

(ZYMV)

***** **** *** ** *

2007\2006-2006\2005

SYR-N2, JOR-N1, (ZYMV) 5
 JOR- .JOR-N3, SYR-N4, SYR-N5,
C. Chenopodium amaranticolor N1, JOR-N3, SYR-N4

SYR-N2, SYR-N5 quinoa

N. benthamiana SYR-N2

(RNA) (Cucurbita pepo)

(RT-PCR) (cDNA)

(ZYMVfor, ZYMVrev) :

(AB004641) JOR-N3
 (JOR-N1, SYR-N2, SYR-N4, SYR-N5) %100

%99-97 1

(SYR-N2) %95-92 (accession no. AF308732)

(%95) (%92)

(SYR-N2) %98-94 %100-96
 (%94)

:

(Potyviridae

(Lisa et al., 1981) 1973

Zucchini

Potyvirus ZYMV) yellow mosaic virus

%99

Desbiez & Lecoq, 1997;)

ZYMV (Albrechtsen, 2006

Mansour &) %55

.2009/12/9

2009/5/5

Mansour, 1981; Al-Musa 1982; Al-Musa, 1989

*
 **

Lisa *et al.*, 1981; Provvidenti *et al.*,) ;2005 ; 2009 ;
 1984b; Huang *et al.*, 1987; De Bon *et al.*, 1992; .(2005
 Boyhan *et al.*, 1994; Wong *et al.*, 1994; Wisler ZYMV (Genome)
 (*et al.*, 1995 (monopartite)
 ZYMV (encapsidated) (+ssRNA)
 9600 (Flexuous filamentous)
 ZYMV
 %100-93 Lisa & Lecoq) %95.5-93 %7-4.5
 (Desbiez *et al.*, 1996) 1984; Lecoq & Pitrat, 1983; Balint *et al.*, 1990;
 12 .(Dougherty & Semler, 1993
 -90.4 ZYMV
 %98.8-78 %98.8 Lisa &) (ZYMV)
 50 Lecoq, 1984; Lecoq, *et al.*, 1981; Provvidenti *et*
 ZYMV (*al.*, 1984a
 Lecoq, *et al.*, 1991, 1994; Bourdin & Lecoq,)
 Glasa & Pittnerová,) (1991; Gal-On *et al.*, 1992; Granier *et al.*, 1993
 .(2006 ZYMV-MZ
 (ZYMV) Sidek *et al.*,) *Aphis gossypii* *Myzus persicae*
 (1999
 (NCBI) ZYMV-CT ZYMV-FL ZYMV-TW
 TW-) %95 Connecticut
 %94 (NT1 .(Wang *et al.*, 1992) ZYMV-WK
 .(Augar *et al.*, 2004)
 (Amino acids) (RNA) (Fn) (melon)
 (Lecoq *et al.*, 1981) (semi-dominant)
 Ward *et*) 22
 .(*al.*, 1992 (ZYMV)
 : line) (pathotypes)
 Lisa &) (414723
 Lecoq, 1984; Risser *et al.*, 1981; Lecoq &
 25 (Pitrat, 1985
 (Guner & Wehner, 2008) (ZYMV)
 (ZYMV)

Nepovirus TRSV) ring spot virus :
 Lettuce (Comoviridae) :
Potyvirus LMV) mosaic virus 2006/2005-2005/2004
 Arabis (Potyviridae (64)
Nepovirus ArMV) mosaic virus)
 (Comoviridae) ()
 ToBRV) Tomato black ring virus ()
 .(Comoviridae *Nepovirus*
 Adgen (ZYMV)
 :
 ZYMV) Zucchini yellow mosaic virus
 (Potyviridae *Potyvirus* °4
 Cucumber green mottle
Tobamovirus CGMMV) mosaic virus
 (Tobamoviridae °20-
Potyvirus PRSV) Papaya ring spot virus :
 (Potyviridae
 WMV) Watermelon mosaic virus (ELISA) ()
 (Potyviridae *Potyvirus* .(Clark & Adams, 1977)
 SLCV) Squash leaf curl virus
 (Geminiviridae *Bigeminivirus* (0.01M)
 7.0
 Tomato : Agritest
Tombusvirus TBSV) bushy stunt virus : Florilab
 (Tombusviridae CMV) Cucumber mosaic virus
 TSWV) Tomato spotted wilt virus (Bromoviridae *Cucumovirus*
 .(Bunyaviridae *Tospovirus* AMV) Alfalfa mosaic virus
 (Bromoviridae *Alfamovirus*
 405 (ELISA Reader) SqMV) Squash mosaic virus
 (Comoviridae *Comovirus*
 MNSV) Melon necrotic spot virus
 (Tombusviridae *Camovirus*
 .
 5 :
 3) 64 *Nepovirus* ToRSV) virus
 Tobacco (Comoviridae

