

Serological Detection of Grapevine Associated Closteroviruses in Infected Grapevine Cultivars

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ABSTRACT

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Western blot immunoassay and enzyme-linked immunosorbent assay using different monoclonal antibodies (MAb) and polyclonal antisera (PA) revealed mixed infections of serologically related and unrelated grapevine leafroll associated viruses (GLRaVs) and grapevine corky bark associated virus (GCBaV) in symptomatic grapevines. A PA designated rootstock-scion incompatibility (RSI)-24 kDa, grapevine corky bark PA, and GLRaV-2b MAb reacted to polypeptides of approximately 24 kDa isolated from grapevines exhibiting rootstock-scion incompatibility, leafroll, and corky bark disease symptoms, suggesting that these isolates are infected with closely related viruses. A PA designated GLRaV-2 US detected virus specific polypeptides of 38, 37, 36, and 24 kDa, while a polyclonal antiserum designated GLRaV-2 FR detected a single virus-specific polypeptide of approximately 24 kDa. The reactivity of GLRaV-2 US to various polypeptides suggests that the immunogen used to produce this antiserum was a mixture of viruses. Apical meristems were excised and cultured to eliminate the infection of viruses in the grapevines showing RSI symptoms and in the cultivar French Colombard infected with GLRaV-1. The elimination of these viruses was confirmed by Western blot assay. These studies show that the Western blot assay can be used to detect and differentiate grapevine disease-associated closteroviruses.

Grapevine leafroll (LR) and corky bark (CB) diseases occur wherever grapevines are grown and are associated with undesirable viticultural effects which include delayed ripening of fruit, reduced yield, altered fruit pigmentation, and reduced accumulation of sugar (6,24). Rootstock-scion incompatibility (RSI) is a disease syndrome characterized by incompatibility at the graft union resulting in slow death of the scion (5).

The etiology of LR disease is not clear. Different kinds of virus particles have been associated with symptomatic variants of LR disease (24), but only closterovirus-like particles have been consistently associated with this disease (14). Viruses found in LR infections are designated grapevine leafroll associated viruses (GLRaVs). Other closterovirus-like particles designated as grapevine corky bark associated viruses (GCBaVs) were found in CB infections (20). Many cultivars often carry latent infections of grapevine viruses and are asymptomatic until they are grafted onto a susceptible rootstock (5).

Several laboratories have purified closterovirus-like particles from diseased vines and prepared monoclonal antibodies (MAb) and polyclonal antisera (PA)

(8,10,11,20,26). Using sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE), the molecular weights of GLRaV-1, -3, -4, and GCBaV coat proteins have been reported to be approximately 38, 43, 36, and 26 kDa, respectively (8,10,11,20). Two different molecular weights (circa 26 and 36 kDa, respectively) have been reported for the coat proteins of the United States and French GLRaV-2 isolates (2,26), suggesting that these are different viruses. In addition, Gugerli and Ramel (8) reported that a MAb designated MCA 29-1 (prepared with GLRaV-2b) reacted to a polypeptide of approximately 26 kDa in a Swiss virus isolate.

With the exception of GLRaV-2, GLRaV-2b, and GCBaV, the reported mass of the capsid proteins of grapevine associated closterovirus-like particles are greater than expected for closterovirus capsid proteins (circa 22 to 26 kDa). This information suggests that these viruses could represent a new virus group.

Presently, LR is diagnosed by biological indexing and enzyme-linked immunosorbent assay (ELISA), while CB is diagnosed by biological indexing only. Symptom evaluation of an indicator host is subjective, non-specific, and time consuming. Discrepancies have been found between ELISA and woody indicator tests (18,22). GLRaV-1 and GLRaV-3 were detected by ELISA and Western blotting in previously registered foundation material that had been determined to be virus free by biological indexing (17,18). ELISA has also been unreliable, since the poor quality of

the PA available would yield a positive reaction in response to non-specific reactivity or infection with one or more viruses. Furthermore, the distribution of GLRaVs is variable in vegetatively grown and dormant grapevines (18), making the choice of sampling strategies critical to successful detection of grapevine associated closteroviruses.

In this study we show that a Western blot (WB) assay using various antibodies permits reliable and accurate detection of viral polypeptides, and hence diagnosis of viruses associated with RSI, leafroll, and corky bark diseases. Furthermore, the WB assay has allowed the characterization of PA reactive to several virus isolates.

