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RESEARCH ARTICLE

Weeds as A Potential Host of *Xanthomonas Oryzae* Inoculum in Irrigated Rice Systems in Burkina Faso

Wonni Issa^{1,*}, Sanou Adama¹, Berthe Habibatou¹, Zougrana Sylvain¹ and Léonard Ouedraogo¹

¹Institut de l'Environnement et de Recherches Agricoles, 01 BP 910 Bobo Dioulasso 01, Burkina Faso

Abstract:

Background:

Bacterial Leaf blight and bacterial leaf streak caused, respectively, by *Xanthomonas oryzae* pv. *oryzae* and *X. oryzae* pv. *oryzicola* are two rice diseases that occur in rice-growing areas in Burkina Faso. For better management of both pathogens, it is essential to integrate the use of resistant varieties with good agronomic practices including weed management which plays a role in their survival and dissemination.

Objective:

This study aims to identify the main weeds that can act as reservoirs of *Xanthomonas oryzae* strains.

Materials and Methods:

Soil was sampled on three rice-growing sites. In the laboratory, we sifted the soil to collect the seeds of the weeds which were put in pots in the greenhouse for germination. After emergence, all weed species were inventoried and then inoculated to determine their host capacity.

Results:

We identified 14 species belonging to dicotyledons against 10 species from the Poaceae and three species from the Cyperaceae genus. This presence is effective in all plots sampled but at a variable level of prevalence. Our results showed that some Poaceae and Cyperaceae species were the main reservoirs of *X. oryzae* pv. *oryzae* and *X. oryzae* pv. *oryzicola* and behave sometimes like epiphytes. We detected a latent infection of *X. oryzae* pv. *oryzicola* from *Oryza sativa*, *Echinochloa colona*, *Eleusine indica*, *Digitaria horizontalis*, and *Rottboellia cochinchinensis*. Except *Oryza longistaminata*, *Sacciolepis Africana*, *Paspalum vaginatum*, *Paspalum polystachyum*, and *Echinochloa colona* which manifested typical symptoms of bacterial leaf streak, the other species showed yellowing, whitish spots and necrosis at the inoculation site.

Conclusion:

The high prevalence of weeds in rice fields, especially Poaceae and Cyperaceae, is one of the main causes of the importance of bacterial leaf streak disease in the irrigated rice system in Burkina Faso.

Keywords: *Xanthomonas oryzae*, Host, Weed, Burkina Faso, Faso, Cyperaceae genus.

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1. INTRODUCTION

In Burkina Faso, cereals (sorghum, millet, and maize) constitute the staple food of the population and rice is the fourth cereal cultivated in terms of area and production [1]. Despite the enormous potential to increase rice productivity, Burkina Faso imports large quantities of rice each year, which accentuates the country's food dependency. Unfortunately, rice productivity is still very low due to several biotic and abiotic

constraints, including pests and the effects of climate change that affect field yields. It has been reported that over 47% of crop production losses are caused by pathogens. Indeed, more than fifteen pathogens have been identified in rice in Burkina Faso [2 - 5]. The most damaging of these are blast (*Magnaporthe oryzae*), Rice yellow mottle virus, and bacterial diseases caused by *Xanthomonas oryzae* (*Xo*). It causes two important diseases of rice; Bacterial Leaf Blight (BLB) due to *X. oryzae* pv. *oryzae* (*Xoo*) that colonizes xylem vessels and Bacterial Leaf Streak (BLS) caused by *X. oryzae* pv. *oryzicola* (*Xoc*) that colonizes spaces between leaf parenchyma. BLB and

* Address correspondence to this author at the Institut de l'Environnement et de Recherches Agricoles, 01 BP 910 Bobo Dioulasso 01, Burkina Faso; Tel: +22670156832; Fax: +22625340271; E-mail: wonniissa@gmail.com

