Characterization of the Zucchini Yellow Mosaic Virus from Squash in Tehran Province

S. Hosseini1*, G. H. Mosahebi1, M. Koohi Habibi1 and S. M. Okhovvat1

ABSTRACT

Zucchini yellow mosaic virus (ZYMV) is a potyvirus with a worldwide distribution. This virus causes serious economic losses in Iran in many cucurbits. During 2002-2003, samples were collected from squash fields in Tehran Province. Five isolates (Z1, Z2, Z3, Z4 and Z5) were inoculated on 27 species of Cucurbitaceae, Chenopodiaceae, Amaranthaceae, Solanaceae, Leguminosae and Ranunculaceae. Chenopodium quinoa and C. amaranthicolor showed chlorotic local lesions. Gomphrena globosa developed necrotic local lesions. Systemic symptoms were produced in the members of Cucurbitaceae and Ranunculus sardous. Z2, Z4 and Z5 caused mosaic symptoms on Phaseolus vulgaris cv. Red Kidney and P. vulgaris cv. Khorram but Z1 and Z3 caused chlorotic local lesions. Virus was purified from Cucurbita pepo. Virus particles in immunoelectron microscopy were filamentous flexuous. The molecular weights of coat protein using SDS-PAGE and western blotting were estimated at 32 kDa. Reverse transcription polymerase chain reaction (RT-PCR) was performed using one primer pairs designed by Desbiez et al. An approximately 458 bp fragment was amplified with a specific primer.

Keywords: ELISA, RT-PCR, SDS-PAGE, ZYMV.

INTRODUCTION

Zucchini yellow mosaic virus (ZYMV), a potyvirus with a worldwide distribution, was first discovered in Italy (Lisa et al., 1981). In Iran, ZYMV is known to occur in squash fields in Tehran Province (Ghorbani, 1988).

This virus can cause severe economic losses in many cucurbits. ZYMV belongs to the genus potyviruses in the Potyviridae, a group of plant viruses characterized by a monoparticle, positive-sense, single stranded RNA genome encapsidated in flexuous rod shape particles. Viral RNA consists of about 9600 nucleotides (Balint et al., 1990) with a 5′ viral protein genome linked (VPG) and a poly(A) tail (Dougherty et al., 1993). Like other potyviruses, ZYMV is efficiently transmitted by aphids in a nonpersistent manner (Lisa et al., 1981). Control of ZYMV is difficult but the use of resistant cultivars, inoculation of mild ZYMV-WK strains for cross protection against severe challenging strains (Walkey et al., 1992; Lecoq et al., 1991) and the use of mineral oil sprays (Makkouk et al., 1986) in association with pyrethroids (Raccah et al., 1985), might provide protection under certain ecological conditions.

MATERIALS AND METHODS

Sample Collection

Samples were collected during the 2002 and 2003 growing seasons from field-grown squash (Cucurbita pepo) in Tehran Province. In this region, squashes are planted during early April and harvested from June

1. Department of Plant Protection, College of Agriculture, University of Tehran, Karaj, Islamic Republic of Iran.

* Corresponding author, email: samin_hf@yahoo.com
to July. Virus infections become visible after the setting of the first fruits. Infected plants show symptoms such as: mosaic, yellowing, leaf distortion, shoestring, fruit deformation and yield reduction. Young leaves from some symptomatic plants were collected at random. All samples were kept in ice chests for transportation to the laboratory. Each plant sample was kept separately in a plastic bag at 4°C until analyzed.

Virus Identification

DAS-ELISA (double antibody sandwich ELISA) as described by Clark and Adams (1977) was used with a polyclonal antiserum against ZYMV (DSMZ-AS0239). A 200 µl aliquot of IgG was added to coat each well of plates. Each step of ELISA was followed by a 4-hr incubation at 37°C or a 12-hr incubation at 4°C. This was followed by three washes with a washing buffer. Ten milliliters of sample buffer, pH 7.4, was added to 1g tissue samples that had been ground in liquid nitrogen, and 200 µl of this extracted was added to each well. The reaction was read using a colorimeter at 405 nm after adding conjugate incubation with substrate for about one hour.

