

INCIDENCE OF SOME SWEET POTATO VIRUSES, THEIR INTERACTION AND FREQUENCY UNDER FIELD CONDITIONS

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ABSTRACT

During 2005-2006 a field survey of sweet potato viruses was carried out and 19216 plants growing in 42 feddans were examined. Symptoms suspected to be due to virus infection reached about 4.5%, 10% and 22% in Menoufeya, Damietta and Kalyoubia Governorates respectively. Symptomatic plants showed mosaic, yellowing or leaf curl accompanied with mosaic. Percentage of each pattern was varied according to the tested cultivar and location. One hundred and forty six symptomatic and sixty nine asymptomatic plants were collected and serologically analyzed by ELISA against the most wide spread sweet potato viruses (i.e. SPFMV, SPMMV, SwPLV, SPCFV, SPCaLV, SPMSV and SPCSV). Data revealed that 13 % of asymptomatic and 68% of symptomatic plants was virus(s)-infected. SPFMV were detected in most infected plants either exhibiting symptoms or not. The virus existed either alone or in combination with the other viruses in double and mixed infections. Plants singly infected with this virus showed mild or no symptoms. Variable data were obtained with the other viruses. In contrast, SPMMV was found to be the lowest detectable virus. It was not detected in any single infection but tended to combine with SPFMV in double and mixed infections. Only three samples were infected with all tested viruses. Due to its wide spread, incidence in high frequency and problems facing its detection and diagnosis, some additional studies were carried out on the most predominant virus, SPFMV. It was identified by NCM-ELISA, immunosorbent electron microscopy and specific primers were used to ensure the identity.

keywords: Sweet potato viruses, Survey, Detection, NCM- ELISA. ISEM- PCR.

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INTRODUCTION

Sweet potato, *Ipomoea batatas* (L.) Lam. is the third most important root crop in the world after potato (*Solanum tuberosum* L.) and cassava (*Manihot esculenta* Crantz) (Anonymous, 1998). During 2004, the global cultivated area with sweet potato was about 8,750,695 hectare that produced 129,403,827 tons (Anonymous, 2005).

Sweet potato requires few inputs, making it appropriate for subsistence farmers with limited resources. Among the major starch staple crops, it has one of the highest rates of production per unit area per unit time (Woolfe, 1992). However, production is greatly constrained, particularly by viral diseases that cause great yield reduction ranging between 56 and 98% (Ngeve, 1990 and Gibson *et al.*, 1997).

Sweet potato feathery mottle potyvirus (SPFMV) occurs in Africa and in everywhere sweet potato is grown (Karyeija *et al.* 1998). Economic

loss may be associated with external cracking and internal corkiness, making the tuberous roots unmarketable. However, there are few reports about this form of losses from Africa. In a small screen house trial in Uganda, a root yield of virus-free clone was about twice of the same inoculated clone (Gibson *et al.*, 1997).

In Egypt the cultivated area increased from 9,130 to 10,000 feddans during 2000-2005 and the production was also increased from 249,548 to 300,000 tones respectively (Anonymous, 2005). Ishack *et al.* (2000) found that SPVD was the most prevalent virus disease in Egypt, which was detected during survey program (1997-2000) in the International of Potato Center in Gharbia.

The present study aimed to a) detect viruses infecting sweet potato in some productive area, b) determine the percentage of incidence of each virus and c) elucidate virus(s) frequencies and combinations in the infected plants.

MATERIALS AND METHODS

Survey of viruses infecting sweet potato was conducted in three productive Governorates i.e. Damietta, Menoufeya and Kalyoubia. A total of 19216 plants growing 42 feddans were examined with X- shape transects stretching between opposing corners of each field (Ateka *et al.*, 2004). Number of tested plants was correlated with the area of each field. Number of plants exhibiting external symptoms suspected to be due to virus infections as well as type of symptoms was recorded for each selected field.

