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Host suitability of weeds to Meloidogyne ottersoni and Meloidogyne graminicola

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ABSTRACT: We evaluated the host suitability of the weeds associated with rice crops regarding Meloidogyne ottersoni and M. graminicola. Both plant-parasitic nematodes can develop in Oryza sativa, but Cyperus ferax plants were resistant to M. ottersoni. Plants of Cyperus iria, Cyperus difformis, Echinochloa crus-galli and Echinochloa colonum were susceptible to M. ottersoni, but resistant to M. graminicola. Besides this, Aeschynomene denticulata and Leersia hexandra were immune to M. graminicola and susceptible (1st assessment) and resistant (2nd assessment) regarding M. ottersoni. The results shed light on the role of hosts of M. ottersoni and M. graminicola, demonstrating that weed management should be included in strategies to control root-knot nematode diseases. Key words: management of diseases, rice, plant-parasitic nematodes.

Reação de plantas daninhas a Meloidogyne ottersoni e Meloidogyne graminicola

RESUMO: Objetivou-se avaliar a reação de plantas daninhas associadas à cultura de arroz em relação a Meloidogyne ottersoni e M. graminicola. Ambos fitonematoides podem se desenvolver em Oryza sativa, mas plantas de Cyperus ferax foram resistentes a M. ottersoni. Plantas de Cyperus iria, Cyperus difformis, Echinochloa crus-galli e Echinochloa colonum foram suscetíveis a M. ottersoni, mas resistentes a M. graminicola. Além disso, Aeschynomene denticulata e Leersia hexandra comportaram-se como imunes a M. graminicola e suscetíveis (1ª avaliação) e resistentes (2ª avaliação) em relação a M. ottersoni. Os resultados ampliam o conhecimento acerca de plantas hospedeiras de M. ottersoni e M. graminicola, demonstrando que controle de plantas daninhas deve ser incorporado nas estratégias de manejo de meloidoginoses. Palavras-chave: manejo de doenças, arroz, fitonematoides.

1 INTRODUCTION

2 3 Rice (Oryza sativa L.) growing is 4 hugely important to produce food, employment 5 and income by millions of people around the 6 world. Among the leading producing countries in 7 2022 were continental China (208.5 million tons), 8 followed by India (196.2 million tons). Brazil was 9 in eleventh place, with an output of 10.8 million 10 tons (FAO, 2024) and Brazil's South region stands 11 out in the production of irrigated rice, with the state 12 of Rio Grande do Sul being the leading producer 13 (7.29 million tons), followed by Santa Catarina 14 (1.12 million tons) (IBGE, 2024).

15 Among the phytosanitary factors that 16 limit the productivity of rice are attacks by pests 17 and encroachment of weeds (SAVARY et al., 2012;

AVILA et al., 2021). The losses caused by weeds to 1 2 rice crops can be direct (competition) and indirect, 3 due to the multiplication/maintenance of various pathogens (FERRAZ et al., 1983; AGOSTINETTO 4 5 et al., 2008; SILVA et al., 2010; CONCENÇO et al., 2014). Globally, M. graminicola GOLDEN & 6 7 BIRCHFIEL, 1965, is the species with the greatest potential to damage irrigated rice crops (DE WAELE 8 9 & ELSEN, 2007). Besides rice, various plants present in fields between harvests can serve as hosts, such 10 as Echinochloa colonum (L.) Link. (GOLDEN & 11 BIRCHFIELD, 1965), E. crus-galli (L.) P. Beauv., 12 1812, Eleusine indica (L.) Gaerth., 1788, and Cyperus 13 14 difformis L., 1756 (BAJAJ & DABUR, 2000; DABUR 15 et al., 2004; NEGRETTI et al., 2014; KUMAR et al., 2019), as well as Juncus microcephalus Kunth, 1816 16 (BELLÉ et al., 2021). 17 2

