## Evaluation of Phytophthora colocasiae Resistance in Taro (Colocasia esculenta) Using Leaf Disc Bioassay

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#### Abstract

Taro leaf blight disease is the most destructive disease affecting taro production worldwide. Identifying resistant genotypes is the most practical means for managing the disease. In this regard, eleven taro genotypes were screened for taro leaf blight disease resistance with four isolates of P. colocasiae (Pc7, Pc12, Pc25 and Pc35) using leaf disc assay. Leaf discs of each genotype was inoculated with approximately  $1 \times 10^4$  zoospores of the P. colocasiae isolates which were arranged in Completely Randomised Design (CRD) with three replications in a factorial experiment. Results of the study showed varied reactions of taro genotypes to the isolates tested. Significant differences (P < 0.05) in lesions size was recorded among the genotypes irrespective of isolate used. Similarly, significant genotype-isolate interactions were observed. Taro genotypes BL/SM/134 and BL/SM/10 inhibited growth of all P. colocasiae isolates. They recorded mean lesion sizes of 16.6 and 17.3 mm compared to 59.9 mm recorded for local genotype (control) at 5days-post-inoculation. The local landrace (check) genotype was susceptible to all P. colocasiae isolates whilst 2 and 7 taro genotypes were categorized as reistant resistant and moderately resistant. It is recommended that the identified resistant genotypes (BL/SM/134 and BL/SM/10) be screened further under natural infestation to confirm results.

Key words: Disease management, Ghana, lesion size, resistance, taro leaf blight

### Reaction Resistante de Genotypes De Taro (Colocasia esculenta) aux Isolats **De Phytophthora colocasiae**

#### Résumé

La rouille des feuilles de taro est la maladie la plus destructive qui affecte la production de taro. Identification des génotypes résistants est important pour une gestion efficace de la rouille des feuilles de taro. Onze génotypes de taro ont été dépistés contre quatre isolats (Pc7, Pc12, Pc25 et *Pc35*) de *P. colocasiae. Le dépistage de la résistance a été effectué en utilisant la méthode de la* feuille détachée. Les résultats de l'étude ont montré une réaction de résistance variée des génotypes aux différents isolats. Une différence significative (p < 0,05) de la taille des lésions a été enregistrée parmi les génotypes indépendamment de l'isolat utilisé pour l'évaluation. De même, des interactions significatives ont été observées entre les génotypes et les isolats. Deux génotypes ont présenté une réaction de résistance à l'isolement de Pc1, six génotypes étaient susceptible, alors que 2 et 1 génotypes étaient modérément résistants et très sensibles à l'isolat, respectivement. Cinq, trois et deux génotypes étaient modérément résistants, susceptibles et résistants à l'isolement de Pc12 respectivement. En ce qui concerne l'isolat Pc25, les génotypes 1,

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7, 2 et 1 ont exprimé respectivement des réactions de résistance, de résistance modérée, sensibles et hautement susceptibles, tandis que 5 génotypes étaient résistants à l'isolat de Pc35, les génotypes 3, 2 et 1 étant respectivement modérément résistants, sensibles et hautement susceptibles.

# Mots-clés: Gestion de la maladie, Ghana, Taille de la lésion, Résistance, Susceptible, La rouille des feuilles de taro.

#### Introduction

Taro (*Colocasia esculenta* (L.) Schott) is an important food security crop staple in the Pacific Islands, West Africa, Asia, the Caribbean and South America (Tarla *et al.*, 2016; Akwee *et al.*, 2015). It is a rich source of carbohydrates, proteins, minerals and vitamins and has medicinal properties to reduce tuberculosis, ulcers, pulmonary congestion and fungal infection (Sharma *et al.*, 2008). In addition, the corms are used for the production of fructose syrup and alcohol (Misra *et al.*, 2008).