Virus detection and identification

One and half month after transplanting, 131 and 15 symptomatic, 64 and 5 asymptomatic sweet potato plants were collected from different locations at Kalyoubia and Damietta respectively. Collected plants were tested against the most wide spread sweet potato viruses i.e. *Sweet potato feathery mottle virus* (SPFMV), *Sweet potato mild mottle virus* (SPMMV), *Sweet potato latent virus* (SwPLV), *Sweet potato chlorotic fleck virus* (SPCFV), *Sweet potato caulimo-like virus* (SPCaLV) and *sweet potato mild speckling virus* (SPMSV). Detection of these viruses was carried out by nitrocellulose membrane enzyme-linked immunosorbent assay (NCM-ELISA) as done by Anonymous (2001), using standard kits obtained from the International Potato Center (CIP). Detection of *sweet potato chlorotic stunt virus* (SPCSV) was done by triple antibody sandwich (TAS)-ELISA in microtitre plates as described by Gibson *et al.*, (1998). Percentages of infection with single double or multiple virus(s) were recorded.

Detection and identification of SPFMV:

Samples of Mabroka sweet potato plants found to be SPFMV infected (as tested by ELISA), were extracted, and their infectious sap was used to inoculate *Ipomoea nil* and *I. setosa*. Three weeks later, inoculated plants were used in the following experiments.

Immunosorbent electron microscopy:

Identification was carried out by immunosorbent electron microscopy (electron transmission microscope, at Al-Azhar University, JEOL, JEM 1010; Japan) as described by Derrick (1973). The shape and size of the virus were determined by using the software analysis (Doc 2.11.005, soft imaging GmbH, 1986-1996).

Reverse transcription polymerase chain reaction (RT-PCR):

Total RNA was extracted from infected and healthy *I. nil* as well as from infected *I. setosa* and Mabroka sweet potato plants, according to Kern *et al.*, (2005). Two specific primers were used i.e. the up stream primer SPFMV1 (5' -ATA GTG GGG GCA TCA TCA AAG G- 3') and the down stream primer SPFMV 2 (5' - CCT AAA AGT AGG CAC TGC ATG G- 3'). RT-PCR was carried out at 42 °C for 45 min in 10 mM Tris- HCL buffer(pH 8.3) containing 4 µl samples RNA, 20 pmol primers, 2.5 mol dNTPs, 30 units reverse transcriptase (RT), 5 units RNase, 1 unit Taq DNA polymerase. The reaction was predenaturated for 2 min at 95°C, annealed for 30 sec at 63°C and then extended for 1 min at 72°C for a total of 40 cycles in DNA thermal cycler (Perkin Elmer Modele 480, USA). PCR amplified products were separated by agarose gel electrophoresis. Aliquots of 10 µl of PCR products were analyzed on 1% agarose gel in TEB A buffer 1X (89 Mm Tris, 89 mM borate and 2.0 mM EDTA, pH 8.3) at 100 volt for 1 h. The gel was stained with ethidium bromide at a concentration of 0.5 µl / ml. DNA molecular weight marker was used to determine the size of RT-PCR amplified cDNA products of SPFMV. Bands of DNA were visualized on a UV transilluminator and photographed using documentation system (Bio/ Doc Analyze, Biometra)

RESULTS AND DISCUSSION

There is a lack of information about ecology aspects of sweet potato viruses under field conditions. Also, there are no available data on the interactions between different viruses affecting sweet potato in Egypt. In the present study, we tried to shed some light on the situation and interaction of sweet potato viruses in some productive areas in Egypt.

Accordingly, field survey was conducted at 42 feddans in three productive Governorates i.e. Kalyoubia, Damietta and Menoufiea using the common cultivar in each Governorate. Infected plants showed mosaic, yellowing and/or leaf curl accompanied with mosaic (Fig. 1 A, B and C). At some locations mosaic appeared on young leaves, while mild chlorotic pattern was prominent on older leaves, Data in Table (1) revealed that, Kalyoubia has the highest percentage of infection (21.95 %), followed by Damietta (9.75%) and then by Menoufiea (4.42%). Among symptomatic plants the most prevalent symptoms in Kalyoubia was mosaic (68.58%) followed by yellowing (31.41%) and no leaf curl was observed on any plant. In Damietta, yellowing recorded 52.15% followed by leaf curl with mosaic (47.89%) whereas in Menoufiea, leaf curl with mosaic raised to 83.33% followed by yellowing (16.66%). No mosaic symptoms were observed neither in Damietta nor in Menoufiea. Incidence of different patterns varied greatly according to the cultivar and the location. The results were in parallel with that obtained by Mukasa *et al.* (2003) and Aritua *et al.* (2007).