(
RNA) (ZYMV)
:(extraction
(RNA) JS58-05 (JOR-:
() JS32-06 (JOR-N3) SS22-06 (SYR-N2) N1
(ZYMV) .SS59-05 (SYR-N5) SS24-06 (SYR-N4)
EZ-10 Spin Column Total ()
RNA Minipreps Super Kit (Peatmoss)
Biotechnology Department Bio Basic Inc.) 10 (pots)
(RNA) (Canada)
(RNase-free H₂O) 10 (pots)
. ° 70-
(RNA)
Spectrophotometer S2100 Diode)
.(Array Biowave WPA
(cDNA) **(DNA)** (ZYMV)
(cDNA)
(ZYMV) (RNA) *Chenopodium amaranticolor*, :
Reverse Transcription) *C. quinoa*, *Nicotiana glutinosa*, *N.*
Reverse) (System *benthamiana*, *N. rustica*, *Gomphrena globosa*,,
Random) (Transcriptase AMV (0.01M) *Cucurbita pepo*.
(RNA) 2 (primer Carborundum pH 7.0
4 (200 mesh)
5mM MgCl₂, 1x Reverse) : (Control)
Transcription buffer (10mM Tris-HCl pH 9.0 at ±24
25° C), 50 mM KCl, 0.1% Triton X-100), 1mM 18 °1
each DNTP, 1U/μl Recombinant RNasin
Ribnuclease Inhibitor, 15u/μg AMV (Avian
myeloblastosis virus) Reverse Transcriptase :
(high conc., 0.5 μg Oligo(dT)15 Primer or
Random Primers per microgram RNA,
50ng/μl1.2kb Kanamycin Positive Control
.(RNA, total RNA) (ELISA) ()
(Cat. No. A3500 kit)
(Promega) (ZYMV)

:
 (Thermal Denaturation) 15 (Thermo-cycular)
 ° 95 5 ° 42
 DNA ° 94 5 ° 5-0
 ° 60 (Annealing) (cDNA)
 (Extension) ° 72 ° 20-
 DNA (cDNA) (ZYMV)
 5 (cDNA)
 (DNA) ° 72
:PCR
Electrophoresis and) (Primers)
:(staining 5'-GCTTCCGACAGGACTACGGCA-3'
 DNA ZYMVrev
 1.5) 5'-GCCGAGGTATGGTTTGCTTCG-3'
 Tris 0.605g, Boric acid) TBE 100 (Prieto, *et al.*, 2001) ZYMVfor
 (0.5X (0.2555g, EDTA 0.185g The Midland Certified Reagent Company)
 4 (Inc.
 9508-9118
 (Gel Documentation System) (3' region of coat protein gene)
(DNA analysis) (ZYMV)
 : (accession no. AF308732)
 (DNA) 395bp (DNA)
 Macrogen 20-10 100
 (DNA sequencing) () (cDNA)
 (DNA) 25 dNTP mixture 10 mM (TE
 DNAMAN-6.0 Demo *Taq*) 25 mM MgCl₂
 (Alignment) 50 (DNA Polymerase
 (DNA) 100) (Reverse Transcription buffer)
 Genebank (ZYMV) mM Tris-HCl pH 9.0 at 25° C, 500 mM KCl,
 -(amino acids) - 100 (1% Triton X-100
 (DNA) (PCR) (Nuclease-Free water)
 accession no.) (PTC-100 Programmable Thermal controller)
 .(AF308732 35

SS24-06 (SYR-N4)
 SS59-05 (SYR-N5) :
 (1 1)
 : (ZYMV)
 JS58-05 (JOR-
 25 (ELISA) () SS22-06 N1)
 .(ZYMV) JS32-06 (SYR-N2)
 (JOR-N3)

