Review Article

The Importance of Potato virus Y Potyvirus

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Abstract

Potato virus Y potyvirus (PVY) is one of the most important aphid transmitted viral pathogen of potato worldwide. PVY affects potato plant (*Solanum tuberosum* L.), which is an important economic crop. PVY infection with potato viruses usually causing severe economic damage affecting both tubers quality and trade. PVY can degrease the production of certified seed and also crops grown for processing or fresh market. Potato is considered as one of the most economically important solanaceous crops cultivated in different regions that (PVY) has been studied in many parts of the world. Wide host plant range is a characteristic of viruses belonging to the Potyviridae. 495 species in 72 genera of 31 families were reported to be infected by PVY. Under field conditions, potato plants are subjected to attack by many viruses. The most cited virus could affect potato in the world is PVY. It has been long recognized as a threat to potato cultivation. Several studies carried out and reported that PVY is the most important virus infecting potato in different localities. Our review article is showed that PVY are the most frequently detected viruses in potato.

Introduction

Potato virus Y Potyvirus (PVY) was described for the first time in the early 1930s as the causal agent of a serious disease in potato [1]. PVY has been studied in many parts of the world [2]. PVY is still major virus of potatoes because it spreads easily and depresses yield so much. It is one of the most important aphid transmitted (non persistent manner) viral pathogen of potato worldwide [3]. Depressing varies with cultivar and strain [4,5]. PVY is one of the most important aphid transmitted viral pathogen of potato worldwide. It can affect the production of certified seed and also crops grown for processing or fresh market (Rykbost, et al. 1999). The virions of the viruses belonging to the Potyvirus genus are rod-shaped flexuous filaments 680-900 nm long and 11-13 nm wide [6]. Molecular genome characterization has identified where recombination events on PVY genomes have occurred to create new strains such as PVY^{N-Wilga} and PVY^{NTN} [7].

Classification and electron microscopy

PVY was described for the first time in the early 1930s as the causal agent of a serious disease in potato [1]. PVY is the type member species of the genus Potyvirus, family Potyviridae. PVY has a long filamentous particle containing a single stranded, positive sense RNA genome of approximately 9.7kb [8]. Transmission electron microscope examination of partially purified preparation from infected *Nicotiana*

More Information

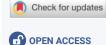
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glutinosa leaves negatively stained with phosphotungistic acid revealed the rod flexible (helical symmetry) particles of PVY with 730 nm (Figure 2) [6]. PVY infects solanaceous plants such as tobacco, tomato, pepper and potato and has a significant impact on potato and other solanaceous plants [9]. PVY has been studied in many parts of the world. It is still major virus of potatoes because it spreads easily and depresses yield so much (up to 80%). Depressing varies with cultivar and strain [4,5]. The potato (S. tuberosum L.) is an important economic crop, however, its infection with potato viruses usually causing severe economic damage affecting both seed quality and trade (Figure 1a) [10]. PVY is one of the most important aphid transmitted viral pathogen of potato worldwide. It can affect the production of certified seed and also crops grown for processing or fresh market (Rykbost, et al. 1999). In Egypt, PVY is widely spread and isolated from potato plants [4,11,12]. PVY was isolated from different crops such as potato, tomato, gladiolus and pepper [13,14]. Also, isolated three isolates of PVY from naturally infected potato plants cvs. Nicola and Diamont at different farms [15].

Approximately 2000 subunits of a single coat protein are organized around the viral genomic RNA in a helical arrangement [16]. Sharma, et al. [17], reported that small drops of purified virus were placed on carbon-coated grids for one minute and dried, then the virus particles existed on grids were stained using 2% uranyl acetate and dried (Figure 2) [6]. The modal length of particles were determined from crude preparation deposited on carbon- formvar coated 38µm (400 mesh) grids negatively stained with 2% uranyl acetate for one minute and dried, then examined using transmission electron microscope [18].