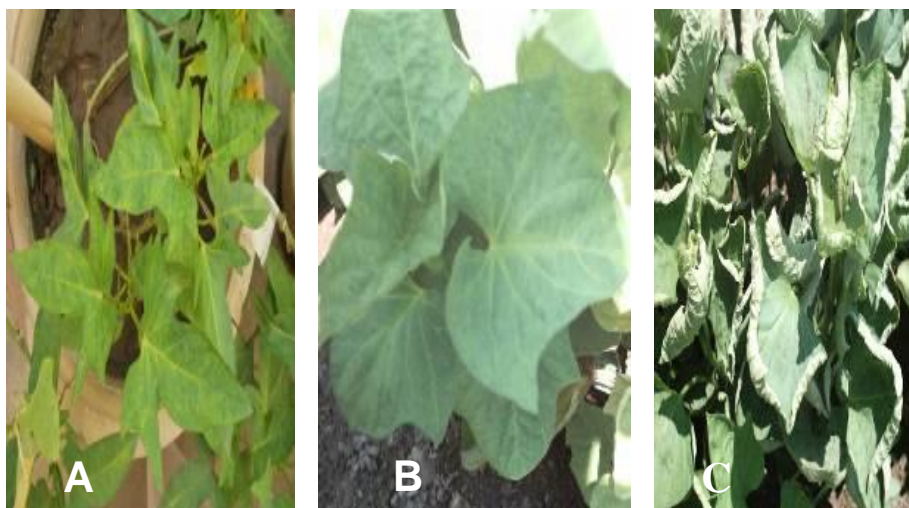


Fig. (1): External symptoms on some naturally infected sweet potato cultivars. (A): Abess showing mosaic. (B): Mabroka showing mosaic. (C): Menoufiea 6 showing leaf curl & mosaic.

Table (1): Percentages of external symptoms on sweet potato cultivars in three Governorates.

Governorate	Cultivar or hybrid	Symptom incidence*			
		S/T	M/S	Y/S	(Lc+M)/S
Kalyoubia %	Mabroka	191/870 21.95	131/191 68.58	60/191 31.41	- -
Damietta %	Abess	190/1948 9.75	- -	99/190 52.1	91/190 47.89
Menoufiea %	M 6	726/16398 4.42	- -	121/726 16.66	605/726 83.33

*Number of:

S= symptomatic plants

M= mosaic

Y= yellowing

L.c.= leaf curl

T= tested plants

Detection and frequency of some sweet potato viruses

Symptomatic (146) and asymptomatic (69) sweet potato plants were collected from the most two productive Governorates i.e. Kalyoubia and Damietta. Infected plants were analyzed serologically with NCM-ELISA against SPFMV, SPMNV, SwPLV, SPCFV, SPCaLV, SPMSV and TAS-ELISA against SPCSV.

In Kalyoubia Governorate Table (2) 86 out of 131 symptomatic samples were found to be virus infected (65.6 %). Single, double and mixed infections represented 58%, 20.9% and 20.9 % respectively. SPFMV was the most prevalent virus either in single (66%), double (88, 9%) or in mixed (100%) infections. These results are in harmony with those obtained by Ateka

et al. (2004) who found that SPFMV was the most common and the most widespread virus in Kenya. In contrast, SPMMV was found to be the lowest detectable virus representing about 5% of all infected plants. In addition, this virus was not detected in any single infection and tended to combine only with SPFMV either in double and mixed infections. SPMSV tended also to combine more frequently with SPFMV either in double or mixed infections. SPFMV was detected in all mixed infections. All the seven tested viruses were detected in three samples only, representing about 3.5% of all infected plants.