BLS are occurring in the tropical and subtropical areas of Asia and Australia. In Africa, BLB was reported in several West African countries and Madagascar in the 1980s; whereas *Xoc* was described in these areas in the 2000s [6, 4]. BLS symptoms occur as small, translucent, and typically yellow lesions along the leaf between the veins that expand lengthwise. At later stages, leaves turn grayish-white and die. *Xoo* induces pale-yellow leaf syndrome and seedling blight or “kresek” at an early stage. In Burkina Faso, BLB induced important yield losses in 1998 and 2004, respectively, in the Bagre area. However, BLS is recurrent on the major rice sites in most rice-cultivated varieties [7, 8]. To control both bacterial diseases, it is needed to combine the use of resistant varieties with good cultural practices including weed control. Weeds are at present the major biotic constraint to increase rice production worldwide. In Africa, weeds induce important yield losses from 28 to 74% in the irrigated rice sites and 48 to 100% in the upland rice systems. The importance of their control has been emphasized in the past by various authors [9 - 11]. Among several weeds affecting rice production, two major weed complexes have been identified which require an integrated approach for their control. They are the complex formed by various *Echinochloa* species and the red/weedy rice complex. Wild rice, *Oryza barthi* and *O. longistaminata*, are among the most important weeds in West Africa and the Sahel but total control is not evident. Unfortunately, chemical control is used today against weeds with its harmful effects on human health and the environment. In addition to weed control, the deployment of resistant varieties is one of the best ways to manage bacterial diseases as well as weeds.

In rice, there were 29 major genes for resistance to bacterial blight, but so far only a few quantitative resistance loci for bacterial leaf streak [12]. However, while it is well known that rice is the main host of both pathogens, it has been reported that a number of weeds, notably Poaceae and Cyperaceae [13], played the role of inoculum reservoir. Indeed, weeds can significantly influence disease incidence and should be considered when endeavoring to manage and control plant pathogens of cultivated plants [14]. It is essential to know all factors which contribute to the BLB and BLS epidemic for better management of pathogens. Hence, the aim of our study is to identify the weed hosts of *Xoc* and *Xoo* in the irrigated rice growing system in Burkina Faso.

2. MATERIALS AND METHODS

2.1. Weed Samples

Soil sampling was performed on the rice-growing sites of Karfiguela, Banzon, and Vallée du Kou based on five plots per site. Five samples of three (03) kg of soil were taken in the first 10-20 cm along the two diagonals of each plot. The samples were then pooled to form a composite sample of 15 kg per plot. The samples are bagged and then labeled by assigning them a code, the name of the site, the date of collection, and the GPS coordinates of the plot. In the laboratory, we sifted the soil to collect the seeds of the weeds which were put in pots in the greenhouse for germination. To determine their host capacity, all weed species were inventoried after emergence and then inoculated.

In addition, during the period of a high prevalence of BLB and BLS in the field (September- October), we performed observations on weeds in the plots on the main irrigated sites: Vallée du Kou, Bazon, Karfiguela, Douna, Di, Niassan, Mogtêdo, and Bagre. Symptomatic and asymptomatic samples were collected to determine the presence/absence of *Xoo* and *Xoc* strains.

2.2. Rice Varieties and *Xo* Strains Tested

Two improved rice varieties, TS2 and FKR62, from National Research Institute were used to test the pathogenicity of the strains on weeds. These varieties were previously identified by Wonni *et al* [8]. are susceptible to *Xoo* and *Xoc* strains representing *Xo* diversity in Burkina Faso. The *Xoo* strains BA13 (Race A1) and *Xoc* Strain BA110 [4, 14] were used to inoculate weed and rice varieties, respectively.

2.3. Weed Cultivation for Greenhouse Tests

To obtain the weed seeds contained in each sample, we sieved them using two (02) sieves of different mesh (5 and 0.2 mm). The grains of sand passed through the sieves and the particles retained by the mesh, called refusals, usually contain weed seeds. The refusals obtained were dried in the shade and then weighed.

The seeds were grown in soil (2/3 soil and 1/3 sand) previously sterilized at 100 °C to avoid any development of unsampled weeds. The refusals of each sample were divided into four batches (or four replicates), and were mixed in the soil of each pot. Fifteen days after weed germination, to avoid competition between species for mineral elements, we applied 0.4 g of NPK in each pot in accordance with the recommended dose per hectare (200 kg/Ha) in rice fertilizing. Before the inoculation tests, we applied the same dose again, to eliminate the doubt that the symptoms would be due to a deficiency in mineral elements.