Symptomatology and host range of Potato virus Y Potyvirus

PVY infects a number of plant species in the family Solanaceae and causes a wide range of symptoms from symptomless to mosaic, mottling, lesions, stunting, necrosis and plant death, depending on the plant species, the cultivar, the virus strain and isolate [16]. Mustafa, et al. [19], showed that pepper isolate of PVY caused vein clearing and mosaic symptoms on Datura metel (Figure 1b) and Capsicum annum, mosaic on *N. glutinosa* (Figure 1c) and *N. repanda* (Figure 1d) as [6] N. tabacum cv. White Barley, N. tabacum cv. Samsun and N. rustica but didn't show any symptoms on Physalis floridana, Chenopodium amaranticolor, C. quinoa and Solanum tuberosum. Datura stramonium is not infectable by PVY [20]. Reaction of potato and other different susceptible plants to PVY was found to be varied from local lesions to systemic symptoms [21], depending on inoculated host, virus strain and environmental conditions [22]. Typical foliar symptoms can include veinal necrosis, leaf drop and a mosaic pattern sometimes accompanied by leaf roughness [23]. Smith [24],

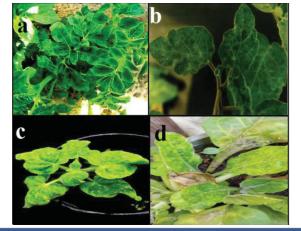


Figure 1: a) Mosaic symptom on potato plants naturally infected with PVY; b) Malformation, blisters and mosaic on *D. metal* caused by PVY; c) PVY induced mosaic symptoms on *N. glutinosa*; d) Mottling symptoms and yellow leaves appears on *N. repanda* after mechanical inoculation.

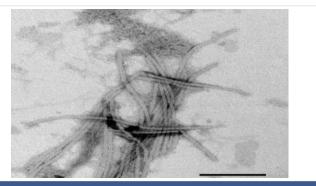


Figure 2: Electron micrograph showing rod flexible particles (helical symmetry) of PVY isolate negatively stained with 2% phosphotungistic acid with magnification power 40000x.

described the differences in disease reaction by number of different potato cultivars. Different patterns of symptoms of PVY infection were recorded by several authors [25,26]. Rich [21], concluded that chronic symptoms on potato plants consist of chlorotic mottling, severe rugose wrinkling, moderate to severe dwarfing and premature death. Tubers are smaller than normal. Current-season symptoms are characterized by the development of brown necrotic streaks along the veins of the leaves, the petioles, and the stems. In severe cases, the leaves and petioles die and hang downward from the stems; hence the name "leaf drop streak" was adopted. Severity of symptoms expression varies considerably from one cultivar to another. If infection takes place later in the season, tubers may become infected without development of plant symptoms [27,28]. Eskarous, et al. [29], reported that PVY caused mottling, vein clearing and leaf abnormalities on S. nigrum var. judaicum. The virus induced systemic symptoms with vein clearing and vein banding, in N. tabacum [21]. In this respect, Park, et al. [30] observed that the virus caused vein banding and necrosis on Burley tobacco plants. Vein clearing, vein necrosis and bronzing were observed on three tobacco cultivars, namely virgin A Mutant; Virginia Aurae and Virginia. These cultivars were previously reported as being immune to PVY^N [31]. Some workers found that C. annum was not susceptible to PVY [4], but other workers stated that extremely severe leaf necrosis followed by leaf drop, occurred in very young chill seedlings, while older plants were only mildly affected by vein banding and mosaic disease [32,33]. El-Hammady, et al. [34], studied the susceptibility of 14 pepper cultivars (C. annum) to PVY (Previously isolated from naturally infected pepper plants) and found that all tested cultivars were susceptible but with different degrees. Ch. amaranticolor was recorded as a local lesion host for this virus [21,34,35]. Ch. quinoa was also used as a local lesion host for PVY [26,36]. In addition, El-Hammady et al. [34] reported that the virus produced local lesions on Ch. album. The potato hybrid A6 (S. demissum x S. tuberosum) is the most important indicator host for PVY detection, that the virus produced local lesions accompanied with systemic symptoms on this indicator host [36]. Wide host plant range is a characteristic of viruses belonging to the Potyviridae. 495 species in 72 genera of 31 families were reported to be infected by PVY in the laboratory [37]. PVY has a wide range of natural host species, some of which are edible crops like potato, tobacco, pepper, and tomato, while others are ornamental plants (Dahlia and Petunia spp.) or weeds such as Datura spp., Physalis spp., S. dulcamara and S. nigrum [38]. It has been reported that the weed S. elaeagnifolium, which is abundant in potato fields in Tunisia and other Mediterranean countries, is infected up to 32% by PVY^N which can provide a source for virus infection of potatoes [39]. There are recent reports of such new hosts. For example, Fletcher (2001) reported that Cotula australis and Capsella bursa-pastoris can be infected naturally by PVY. In Europe, Kaliciak and Syller [40], reported other new wild hosts for PVY including Erodium cicutarium, Geranium