Table (2): Natural occurrence and frequency of seven viruses in diseased sweet potato plants grown under field conditions as indexed by NCM-ELISA

Location	Frequency of the tested viruses in single, double and mixed infections		
	Single	Double	Mixed
Kalyoubia	SPFMV(33)	SPFMV+SPMMV(2)	SPFMV + SPCSV + SPCaLV (1)
	SPCSV(5)	SPFMV + SPCFV (2)	SPFMV + SPCaLV + SPMSV (2)
	SwPLV(2)	SPFMV + SwPLV (4)	SPFMV + SPCaLV + SwPLV (1)
	SPMSV(2)	SPFMV+ SPCaLV (2)	SPFMV + SPCFV + SPCaLV + SPMSV (2)
	SPCFV(4)	SPFMV + SPCSV (2)	SPFMV + SPCaLV + SwPLV + SPMSV (1)
	SPCaLV(4)	SPFMV + SPMSV (4)	SPFMV + SPCSV + SPCaLV+SwPLV+SPMSV (1)
		SwPLV + SPMSV (2)	SPFMV + SPCSV + SPMSV (1)
			SPFMV + SwPLV + SPMSV (3)
			SPFMV +SPMMV+ SPMSV (2)
			SPFMV + SPCFV + SPMSV (1)
			All tested viruses(3)
total	50	18	18
Damietta	SPFMV(5)	SPFMV + SwPLV (2)	SPFMV + SPCSV + SwPLV (6)

In Damietta Governorate, only three viruses (SPFMV, SwPLV and SPCSV) were detected (Table, 2). Thirteen out of fifteen plants were found to be infected (86.6%). Single, double and mixed infection represented 38.4, 15.3 and 46.2% respectively. In accordance with results obtained with Damietta plants, SPFMV was found in all samples (single, double and mixed infections).

Regarding asymptomatic plants, 9 out of 69 plants were virus(s)-infected, among these 9, 8 were singly infected with SPFMV and one was doubly infected with SPFMV +SPMSV infection. SPFMV was found in all infected plants, but without any external symptoms. Clark and Moyer (1988) and Gibson *et al.* (1997) reported that SPFMV caused mild or no symptoms in sweet potato when it found alone.

As shown in Table (3) SPFMV was detected in 54.8% of symptomatic sweet potato plants, representing the most prevalent virus. This virus was followed by SwPLV 17.1%, SPMSV(16.4), SPCSV(13%), SPCaLV (11.6%), SPCFV(8.2%) and SPMMV(4.8%). In symptomatic sweet potato plants, SPFMV was reported to occurrence in 67.5% by Mukasa *et al.* (2003) and in 74% by Ateka *et al.* (2004).

Sweet potato feathery mottle virus (SPFMV):

SPFMV-infected Mabroka sweet potato plants (as detected by ELISA) were used to isolate this virus. Infectious sap was extracted and used

to inoculate 2 indicator hosts (*I. nil* and *I. setosa*). Three weeks after inoculation, vein clearing and mild mosaic accompanied with malformation were observed on the new leaves. These symptoms are similar to those observed on the indicator hosts by Cadena-Hinojosa and Campbell (1981) and Moyer and Salazar(1989). However, identity of SPFMV was ensured by ELISA test. The inoculated indicator hosts were used as a source of the virus in the following studies.

Table (3): Incidence of different viruses in symptomatic sweet potato plants grown in Kalyoubia and Damietta as indexed by NCM-ELISA

Governorate	No. of samples	SPFMV		SPCSV		SPMMV		SPMSV		SwPLV		SPCaIV		SPCFV	
		+	%	+	%	+	%	+	%	+	%	+	%	+	%
Kalyoubia	131	67	51.1	13	10	7	5.3	24	18.3	17	13	17	13	12	9.1
Damietta	15	13	86.6	6	40	-	-	-	-	8	53.3	-	-	-	-
Total	146	80	54.8	19	13	7	4.8	24	16.4	25	17.1	17	11.6	12	8.2

+ = virus infected plants

Immunosorbent electron microscopy(ISEM)

EM-serology is highly sensitive and have become an important method for virus identification. (Derrick, 1973; Milne and Luisoni, 1975). This test was done using partially purified SPFMV. Negatively stained preparations (after trapping with specific SPFMV antibody with 2% uranyl acetate) were examined. Electron micrographs revealed the presence of flexuous particles with about 830 nm in length (Fig. 2). Shape and size of these particles are in accordance with those reported by some authors(McLean 1959; Campdell *et al.* 1974; Nome *et al.* 1974; Moyer and Kennedy 1978 and Cali and Moyer, 1981). They mentioned that, particle length of SPFMV ranged between 800 and 850 nm in the infected plants.