Production of the crop is however constrained by high incidence of taro leaf blight disease caused by Phytophthora colocasiae Racib. It is the most destructive disease infecting the crop in major producing countries (Gadre and Joshi, 2003). It is highly prevalent in Ghana (Adomako et al., 2016), although its impact has not been fully established in the country. In countries such as Hawaii, the disease has been associated with heavy yield decline in taro for over 30 years (Miyasaka et al., 2012) and has compelled farmers to abandon their fields or shifted to production of other staple crops. Several management strategies have been employed to manage taro leaf blight disease viz: crop rotation, removal of diseased leaves and use of fungicides. Although metalaxyl-based fungicides have proven effective, like most synthetic chemicals, their detrimental effect on the environment, animals and underground water bodies (Luc et al., 2005) makes them inappropriate option in disease management strategies. Also high incidence of the disease renders it too expensive, as repeated applications are required to protect the crop.

Identification and use of resistant cultivar appears the most sustainable, efficient and cost effective way of managing plant diseases. Screening of taro cultivars for P. colocasiae resistance has traditionally been conducted in disease hot spot zones. Although this method according to Nath et al. (2016) remains the benchmark for evaluating taro cultivars for leaf blight resistance, it relies on the presence and evenly distribution of pathogen inoculum and conducive environmental conditions for disease development. In view of this, previous studies (Tyson and Fullerton, 2015; Brooks, 2008) successfully screened and identified taro cultivars resistance to P. colocasiae using leaf disc bioassay. This method curtails the limitations of uneven inoculum distribution and inconducive environmental conditions. It ensures standard inoculum pressure and uniform disease development on the test material (Nath et al., 2016). Leaf disc bioassay also allows the use of multiple isolates to screen host materials to determine host-isolate interaction over a short period of time. In the present study, 11 taro genotypes were evaluated for their response to four P. colocasiae isolates infection using leaf disc assay.

#### Materials and Methods Preparation of *P. colocasiae* inoculum Diseased taro leaves were collected from taro

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farms in Nkawkaw (Pc1), Asukese (Pc35), Dumasua (Pc25) and Kwamo (Pc12) in the semi-deciduous forest zone of Ghana and brought to the Plant Pathology Laboratory of CSIR - Crops Research Institute, Kumasi where isolations were made on V8 juice ajar medium following standard isolation techniques. Phytophthora colocasiae isolates were maintained on V8 juice agar medium amended with 20 mgL<sup>-1</sup> Nystatin and 250 mg  $L^{-1}$  Ampicillin for 21 days at 26±2 °C in the incubator. Sporangia suspensions were produced by adding 10ml distilled water to the plate of each isolate. The surface of their mycelia were gently scrapped with the edge of a sterilised glass rod to dislodge sporangia. The mycelia-sporangia suspension was filtered through double layered cheesecloth to remove mycelia fragments before chilling at 4°C for 2h to induce zoospore release (Fontem et al., 2005).

#### Sources of taro genotypes

Ten taro genotypes *viz;* CE/IND/16, CE/MAL/32, BL/SM/132, BL/SM/116, BL/SM/134, BL/SM/10, BL/SM/16, BL/SM/115, BL/SM/80 and KAO 022, were obtained from the CSIR-Plant Genetic Resources Research Institute (CSIR-PGRRI), Bunso whilst the check, a local landrace was obtained from a farmer at Kwamo in the Ejisu-Juaben Muncipal of the Ashanti Region of Ghana.

## Evaluation of taro genotypes for resistance to *P. colocasiae*

Screening of taro leaves against *P. colocasiae* was done by detached leaf method (Nath *et al.*, 2016; Tyson and Fullerton, 2015). Leaves of the 11 taro genotypes were detached from 12 week old plants and thoroughly washed under running water. Sampled leaf of each genotype was cut into leaf disc of 80 mm and surface sterilized with 70% ethanol for 60 sec. Ethanol-sterlized leaves were rinsed separately in sterile distilled water to wash off

the ethanol. Single leaf disc of each genotype was placed on adaxial side in a different Petri dish lined with moistened Whatman No 9 filter paper. A single drop (0.5 ml) of inoculum of each P. colocasiae isolate containing approximately 10,000 zoospores was placed at the centre of the leaf disc of each genotype in a Petri dish. The set up was incubated at 26±2 °C for 5 days. Leaf discs inoculated with 0.5 ml distilled water without zoospores served as control for the experiment. The study was a factorial experiment mounted on a Complete Randomized Design (CRD) with three replications. The factors consisted of 11 taro genotypes and four P. colocasiae isolates. Leaf discs were visually examined daily for Taro leaf blight disease symptoms.