2.4. Weed Population by Sample

We carried out a visual observation and a count of each species. The inventory is updated every 15 days after sowing for three (03) months. We used the Adventrop guide (Weeds of African rice fields and Weeds of subtropics) to identify the species even in the juvenile state. abundance index [15] was used to assess the frequency of individuals of a species within the same family. The frequency of occurrence (number of pots where the species is present) and the maximum abundance per station (maximum score obtained by the species) were then recorded.

Weed diversity is assessed using three diversity indices namely: the Shannon-Weave Index (Ish) which is an indicator of diversity taking into account not only the species richness but also the proportion of each species represented within the community, it varies between 1 and 5,

$$Ish = -\sum (ni/N) \log_2 (ni/N)$$
, where ni is the number of species i and N the total number of species;

Simpson's index (SI), expresses the probability that two (02) individuals chosen at random in an infinite population, belong to the same species, it varies between 0 and 1,

$IS = \sum (ni/N)^2$, where ni is the number of species i and N the total number of species;

Pielou Equitability Index (Eq), measures the degree of diversity achieved by a stand in relation to its maximum value and allows comparison between two groups that do not have the same number of species. It is between 0 and 1,

$Eq = ISh/(\log_2 N)$, where N is the total number of species.

2.5. Evaluation of Weed Epiphytic Capacity

The asymptomatic leaves of the weeds after growing, were collected for *Xo* strain isolation on Peptone-Sucrose Agar medium. Then, Multiplex PCR specific to both pathogens, described by Lang *et al.* [16] was achieved from the leaf's suspension.

Multiplex PCR combines four pairs of primers. It can be performed directly on bacterial colonies or from genomic DNA. The reaction volume of the PCR is 25 μ l composed of 5 μ l buffer 5X containing $MgCl_2$, 0.5 μ l dNTPs, 2.5 μ l primers, 0.05 μ l Taq polymerase, 5 μ l of leaf's suspension. Samples were initially denatured for 5 min at 95 °C, then 35 reaction cycles including (i) denaturation for 30s at 94 °C, (ii) hybridization for 30 s at 60 °C, (iii) elongation for 2mn at 68 °C, followed by a final elongation phase for 10 mn at 68 °C. The amplicons were separated by electrophoresis on 0.8%.

2.6. Identification of Weeds as Hosts for *Xo* Strains

To determine the capacity of the weed species to harbor *Xoo* and *Xoc* strains, we inoculated the weed leaves 30 days after sowing with 108 bacteria/ml suspension. The leaf reactions after inoculation were assessed as follows: (i) presence of typical symptoms after inoculation; (ii) no typical symptoms, but the presence of necrosis at the site of inoculation; (iii) presence of yellowish or whitish lesions at the point of inoculation.

Two weeks after inoculation, the inoculated leaves are collected for the detection of *Xoo* and *Xoc* by isolation or multiplex PCR.

2.7. Evaluation of the Survival and Dynamics of *Xanthomonas Oryzae* Strains after Inoculation

2.7.1. Survival Tests

To assess the survival of *Xo* strains in the leaves of weeds, we used suspensions obtained from ground leaves inoculated and collected at 7, 14, and 21 JAI; to inoculate the susceptible rice varieties TS2 and FKR62N. Symptom induction on one of these two varieties indicates the presence or survival of the bacteria in the weed's leaves inoculated.

2.7.2. Assessment of Bacterial Numbers

The ability of the bacterium to multiply in the plant tissues was assessed at 7, 14, 21, and 30 days after inoculation of one-month-old plants following the method described by Wonni *et al.* (2016). For *Xoo*, the number of bacterial cells in the leaves

was determined within the entire leaf. *Xoc* cell number was assessed from the portion of the leaf where symptoms or necrosis developed, including the site of inoculation and surrounding tissues. Leaf sections were surface-sterilized in 75% ethanol for 10 s, followed by submersion in sterile water for 30 s, and crushed and resuspended in 1ml sterile distilled water. Leaf homogenates were serially diluted (up to 10-12) and plated onto a PSA medium amended with cephalixin (40 ug/ml), kasugamicyn (20 ug/ml), and cycloheximide (50 ug/ml) to avoid contamination. The plates were incubated at 28 °C until single colonies could be counted. The experiment was repeated independently three times.