pusillum, Lactuca serriola and Lamium purpureum. Chikh-Ali, et al. [41], reported that PVY is very common in S. nigrum and *Physalis spp.* which are abundant at the field margins in potato crops in Syria. They speculated that these weeds may serve as a virus source for aphids alighting on potato crops. This growing list of alternative host species is very challenging when designing virus control strategies, particularly because of the important role of these hosts in providing virus sources for aphid vectors, several investigators recorded that PVY can infect many hosts belonging to different families. These families are about eight of the angiosperm [21]. S. tuberosum is the main host of PVY [1,25]. Singh, et al. [42], found that PVY causes diseases and production loss in tobacco crops, pepper (C. annuum L.), tomato (S. Lycopersicum L.) [43] and Ch. Amaranticolor [24,34], D. metel, N. debneyii [22,34,36], N. glutinosa [26,34], N. tabacum [21,26,34,35], S. dulcamora [21], S. melongena [44], S. nigrum [21,25,34] and Petunia spp [21,34,36]. D. stramonium was recorded as immune host for PVY infection [4,25,34,35,45,46]. In addition, Gomphrena globosa, Ch. amaranticolor, Ch. quinoa, Ch. Murale and Ch. album were also found to be immune for PVY [4,22,34,45,47].

Transmission of Potato virus Y Potyvirus (PVY)

PVY is transmitted easily by mechanical inoculations [4,6,29,45,48-51]. The virus was found to be stylet-borne in many species of aphids [21]. The green peach aphid (*Myzus persicae* Sulz.) is the most common and efficient vector [36,48,52,53]. Other aphid vectors include *Aphis compositae, A. craccivora* [29], *Rhopalosiphum padi* [52,53] and *A. nerii* [6]. According to [53], *M. persicae* and *Acyrthosiphon pisum* were the most effective vectors, infecting 26% and 25% of the test plants respectively. *A. fabae, A. nasturtii* and *Rhopalosiphum padi* infected only 1% - 7% of the test plants. *Sitobion avenae* and *Brevicoryne brassicae* did not transmit PVY°. The virus is transmissible through infected tubers [11,21,54].

Purification of Potato virus Y Potyvirus

The methods used in purification of PVY by several investigators were different from each other mainly in the type of extraction buffer and the type of organic solvent used to clarify the leaf extract [55]. Sankari et al. [56], reported that Virus purification was carried out according to [57] with some modifications by [6]. In brief, infected leaves were triturated with four volumes (v/w of leaf tissue) of 0.1M Citrate buffer (pH 7.2) containing 0.02 M EDTA, 0.04 M Sodium sulfite, 0.2% mercaptoethanol, and 0.1% Triton X-100. Then this was mixed with cold chloroform : charbontetrachloride (1:1) to 50% of total tissue and buffer volume. Mixture was centrifuged at low speed (10000 g) and aqueous phase was stirred with .4% PEG and 0.4 M Na Cl for one hour followed by low speed centrifuge (10000 g). Pellet was dissolved in 0.02 M Tris-HCl (pH 7.2) containing 0.1% mercaptoethanol, 0.01 M EDTA and 0.1% Triton X100 and mixed with cold chloroform and finally centrifuged at low speed (10000 g). Aqueous phase was centrifuged at high speed (110000 g) and the resultant pellet was dissolved in 0.02 M Tris-HCl, pH 7.2 containing 0.01 M EDTA. Dissolved pellet was layered on (10-40%) sucrose gradients and recovered virus zone was diluted in 7.2 M Tris-H Cl, pH 7.2 containing 0.01 M EDTA and centrifuged at high speed (112000 g) for 90 min. at 4^oC Purified virus was dissolved in Na Cl (0.85%) solution and stored at -80 °C. Nerway and Kassim [58], Modified procedures of [59,60] to purify PVY using differential centrifugation, precipitation and filtration technique.

Spectrometrical tests were used to assess the purity of PVY suspension and the efficiency of procedure used to purify the virus. This test was done using UV-spectrophotometer for scanning the purified solution at a wave length ranged between 200-300 nm, at 10 nm intervals [58] to plot the absorbency curve, estimate A260/A280 ratio (nucleic acids : nuclei protein) and PVY concentration in potato tissues. The virus yield was calculated according to [61] as follows: Virus yield = 0.D. (260) X Dilution factor/ E.C. Whereas: 0.D. = optical density at 260 nm; E.C. = extinction coeffcient 2.4 (mg/ml⁻¹ cm⁻¹).

