAI	8.6	ns	***	
				5DAI	14.3	ns	***	S
				7DAI	31.4	ns	***	
WRC51	Urasan 1	Japan	Landrace	3DAI	6.2	ns	***	
				5DAI	6.2	ns	***	S
				7DAI	11.5	ns	***	
WRC52	Khau Tan Chiem	Vietnam	Landrace	3DAI	2.9	ns	***	
				5DAI	9.0	ns	***	S
				7DAI	11.9	ns	***	
WRC53	Tima	Bhutan	Landrace	3DAI	6.2	ns	***	
				5DAI	9.0	ns	***	S
				7DAI	9.0	ns	***	
WRC68	Khao Nam Jen	Laos	Landrace	3DAI	5.4	ns	***	
				5DAI	5.4	ns	***	S
				7DAI	7.9	ns	***	
				3DAI	2.5	-	***	
165				5DAI	5.0	-	***	S
				/DAI	5.0	- ***		
DUOS				5DAI	91.0	***	-	TID
DV85				5DAI	97.1	***	-	нк
				7DAI	100.0	***	-	

Name	Cultivar name	Country of origin	Туре		ANM (%)	Different from T65	Different from DV85	Classification
WRC2	Kasalath	India	Landrace	3DAI	54.8	***	ns	
				5DAI	62.2	***	ns	HR
				7DAI	65.6	***	ns	
WRC3	Bei Khe	Cambodia	Landrace	3DAI	93.3	***	*	
				5DAI	93.3	***	ns	HR
				7DAI	93.3	***	ns	
WRC5	Naba	India	Landrace	3DAI	44.4	***	ns	
				5DAI	51.1	***	ns	HR
				7DAI	64.4	***	ns	
WRC10	Shuusoushu	China	Landrace	3DAI	38.9	***	*	
				5DAI	55.2	***	ns	HR
				7DAI	55.2	***	ns	
WRC14	IR 58	Philippines	Landrace	3DAI	80	***	ns	
				5DAI	83.3	***	ns	HR
				7DAI	83.3	***	ns	
WRC16	Vary Futsi	Madagascar	Landrace	3DAI	54.4	***	ns	
				5DAI	65.2	***	ns	HR
				7DAI	65.2	***	ns	
WRC17	Keiboba	China	Landrace	3DAI	93.3	***	*	
				5DAI	100	***	ns	HR
				7DAI	100	***	ns	
WRC18	Qingyu (Seiyu)	China	Landrace	3DAI	96.7	***	*	
				5DAI	100	***	ns	HR
				7DAI	100	***	ns	
WRC28	Jarjan	Bhutan	Landrace	3DAI	36.7	**	*	
				5DAI	60	***	ns	HR
				7DAI	60	***	ns	
WRC35	ARC 5955	India	Landrace	3DAI	83.3	***	ns	
				5DAI	95.8	***	ns	HR
				7DAI	100	***	ns	
WRC36	Ratul	India	Landrace	3DAI	85.2	***	ns	
				5DAI	92.6	***	ns	HR
				7DAI	96.3	***	ns	
WRC39	Badari Dhan	Nepal	Landrace	3DAI	44 2	***	ns	
	Dunni Dinii	pu	Lundruce	5DAI	52	***	ne	HR
					52	***	115 DC	
				/DAI	52		115	