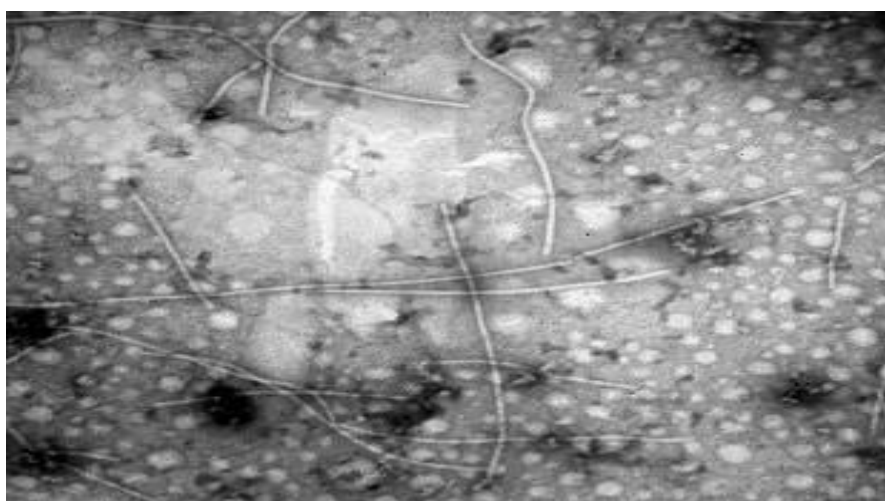


Fig. (2): Electron micrograph of partially purified virus preparation stained with uranyl acetate, x = 60000.

Reverse transcription- polymerase chain reaction (RT-PCR):

The total RNAs prepared and purified from infected- *I nil*, *I setosa* and Mabroka sweet potato plants as well as from healthy *I nil* plants were reverse transcribed using one step PCR mix as mentioned before. This reaction was done by using the specific primers for coat protein gene of SPFMV. The up stream primer SPFMV1 was corresponded to nucleotides 202 to 223 whereas the down stream primer SPFMV2 was complementary to bases 591 to 612 of the viral RNA as mentioned by Jeong *et al.* (2003). Both primers (SPFMV1 & SPFMV2) successfully amplified the cDNA Fig.(3). The size of PCR product amplified from infected plants was 411 bp. This was in agreement with the expected size calculated from the position of the primers and also as published for nucleotide sequence of coat protein gene. No signal was detected in the negative controls (non infected plants).

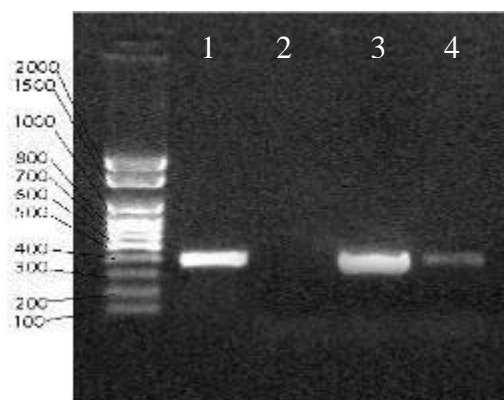


Fig (3): Agarose gel electrophoresis(1%) showing the results of RT-PCR products, using SPFMV1 and SPFMV2 primers to amplify the full length(411 bp),using RNA extracted from infected *Ipomoea nil* (lane 1), healthy *I nil* (lane 2), infected *Ipomoea setosa* (lane3), infected sweet potato Mabroka (lane 4) M: DNA Molecular Weight Marker (Bioron).

This result was in agreement with that of Jeong *et al.* (2003) who described RT-PCR protocol using two specific 22- mer primers located in coat protein gene of SPFMV {the up stream primer SPFMV1(5'-ATA GTG GGG GCA TCA TCA AAG G-3')and the down stream primer SPFMV2(5'-CCT AAA AGT AGG CAC TGC ATG G-3')}. A 411 bp PCR- product was detected in virus infected plants but not in healthy ones.