#### Data collected and analysis

Lesion size was recorded five days post inoculation (dpi). Lesion size was measured using a leaf area meter. Relative lesion size was determined as a proportion of damaged area of each leaf disc and rated on a 0-5 rating scale (Table 1) to determine disease severity. Based on the disease severity, taro genotypes were categorised as immune, highly resistant, resistant, moderately resistant or susceptible. Disease incidence was determined as the number of taro leaflets infected by an isolate compared to total number of leaflets inoculated. Data collected on lesion size was subjected to Analysis of variance (ANOVA) using the Genstat statistical package version 12. Means were separated using Tukey's Honest significant difference test (HSD) at *P*<0.05.

#### Results

Lesions were found on *P. colocasiae* inoculated sites on all taro leaf discs (Plate 1A) compared to control (Plate 1B) which showed no lesions.

At five days post inoculation, lesion size

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blight disease using leaf disc essays						
Rating Scale	Leaf Area Damage	Host Reaction				
0	0.00	Immune				
1	0.01-10	Highly Resistant				
2	10.01-25	Resistant				
3	25.01-40	Moderately Resistant				
4	40.01-60	Susceptible				
5	>60.01	Highly susceptible				

Table 1: Disease rating scale of Tare last

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Plate 1. Lesion symptoms (A) and assymptomatic (B) taro leaflets following *P. colocasiae* infection

on leaf discs, was significantly different (P < 0.05) among taro genotypes irrespective of *P. colocasiae* isolate used (Table 2). For taro genotypes screened with isolate Pc1, mean lesion size ranged from 13.4 mm in genotype BL/SM/10 to 60.4 mm in the local material (check). Similarly, the local taro genotype (check) recorded the highest mean lesion size of 55.2 mm compared to 10.4 mm recorded in BL/SM/134 when the taro genotypes were screened with isolate Pc12. The local taro genotype (check) again recorded the highest mean lesion size of 62.2 mm and 61.9 mm when taro genotypes were screened with isolates Pc 25 and Pc35 respectively . For the same isolates mean lesion size of 19.4 mm and 18.3 mm were recorded for genotypes CE/IND/16 and BL/SM/115 respectively. Genotype x isolate interaction was significantly different (P<0.05) indicating that taro genotypes responded differently to P. colocasiae isolate used (Table 2).

None of the taro genotypes screened was found to be immuned or highly resistant to any of the *P. colocasiae* isolates used (Table 3). Taro genotypes BL/SM/134 and BL/SM/10 showed resistant reaction to three isolates-Pc1, Pc12 and Pc35 (Table 3). Seven taro genotypes (BL/SM/132, BL/SM/116, BL/SM/80, BL/SM/16, CE/MAL/32, BL/SM/115 and CE/IND/16), showed moderately resistant reaction to isolate Pc12 with two genotypes (KAO 022 and the local taro genotype) being

Taro	Isolates					
Genotypes	Pc1	Pc12	Pc25	Pc35	Mean (Genotypes)	
BL/SM/10	13.4	12.5	31.7	11.7	17.3	
BL/SM/115	28.3	38.1	30.0	18.3	28.7	
BL/SM/116	36.6	32.0	33.0	30.0	32.9	
BL/SM/132	32.4	32.2	25.5	32.3	30.6	
BL/SM/134	13.7	10.4	25.7	16.6	16.6	
BL/SM/16	37.3	31.2	37.9	16.6	30.8	
<b>BL/SM/80</b>	37.9	38.9	38.6	17.7	33.3	
CE/IND/16	27.4	33.5	19.4	32.7	28.3	
CE/MAL/32	38.4	37.7	27.3	27.5	32.7	
KAO 022	42.8	52.7	50.2	46.4	48.0	
CHECK	60.4	55.2	62.2	61.9	59.9	
Mean (Isolates)	33.5	34.0	34.7	28.3		
HSD (P<0.05)						
HSD (Isolates)	1.0					
HSD Genotypes)	1.7					
HSD (GxI)			3.5			