### Table S4. The nymph mortality of the green rice leafhopper on the rice landrace accessions at booting stage.

Continued								
WRC42	Local Basmati	India	Landrace	3DAI	68.9	***	ns	
				5DAI	86.3	***	ns	HR
				7DAI	83	***	ns	
WRC57	Milyang 23	Korea	Modern	3DAI	93.3	***	*	
				5DAI	100	***	ns	HR
				7DAI	100	***	ns	
WRC58	Neang Menh	Cambodia	Landrace	3DAI	89.2	***	ns	
				5DAI	100	***	ns	HR
				7DAI	96.7	***	ns	
WRC59	Neang Phtong	Cambodia	Landrace	3DAI	60	***	ns	
				5DAI	63.3	***	ns	HR
				7DAI	66.7	***	ns	
WRC61	Radin Goi Sesat	Malaysia	Landrace	3DAI	70	***	ns	
				5DAI	90	***	ns	HR
				7DAI	90	***	ns	
WRC66	Bingala	Myanmar	Landrace	3DAI	93.3	***	*	
				5DAI	96.7	***	ns	HR
				7DAI	96.7	***	ns	
WRC22	Calotoc	Philippines	Landrace	3DAI	43.3	***	*	
				5DAI	46.7	***	*	CR
				7DAI	50.8	***	*	
WRC63	Bleiyo	Thailand	Landrace	3DAI	20	ns	***	
				5DAI	36.7	**	**	CR
				7DAI	46.7	***	*	
WRC65	Rambhog	India	Landrace	3DAI	46.7	***	ns	
				5DAI	46.7	***	*	CR
				7DAI	53.3	***	*	
WRC19	Deng Pao Zhai	China	Modern	3DAI	0	ns	***	
				5DAI	6.7	ns	***	MR
				7DAI	30	*	***	
WRC20	Tadukan	Philippines	Landrace	3DAI	30	**	**	
				5DAI	36.7	**	**	MR
				7DAI	40	**	**	
WRC24	Pinulupot 1	Philippines	Landrace	3DAI	23.3	*	***	) (D
				5DAI	26.7	*	***	MR
WRC26	Ihona ?	India	Landrace		21.8	ne	***	
¥¥ IXC20	Juona 2	mula	Lanulace	50 11	34 5	**	**	MR
					37.0	**	**	IVIIX
				/DAI	57.7			

Continued								
WRC30	Anjana Dhan	Nepal	Landrace	3DAI	22.7	*	***	
				5DAI	22.7	*	***	MR
				7DAI	22.7	*	***	
WRC1 ^a	Nipponbare	Japan	Modern	3DAI	0	ns	***	
				5DAI	0	ns	***	S
				7DAI	3.7	ns	***	
WRC4	Jena 035	Nepal	Landrace	3DAI	0	ns	***	
				5DAI	0	ns	***	S
				7DAI	0	ns	***	
WRC6	Puluik Arang	Indonesia	Landrace	3DAI	6.7	ns	***	
				5DAI	10	ns	***	S
				7DAI	13.3	ns	***	
WRC7	Davao 1	Philippines	Landrace	3DAI	0	ns	***	
				5DAI	3.3	ns	***	S
				7DAI	6.7	ns	***	
WRC9	Ryou Suisan Koumai	China	Landrace	3DAI	3.3	ns	***	
				5DAI	10.4	ns	***	S
				7DAI	3.3	ns	***	
WRC11 ^a	Jinguoyin	China	Landrace	3DAI	0	ns	***	
				5DAI	0	ns	***	S
				7DAI	10	ns	***	
WRC13	Asu	Bhutan	Landrace	3DAI	6.7	ns	***	
				5DAI	10	ns	***	S
				7DAI	10	ns	***	
WRC15	Co 13	India	Landrace	3DAI	4.8	ns	***	
				5DAI	8.1	ns	***	S
				7DAI	8.1	ns	***	
WRC21 ^a	Shwe Nang Gyi	Myanmar	Landrace	3DAI	3.3	ns	***	
				5DAI	3.3	ns	***	S
				7DAI	7.5	ns	***	
WRC23 ^a	Lebed	Philippines	Landrace	3DAI	9.4	ns	***	
				5DAI	12.4	ns	***	S
				7DAI	12.4	ns	***	
WRC25	Muha	Indonesia	Landrace	3DAI	3.3	ns	***	
				5DAI	3.3	ns	***	S
				7DAI	3.3	ns	***	
WRC27 ^a	Nepal 8	Nepal	Landrace	3DAI	0	ns	***	
				5DAI	3.3	ns	***	S