2.8. Data Analysis

The data were entered into Excel 2010 software. To compare the level of weed and bacterial populations, histograms were constructed and variance analyzes were performed with SPSS20 software. The phytosociological and ordination analyses were performed with PC-ORD version 5.0 ordination software [17].

3. RESULTS

3.1. Weeds Inventoried from Soil

The inventories performed in the three prospected sites identified 24 species including the three (03) major families of weeds, namely, *Poaceae*, *Cyperaceae*, and dicotyledons. In fact, we have counted 12 species belonging to the broad leaf weed family with nine (09) species of the *Poaceae* family and three (03) species of the *Cyperaceae* family. The *Poaceae* were represented by *Pennisetum pedicellatum* Trin, *Setaria pallidifusca* (Schumach.) Stapf & C.E. Hubbard, *Paspalum scrobilatum* L., *Echinochloa colonum* (L.) Link, *Echinochloa stagnina* (Retz.) P. Beauv., *Rottboellia cochinchinensis* (Lour.) W.D. Clayton, *Oryza longistaminata* A. Chev. & Roehr, *Eleusine indica* (L.) Gaertn., *Eragrostis tenela* (L.) Roem. & Schult., and *Digitaria horizontalis* Willd. As for *Cyperaceae*, we identified *Cyperus esculentus* L., *Cyperus difformis* L., and *Fimbristylis littoralis* Gand.

The broad-leaved species inventoried were *Phyllanthus amarus* Schum. & Thonn., *Corchorus tridens*, L., *Portulaca oleracea* L., *Melochia corchorifolia*, L., *Physalis angulate* L., *Cleome viscosa* L., *Euphorbia hirta* L., *Stachytarpheta angustifolia* (Mill.) Vahl, *Triumfetta rhomboidea* Jacq., *Ammania priureana* Guill. & Perr., *Ludwigia hyssopifolia* (G. Don) Exell, *Mollugo nudicaulus* Lam., *Sphenoclea zeylanica* Gaertn. and *Hyptis spicigera* Lam..

3.2. Weed Frequency and Dynamics

Three major families of weeds, namely, *Poaceae*, *Cyperaceae*, and Dicotyledonous, were inventoried at the sites of Bama, Karfiguèla, and Banzon (Fig. 1). However, broadleaf (dicotyledonous) weeds were most abundant at all three sites with 51% at Banzon, followed by *Cyperaceae* (41.02%) at the Karfiguèla site and *Poaceae* Bama (33.51%).



Fig. (1). Weed populations inventoried in soil form Bazon, Bama, and Karfiguèla.

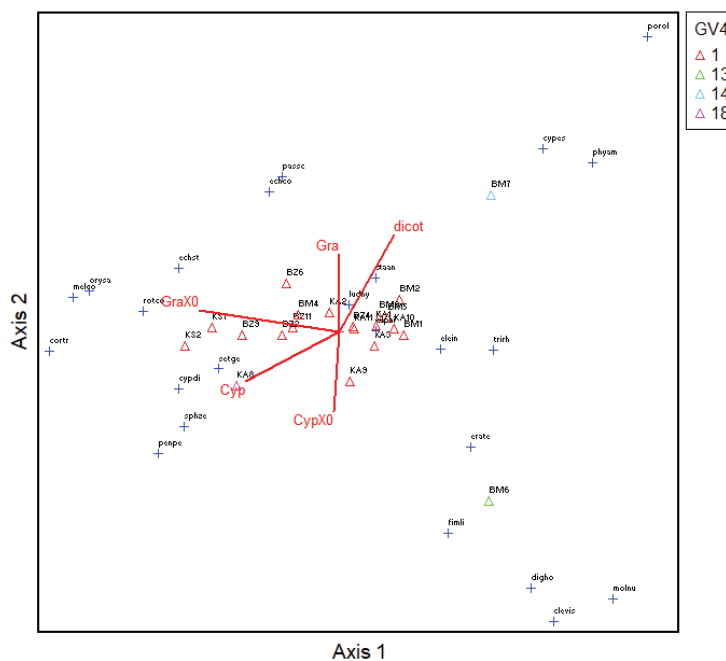


Fig. (2). Dynamic of weeds populations inventoried in soil per site.