Host Range Studies

Five original isolates (Z1, Z2, Z3, Z4 and Z5) recovered from infected squash plants in the host range studied (from different locations) were maintained in Cucurbita pepo by sap inoculation. For plant assays, 27 species from 6 families were inoculated with the virus isolates. Sap prepared from leaves which were in 0.01 M Sodium phosphate buffer, pH 7, was rubbed onto leaves dusted with carborundum powder. The Leaves were then rinsed with water, and plants were maintained in an insect–proof screen house for observation. Symptoms on both inoculated and upper, uninoculated leaves were recorded. Tests for latent infection were conducted by back–inoculation to either Chenopodium amaranticolor Cost and Reyn or C. quinoa Wild.

Virus Purification

The virus was purified according a methods described by Purcifull and Hiebert (1979) and was purified from systematically infected tissue of Cucurbita pepo. All stages of the purification were completed at 4°C. Tissue was homogenized (1:2w/v 400g/800ml) in a 0.5 M potassium phosphate buffer, pH 7.2, containing 2 g Na2SO3 and a 1:1(v/v) mixture of chloroform and carbon tetrachloride.

The filtrate of the homogenate after passing it through eight layers of cheese cloth, was given low–speed centrifugation at 4000 g for 5 minutes. The aqueous phase was collected and centrifuged at 12000 g for 20 minutes to the resultant supernatant. Polyethylene glycol (PEG) (mol wt 6000) to concentration was added to 8%, and the preparation was stirred for one hour. The mixture was centrifuged at 12000 g for 10 minutes. The precipitate was resuspended in small volume of buffer. Virus purity was determined by UV spectroscopy. A260/A280 ratio was determined.

Protein Analysis

The molecular weight of the viral coat protein was estimated by SDS PAGE (Sodium dodecyl sulfate poly acryl amid gel electrophoresis) as described by Lamemmli (1970).

A purified virion suspension and plant samples infected with the virus were mixed with the sample buffer, boiled at 100 °C for 5 minutes and subjected to electrophoresis on 12% poly acryl amid gel along with poly peptide size standard. The gel was stained with coomassie blue. The molecular weight of polypeptides from ZYMV was determined by comparison with the size standards.

For western blot analysis, the purified virus and infected plant samples were electro-
phoresed as described previously. The separated proteins were transferred to nitrocellulose using the buffer styles of Sambrook et al. (1989). Probing of the western blots was done as described by Burgermeister and Koeing (1984) using the ZYMV antiserum to ZYMV (DSMZ DAS-0234) at a dilution of 1:1000 (v/v).

**Electron Microscopy**

Immunoelectron microscopical decoration tests was carried out at DSMZ (German collection of microorganism and cell cultures plant collection) in Germany.

**Reverse Transcription Polymerase Chain Reaction (RT-PCR)**

Total RNA from ZYM-infected plants was extracted using a phenol/chloroform protocol (Wadsworth et al., 1988). Three μl of RNA were submitted to reverse transcription in a final volume of 20 μl, using 2μl PCR buffer 10x (0.5 M Tris-HCl, 0.7 M KCl, 0.1 M MgCl₂, pH 8), 1μl DTT (100 mmol/μl), 1μl dNTPs (10 mmol/μl), 0.5 μl RNase inhibitors enzymes (10 mmol/μl) and 2 μl Reverse-DAG primer (100 pmol/μl) (5‘-GCG TGG CAA TGA CAT-3‘: nucleotide position 8735-8749 on sequence L 31350) for one hour at 42°C with 0.5 μl MMLV reverse transcriptase (200 mmol/μl). 5 μl of the RT reactions were used for PCR using a 5 μl PCR buffer 10x, 2 μl MgCl₂, 1 μl dNTPs (10mmol/μl), 0.5 μl Taq polymerase (5 unit/μl), 1 μl Reverse-DAG (100 pmol) and 1 μl Forward-DAG (100 pmol) (5‘-ATT TGC GCT GCG ATG-3‘: 8291-8305 on sequence L 31350) oligonucleotides encompassing the N-terminal part of the coat protein coding region and the C-terminal part of the polymerase (Nlb) (primers designed by Desbiez et al., 2002). PCR reactions were performed by a first denaturation of the samples at 94°C for 3 minutes followed by 35 cycles at 94°C for 30 seconds, 43 °C for 30 seconds and 72°C for 30 seconds and a final elongation step at 72°C for 7 minutes. PCR products were controlled by electrophoresis on 1% agarose gel (Desbiez et al., 2002).