(ELISA) () :1

			Accessions		
			no.		
0.825	19/4/2005	/	EU999761	JS58-05 ** (JOR-N1)	1
1.119	8\8\2006	/	EU999762	SS22-06 * (SYR-N2)	2
0.871	20/12/2006	/	EU999763	JS32-06 (JOR-N3)	3
0.544	6/8/2006	/	EU999764	SS24-06 (SYR-N4)	4
0.81	9/7/2005	/	EU999765	SS59-05 (SYR-N5)	5

*SS: Syrian isolate **JS: Jordanian isolate

Cucurbita :
Nicotiana pepo
Gomphrena globosa rustica (ZYMV)
 (ELISA) () C.
 (ZYMV) JOR-N3 JOR-N1 *amaranticolor*
 SYR-N4
 .2 2 SYR-N5 SYR-N2
C. quinoa



.1JOR N1
.2SYR-N2
.3
JOR-N3
.4SYR-N4
.5SYR-N5

(N1, N2, N3, N4, N5)
.(ZYMV)

:1

(ZYMV) 2 :

	* Isolates				
	** (Symptoms and ELISA results)				
	JOR-N1	SYR-N2	JOR-N3	SYR-N4	SYR-N5
<i>Chenopodium amaranticolor</i>	CLL (+)	RLL (+)	CLL (+)	CLL (+)	RLL (+)
<i>C. quinoa</i>	CLL (+)	CLL (+)	CLL (+)	CLL (+)	CLL (+)
<i>Cucurbita pepo</i>	Mm, Vy (+)	Sm, B (+)	M, Ld (+)	M, CLL (+)	Ld, B (+)
<i>Nicotiana rustica</i>	-	-	-	-	-
<i>N. benthamiana</i>	-	M	-	-	-
<i>Gomphrena globosa</i>	-	-	-	-	-
<i>N. glutinosa</i>	-	-	-	-	-

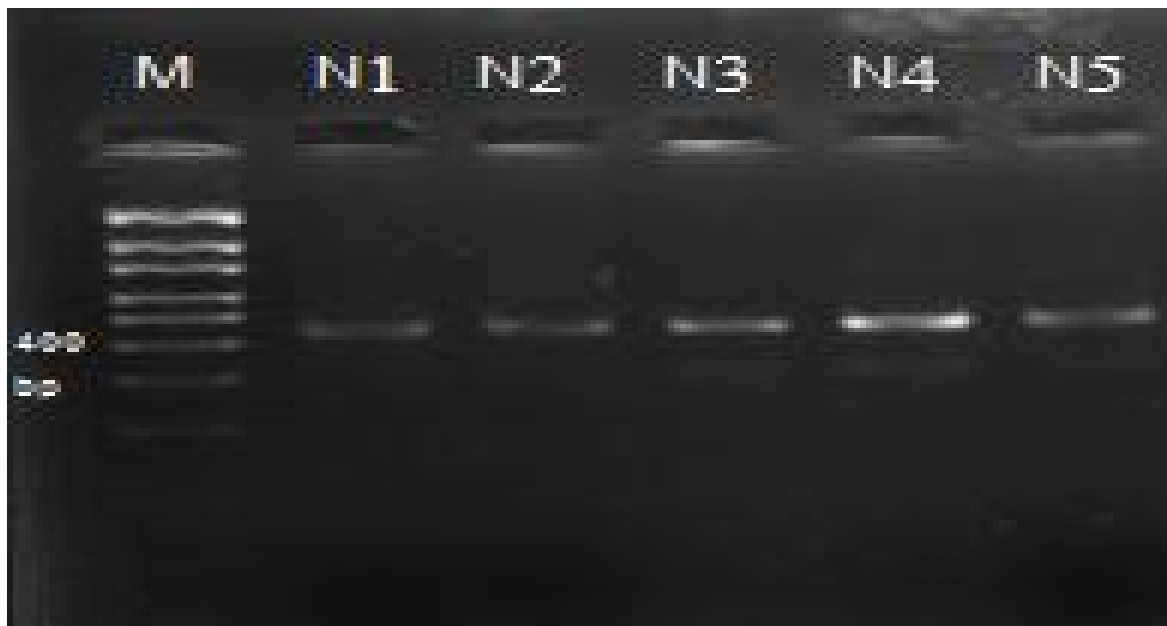
JS58-05(JOR-N1)*, SS22-06 (SYR-N2), JS32-06 (JOR-N3), SS24-06 SYR-N4), SS59-06 (SYR-N5), **Mm: mild mosaic, M: mosaic, Sm: severe mosaic, Vy: vein yellowing, -: not infected, CLL: chlorotic local lesions, RLL: red local lesions, Ld: leaf deformation, B: Blister, + positive ELISA, - negative ELISA