GV4: Group of weeds where the effect of the treatments is the most significant;

1; 13; 14; 18: Sub-group of species according to the effect of the most significant treatments.

Concerning the dynamics of weeds, the ordination diagram (Fig. 2) shows that there is a slight link (13%) between the classes, the capacity to be a reservoir of *Xoo* and *Xoc* of weeds, and the distribution of collection sites. Indeed, regarding the capacity to be a *Xo* reservoir, it appears that dicotylodonous (Dicot) species do not host *Xo* strains at any site. However, *Cyperaceae* (Cyp), reservoirs of *Xo* are mainly present in Karfiguèla. As for the *Poaceae* (Gra) reservoirs of *Xo*, they are more present in Bazon and Karfiguèla.

The *Cyperaceae* genus are dominated by two species, *Cyperus difformis* L. with a sum of 99.4 individuals per sample, and *Fimbristylis littoralis* Gand. (Sum = 24.3) (Table 1). The two most important species of *Poaceae* are

Echinochloa colona (L.) Link (sum=16.3) and *Eleusine indica* (L.) (sum= 4.9). Regarding the broad-leaved species, the most inventoried species are *Stachytarpheta angustifolia* (Mill.) Vahl (sum=28.2), and *Ludwigia hyssopifolia* (G. Don) Exell (sum=22.8).

The *Cyperaceae* genus obtained the highest specific diversity with *F. littoralis* (Shannon’s diversity index: H = 1.03) but also the lowest for *C. difformis* (H =0.41).

3.3. Latent Infections due to *Xanthomonas Oryzae* Strains

In order to determine the possible transmission of BLB and BLS by grasses and *Cyperaceae* seeds, we evaluated by PCR and biological tests, the presence/absence of *Xoo* and *Xoc*

strains in the weed's leaves. Of all *Poaceae* and *Cyperaceae* leaves collected, no weed species exhibited symptoms of BLB and BLS after seed emergence and plant development. PCR analysis of the leaf suspension collected from each species of *Poaceae* and *Cyperaceae* revealed the absence of *Xoo* and *Xoc* strains.

Surprisingly, foliar infections performed with foliar suspensions from *Poaceae* and *Cyperaceae* species collected in experimental conditions, on the two susceptible rice varieties, induced apparent BLS symptoms. These are samples of *Echinochloa colona* and *Rottboellia cochinchinensis* from Bazon; *Eleusine indica* from Bama and *Digitaria horizontalis* from Karfiguela. PCR analysis of infected rice leaf samples after inoculation of foliar suspensions of weeds revealed the presence of *Xoc*.

During our surveys, we collected weeds showing typical

symptoms of BLS on *Oryza longistaminata*, *Sacciolepis Africana*, *Paspalum vaginatum*, *Paspalum polystachyum*, and *Echinochloa colona*. However, some weeds exhibited non-specific symptoms of BLB and BLS, which are leaf blight-like. Diagnosis by isolation and multiplex PCR tests has been made to identify mainly *Xoc* strains from these samples which are *Paspalum vaginatum*, *Brachiaria lata*, and *Pennisetum pedicellatum* sp.

Under the infiltration tests in the greenhouse, all species of *Poaceae* and *Cyperaceae* have the capacity to harbor *Xo* strains, without exhibiting typical symptoms of BLB or BLS. These are identified as hosts following reactions:

(i) typical symptoms of BLS (Fig. 3A - C) on *Oryza longistaminata*, *Sacciolepis africana*, *Paspalum vaginatum*, and *Echinochloa colona*;



Fig. (3). Foliar reaction of weed at 7 days after inoculation with *Xoc* BAI10 strain. (A): *Echinochloa colona*; (B): *Paspalum vaginatum*; (C): *Paspalum vaginatum*; (D): *Pennisetum alopecuroides*; (E): *Brachiaria lata*.

