

International Journal of Current Research in Biosciences and Plant Biology

Volume 6 • Number 8 (August-2019) • ISSN: 2349-8080 (Online)

Journal homepage: www.ijcrbp.com



Original Research Article

doi: https://doi.org/10.20546/ijcrbp.2019.608.005

Weed hosts of major legume viruses in Niger State, Southern Guinea Savanna of Nigeria

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Article Info	ABSTRACT
<i>Date of Acceptance:</i> 26 March 2019	Field surveys were conducted during the 2015 cropping season to identify and determine the distribution of legume viruses in Niger State, Nigeria. A total of 27
<i>Date of Publication:</i> 06 August 2019	locations were visited during the surveys. Leaves were collected from weed plants showing virus and virus-like symptoms in fields of cowpeas, groundnuts and soyabean. Samples were also collected from asymptomatic plants within the vicinity of infected plants. The antion control plats are supplied in a symptomatic plants within the vicinity of infected
Keywords ACP-ELISA Legume Survey Symptoms Virus disease Weed host	plants. The antigen coated plate - enzyme linked immunosorbent assay (ACP –ELISA) method was employed for virus detection in the collected leaf samples. Results showed that <i>Blackeye cowpea mosaic virus</i> (BICMV), <i>Cowpea mild mottle virus</i> (CPMMV) and <i>Cowpea mottle virus</i> (CPMoV) were the viruses detected. Thus <i>Aeschynomene indica</i> , <i>Amaranthus caudatum</i> and <i>Centrosema pubescens</i> were positive to BICMV; while <i>Aspilia africana</i> , <i>Cleome viscera</i> , <i>Euphorbia hirta</i> and <i>Heterotis rotundifolia</i> were hosts of CPMoV and CPMMV was detected in <i>Chenopodium amaranticolor</i> , <i>Desmodium scorpiurus</i> , and <i>Vicia faba</i> . The detection of these viruses in weed species in the surveyed areas indicates their importance in the ecology, survival and the significant role they play in the epiphytology of the various virus diseases. The occurrence of BICMV, CPMoV and CPMMV in these weed species is believed to be the first report in the study area.

Introduction

Cowpea (*Vigna unguiculata* [L.] Walp), groundnut (*Arachis hypogaea* L.) and soyabean (*Glycine max* [L.] Merril) are the major legumes in sub-Saharan Africa (Ndiaye, 1993; Batiano, 2011). The crops contribute bulk of the protein in the diets of millions of people (Vesper et al., 1999; Lokuruka, 2010). The grains of these crops are a major source of plant proteins for man, feed for animals, and a source of income (Diaz, 2005). Moreover, these plants play an important role in providing soil nitrogen to cereal crops such as maize, millet, and guinea when grown in rotation, especially in areas where poor soil fertility is a problem (Deacon, 1998; Herridge et al., 2008).

Virus diseases are considered a major limiting factor for the productivity of legumes in the tropical and sub-tropical countries (Bashir, 2008; Vincent et al., 2014; Palanga et al., 2016). One of the principal avenues by which these viruses are perpetuated is their over seasoning in weeds (Meliansyah et al., 2012; Makkouk et al., 2015). For instance, it has been documented that perennial grasses in the lowland ecology or *fadama*, might be the original host of *Rice yellow mottle virus* from which it spreads to cultivated rice when conditions become favourable, especially after intensive cultivation (Salaudeen et al., 2008).

In addition, these weed species harbour plant viruses during the growing season and serve as sources of inocula for their secondary spread (Cooper and Jones, 2006; Meliansyah et al., 2012). Odedara et al. (2011) had reported that broad leaves and grasses which occur abundantly around legume fields could serve as alternative hosts and possibly be responsible for inoculum carry-over. Besides weeds competition with crop plants for space, nutrient and light and as host to insect pests, various weed species have been implicated in the epiphytology of viruses of some notable crops (Palanga et al., 2016). Salaudeen et al. (2008) reported that the Rice yellow mosaic virus which causes yield losses of between 25 and 100% was harboured by weed species in the family Cyperaceae. Information on virus types and their weed hosts has a lot of implication on virus survival and epiphytology. The ultimate goal of such information is for designing sustainable management strategies against legume viruses. The objective of this study was, therefore, to identify the weed hosts of major legume viruses in Niger State, Southern Guinea Savanna of Nigeria.

Materials and methods

Description of the study area

Niger State is located in the Southern Guinea Savannah agro-ecological zone of Nigeria and lies between latitude 6° 8′ E and longitude 8° 44′ N of the equator. The site experiences distinct dry and wet seasons with an annual rainfall ranging from 1100 mm in the northern part to 1600 mm in the south with a mean of 1350 mm. The rainfall which peaks in September normally begins in April and ends in October. The temperature ranges between 35 and 37.5°C with relative humidity between 40 and 80% in January.