1 Various studies have been carried out to 2 investigate weeds commonly found in rice fields as 3 hosts of M. graminicola (RUSINQUE et al., 2021). 4 In Brazil, the ability of weeds to host M. graminicola 5 was initially reported by MONTEIRO & FERRAZ (1988), in C. ferax L.C. Rich., 1792, but it was only 6 7 in the 1990s that this plant-parasitic nematode was reported in various species of native and cultivated 8 9 plants in the state of Rio Grande do Sul (SPERANDIO 10 & MONTEIRO, 1991; SPERANDIO & AMARAL, 11 1994). Several Meloidogyne species have been found 12 in rice-growing areas of Rio Grande do Sul, Santa 13 Catarina and Paraná, among them M. graminicola 14 (SOARES et al., 2020), M. ottersoni (Thorne, 1969) Franklin, 1971 (LEITE et al., 2020), M. javanica 15 16 (Treub, 1885) Chitwood, 1949, and M. oryzae Mass, 17 Sanders and Dede, 1978 (MATTOS et al., 2017). In 18 other studies, conducted in assays under greenhouse 19 conditions, reproduction of M. graminicola has been 20 reported in E. crus-galli, C. difformis and C. iria L. 21 1753, (good hosts) in Rio Grande do Sul and Santa 22 Catarina (NEGRETTI et al., 2014). Furthermore, 23 also with artificial inoculation under greenhouse 24 conditions, M. ottersoni was confirmed in E. crus-25 galli, E. colonum, and Phalaris canariensis L., 1753, 26 but little information is available about the range 27 of hosts of this species associated with rice crops 28 (LEITE et al., 2020).

Due to the scenario described above, this study characterized, in greenhouse conditions, the reaction of weeds associated with irrigated rice crops in relation to the species *M. ottersoni* and *M. graminicola*.

34 MATERIALS AND METHODS

The host suitability of weeds that occur in 36 37 flooded rice fields to M. ottersoni and M. graminicola 38 was evaluated under greenhouse conditions at 39 Embrapa Clima Temperado, Pelotas, Brazil. 40 Previously, we applied electrophoresis to confirm the 41 purity of the inocula (CARNEIRO & ALMEIDA, 42 2001). The experiments with M. ottersoni were 43 conducted from December 15, 2020, to February 44 26, 2021 (#1) and again from February 23, 2022, to 45 May 4, 2022 (#2). In the case of M. graminicola, the experiments were carried out from December 20, 46 47 2020, to March 3, 2021 (#3) and from February 24, 2022, to May 5, 2022 (#4). 48

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50 Inoculum origin and identification

51 Isolates were obtained from samples 52 collected in flooded rice fields located in Capão 53 do Leão (*M. ottersoni*) and Uruguaiana (*M.* graminicola), Rio Grande do Sul state, Brazil. The isolates (one eggs mass) were routinely multiplied on rice plants under greenhouse conditions (25 ± 5 °C). Both *Meloidogyne* species were identified based on esterase phenotypes as *M. ottersoni* (Est Ot0; Rm=0) and *M. graminicola* (Est G2; Rm: 0.85, 0.91), according to LEITE et al., (2020). For this purpose, protein extract from both nematodes were individually submitted to a horizontal (continuous) electrophoresis system with polyacrylamide gel (7%) (CARNEIRO & ALMEIDA, 2001) using *M. javanica* [Est J3 (Rm: 1.0, 1.20, 1.35)] as reference.

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Weed seeds: collection, treatment, and sowing

Seeds of weeds were collected from a lowland rice field at the Palma Agricultural Center/ UFPel, located at Capão do Leão, Rio Grande do Sul, Brazil. Seeds collected from *C. ferax*, *C. iria*, and *C. difformis* were submitted to thermal treatment at 40 °C for 3 days to break dormancy (DERAKHSHAN & GHEREKHLOO, 2013). Seeds with fast germination but slow emergence (Cyperaceae) were firstly sown, while seeds with slow germination and fast emergence (Poaceae) were sown later (3 days) in a commercial substrate (Germina Plant Horta Turfa Fértil[®]) and maintained under greenhouse conditions (25 ± 5 °C).

Experimental design

The experiments with both nematodes were performed twice under greenhouse conditions $(25 \pm 5 \text{ °C})$. In both experiments, the design was randomized blocks with 6 (#1, 2 and 3) and 5 (#4) repetitions with 10 treatments (weeds species + control). The weeds tested were *C. ferax*, *C. iria*, *C. difformis*, *Spergula arvensis* L., *O. sativa* (red rice), *E. crus-galli*, *E. colonum*, *Aeschynomene denticulata* Rudd, and *Leersia hexandra* Sw. There was only one assessment for the species *C. iria* and *C. difformis*, since the seeds did not germinate in the first and second periods, respectively.

For the experiments with *M. graminicola*, seedlings with two leaves were transplanted to pots containing 1 L of sterile substrate (18% clay). Experiments with *M. ottersoni* had seedlings with two leaves transplanted to pots with 3 L of the same sterilized substrate (18% clay). *Oryza sativa* cv. BRS Querência (*M. graminicola*) and *O. sativa* cv. IRGA 424 (*M. ottersoni*) were used as susceptible control.