**RESULTS**

**Host Range**

The virus was isolated from infected squash species from the families Cucurbitaceae, Amaranthaceae, Chenopodiaceae and Leguminosae but it did not infect the numbers of Solanaceae that were tested. Mosaic, blistering, leaf distortion and stunted growth were observed on *Cucurbita pepo* cv. Zucchini (figures 3 and 4), *C. Pepo* cv. Maragheh, cv. Khoy, *Cucumis melo* cv. Asgrown. Necrotic local lesions were found on *Gompherena globosa* and chlorotic local lesions with symptoms of systemic infection on *Chenopodium quinoa*, *C. amaranticolor* and *Phaseolus vulgaris* cv. Khorram and cv. Red kidney were all observed (Figures 1 and 2). Latent infection was detected in *Ranunculus sardous* (Table 1).

**Figure 1**. Symptoms of mosaic on *Phaseolus vulgaris* cv. Red Kidney.

**Figure 2**. Symptoms of chlorotic local lesions on *Phaseolus vulgaris* cv. Red Kidney.
Virus Purification and Characterization

Greenhouse plants with the highest absorbance values determined in ELISA were chosen to purify the putative ZYMV. Approximately 16.2 mg of purified ZYMV was obtained from every 1 g of 8 day-old infected pumpkin leaves. Purification of the virus using the protocol described by Purcell and Hiebert (1979) was successful. Its $A_{260}/A_{280}$ ratios were 1.28. Electron microscopy analysis showed flexuous filamentous particles typical of the Potyviridae morphology (Figure 5).

### Table 1. Reaction of selected indicator plant species to ZYMV.

<table>
<thead>
<tr>
<th>Families</th>
<th>Test plants</th>
<th>Symptoms in leaves</th>
</tr>
</thead>
<tbody>
<tr>
<td>Cucurbitaceae</td>
<td><em>Cucurbita pepo</em> cv. Zucchini</td>
<td>m, b, Id</td>
</tr>
<tr>
<td></td>
<td><em>C. pepo</em> cv. Khoy</td>
<td>m, b, Id, ss</td>
</tr>
<tr>
<td></td>
<td><em>C. pepo</em> cv. Maragheh</td>
<td>m, b, Id, ss</td>
</tr>
<tr>
<td></td>
<td><em>Cucumis melo</em> cv. Asgrown</td>
<td>b, m</td>
</tr>
<tr>
<td></td>
<td><em>C. melo</em> cv. Melon Seed</td>
<td>b</td>
</tr>
<tr>
<td></td>
<td><em>C. melo</em> Local cultivar</td>
<td>b</td>
</tr>
<tr>
<td></td>
<td><em>C. sativus</em> cv. Dominus</td>
<td>m</td>
</tr>
<tr>
<td></td>
<td><em>C. sativus</em> cv. PS</td>
<td>m</td>
</tr>
<tr>
<td></td>
<td><em>Citrullus lanatus</em> cv. Crimson Sweet</td>
<td>s, m, b</td>
</tr>
<tr>
<td></td>
<td><em>Luffa acantangula</em></td>
<td>m</td>
</tr>
<tr>
<td></td>
<td><em>Gomphrena globosa</em></td>
<td>nll</td>
</tr>
<tr>
<td>Amaranthaceae</td>
<td><em>Amaranthus paniculatus</em></td>
<td>-</td>
</tr>
<tr>
<td>Chenopodiaceae</td>
<td><em>Chenopodium quinoa</em></td>
<td>cll</td>
</tr>
<tr>
<td></td>
<td><em>C. amaranticolor</em></td>
<td>cll</td>
</tr>
<tr>
<td></td>
<td><em>Spinacia oleracea</em> cv. Keshtzar</td>
<td>-</td>
</tr>
<tr>
<td></td>
<td><em>Phasesolus vulgaris</em> cv. Khorram</td>
<td>cll , m</td>
</tr>
<tr>
<td></td>
<td><em>P. vulgaris</em> cv. Daneshkadeh</td>
<td>-</td>
</tr>
<tr>
<td>Leguminoseae</td>
<td><em>P. vulgaris</em> cv. Red Kidney</td>
<td>cll , m</td>
</tr>
<tr>
<td></td>
<td><em>P. vulgaris</em> cv. Bountiful</td>
<td>-</td>
</tr>
<tr>
<td></td>
<td><em>Vigna unguiculata</em></td>
<td>-</td>
</tr>
<tr>
<td></td>
<td><em>Vicia faha</em></td>
<td>-</td>
</tr>
<tr>
<td></td>
<td><em>Pisum sativum</em></td>
<td>-</td>
</tr>
<tr>
<td>Solanaceae</td>
<td><em>Dautura metel</em></td>
<td>-</td>
</tr>
<tr>
<td></td>
<td><em>D. stramonium</em></td>
<td>-</td>
</tr>
<tr>
<td></td>
<td><em>Nicotiana tabacum</em></td>
<td>-</td>
</tr>
<tr>
<td></td>
<td><em>N. glutinosa</em></td>
<td>-</td>
</tr>
<tr>
<td></td>
<td><em>N. benthamiana</em></td>
<td>-</td>
</tr>
<tr>
<td>Ranunculaceae</td>
<td><em>Ranunculus sardous</em></td>
<td>-</td>
</tr>
</tbody>
</table>