Collection of samples

A field survey was conducted in major legume

producing areas of Niger State with crops aged 6 – 9 weeks after planting in September and October, 2015. The state has 25 Local Government Areas (LGAs), grouped into three Agricultural zones. A multi-stage sampling procedure was employed to select three LGAs from each of the three zones, giving a total of 9 LGAs. Three villages/locations from the nine LGAs were randomly selected for the survey to give a total of 27 locations. Twenty leaf samples with symptoms such as leaf mottling, mosaic, leaf curling, distortion, chlorotic spots and stunting were collected from weed species growing from the edges and within the legume fields.

The weed species were botanically identified and classified by their morphological characteristics as described by Akobundu and Agyakwa (1987). The leaf samples were preserved over silica gels in airtight vial tubes. In each location, samples were collected at approximately 5 to 10 km from the previous sampled farm. Geographical location of each field denoting longitude and latitude was recorded using the Global Positioning System (GPS- 4300) equipment (Ethrex Garmin GPS, Taiwan) (Table 1).

Serological detection of legume viruses

The sampled leaves were subjected to antigen coated plate enzyme-linked immunosorbent assay (ACP - ELISA) as described by Kumar (2009) and absorbance values were quantified at 405 nm using a microplate reader (MRX, Dynex Technologies, Inc., USA) after overnight. Values were accepted to be positive when the optical density reading was at least twice that of the mean for the negative control.

Results

Prevalence of characteristic virus symptoms

The results showed that mosaic was the most prevalent symptom in all the locations surveyed. Leaf mottling was the next most rampant virus symptom observed during the survey. Leaf curl was another frequently encountered symptom followed by chlorosis, but not as rampant as the mosaic and leaf mottling symptoms. Stunting of plants, leaf puckering and deformation, dead of plants and others were also recorded but at specific locations only (Fig. 1).

Table 1. GPS coordinates for locations where farms were surveyed	Table 1. GPS	coordinates f	or loca	tions whe	ere farms	were surve	yed.
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Town	Longitude/Latitude	Elevation	
Paiko Jazu	09 º 25.672 N/ 006 º 39.447 E	353.0 m	
Tungan Makum	09 º 29.291 N/ 006 º 37.384 E	277.4 m	
Kuta	09 º 51.479 N/ 006 º 42.740 E	320.2 m	
Shiroro Dam	09 ° 55.582 N/ 006 ° 48.797 E	379.0 m	
Zungeru	09º 46.300 N/ 006º 08.846 E	121.0 m	
Dabiri	09 ° 35.895 N/ 006 ° 01.129 E	97.1 m	
Lemu	09º 22.290 N/ 006º 01.703 E	136.9 m	
Gidan Gwari	09º 14.440 N/ 006º 09.054 E	118.5 m	
Zachinta	09º 07.678 N/ 005º 54.765 E	153.3 m	
Kosteni	09º 11.925 N/ 005º 25.464 E	144.4 m	
Awuru	09° 49.977 N/ 004° 35.019 E	162.3 m	
Ba'aburasa	09º 53.462 N/ 004º 23.749 E	254.2 m	
Manigi	09º 44.196 N/ 005º 28.901 E	215.4 m	
Makera	09º 37.281 N/ 0050 21.621 E	204.7 m	
Zugurma	09º 28.909 N/ 004º 57.315 E	148.4 m	
Bokani	09º 26.958 N/ 005º 93.490 E	163.1 m	
Mokwa	09º 18.884 N/ 005º 07.122 E	206.9 m	
Rafin Gora	10°06.304 N/ 005°24.155 E	288.2 m	
Farin shenge	10 ° 24.453 N/ 005 ° 30.859 E	346.3 m	

Table 2. Reaction of weed species in Enzyme Linked Immunosorbent Assay (ELISA) in Niger State Southern Guinea

 Savanna of Nigeria in the year 2015.