REFERENCES

- Anonymous (1998). FAO production Year Book for 1996. Food and Agriculture Organization of the United Nation, Rome, Italy. pp. 91-92. (c.f. Mukasa *et al.*, 2003).
- Anonymous (2001). Techniques in plant virology in CIP. Training Manual Version: Lima, Peru: International Potato Centre.

- Anonymous (2005). FAO. <http://faostat.fao.org/DesktopDefault.aspx?pageID=336&lang=en>
- Aritua, V., Bua, B., Barg, H., Vetten, J., Adipala, E. and Gibson, W. (2007). Incidence of five viruses infecting sweet potatoes in Uganda; the first evidence of sweet potato caulimovirus-like virus in Africa. *Plant pathology* 56: 324-331.
- Ateka, E.M., Njeru, R.W., Kibaru, A.G., Kimenu, J.W., Barg, E., Gibson, R.W. and Vetten, H.J. (2004). Identification and distribution of viruses infecting sweet potato in Kenya. *Annals of applied Biology* 144: 371-379.
- Cadena-Hinojosa, M.A. and Campbell, R.N. (1981). Serologic detection of feathery mottle strains in sweet potatoes and *Ipomoea incarnata*. *Plant Dis.* 65: 412-414.
- Cali, B.B. and Moyer, J.W. (1981). Purification, serology, and particle morphology of two russet crack strains of sweet potato feathery mottle virus. *Phytopathology* 71:302-305.
- Campbell, R.N., Hall, D.H. and Mielins, N.M. (1974). Etiology of sweet potato russet crack disease. *Phytopathology* 64: 210-218.
- Clark, C.A. and Moyer, W. (1988). *Compendium of Sweet Potato Diseases*. The American phytopathological society. Minnesota, USA. 74pp.
- Derrick, K.S. (1973). Quantitative assay for plant viruses using serologically specific electron microscopy. *Virology* 56: 652-653.
- Gibson, R. W., Mpembe, I., Alicai, T., Carey, E. E., Mwanga, R. O. M., Seal, S. E., and Vetten, H. J. (1998). Symptoms, etiology and serological analysis of sweet potato virus disease in Uganda. *Plant Pathology* 47:95-102.
- Gibson, R.W., Mwanga, R.O.M., Kasule, S., Mpembe, I. and Carey, E.E. (1997). Apparent absence of viruses in most symptomless field grown sweet potato in Uganda. *Annals of Applied Biology* 130: 481-490.
- Ishack, J., Bekheit, H., Ibrahim, L., El-Bedewy, R. and Abo El- Abbas, F. (2000). Occurrence of sweet potato virus disease in Egypt: Detection and epidemiological aspects. Fifth Triennial Congress of the African Potato Association. Pp.35(Abstr.)
- Jeong, J.H., Chakrabarty, D., Kim, Y.S., Eun, J.S., Choi, Y.E. and Paek, K.Y. (2003). A simple detection of sweet potato feathery mottle virus by reverse transcription polymerase chain reaction. *J. Plant Biotechnology* 5:283-286.
- Karyeija, R.F., Gibson, R.W. and Valkonen, J.P.T. (1998). The significance of sweet potato feathery mottle virus in subsistence sweet potato production in Africa. *Plant Dis.* 82: 4-15.
- Kern, A.J., Chaverre, M.E. and Dyer, W.E. (2005). Dicamba-responsive genes in herbicide-resistant and susceptible biotypes of Kochia (*Kochia scapania*). *Weeds Science* 53:139-145.
- McLean, D. D. (1959). Some aphid vector-plant virus relationships of the feathery mottle virus of sweet potato. *Journal of Economic Entomology* 52: 1057-1062
- Milne, R.G. and Luisoni, E. (1975). Rapid high-resolution immune electron microscopy of virus preparations. *Virology* 68: 270-274.