Table 2: Reaction of taro genotypes to four P. colocasiae isolates

Data are means of three replications

susceptible and highly susceptible respectively (Table 3). Seven taro genotypes (BL/SM/134, BL/SM/10, B L / S M / 1 6, B L / S M / 8 0, BL/SM/115,CE/MAL/32 BL/SM/132) and one taro genotype (CE/IND/16) were moderately resistant and resistant to isolate Pc25 (Table 3). Taro genotype KAO 022 and the local genotype were susceptible to isolate Pc 25. With respect to *P. colocasiae* isolate Pc35, taro genotypes, BL/SM/10, BL/SM/134, BL/SM/16, BL/SM/10, BL/SM/134, BL/SM/16, BL/SM/80 and BL/SM/115 were found to be resistant (Table 3).

#### Discussion

The objective of the study was to screen for resistant taro genotypes with different *P. colocasiae* isolates. It was observed that taro genotypes reacted differently to *P. colocasiae* isolates used. The observed variations in genotypes confirm the need to use numerous pathogen isolates to screen for resistance as single isolate may not adequately confirm the susceptibility or resistance of a host genotype. The significant interaction observed between *P. colocasiae* isolates and taro genotypes demonstrates the existence of physiological races in *P. colocasiae*. Identification of resistant taro genotypes to different isolates of

to different P. colocasiae isolates								
	Isolates							
Taro Genotypes	Pc1	Pc12	Pc25	Pc35				
BL/SM/10	R	R	MR	R				
BL/SM/115	MR	MR	MR	R				
BL/SM/116	MR	MR	MR	MR				
BL/SM/132	MR	MR	MR	MR				
BL/SM/134	R	R	MR	R				
BL/SM/16	MR	MR	MR	R				
BL/SM/80	MR	MR	MR	R				
CE/IND/16	MR	MR	R	MR				
CE/MAL/32	MR	MR	MR	MR				
KAO 022	S	S	S	S				
CHECK	HS	S	HS	HS				

Table 3: Reaction level of taro genotypes

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R - Resistant, MR - Moderately resistant,

S - Susceptible, HS - Highly susceptible

P. colocasiae using leaf disc assay agrees with Padmaja (2013) and Brooks (2008) who effectively evaluated and identified resistant taro genotypes using the leaf disc method. Taro genotypes resistant to P. colocasiae have been reported in previous studies by Bassey et al. (2016) and Singh et al. (2006) in Nigeria and Papua New Guinea respectively indicating the presence of resistant genotypes in taro populations. Also, the identification of moderately resistant genotypes identified in this study corroborates findings of similar studies by (Amadi et al., 2015; Singh and Okpul, 2000) who identified moderate resistance of improved genotypes to isolates of P. colocasaie in Nigeria, and Papua New Guinea respectively. The high susceptibility of the local genotype to local P. colocasiae agrees with Ackah et al., (2014) and Amadi et al., (2 0 1 5) who observed that landrace genotypes of taro were highly susceptibility to local pathogen compared to improved

genotypes. Genotypes BL/SM/134 and BL/SM/10 exhibited resistance to multiple P. colocasiae isolates. This confirms Singh et al, (2012) assertion that resistance in taro is controlled by horizontal or partial resistant traits. Horizontal resistance is polygenic and controlled by several genes. It is considered to be more stable and according to Agrois (2005) horizontal resistance is relatively difficult for a pathogen of interest to overcome numerous genes which individually may provide only minor effect against the pathogen. The practical method to manage taro leaf blight disease is to identify and use resistant cultivars which is the most sustainable and cost effective way of managing plant diseases. Screening for resistance have traditionally been in field trials. However, the success of it depends on conducive environmental conditions, inoculum pressure and evenly distribution of the pathogen inoculum on the field (Iramu et al., 2004). Selecting resistant genotypes based on the leaf disc approach is faster and overcomes the challenges associated with field evaluations. In conclusion, the study observed varied resistant reactions among the genotypes. Two genotypes, BL/SM/134 and BL/SM/10 were found to be resistant to three isolates of P. colocasiae used in the study. Eight of the genotypes were moderately resistant to the pathogen whilst the local landrace was highly susceptible to all isolates of the pathogen. The promising genotypes, BL/SM/134 and BL/SM/10 should be screened further under field conditions to validate the laboratory results.

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