Polyclonal antiserum production from Potato virus Y Potyvirus

Nerway and Kassim [58], mentioned that antiserum was produced according to that of [61], as follows. 2 ml of the purified viral preparation was injected intravenously at interval of one week into the ear vein of albino New Zealand male rabbit. After two weeks, the injection was repeated at the same quantity (booster injection) intramuscularly into the muscle of rabbit, after mixing the purified virus solution with sterilized alum solution (1:1). Kassim and Ali, [62], reported that the injected rabbit with PVY was bleed 7 days after final injection, 15 ml of the blood was taken from the injected rabbit. The antiserum was obtained and purified as reported by [63]. Blood was allowed to clot at room temperature, and the clot was removed. Clotted blood was centrifuged at 5000 rpm for 10 minutes. The clear antiserum (supernatant) was collected and discarded the pellet. The antiserum (serum plus antibodies) was poured into a small vial and kept frozen at -18°C [6,58].

Serological detections

Serological techniques are useful and accurate test for identifying PVY [64]. ELISA was the most common serological test used to diagnose viral diseases in potato [5,50,65]. Lin, et al. [66], described a technique of tissue-blot immunoassay (TBIA) on nitrocellulose membranes for detection of plant viruses and micoplasma-like organisms (MLO) in infected plants. Tissue blots were made by pressing, with a firm and gentle force, the freshly cut tissue surface on nitrocellulose membranes. The reliability of ELISA when used on leaves is efficient and reliable as using nucleic acid based technique [67]. In certification programs, it is the most practical diagnostic method which is performed using leaf sap for routine testing of large number of potato plants [68]. Testing of potato tubers using ELISA has not been shown to be accurate and reliable [47,69-71]. Antigens in tissue blots on nitrocellulose membranes were detected by enzyme-labeled immunological probs [66]. Indirect immunological methods, the blots were reacted with antigen-specific primary antibodies and detected with enzyme-labeled species-specific secondary antibodies [2]. Alternatively, the blots were reacted with antigenspecific biotinylated primary antibodies and detected with avidin-enzyme conjugates. Indirect immunological methods, the blots were reacted and detected with enzyme-labeled antigen-specific antibodies [66]. Samson, et al. [72] pointed out that serological detection of potato viruses was more sensitive, faster and less expensive with direct tissue blotting (DTBA) than with ELISA. Positive reaction were obtained with Tobacco rattle Tobravirus (TRV), Potato virus S Carlavirus (PVS), PVY, Potato virus M Carlavirus (PVM), Potato virus A Potyvirus (PVA), Potato virus X Potexvirus (PVX) and Potato leaf roll Luteovirus (PLRV). Viruses were detected in leaves, petioles, stems, roots and tubers [73].

Molecular detection of *Potato Virus Y Potyvirus* by reverse transcription polymerase chain reaction (RT–PCR)