Weed species	Growth	Growth Polyclonal antibody						
weed species	habit	CABMV	BICMV	CMV	SBMV	CPMoV	CYMV	CPMMV
Aeschynomene indica (Linn.)	А	0.113	0.613*	0.121	0.204	0.139	0.250	0.211
Ageratum conyzoides (Linn.)	А	0.230	0.182	0.214	0.119	0.291	0.244	0.117
Amaranthus caudatum L.	Α	0.210	0.839*	0.118	0.299	0.237	0.201	0.481*
Aspilia africana (Pers.) C.D Adams	Α	0.221	0.231	0.091	0.201	0.382*	0.110	0.291
Cenchrus biflorus Roxb.	А	0.229	0.197	0.239	0.103	0.154	0.249	0.102
Centrosema pubescens Benth.	А	0.211	0.892*	0.101	0.199	0.127	0.102	0.110
Chenopodium amaranticolor (Coste &	Α	0.221	0.710*	0.229	0.120	0.101	0.212	0.429*
Reyn)								
Cleome viscosa L.	А	0.091	0.118	0.197	0.217	0.451*	0.228	0.102
Desmodium scorpiurus (Sw.) Desv.	Α	0.110	0.401*	0.120	0.193	0.207	0.229	0.494*
Euphorbia hirta (Linn.)	Α	0.211	0.291*	0.111	0.129	0.429*	0.221	0.098
Heterotis rotundifolia (SM)	Р	0.162	0.221	0.102	0.226	0.562*	0.081	0.170
Hyptis suaveolens (Poit)	Α	0.110	0.129	0.298*	0.190	0.281	0.153	0.461*
Laportea aestuans (L.) Chew.	Α	0.211	0.105	0.203	0.214	0.221	0.092	0.117
Physalis angulata (Linn.)	Α	0.119	0.049	0.106	0.117	0.162	0.114	0.215
Talinum triangulare (Jacq.) Willd.	Р	0.225	0.104	0.217	0.182	0.135	0.139	0.231
Tridax procumbens Linn.	А	0.219	0.193	0.221	0.210	0.107	0.183	0.216
Vicia faba L.	Α	0.238	0.249	0.148	0.292	0.119	0.232	0.459*
Diseased		2.138	2.377	2.586	2.140	2.584	0.262	0.452
Healthy		0.138	0.124	0.132	0.154	0.142	0.127	0.191
Buffer		0.128	0.141	0.138	0.167	0.141	0.165	0.214

CABMV = Cowpea aphid borne mosaic virus; BICMV = Blackeye cowpea mosaic virus; CMV = Cucumber mosaic virus; SBMV = Southern bean mosaic virus; CPMMV = Cowpea mild mottle virus; CMoV = Cowpea mottle virus; CPYMV = Cowpea yellow mosaic virus. A = annual: P = perennial: * positive.

Weed host of legume viruses

Results obtained from the identification of weeds that are hosts to legume crops in Niger State, Southern Guinea Savanna of Nigeria using ACP -ELISA are shown in Table 2. The results show the presence of viruses in some specific locations. Blackeye cowpea mosaic virus (BICMV), cowpea mild mottle virus (CPMMV) and cowpea mottle virus (CPMoV) were the only viruses detected in weeds. These viruses occurred in both single samples and mixtures of two or more collected leaves at the different locations surveyed. Specifically, BICMV was found in Dabiri, Farin Shinge, Gidan Kwano and Tatiko, CPMMV was detected in Farinn Shinge and Manigi while CPMoV was found in Awuru, Gidan Kwano, Lemu, Manigi, Mokwa and Tatiko. Similarly, *Aeschynomene indica* (Linn.), *Amaranthus* caudatum L., Centrosema pubescens Benth., Corchorus spp. (Jacq.) Willd. and Aspilia africana (Pers.) C.D Adams species, BICMV and Cleome viscosa L. were detected in the area hosting BICMV, CPMMV and CPMoV respectively.



Fig. 1: Some weed samples with symptoms of virus infections during the 2015 survey.

Discussion

The virus symptoms which include leaf mottling, mosaic, leaf curling, distortion, chlorotic spots and stunting observed on the sampled weeds are in agreement with reports on viral infected legumes in the tropics by several workers (Shoyinka et al., 1997; El-Muadhidi et al., 2001; Najar et al., 2011; Makkouk et al., 2014). Occurrence, distribution and spread of these viruses could probably be influenced by environmental factors, presence of suitable and susceptible hosts and presence and activities of vectors. Incidence and distribution of the viruses were natural and may have stemmed primarily from seed infection and weed hosts as observed from the cropping pattern in the present study areas and as reported by Alabi et al. (2010).

The BICMV and CPMoV incidences recorded at Lemu and Dabiri which are separated by less than 5 km collaborate the findings of Aliyu et al. (2012) who discovered two different viruses from the same genus co-existing in the nearby field in Kwara State, Nigeria. This implies that subsequent mutation and replication of the viruses could simply result in several serotypes with varying degrees of virulence on the one hand and multiple infections of legume crops on the other. The occurrence of CPMoV and CPMMV in naturally infected cowpea witnessed from this study is believed to be the first report from Niger State. BICMV was detected at Awuru and Gidan-Kwano in mixture with CPMoV.