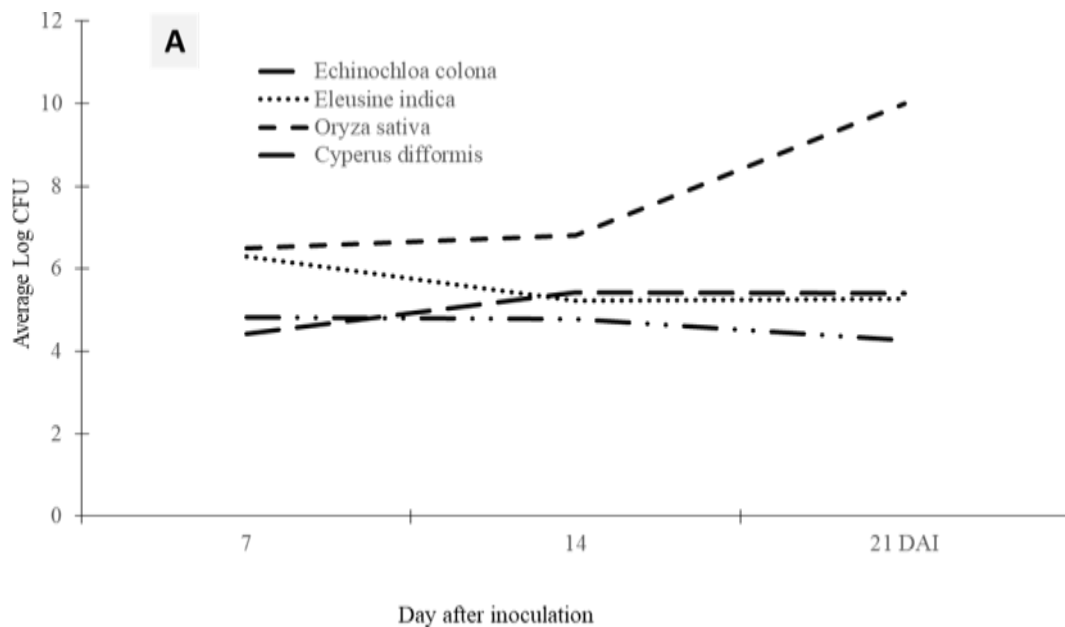


Fig. 4 contd.....

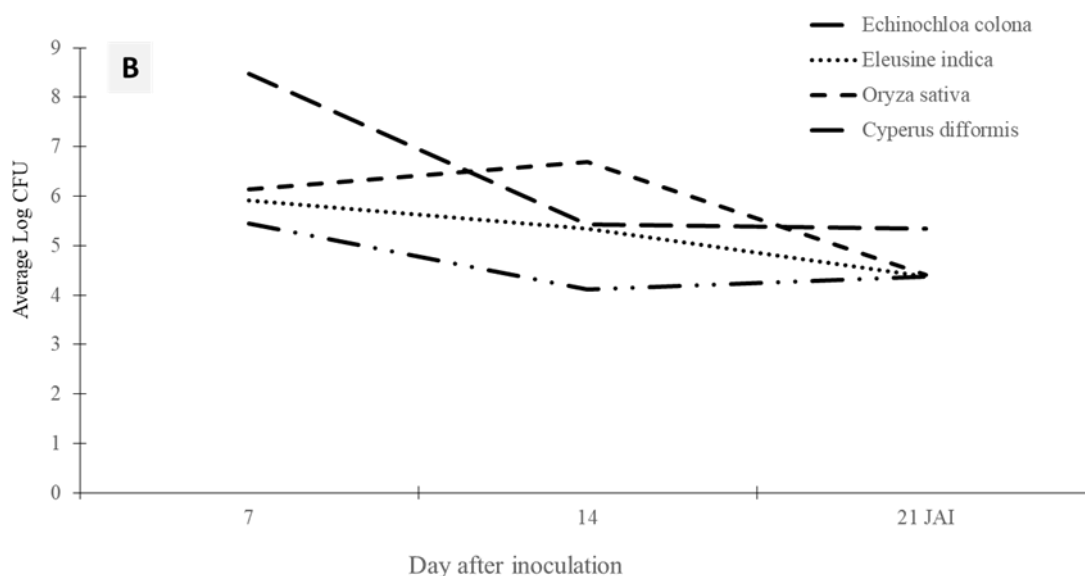


Fig. (4). Dynamic of *Xanthomonas oryzae* strains in weed leaves. (A): *Xanthomonas oryzae* pv. *oryzae*; (B): *Xanthomonas oryzae* pv. *oryzicola*.