Inoculation of M. ottersoni and M. graminicola and evaluation criteria

Inoculum of M. ottersoni and M. 52 graminicola was extracted from the roots of rice 53

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1 plants, according to the method proposed by 2 HUSSEY & BARKER (1973), using a blender 3 instead of manual shaking for 30 seconds with sodium hypochlorite solution (BONETI & FERRAZ, 4 5 1981). The suspension obtained was then poured into attached sieves and the specimens were collected 6 7 on the 500-mesh sieve. After 10 days, these plants were inoculated with approximately 5,000 specimens 8 9 (eggs plus J2s) (initial population - IP), with the 10 inoculum being deposited at an approximate depth 11 of 2 cm around each plant (two holes). Ten days after inoculation (DAI), the water level was adjusted 12 13 at 1 cm above the soil and maintained during the 14 experimental period. The plants inoculated with M. 15 ottersoni were evaluated at 71 DAI (first evaluation) 16 and 73 DAI (second evaluation), while those 17 inoculated with M. graminicola were evaluated at 70 (first evaluation) and 73 DAI (second evaluation). 18

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20 Evaluation of nematological variables

21 Plant root systems were examined 22 regarding the number of galls (NG) and then were 23 separated from the shooting part, washed, weighed, ground, and processed for extraction of eggs and 24 25 second-stage juveniles (J2s), according to the method described by HUSSEY & BARKER (1973), using a 26 27 blender instead of manual shaking for 30 seconds with sodium hypochlorite solution (BONETI & FERRAZ, 28 29 1981). The suspension obtained was then poured into attached sieves and the specimens were collected on 30 31 the 500-mesh sieve. The extracted specimens (Final 32 population - FP) were counted nematodes on Peter's 33 slide and used to calculate the reproduction factor 34 (RF=FP/IP), according to OOSTENBRINK (1966). 35

36 Statistical analysis

37 The data were analyzed using the R 38 software (version 4.2.1) (R DEVELOPMENT CORE 39 TEAM, 2022). The data referring to the variables NG and RF were transformed by CenterScale, (x+1)1/2 and 40 41 $(x+0.5)^{1/2}$, when necessary to satisfy the assumptions 42 for analysis of variance (ANOVA), with the 43 bestNormalize package version 1.8.3 (PETERSON, 44 2021). The Shapiro-Wilk and Bartlett tests were applied 45 to assess the normal distribution of the residuals and homoscedasticity of the variances, respectively. 46

47 When the assumptions of ANOVA were 48 satisfied, the data were submitted to the Scott-Knott 49 test for comparison of the means ($P \le 0.05$). When 50 the assumptions were not satisfied, even after the 51 transformations, the nonparametric Friedman test 52 was used to analyze the data, with the separation 53 of the means accomplished by the method of Bonferroni adjusted to a confidence interval of 0.05. We considered the weeds to be resistant (poor hosts) when plants showed RF < 1.00; susceptible (good hosts) when RF \ge 1.00; and immune (non-hosts) with RF=0,00 (OOSTENBRINK, 1966).

RESULTS AND DISCUSSION

The reactions of weeds to M. ottersoni are presented in table 1. Differences between the variables were observed between treatments (P ≤ 0.05): A. denticulata did not have galls in any of the evaluations, while S. arvensis presented only a small number in the first evaluation (0.04 ± 0.05) and L. hexandra presented only a small number in the second evaluation (3.17 \pm 2.92) (Table 1). The species *E. colonum* presented intermediate results in both assessments (4.83 ± 0.98 and 15.83 ± 7.57). The greatest NG results were presented by C. ferax (12.16 ± 2.13 and 0.83 ± 0.98), C. iria $(7.83 \pm 1.47 \text{ and } 30.00 \pm 19.96)$, E. crus-galli $(6.84 \pm 2.22 \text{ and } 18.17 \pm 10.26) \text{ and } C. differmis (11.00)$ \pm 1.67). Nevertheless, in comparison with the rice cultivar IRGA 424, this number was very small for all species (129.33 ± 27.74) .