In 1989, scientists selected polymerase chain reaction as the major scientific development of the year [74]. Nowadays RT-PCR is considered to be the most powerful tool in molecular biology. In the past, serology was routinely used for the detection of plant viruses from infected tissues. However, this method is less sensitive than some other techniques such as nucleic acid hybridization or PCR, which are becoming more popular [75]. PVY is a rapidly evolving virus [10], where both variation within strains and recombination events between strains occurs. Analysis of the genome using molecular biology methods has brought a new understanding of PVY at a nucleotidic level. Rapid developments in molecular biology and sequencing of isolates since the late 1980s have made it possible to characterise and classify PVY strains based on molecular characteristics [76-78] which may or may not correspond to traditional PVY strain groups. Overall, at the nucleotidic level, multiple isolates of the PVY^0 and PVY^N groups differ from each other by approximately 8% along their genomes [79]. Molecular genome characterization has identified where recombination events on PVY genomes have occurred to create new strains such as PVY^{N-Wilga} and PVY^{NTN} [7]. Molecular characterisation has also revealed that in many countries the majority of isolates are now recombinants between PVY⁰ and PVY^N [80,81]. Takacs [82], studied the main characteristics and diagnostic methods of PVY and its strains, PVY⁰ (normal strain), PVY^C (strain C), PVY^{An} (anomalous strain), PVY^N (tobacco vein necrosis strain), and PVY^{NTN} (Potato ring spot strain) were identified using PCR techniques. Several authors used RT-PCR for the detection of PVY and other viruses. The application of PCR for potato viruses detection has already been reported [83]. This method followed by an additional restriction enzyme digestion of the amplification product has potential for rapid identification of PVY isolates [84]. Rapid molecular methods have recently been developed for virus strain differentiation, based on restriction fragment length polymorphism (RFLP) [84] and single-stranded conformation polymorphism (SSCP) [85] of PCR products. The RT-PCR is used to characterize isolates of potato viruses by converting RNA into cDNA followed by amplification and sequencing [86] and Chikh Ali, et al. 2014). RT-PCR method has been also successfully utilized for the detection of PVY in potato tubers [87]. The RT-PCR method has been successfully utilized to detect viruses from infected potato leaves [2,88]. Sequencing techniques have enabled molecular taxonomy in closely related species. Using the amino acid sequences of the CP of virus species, the genus Potyvirus exhibit 38% - 71% similarity, while strains share 90-99% similarity [8]. The polymerase chain reaction PCR technology has become the most powerful tool for plant pathologists. Since the adaptation of ELISA to plant virology by [63]. The first published account of PCR in 1985 was its application to the prenatal diagnosis of Sickle cell anemia [89]. PCR is an in vitro method in which by DNA sequences or transcripts are amplified rapidly with very high specificity and fidelity using oligonucleotide primers and Taq DNA polymerase in simple automated reaction. The PCR was developed to use it in the areas of disease diagnosis, detection of pathogens, detection of DNA in small samples, DNA comparison, high efficiency cloning of genomic sequences and gene sequencing. PCR is a powerful technique used for the amplification of a specific DNA sequences and is capable of enrichment by factor of 10⁶ to 10⁷, enabling the detection of a few target nucleotides [89]. Several investigations in their research work used specific method for extraction of total nucleic acid (TNA) from plant tissue; this procedure was used successfully for the extraction of numerous samples for RT - PCR detection and diagnosis of several viroids and viruses. RT-PCR amplification has been successfully utilized for the detection of RNA plant viruses from infected tissue. Several researches used this technique to detect CMV in infected samples. Proper virus identification is always the key in developing appropriate practical solutions to manage plant virus diseases [10,90]. Recent advances in biotechnology and molecular biology have played a significant role in the development of rapid, specific and sensitive diagnostic tests. Reverse transcriptase PCR (RT-PCR) can be used for viral disease diagnosis because it is a quick and more reliable method in comparison to ELISA, and PCR can be used for further characterization of cowpea plant viruses. Combining PCR with molecular hybridization can detect even pictogram qualities of virus, and this combination is 4 - 5 orders magnitudes superior to direct molecular hybridization (Vunsh et al., 1999). Diagnostic approaches based on nucleic acid hybridization are not only highly specific but also applicable for routine testing of large number of samples. The development of nucleic acid-based tools was another new dimension of virus detection. The most common among these techniques are cDNA hybridization and

polymerase chain reaction (PCR). In addition, PCR can be used as a confirmatory test for TBIA, where processed blots can be cut individually and tested by PCR. This proved to work well with both DNA and RNA plant viruses. Furthermore, unprocessed plant tissue blots on nitrocellulose membrane represent a good sample for PCR amplification. PCR products can also be used for cloning and subsequent sequencing which is extremely useful for identification of new viruses or virus strains. Gibbs and Mackenzie [91], mentioned that the sequence analysis was used to design a pair of degenerate oligonucleotide primers that amplified a 1.6-2.1 kbp fragment from the 3' end of the genome (virion protein gene and part of the Nib gene) of 17 species of the Potyviridae ('potyvirids'); 11 potyviruses, 2 bymoviruses, 2 macluraviruses, an ipomovirus and a rymovirus. The 'potyvirid primer 1' hybridizes to the 3' terminal poly-A region of the genome, and 'potyvirid primer 2' to the genomic region encoding the-GNNSGQ-motil' of the Nib protein. Database searches showed that the potyvirid 2 primer is specific for potyvirids. Associated analyses indicated that the published amino acid sequence of part of the wheat streak mosaic rymovirus Nib protein is probably incorrect in part. Kumari [92], mentioned the A one-step reverse transcription-polymerase chain reaction (RT-PCR) protocol was developed and used for the detection of *Cherry leaf roll* virus (CLRV) and Strawberry latent ring spot virus (SLRSV). The protocol was used to test infected screen house plants. The one-step RT-PCR protocol is rapid and sensitive and has the potential to be used for the diagnosis of CLRV and SLRSV in routine diagnostic laboratories [2,92].

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