(ii) No specific symptoms such as necrosis, whitish and yellowish lesion, or total absence of reaction at the inoculation point were induced by the two pathovars. However, the presence of *Xoo* and *Xoc* has been detected by multiplex PCR or after infiltration of left suspension from infected weeds with FKR 62N and TS2.

Furthermore, to assess the bacteria's ability to multiply in weeds and leaves, we quantified the bacterial populations at 7, 14, and 21 days after inoculation. These quantifications made it possible to confirm the host nature of some weed species.

The results showed that *Xoo* and *Xoc* strains survive in the asymptomatic leaves beyond one (01) month after inoculation. Indeed, high bacterial growth is observed during the first week of inoculation and decreases gradually from 14 days after inoculation (Fig. 3A and B). In general, bacterial growth is greater in grasses than in Cyperaceae. On the other hand, *Xoo* grows more abundantly in the leaves of *O. sativa* than those of weeds. Dicotyledons inoculated with both strains show necrosis and bleaching on the inoculation.

4. DISCUSSION

4.1. Importance and Dynamics of Weeds in the Studied Sites

Three major families of weeds, namely, *Poaceae*, *Cyperaceae*, and Dicotyledons, have been inventoried on the sites of Bama, Karfiguela, and Banzon. However, broadleaf (dicotyledon) weeds were the most abundant, followed by *Cyperaceae* (28%) and *Poaceae* (23%) at all three sites. These results agree with those [18] who showed that broadleaf weeds constitute the most important group of weeds in irrigated rice and lowland areas followed by *Cyperaceae* and *Poaceae*.

Similar results are obtained in irrigated rice with the same weed species by [19], where the species such as *Cyperus difformis*, L., *Fimbristylis littoralis* Gand., *Echinochloa colona* (L.) Link, *Oryza longistiminata* A. Chev. & Roehr. *Ludwigia abyssinica* A. Rich., *Spilanthus uliginosa* Sw. and are some major species found in the rice plains of Bama and Karfiguela. Fifty weed species was inventoried in the rice fields in Ivory Coast, that half of them belong to dicotyledons, the rest are distributed equally between *Cyperaceae* and *Poaceae* [20]. Studies achieved possible after it [20 - 22], in the valley of the Senegal river and in Wianga made it to inventory 90 species of weeds, distributed in 27 families where *Poaceae* and *Cyperaceae* are the most important with respectively 29% and 16%. Given the importance of *Poaceae* and *Cyperaceae*, which represent 50% of the weeds present in the rice fields surveyed in our study, they could constitute potential reservoirs for the two pathovars of *X. oryzae*. These results agree with studies by Sanou et al. [23]. which showed that farmers consider *Poaceae* and *Cyperaceae* to be the most harmful weeds in the irrigated rice system.

The ability of weeds to be a reservoir of *X. oryzae* pv. *oryzae* and *X. oryzae* pv. *oryzicola* would depend on the shape of the leaves, stems, and root system. Indeed, from the distribution of species according to these parameters, it appears that broad-leaved weeds (dicotyledones) are not the host of *Xoo* and *Xoc*. This classification of weeds by Defoer et al. [24], allowed rice farmers to identify *Cyperaceae* and *Poaceae* as the most damaging weed weeds [23].

Cyperaceae and *Poaceae* have been identified as the host group of *Xoo* and *Xoc*. In addition, they contained the greatest specific diversity, in particular with the species *Fimbristylis littoralis*. Their ability to host *Xoo* and *Xoc* would be explained by their morphology; but also their proliferation capacity.