With regard to RF, the lowest values were observed for *C*. *ferax* (0.23 ± 0.08 and 0.07 ± 0.09), *S*. *arvensis* (0.05 ± 0.04 and 0.00 ± 0.00), *A*. *denticulata* (5.46 ± 1.90 and 0.02 ± 0.03) and *L*. *hexandra* (3.53 ± 1.02 and 0.20 ± 0.49), which were significantly lower ($P \le 0.005$) in comparison with *C*. *iria* (8.59 ± 0.76 and 29.90 ± 11.54), *E*. *crus-galli* (7.86 ± 0.96 and 25.22 ± 10.46) and *E*. *colonum* (7.26 ± 1.23 and 26.63 ± 9.62), so they were considered to be good hosts (Table 1). The *C*. *difformis* plants, although only evaluated once, were classified as susceptible (5.33 ± 1.74).

Despite the low NG value, *C. iria*, red rice (7.17 \pm 0.75), *E. crus-galli* and *E. colonum* were classified as good hosts of *M. ottersoni* (RF > 1.0). The species *A. denticulata* (0.00 \pm 0.00) and *L. hexandra* (0.00 \pm 0.00) were judged susceptible in the first evaluation (RF > 1.00), while they were classified as resistant (RF < 1.00) in the second assessment. The species *C. ferax* (0.83 \pm 0.98) and *S. arvensis* (0.00 \pm 0.00) were considered resistant (RF < 1.00).

For M. graminicola, there were significant 46 differences between the treatments regarding the 47 variables assessed ($P \le 0.05$) (Table 2). In the first 48 experiment (2021), the highest NG value was 49 observed for the cultivar BRS Querência (139.60 50 \pm 27.67), followed by red rice (4.20 \pm 0.50) and C. 51 difform is (6.00 ± 0.95) . No galls were detected in 52 the other species. In the second experiment (2022), 53

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Treatments		NG				RF				Reaction ¹	
	2021*		2022**		2021***		2022		2021	2022	
Oryza sativa 'IRGA 424'+	-		129.33	а	-		32.24	а	-	S	
Cyperus ferax	12.16	а	0.83	с	0.23	d	0.07	d	R	R	
Spergula arvensis	0.04	d	0.00	с	0.05	d	0.00	d	R	Ι	
Cyperus iria	7.83	b	30.00	b	8.60	а	29.90	а	S	S	
Oryza sativa (red rice)	7.17	b	19.00	b	5.13	b	12.30	с	S	S	
Echinochloa crus-galli	6.83	b	18.17	b	7.86	а	25.22	b	S	S	
Echinochloa colonum	4.83	с	15.83	b	7.26	а	26.63	b	S	S	
Aeschynomene denticulata	0.00	d	0.00	с	5.46	b	0.02	d	S	R	
Leersia hexandra	0.00	d	3.17	с	3.53	с	0.20	d	S	R	
Cyperus difformis	11.00	а	-	-	5.33	b	-	-	S	-	
CV%	3.89		30.04		10.53		34.49				

Table 1 - Reaction of common weeds in rice fields with Meloidogyne ottersoni.

Means followed by the same letter in the column do not differ significantly based on the Scott-Knott test at 5%; * original values transformed with CenterScale, ** (x+1)1/2 e, ** (x+0,5)1/2; CV = Coefficient of variation; NG = Number of galls; RF = reproduction factor; Sign - = Missed treatment.+ Control.

¹Resistance/susceptibility reaction according to OOSTENBRINK (1966). R - Resistant or poor host; S - Susceptible or good host, I -Immune or non-host.

the species with the highest NG value was 'BRS 1

2 Querência' (36.8 ± 15.51) , a significantly higher 3 result (P \leq 0.05) in comparison with red rice (1.4 \pm

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1.14) and *E. crus-galli* (1.4 ± 1.67) .

5 Although red rice and E. crus-galli 6 presented low NG values, O. sativa was a good host to

M. graminicola in both experiments (RF = 1.08 ± 0.09 7

and 5.42 \pm 1.85), while *E. crus-galli* was classified as resistant in both (RF = 0.56 ± 0.05 and 0.23 ± 0.08) (Table 2). The species C. ferax presented higher NG in the second evaluation (18.0 ± 12.28) and variable RF (0.00 ± 0.00 and 3.76 ± 1.32), thus being classified as a good host. The other species were classified as immune or non-hosts. In this respect, the immunity/

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Table 2 - Reaction of common weeds in rice fields with Meloidogyne graminicola.