- Moyer, J.W. and Kennedy, G.G. (1978). Purification and properties of sweet potato feathery mottle virus. *Phytopathology* 68: 762-767.
- Moyer, J.W. and Salazar, L. F. (1989). Virus and virus-like diseases of sweet potato. *Plant Dis.* 75:451-455.
- Mukasa, S.B., Rubaihayo, P.R. and Valkonen, J.P.T. (2003). Incidence of viruses and virus-like diseases of sweet potato in Uganda. *Plant Dis.* 87:329-335.
- Ngeve, J. M. (1990). Yield stability and yield depression in sweet potato cultivars susceptible to the sweet potato virus disease. *J. Hort. Sci.* 65:225-230.
- Nome, S.F., Shalla, T.A. and Petersen, L.J. (1974). Comparison of virus particles and intracellular inclusions associated with vein mosaic feathery mottle and russet crack diseases. *Phytopathol. Z.* 79: 169- 178.
- Woolfe, J. A. (1992). *Sweet Potato, an Untapped Food Resource.* Cambridge University Press, New York. 643 PP.

تواجد و تفاعل بعض فيروسات البطاطا وتكرارها تحت ظروف الحقل

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نظرا لأنة لا توجد دراسة في مصر تشير إلى التواجد الطبيعي للفيروسات التي تصيب نباتات البطاطا في الحقل ومدى انتشار كل منها وكيفية تواجدها في النبات الواحد و مدى قابلية كل منها للارتباط بفيروسات معينة وعدم الارتباط بفيروسات أخرى فقد كان من الضروري عمل حصر لتلك الفيروسات في الحقل وتدارس التداخلات المختلفة وتكرار تواجدها كل منها مع التركيز على تدارس تلك الفيروسات انتشارا وارتباطها بالفيروسات الأخرى في الإصابات المشتركة. نتائج مثل هذه الدراسة يمكن استخدامها في التعامل مع تلك الفيروسات والتعرف على النباتات لمصابة وفى مكافحتها خاصة أن تدخلها مع بعضها يسبب مشاكل في التشخيص.

خلال أعوام ٢٠٠٦/٢٠٠٥ تم إجراء حصر حقل للنباتات المنزرعة في ٤٢ فدان موزعة على ثلاث محافظات منتجة للبطاطا حيث تم عمل فحص ظاهري لعدد ٨٧٠ نبات صنف مبروكة في القليوبية. ١٩٤٨ نبات صنف أبيض في دمياط و١٦٣٩٨ نبات هجين في المنوفية. تراوحت نسبة للأعراض الفيروسية (موزايك، اصفرار، تجعد للأوراق مصحوبا بتبرقش) على هذه النباتات بين ٤,٥% في المنوفية ١٠% في دمياط و ٢٢% في القليوبية.

أخذت ١٤٦ عينة عليها مظاهر إصابة , ٦٩ عينة خالية من أي مظهر من مظاهر الإصابة وقد أثبت الفحص السيرولوجي باستخدام الاليزا على الأغشية النيتروسيلولوزية (NCM-ELISA) وجود سبعة فيروسات في النباتات المصابة مع اختلاف في التوزيع والارتباط ببعضها البعض حيث كانت ٦٨% من النباتات مصابة بفيروس واحد و ١٦% مصابة بفيروسين بالإضافة إلى ١٦% مصابة بأكثر من فيروسين اتضح من الدراسة أن ١٣% من النباتات الخالية تماما من أية أعراض ظاهرية مصابة ب

SPFMV و إن الإصابات الفردية في تلك النباتات بلغت ٩٠% من اجمالي النباتات المصابة ونظرا لأن الدراسة أثبتت أن هذا الفيروس يتواجد بصورة متخفية في الإصابات الفردية ولا يعطى أعراض واضحة وأنة يتواجد مع الإصابات المختلفة وأن نسبة تواجده بلغت ٦٦% في الإصابات الفردية , ٨٩% في الإصابات المزدوجة , ١٠٠% في الإصابة بأكثر من فيروسين لذلك كان من الضروري عمل بعض الدراسات الإضافية علي حدة حيث تم عزلة والتعرف علي بدقة سيرولوجيا وبواسطة الميكروسكوب الاليكتروني حيث وجد أنه خيطي وبطول ٨٣٠ نانومتر كما تم عمل بعض الدراسات الجزيئية علي حدة حيث تم عزل حامضة النووي وباستخدام بادئين متخصصين في تفاعل البلمرة المتسلسل أمكن الكشف الدقيق عنه.