The CPMoV incidence was prevalent in Tatiko, Manigi, Awuru, Lemu and Mokwa Dabiri, locations. Although, Awuru is located at the riverine area which partially agrees with the report of Alegbejo (2015), who recorded high incidence of the pathogen in riverine areas of the middle belt of Nigeria which has a Southern savanna climate and where a lot of bambara groundnut is grown. Furthermore, this finding shows that the pathogen can also be spread to other areas within the Southern savanna which are not riverine. The seed borne nature of CPMoV and its recent detection among the surveyed legume crops, suggest that the virus could be spreading through seeds to other parts of the Southern guinea Agro-ecological zone of Nigeria.

Cowpea mild mottle virus has been reported to occur naturally in the middle belt of Nigeria which includes Niger State (Alegbejo, 2015). Odedara (2011) reported CPMoV as a seed-borne virus that is considered a major constraint to yield in legume fields. Due to emerging plants which quickly get exposed to virus inocula with greater damage at the early stages of crop plant development shows how important the virus can be in the study area.

The negative reaction of some of the weed leaf samples to ELISA in the present study implies that they belong to entirely different virus types or are

non-legume viruses. The detection of CPMoV in some of the weed species indicates its importance in the ecology as well as its survival. Odedara (2011) reported that non-crop hosts of viruses can be particularly troublesome if they act as over seasoning hosts and remain unknown since bioassay must be carried out before to confirm their presence. The weed species which were implicated in the leaf samples from the surveyed legume fields show that they can harbour these pathogens during the growing season and serve as sources of inocula for their secondary spread. This is in agreement with report by Naimuddin, et al. (2014) who found Ageratum conyzoides harbouring Mungbean yellow mosaic India virus and thus acting as an important source of primary inoculum of the virus for the recurrence of yellow mosaic disease in grain legumes in northern India. Thus, these viruses can survive on these weed species during the off season and then serve as sources of primary inocula at the beginning of the new season.

Alegbejo (2015) earlier reported that the more important natural reservoirs of these pathogens are the annual weeds from theLeguminosae which can host the organisms all the year round and for extended numbers of years. Odedara (2011) also maintained that the viruses can also survive in infected dry leaves of the susceptible annual leguminous weed species, resulting in accumulation of virus inocula.

In the present study, Aeschynomene indica (Linn.). Amaranthus caudatum L., and Centrosema pubescens Benth. were among the common weed species in which viruses were detected in the study area. Hampton and Thottappilly (2003) reported that some of these forage legume species are naturally infected by these viruses. That CPMMV and CPMoV were detected in Corchorus spp. (Jacq.) Willd., and Aspilia africana (Pers.) C.D Adams, which is contrary to the work of Alegbejo (2015), who reported them as non-hosts. This might be due to differences in the virulence of the virus strains involved.

The detection of BICMV in *Cleome viscera* L., contradicts the findings of Odedara et al. (2011) who reported the weed as non-host of the virus and this can also be attributed to the virulence of the

viruses strains assayed. Infection of *Euphorbia hirta* (Linn.) and *Heterotis rotundifolia* (SM) CPMMV in this study, is at variance with the report of (Alegbejo, 2015) who found them as only hosts of BICMV.

The infection of *Hyptis suaveolens* (Poit) by CPMMV, *Aeschynomene indica* L. and *Centrosema pubescens* Benth. by BICMV and *Cleome viscera* L. by CPMoV viruses respectively under natural conditions as hosts in the field in the present study is a first report on them.

Conclusion and recommendation

Legume production in Niger State suffers from high virus infection with resultant substantial loss in grains. Virus diagnosis showed that three important legume viruses, BICMV, CPMoV and CPMMV were prevalent in some specific locations surveyed. Aeschynomene indica (Linn.), Amaranthus caudatum L., Centrosema pubescens Benth., Corchorus spp. (Jacq.) Willd., and Aspilia africana (Pers.) C.D Adams species and Cleome viscera L were detected in the area hosting BICMV, CPMMV and CPMoV respectively. There is, therefore, need for constant monitoring of legume crops fields through regular field disease surveillance and surveys to identify new and emerging weeds as starting point for legume viruses' management in the study area. Further studies should be carried out on the weed samples that showed weak positive and negative reactions to the identified viruses from this study in order to validate their roles in the virus-weed host-legume crops complex for their increased production to ensure food security.

Conflict of interest statement

Authors declare that they have no conflict of interest.

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How to cite this article:

Ahmed, A. A., Kolo, M.G.M., Salaudeen, M.T., Wada, A. C., 2019. Weed hosts of major legume viruses in Niger State, Southern Guinea Savanna of Nigeria. Int. J. Curr. Res. Biosci. Plant Biol. 6(8), 30-36. **doi:** <u>https://doi.org/10.20546/ijcrbp.2019.608.005</u>