Treatments	NG			RF				Reaction ¹		
	2021*		2022*	*	2021*		2022^{*}		2021	2022
'BRS Querência'+	139.60	а	36.80	а	4.50	а	2.13	а	S	S
Oryza sativa (red rice)	4.20	ab	1.4	с	1.08	ab	5.42	а	S	S
Echinochloa crus-galli	0.00	b	1.4	с	0.56	ab	0.23	ab	R	R
Leersia hexandra	0.00	b	0.00	с	0.00	b	0.02	ab	Ι	R
Spergula arvensis	0.00	b	0.00	с	0.00	b	0.04	ab	Ι	R
Cyperus ferax	0.00	b	18.0	b	0.00	b	3.76	а	Ι	S
Aeschynomene denticulata	0.00	b	0.00	c	0.00	b	0.00	b	Ι	Ι
Echinochloa colonum	0.00	b	0.00	c	0.00	b	0.43	ab	Ι	R
Cyperus iria	-	-	11.80	b	-	-	0.11	ab	-	R
Cyperus difformis	6.00	ab	-		0.38	ab	-		R	-
CV %	64.81		30.40		8.11		35.73			

Means followed by the same letter in the column do not differ significantly, * original values analyzed with the nonparametric Friedman test and ** original values transformed by $\sqrt{(x+1)}$ and means analyzed based on the Scott-Knott test at 5%; CV = Coefficient of variation; NG=Number of galls; RF = Reproduction factor. Sign - =Plant not evaluated, +Control.

¹Resistance/susceptibility reaction according to OOSTENBRINK (1966). R – Resistant or poor host; S – Susceptible or good host, I – Immune or non-host.

1 resistance of the species L. hexandra (RF = 0.00 ± 0.00 2 and 0.02 ± 0.01), S. arvensis (RF = 0.00 ± 0.00 and 3 0.04 ± 0.01), A. denticulata (RF = 0.00 ± 00 and 0.004 \pm 0.00) and *E. colonum* (RF = 0.00 \pm 0.00 and 0.43 \pm 5 0.43) was verified, with the cultivar BRS Querência 6 $(RF = 4.50 \pm 0.14 \text{ and } 2.13 \pm 0.60)$ differing.

7 Similar results were observed when 8 weeds were inoculated with M. graminicola, where 9 A. denticulata, L. hexandra and S. arvensis did not 10 present symptoms or reproduction (NEGRETTI et al., 11 2014). Although those authors classified S. arvensis 12 as immune, DABUR et al. (2004) considered it to be 13 a host to the same plant-parasitic nematode.

14 Although the susceptibility of O. sativa and 15 E. crus-galli to M. graminicola has been reported in 16 previous studies (NEGRETTI et al., 2014; KUMAR 17 et al., 2019), in our study E. crus-galli presented as 18 poor host, while O. sativa was a good host. In the 19 study carried out by NEGRETTI et al. (2014), red rice 20 had a higher NG value (38.0) and similar RF value 21 (3.67) in comparison with our results. The low NG 22 and RF results of E. crus-galli in our study did not 23 agree with those found by NEGRETTI et al. (2014) 24 and SOARES et al. (2022), who also classified this 25 species as a good host for M. graminicola, with high 26 RF values in irrigated (5.4 and 110.2) and rainfed 27 conditions (16.20 and 20.30).

28 We also observed differences regarding 29 the RF values of the Poaceae species inoculated with 30 M. ottersoni. We found that E. colonum, E. crus-31 galli and red rice were susceptible to M. ottersoni, 32 corroborating the observations of LEITE et al. 33 (2020), who reported the ability of these plants to 34 host M. ottersoni, with high RF values for E. colonum 35 (110.77) and E. crus-galli (61.56).

Regarding the sedge species evaluated, 36 37 there was variation in relation to the RF of M. 38 graminicola (Table 2). The species C. iria and C. 39 difformis were only evaluated once, as resistant, 40 unlike C. ferax, which was immune and susceptible 41 in the first and second evaluations, respectively. These results differed from those described by 42 43 NEGRETTI et al. (2014), who observed RF values 44 higher than 1.0 for the first cited species. DABUR et 45 al. (2004) also confirmed the ability of C. iria to host 46 M. graminicola, while C. difformis was considered a 47 good host since it can multiply in the plants in rice-48 wheat crop sequences. Likewise, for M. ottersoni, 49 the susceptibility of C. difformis and C. iria was 50 verified, but C. ferax was classified as resistant in both assessments.