7DAI

0

ns



***

Continued								
WRC29 ^a	Kalo Dhan	Nepal	Landrace	3DAI	3.3	ns	***	
				5DAI	3.3	ns	***	S
				7DAI	3.3	ns	***	
WRC31 ^a	Shoni	Bangladesh	Landrace	3DAI	0	ns	***	
				5DAI	0	ns	***	S
				7DAI	0	ns	***	
WRC32 ^a	Tupa 121-3	Bangladesh	Landrace	3DAI	0	ns	***	
				5DAI	3.3	ns	***	S
				7DAI	3.3	ns	***	
WRC33 ^a	Surjamukhi	India	Modern	3DAI	0	ns	***	
				5DAI	0	ns	***	S
				7DAI	3.3	ns	***	
WRC34	ARC 7291	India	Landrace	3DAI	0	ns	***	
				5DAI	0	ns	***	S
				7DAI	6.7	ns	***	
WRC37	ARC 7047	India	Landrace	3DAI	3.3	ns	***	
				5DAI	3.3	ns	***	S
				7DAI	6.7	ns	***	
WRC38 ^a	ARC 11094	India	Landrace	3DAI	3.3	ns	***	
				5DAI	3.3	ns	***	S
				7DAI	3.3	ns	***	
WRC40	Nepal 555	India	Landrace	3DAI	0	ns	***	
	1			5DAI	4.2	ns	***	S
				7DAI	15.3	ns	***	
WRC41 ^a	Kaluheenati	Sri Lanka	Landrace	3DAI	3	ns	***	
				5DAI	6.1	ns	***	S
				7DAI	6.1	ns	***	
WRC43	Dianyu 1	China	Modern	3DAI	6.7	ns	***	
				5DAI	6.7	ns	***	S
				7DAI	6.7	ns	***	
WRC44	Basilanon	Philippines	Landrace	3DAI	0	ns	***	
				5DAI	0	ns	***	S
				7DAI	7.4	ns	***	
WRC45	Ma sho	Myanmar	Landrace	3DAI	0	ns	***	0
				5DAI	0	ns	***	8
WDC14	Khao Nob	Looc	Landraca		2 2	115	***	
VV NC40	NHAU INOK	Laus	Lanuface	5DAI	3.5 10.4	115	***	S
					10.4	115	***	3
				/DAI	15./	115		

Continued								
WRC47 ^a	Jaguary	Brazil	Landrace	3DAI	0	ns	***	
				5DAI	0	ns	***	S
				7DAI	3.3	ns	***	
WRC48	Khau Mac Kho	Vietnam	Landrace	3DAI	3.3	ns	***	
				5DAI	6.7	ns	***	S
				7DAI	10	ns	***	
WRC49 ^a	Padi Perak	Indonesia	Landrace	3DAI	0	ns	***	
				5DAI	0	ns	***	S
				7DAI	0	ns	***	
WRC50	Rexmont	USA	Modern	3DAI	0	ns	***	
				5DAI	10	ns	***	S
				7DAI	10	ns	***	
WRC51 ^a	Urasan 1	Japan	Landrace	3DAI	0	ns	***	
				5DAI	0	ns	***	S
				7DAI	0	ns	***	
WRC52 ^a	Khau Tan Chiem	Vietnam	Landrace	3DAI	0	ns	***	
				5DAI	0	ns	***	S
				7DAI	0	ns	***	
WRC53 ^a	Tima	Bhutan	Landrace	3DAI	3.3	ns	***	
				5DAI	10	ns	***	S
				7DAI	6.7	ns	***	
WRC55	Tupa729	Bangladesh	Landrace	3DAI	0	ns	***	
				5DAI	3.3	ns	***	S
				7DAI	3.3	ns	***	
WRC68 ^a	Khao Nam Jen	Laos	Landrace	3DAI	0	ns	***	
				5DAI	7.4	ns	***	S
				7DAI	3.7	ns	***	
				3DAI	0	-	***	
T65				5DAI	0	-	***	S
				7DAI	0	-	***	
				3DAI	66.7	***	-	
DV85				5DAI	73.3	***	-	HR
				7DAI	80.0	***	-	

^aSusceptible at the seedling stage.



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