:PCR

:RT-PCR

ZYMVrev ZYMVfor
(ZYMV) 395bp
(DNA)
accession) (ZYMV)
(1) (Prieto *et al.*, 2001) (no. AF308732

Reverse Transcriptase)
2 (Random primer) (AMV)
(RNA)
(ZYMV)
(cDNA)
()
(cDNA)
2.000 1.600 280/260
(0.297 µg/µl) (cDNA)
(0.447µg/µl)



ZYMV)

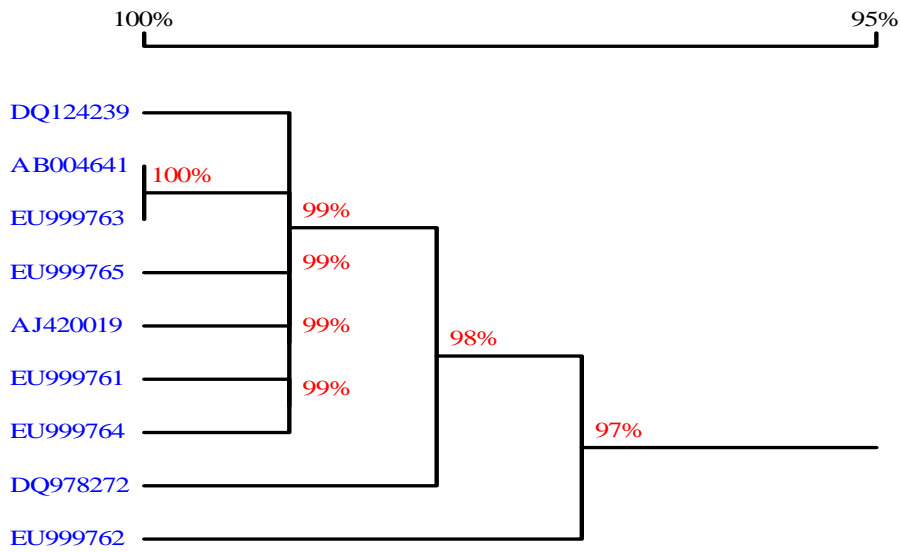
RT-PCR

:1

(Accession no. DQ124239) (Kuchyna) **sequence)**
 (Accession no. AJ420019) Berlin 1 **:(analysis**
 SS22- %97 %99
 (Accession no. EU999762) 06 (SYR-N2) DNA) (DNA) ()
 (ZYMV-KZN) %98 (sequencing
 (Accession no. DQ978272) (ZYMV)
 SS59-05 (SYR- (EU999764) (SYR-N4) (alignment)
 %99 (EU999765) N5) Kuchyna DQ124239 : GenBank
 SS22-06 %2 Berlin 1 AJ420019 AB004641 M
 .(Accession no. EU999762) (SYR-N2) 2) DQ978272 ZYMV-KZN
 SYR-N2, JOR-N3,) (Appendix 1 4
 (SYR-N4, SYR-N5 , JOR-N1, JS32-06 (JOR- (ZYMV)
 accession) %95-92 (Accession no. EU999763) N3)
 (no. AF308732 Accession no.) (ZYMV) (M)
 (SYR-N2) %100 (AB004641
 (%92) () (EU999761) JS58-05(JOR-N1) :
 .(3) (%95) SS59- () (EU999764) SS24-06 (SYR-N4)
 () (EU999765) 05 (SYR-N5)

(accession no. AF 308732) DNA :3

	ZYMV (AF 308732)	ZYMV (JOR-N1)	ZYMV (SYR-N4)	ZYMV (JOR-N3)	ZYMV (SYR-N2)	ZYMV (SYR-N5)
ZYMV (AF 308732)	100	95	95	95	92	95
ZYMV (JOR-N1)			99	98	97	99
ZYMV (SYR-N4)				98	96	99
ZYMV (JOR-N3)					97	99
ZYMV (SYR-N2)						97
ZYMV (SYR-N5)						100



Accession no.		Accession no.	
EU999761	JS58-05(JOR-N1)	DQ124239	Kuchyna
EU999762	SS22-06 (SYR-N2)	AB004641	M
EU999763	JS32-06 (JOR-N3)	AJ420019	Berlin 1
EU999764	SS24-06 (SYR-N4)	DQ978272	ZYMV-KZN
EU999765	SS59-05 (SYR-N5)	-	

(ZYMV) (DNA) 5 .2
GenBank 4 ZYMVrev ZYMVfor

...
 (AF 308732 :
 N1, N2, N3, N4,) (ZYMV)
 (4) %100-96 (N5 (DNA)
 %98-94 (alignment) (ZYMV)
 .(%94) (SYR-N2) Accession no.)