The different host reactions found can result from intraspecific variability of the plants and/or physiological variation of the plant-parasitic nematodes (POKHAREL et al., 2010), as well as climate factors (KUMAR et al., 2021). In the case of weeds, the differences can be presumably attributed to the natural variability of the species studied. On the other hand, we could certainly theorize about the variability of RKN populations as well. SOARES et al. (2022) verified that different plants have different responses according to the plant-parasitic nematode, because when analyzing the effect of different variants of M. graminicola within each plant species, they observed significant differences in the most susceptible plants, among them E. crus-galli and E. colonum, with 13 the G1 variant being most aggressive, followed by G3 and the G2 population. Indeed, some authors have also suggested the possibility that different biotypes (races) of *M. graminicola* share unique physiological traits, 17 which can affect the reproductive capacity in specific 18 hosts (SASSER, 1979).

Another factor that can influence the 20 reproduction of plant-parasitic nematodes is soil 21 temperature (ROBERTS et al., 1981) between our 22 research and those described in literature. Studies have 23 demonstrated low initial infection by plant-parasitic 24 nematodes, so it is likely that the combination of low 25 soil temperature and low reproductive potential of the 26 plants results in little or no increase in the number of 27 plant-parasitic nematodes during the evaluation cycle 28 (PLOEG & MARIS, 1999; TIMPER et al., 2006). 29 However, temperatures between 29 °C and 38 °C 30 favor the development of plant-parasitic nematodes 31 (DEVARAJA et al., 2022). The temperature can explain, 32 at least partially, any discrepancies observed in our 33 experiments, since the maximum reached in a greenhouse 34 is 25 °C (greenhouse conditions), but the average minimum 35 temperature in the region during the experimental period was 36 between 17 (experiments 2 and 4) and 23 °C (experiments 37 1 and 3). This temperature range was slightly lower 38 39 than those found in the literature specifically for M. graminicola (MANTELIN et al., 2017), in which some 40 authors also report temperature ranges between 22 and 41 29 °C and between 27 and 37 °C (RUSINQUE et al., 42 2021). The variation of infection can also be associated 43 with temperature changes (RAVINDRA et al., 2017). 44 Our experiments were carried out in different periods 45 when variations in the average temperatures might 46 have influenced the life cycle of the plant-parasitic 47 nematodes. Studies have demonstrated that the cycle 48 of M. graminicola can vary from 19 to 65 days, 49 depending on the temperature. Hence, the number of 50 generations of plant-parasitic nematodes can differ 51 greatly in the same vegetative cycle of the infected 52 53 plant (RAVINDRA et al., 2017).

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1 Similar results were found for the weed 2 host status of *M. ottersoni*, where LEITE et al. (2020) 3 found higher RF values for E. crus-galli and E. 4 colunum at higher temperatures (15 - 25 °C). Perhaps 5 the lack of flooding could explain the higher RF values in this study. Unfortunately, little research has been 6 7 done on this nematode. Its distribution is probably 8 underestimated because it is difficult to detect, and few 9 studies have been carried out on its biology.

10 We observed that the weeds with RF >1.0 can act as important multiplier agents of M. 11 graminicola and M. ottersoni. Our findings are 12 13 important by contributing to knowledge of the wide range of weeds that can serve as hosts of both plant-14 15 parasitic nematodes. Therefore, these results can be 16 utilized as tools to monitor these crop pathogens, to make recommendations for more effective 17 management seeking to eliminate these plants through 18 19 the application of herbicides or the use of cover plants 20 to suppress plant-parasitic nematodes and minimize 21 crop losses (RICH, 2009; JAIN et al., 2012). 22

23 CONCLUSION

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25 Of the weed species that occur between irrigated rice crops, S. arvensis was found to be a 26 27 poor host of M. ottersoni, while L. hexandra and 28 A. denticulada are good hosts. Among the species tested, all except C. ferax were able to serve as hosts 29 for plant-parasitic nematodes. The presence of these 30 31 species in cropland can serve as alternative hosts, so 32 knowledge in this respect is useful to plan measures to control nematodes and eliminate weeds. 33

The species *L. hexandra* and *S. arvensis* are poor hosts of *M. graminicola*, and *A. denticulada* was immune to the nematode.

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46	DECLARATION	OF	CONFLICT	OF
47	INTEREST			

The authors declare no conflict of interest

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50 51 AUTHORS' CONTRIBUTION

51 AUTHORS' CONTRIBUTIO

Conceptualization: KJAL, CBG, DA and JVAF. Data
 acquisition: KJAL, DGRL and DMC. Design of methodology
 and data analysis: CBG and KJAL. KJAL prepared the draft of

the manuscript. All authors critically revised the manuscript and approved of the final version.

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