Ss= Shoe string, id= Leaf distortion, b = Blistering, s= Stunted growth, m= Mosaic
1= Latent infection, cll= Chlorotic local lesion, nll= Necrotic local lesion.
Molecular weight of the capsid protein was determined as 32 kDa. Western blot analysis also revealed one band of approximately 32 kDa in the purified virus preparation and infected plant samples, while no band was found in the healthy plant extracts (Figures 6 and 7).

**RT-PCR**

RT-PCR was carried out using the primers DAG/F and DAG/R (described previously) which resulted in a fragment of 458 nts (Figure 8).

**DISCUSSION**

Squash (*Cucurbita pepo*) belongs to the Cucurbitaceae family and is one of the most important economic crops in the world. Cucurbitaceae are threatened by viruses and there are many viruses that damage this family. ZYMV is a *potyvirus* with a worldwide distribution that causes serious economic losses in many cucurbits.

ZYMV was first found in squash in Tehran (Ghorbani, 1988) and then was discovered in 2000 in Mazandaran (Delkhosh *et al.*).

The results of the host range studies, ELISA, electron microscopy and RT-PCR indicate that ZYMV isolates from Iran share the reported properties of the ZYMV. In direct caparison with ZYMV, very similar reactions were recorded amongst the isolates of ZYMV from Iran. The isolates induced systemic shoestring symptoms without local...
lesions.
However, some differences in the host range of Iranian isolates were observed in which Z1 and Z3 isolates induced local chlorotic lesions on Phaseolus vulgaris cv. Khorram and cv. Red Kidney. This is rare and only ZYMV-Conecticut, - France, - Taiwan induced these kinds of symptoms. Z2, Z4 and Z5 isolates induced mosaic in Phaseolus vulgaris cv. Khorram and cv. Red Kidney and these symptoms are different from other known strains.

These isolates could systemically infect Cucurbitaceous plants without induced local lesions. ZYMV isolates induced shoestring symptoms in Cucurbata pepo. This symptom has also been described by Wong (1994) in Singapore.

Nicotiana benthamiana, N. tabacum, N. glutinosa and Pism sativam were immune to ZYMV isolates. Ranunculus sardous induced latent infection and was used to as differential host (Wong, 1994).

Our SDS-PAGE pattern showed the presence of a protein of molecular mass 32 kDa, the expected size for the CP from ZYMV previously reported (Wong et al., 1994). This 32 kDa protein was recognized by anti-ZYMV antibodies in Western blot assays, confirming the identity of the isolated viral particle as a potyvirus immunogenically related to ZYMV.

ACKNOWLEDGEMENTS

We thank Dr. S. Winter, DSMZ Plant Virus Department (Germany) and his assistant for conducting the electron microscopy.

REFERENCES

Zucchini yellow Mosaic Virus from Squash

143


Tebien خصوصیات ویروس موزانیک زرد کدو مزارع کدو مسماپی استان تهران

چکیده

ویروس Zucchini yellow mosaic virus (ZYMV) یکی از ویروس‌های مهم مخرب کدو می‌باشد. طی دو سال 1361-1362 از مزارع کدو مسماپی استان تهران تعدادی نمونه جمع آوری شد. نمونه‌ها به چهار گروه متوسط و بزرگ گفته شد. نتایج آزمون ZYMV Zucchini yellow mosaic virus DAS-ELISA نشان داد که ۷۳٪ از نمونه‌ها با این ویروس نفوذ گرفته‌اند.

لیست عبنان (Z5 و Z4، Z3، Z2، Z1) با استفاده از ZYMV، RT-PCR و PAGE روش‌های ایجاد کردند. آنزیم کلون بیشترین تولیدی در نمونه‌های Z1 و Z2 ثبت شده است. روش RT-PCR با استفاده از یک جفت پرتاب که توسط Desbiez (2002) طراحی شده بود، آنالیز شد و قطعه‌ای به طول 458 جفت باز تکثیر شد.