.(accession no. AF 308732)

:4

	ZYMV (AF 308732)	ZYMV (JOR-N1)	ZYMV (SYR-N4)	ZYMV (JOR-N3)	ZYMV (SYR-N2)	ZYMV (SYR-N5)
ZYMV (AF 308732)	100	98	97	98	94	98
ZYMV (JOR-N1)			99	100	97	100
ZYMV (SYR-N4)				99	96	99
ZYMV (JOR-N3)					97	100
ZYMV (SYR-N2)						97
ZYMV (SYR-N5)						100

:

Lisa & Lecoq, 1984;) (ZYMV)
 Risser *et al.*, 1981; Lecoq & Pitrat, 1985; Wang
et al., 1992; Choi *et al.*, 2003; Hosseini *et al.*,
 (2007 (DNA sequence)
 (*Cucurbita pepo*) (Coat protein) Cp
 (SYR-N2) .()
 . (SYR-N4) (ZYMV)

JOR-N1, JOR-N3, SYR-N4, : (ZYMV) :SS.24-06 (SYR-N4)
C. quinoa C. amaranticolor

(Coat protein)	(analysis	Sidek <i>et al.</i> , 1999; Lisa &)
	(ZYMV)	Lecoq, 1984; Lecoq <i>et al.</i> , 1981; Provvidenti <i>et</i>
(Desbiez <i>et al.</i> , 1996)	%99-97	(<i>al.</i> , 1984a; Sidek <i>et al.</i> , 1999
	(ZYMV)	
%100-98	(ZYMV)	(Wong <i>et al.</i> , 1994)
	(ZYMV)	
	(Glasa & Pittnerová, 2006)	Lisa & Lecoq, 1984; Lecoq <i>et al.</i> , 1981;)
		Provvidenti <i>et al.</i> , 1984a; Lecoq & Pitrat, 1985;
	(ZYMV)	(Lecoq & Purcifull, 1992
%100-96		(ZYMV)
SYR-)	%98-94	
(%94)	(N2	(Desbiez <i>et al.</i> , 1996)
		DNA Sequence)

7 . .112-84 :2 .2005 . .2009 . 23 . .6-1 :1 .2005 .2 5

Albrechtsen, S. E. 2006. Testing methods for seed-transmitted viruses: principles and protocols. CABI Publishing, ISBN. pp 2.

Al-Musa, A. M. 1989. Severe mosaic caused by *Zucchini yellow mosaic* in Cucurbits from Jordan. *Plant Pathology*, 38: 541-546.

Augar, J., Martinez, C., Esterio, M. and Prieto, H. 2004. Identification and transmission of a strain of *Zucchini yellow mosaic virus* (ZYMV) causal agent of fruit deformation in squash (*Cucurbita maxima*) in Chile. *Fitopatologia*, 39 (3): 133-143.

Balint, R., Plooy, I. and Steele, C. 1990. The nucleotide sequence of *zucchini yellow mosaic potyvirus*. *Abstr. VIIIth Int. Congress Virology*. 8: 84-107.

Bourdin, D. and Lecoq, H. 1991. Evidence of heteroencapsidaion between two *potyviruses* is involved in aphid transmission of a non-aphid-transmissible isolate from mixed infections. *Phytopathology*, 81 (11): 1459-1464.