Table 1. Characteristics of the groups of weeds inventoried.

Class	Name	Mean	Stand. Dev.	Sum	Minimum	Maximum	E	H	D'
cyp	cypdi	33.15	46.93	99.44	1	87	0.375	0.412	0.2213
Cyp	fimli	8.11	3.27	24.33	4.333	10	0.945	1.038	0.6305
dicot	staan	9.43	7.65	28.28	2	17.278	0.776	0.853	0.5204
dicot	ludhy	7.61	4.43	22.83	3	11.833	0.887	0.975	0.5914
Gra	elein	4.44	4.23	13.33	2	9.333	0.745	0.819	0.465
Gra	echco	1.63	2.02	4.89	0	3.889	0.731	0.507	0.3254

Note: H: Indices de Shannon; D: Indice de Simpson; E: Indice d'Équitabilité de Pielou

4.2. Poaceae Seeds are likely to Transmit *Xoc*

We identified latent infections due to *Xoc* strains from the leaves of *Oryza sativa*, *Echinochloa colona*, *Eleusine Indica*, *Digitaria horizontalis*, and *Rottboellia cochinchinensis* following inoculation of leaf suspensions of these species on two susceptible rice varieties including FKR62N and TS2.

However, PCR analysis of the leaf's suspension from asymptomatic weeds did not reveal the presence of *Xoo* and *Xoc*. This result could be explained by a low concentration of inoculum in the leaves, therefore, a small amount of amplifiable DNA. Indeed, rice seeds are the main source of *Xoc* inoculum [25 - 28], which can be transmitted to the plant. *Xoc* may also infect *Leersia* sp (p), *Leptochloa filiformis*, wild and cultivated species of *Oryza*, *Paspalum orbiculare*, *Zizania palustris*, and *Zoysia japonica* [29]. *Bracharia lata*, *Oryza longistaminata*, *Pennisetum* sp, and *Paspalum vaginatum* have been reported as possible *Xoc* hosts [4, 15]. Consequently, the offending species weeds are *Poaceae* such as rice, which could support the hypothesis of *Xoc* transmission by the seeds of these weeds.

All species of *Poaceae* and *Cyperaceae* retain *Xoc* after inoculation except for *Fimbristylis littoralis*. Indeed, this species is characterized by hard and thin leaves, difficult to infiltrate. Our results confirm those [30], who indicated that *Xoo* and *Xoc* strains survive on wild or low-cultivated *Poaceae*. Analysis of the bacterial population dynamics in the leaves reveals *Xoc* strain's presence of 21 JAI with or without induced necrosis at the inoculation site [31]. Reported epiphytic survival of *Xoo* on wild grass leaves for 140 days. In our study, we noticed a larger *Xoc* population in the leaves of *Poaceae* than that of *Cyperaceae*. Indeed, *Cyperaceae* by their leaf structure can limit the penetration of bacteria and consequently their growth. They have a cylindrical shape at the level of their rachis, which gives them resistance to attacks from both external and internal factors.

CONCLUSION

Our study revealed that 50% of the weeds present in the plains of Bama, Banzon, and Karfiguela are *Poaceae* and *Cyperaceae*. Evaluation of their ability to harbor pathogens shows that all of them retain the bacteria without typical symptoms of BLB and BLS.

Very interestingly, we were able to highlight latent infections due to *Xoc* from *Echinochloa colona*, *Eleusine indica*, *Digitaria horizontalis*, and *Rottboellia cochinchinensis*, proof that weed seeds are potential reservoirs of *Xoc*.

LIST OF ABBREVIATIONS

<i>Xo</i>	=	<i>Xanthomonas oryzae</i>
BLB	=	Bacterial Leaf Blight
BLS	=	Bacterial Leaf Streak

CONSENT FOR PUBLICATION

Not applicable.

RESEARCH INVOLVING PLANTS

The plant species used in this research were not endangered.

AVAILABILITY OF DATA AND MATERIALS

All the data and supporting information are provided within the article.

FUNDING

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CONFLICT OF INTEREST

The authors declare no conflict of interest, financial or otherwise.

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