Boyhan, G. E., Gudauskas, R. T., Norton, J. D. and Abrahams, B. R. 1994. Evaluation of watermelon and germ plasm for resistance to the Egyptian strain of *Zucchini Yellow Mosaic*

-
- virus. Plant Disease*, 78 (1): 100.
- Choi, S. K., Yoon, J. Y., Choi, S. H. and Ryu, K. H. 2003. Movement of *Zucchini yellow mosaic virus* involved in symptoms severity on Zucchini Squash. *Plant Pathology Journal*, 19 (4): 217-220.
- Clark, M. F. and Adams, A. N. 1977. Characteristics of the microplate method of enzyme-linked immunosorbent assay for the detection of plant viruses. *Journal of General Virology* 34: 475 - 483.
- De Bon, H., Boissot, N., Girard, J. C., Michellon, R., Reynaud, B. and Vercambre, B. 1992. A note on research on vegetable crops in Reunion. In: Ninth STASM (Societe de Technologie Agricole et Sucriere de Maurice) Conference. Reunion, Mauritius: *Revue Agricole et Sucriere de l'Ille Maurice*, 71:148-152.
- Desbiez, C. and Lecoq, H. 1997. Review *zucchini yellow mosaic virus*. *Plant Pathology* 46: 809-829.
- Desbiez, C., Wipf-Scheibel, C., Granier, F., Robaglia, C., Delaunay, T. and Lecoq, H. 1996. Biological and molecular variability of *Zucchini yellow mosaic virus* on the Island of Martinique. *Plant Disease*, 80: 203-207.
- Dougherty, W. G. and Semler, B. L. 1993. Expression of virus encoded proteinases, functional and structural similarities with cellular enzymes. *Microbiology Review*, 57: 781-822.
- Glasa, M. and Pittnerová, M. S. 2006. Complete genome sequence of a Slovak isolate of *Zucchini Yellow Mosaic Virus* (ZYMV) provides further evidence of a close molecular relationship among central European ZYMV isolates. *Journal of Phytopathology*, 154: 436-440.
- Granier, F., Durand, T. M., Casse, D. F., Lecoq, H. and Robaglia, C. 1993. Mutations in *Zucchini yellow mosaic virus* helper component protein associated with loss of aphid transmissibility. *Journal of General Virology*, 74: 2737-2742.
- Huang, C. H., Chao, Y. J., Chang, C. A., Hseu, S. H. and Hsiao, C. H. 1987. Identification and comparison of different viruses on symptom expression in loofah. *Journal of Agricultural Research of China*, 36 (4):413-420.
- Hosseini, S., Mosahebi, G. H., Koohi Hbib, M. and Okhovvat, S. M. 2007. Characterization of the *Zucchini yellow mosaic virus* from squash in Tehran Province, *Journal Agriculture Science Technology*, 9: 137-143.
- Lecoq H., Bourdin D., Raccach, B., Hiebert, E. and Purcifull, D. E. 1991. Characterization of a *Zucchini yellow mosaic virus* isolate with a deficient helper component. *Phytopathology*, 81: 1087-1091.
- Lecoq, H. and Pitrat, M. 1983. Field experiments on the integrated control of aphid-borne viruses in muskmelon. In: Plumb RT, Tresh JM, eds *Plant virus Epidemiology*. Oxford, UK: Blackwell Scientific Publications, pp 169-176.
- Lecoq, H. and Pitrat, M. 1985. Specificity of the helper-component-mediated aphid transmission of three potyviruses infecting muskmelon. *Phytopathology*, 75: 890-903.
- Lecoq, H., Pitrat, M. and Clement M. 1981. Identification et caracterisation d'un potyvirus provoquant la maladie du rabougrissement jaune du melon. *Agronomie*, 1: 827-34.
- Lecoq, H. and Purcifull, D. E. 1992. Biological variability of potyviruses, an example: *Zucchini yellow mosaic virus*. *Archives of Virology Vienna*, Austria; Springer-Verlag/Wien, Supplementum, 5:229-234.
- Lisa, V. and Lecoq, H. 1984. *Zucchini yellow mosaic virus* CMI/AAB. *Description of Plant viruses*. No. 282.
- Lisa, V., Boccardo, G., D'Agostino, G., Dellavalle, G., D'Aquilio, M. 1981. Characterization of a *potyvirus* that causes *zucchini yellow mosaic virus*. *Phytopathology*, 71: 667-672.
- Mansour, A. N. 1981. Isolation and Identification of plant Viruses affecting Squash (*Cucurbita pepo*) in Jordan. M.Sc. Thesis, University of Jordan, Amman, Jordan.
- Mansour, A and AL-Musa, A. 1982. Incidence, economic importance and prevention of *watermelon mosaic virus-2* in squash (*Cucurbita pepo*) fields in Jordan *Phytopathology*. Z. 103: 35-40.
- Prieto, H., Bruna, A., Hinrichsen, P. and Muñoz, C. 2001. Isolation and molecular characterization of a Chilean isolate of *zucchini yellow mosaic virus*. *Plant Disease*, 85:

- 644-648.
- Provvidenti, R., Gonsalves, D. and Humaydan, H. S. 1984a. Occurrence of *Zucchini yellow mosaic virus* in cucurbits from Connecticut, New York, Florida and California. *Plant Disease*, 68: 443-6.
- Provvidenti, R., Munger, H. M., Paulus, M.O. 1984b. Epidemics of *Zucchini yellow mosaic virus* in Egypt in the spring of 1983. *Cucurbit Genetics Cooperative*, 7, 78-79.
- Risser, G., Pitrat, M., Lecoq, H. and Rode, J. C. 1981. Sensibilité variétale du melon (*Cucumis melo*) au virus du rabougrissement jaune du melon (MYSV) et à sa transmission par *Aphis gossypii*, Héredite de la réaction de flétrissement. *Agronomie*, 1: 835.
- Sidek, Z., Sako, N. and Ohshima, K. 1999. Characterization and coat protein sequence of a Malaysian isolate of *Zucchini yellow mosaic virus*. *Journal of Plant Protection in Tropics* 12 (2): 92-105.
- Wang H. L., Gonsalves D., Provvidenti R. and Zitter, T. A. 1992. Comparative biological and serological properties of four strains of *Zucchini yellow mosaic virus* *Plant Disease* 76 (5): 530-535.
- Wisler, G., Purcifull, D. and Hiebert, E. 1995. Characterization of P1 protein and coding region of *zucchini yellow mosaic virus*. *Journal of General Virology*, 76: 37-45.
- Wong, S., Chng, C. G., Chng, C. Y. and Chong, P. L. 1994. Characterization of an isolate of *Zucchini yellow mosaic virus* from cucumber in Singapore. *Journal of Phytopathology*, 141: 355-368.

Biological and Molecular Characterization of Some *Zucchini yellow mosaic* Virus Isolates from Southern Syria and Jordan Valley

*Al-Tamimi**, *N.*, *Kawas***, *H.*, *Mansour****, *A.*, *Arabi*****, and *Nida Salem******

ABSTRACT

Zucchini yellow mosaic virus (ZYMV) is an important viral pathogen of squash plants. This study was conducted during the growing seasons of (2005/2006-2006/2007) to differentiate between 5 ZYMV isolates obtained from southern Syria (3) and Jordan Valley (2) using biological and molecular techniques. The nomenclature of the virus isolates was as follows: JOR-N1, SYR-N2, JOR-N3, SYR-N4 SYR-N5. Indicator plants reacted differently when they were inoculated with these isolates. Isolates JOR-N1, JOR-N3 and SYR-N4 induced chlorotic local lesions on *Chenopodium amaranticolor* and *C. quinoa*, whereas red to pink local lesions were observed on plants inoculated with SYR-N2 and SYR-N5 isolates. On *Cucurbita pepo*, JOR-N1 isolate caused mild mosaic and vein yellowing while, isolate SYR-N2 induced severe mosaic and blister. Isolate JOR-N3 caused mosaic and leaf deformation. Squash plants inoculated with SYR-N4 developed chlorotic local lesions and mosaic symptoms while, SYR-N5 induced leaf deformation and blister. Total RNA isolation was extracted from squash plants infected with JOR-N1, SYR-N2, JOR-N3, SYR-N4 SYR-N5 isolates and oligonucleotide primer pair ZYMVfor/ZYMVrev was used to amplify a fragment of about 395bp of the coat protein gene of ZYMV. Sequence analysis of the amplified PCR product showed that JOR-N3 isolate shared high (100%) nucleotide identity with the Japanese isolate (AB004641) of ZYMV, while, the nucleotide similarity of the other isolates ranged between 97% and 99%. On the other hand, the nucleotide sequence similarities between the five isolates studied in this work and ZYMV isolates from Slovakia, Germany and South Africa ranged between 97% and 99%. The lowest sequence similarity of the five isolates was recorded with ZYMV isolate from Chili. Sequence analysis of amino acids indicated that the five isolates shared high (96-100%) identities with each other, and 94-98% when compared with the Chili isolate.

Keywords: ZYMV isolates, southern Syri, Jordan Valley, squash viruses.

*

**

Received on 5/5/2009 and Accepted for Publication on 9/12/2009.