MATERIALS AND METHODS

Plant materials. The isolates used in this study are described in Table 1. Biological indexing confirmed the infection status of all positive and negative controls, with the exception of the French Colombard (FC/2) cultivar that tested negative by grafting onto indicator hosts but tested positive using ELISA (18). Dormant cuttings of LR 101, LR 102, LR 105, LR 106, LR 109, and CB 100 (4) were provided by Deborah Golino (University of California, Davis); positive controls designated in our laboratory as LR-A, LR-B, LR-C, and LR-D were provided by Adib Rowhani (University of California, Davis). Grapevines with RSI, and field selections that reacted with corky bark and GLRaV-2b antibodies (17) were obtained from California vineyards. The negative controls (FPMS certified rootstock cultivars) were provided by Vinifera, Inc. (Petaluma, CA). Cuttings from the above-mentioned vines were rooted and grown in our research greenhouse in Beaverton, Oregon. The vines were grown from March until December at approximately 25 to 28°C and supplemented with 4 h of light. Vines were allowed to go dormant during the period between December and March in a cold frame greenhouse (no temperature or light adjustments).

Serological reagents. The PA and MAb used in this study are described in Table 2. GLRaV-2 US, -3, -4, and GCBaV PA were cross-absorbed (7) with virus-free *Vitis riparia* Gloire (RG) extracts prior to immunoglobulin purification by protein A chromatography and conjugation with alkaline phosphatase (9). GLRaV-1 MAb and alkaline phosphatase-labeled GLRaV-1 MAb used in double antibody sandwich

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ELISA (DAS-ELISA) were purchased from BIORBEA AG (Basel, Switzerland). Phosphatase-labeled goat anti-rabbit and goat anti-mouse IgG used in the WB assay were purchased from Kirkegaard & Perry Laboratories, Inc. (Gaithersburg, MD).

Production of polyclonal antisera.

Polyclonal antisera were produced in duplicate New Zealand rabbits using gel purified SDS-PAGE viral coat proteins. Concentrated extracts from healthy RG grapevine were used to immunize rabbits. The RG extracts were verified to be free of viruses using immunosorbent electron microscopy (ISEM), ELISA, and WB.

GLRaV-1 and RSI-associated virus particles (17) were purified from FC/2 and RSI-symptomatic grapevine cultivars, respectively. Healthy RG extracts were concentrated with the same method used for virus purification. Briefly, stem and petiole tissue were cut in small portions and ground in liquid nitrogen, the tissue was homogenized in virus extraction buffer, and the virus or host proteins were concentrated by two series of differential centrifugation (25). Preparative SDS-PAGE were used to separate the viral proteins from contaminating host proteins. Purified virus preparations (250 µl) were resolved by SDS-PAGE (14% polyacrylamide) (13). The region of the gel containing the viral capsid protein was identified by blotting one lane of each slab gel and testing the reactivity of the transferred polypeptides with anti-GLRaV-2 US PA and GLRaV-1 MAb (38 kDa) or anti-GCBaV PA and GLRaV-2b (24 kDa) in WBs. The WB was aligned with the remaining lanes of the gel and the region corresponding to the 24 or 38 kDa polypeptide bands were excised and eluted using an Elutetrap apparatus (Schleicher & Schuell, NH). To ensure purity of the protein preparation, the eluted protein was run in a second SDS-PAGE and eluted a second time. The identity of the purified proteins was verified by WB using GLRaV-2 US PA and GCBaV PA before the rabbits were immunized; the PA were used in the WB because they would have revealed the contamination of host proteins in the preparation if present. Healthy RG extracts were used as a negative control. A single protein band of 24 or 38 kDa was observed in all protein preparations of RSI and FC/2, respectively. Multiple bands observed in the healthy extract lanes reacted with GLRaV-2 US and GCBaV PA.

Prior to immunizations, rabbits were bled for pre-immune sera. Rabbits that had the lowest reactivity to healthy grapevine extracts were chosen for immunizations with gel-purified 38-kDa (GLRaV-1) and 24-kDa (RSI) polypeptides and concentrated extracts of healthy proteins. All protein extracts were emulsified using complete Freund's adjuvant prior to primary immunizations (9). Three additional injections containing Freund's incomplete

adjuvant were performed 3, 6, and 9 weeks after the primary injection. Test bleeds were taken 1 week after each booster injection to determine the reactivity of the antiserum to the virus and plant protein

extracts using the Western blot assay. The total serum was collected 1 month after the final booster injection. The antisera produced will be referred to as RSI-24 kDa, GLRaV-1, and RG PA.

Table 1. Virus-infected and noninfected grapevine cultivars

Isolate designation	Grapevine variety	Symptoms on indicator
LR A	Unspecified field selection	Leafroll
LR B	Cabernet Sauvignon	Leafroll
LR C	Italia	Leafroll
LR D	Cabernet Sauvignon	Leafroll
LR 101	Thompson Seedless	Leafroll
LR 102	Thompson Seedless	Leafroll
LR 105	Teroldigo-1	Leafroll
LR 106	Thompson Seedless	Leafroll
LR 109	Thompson Seedless	Leafroll
FC/2	French Colombard	None
RSI	Malbec	Incompatibility
CB100	Semillion	Corky bark
SJ	Unspecified field selection	Incompatibility/corky bark
MBF	Merlot	Incompatibility
PS	Petit Shirah	Incompatibility
PS3	Petit Shirah	Leafroll/incompatibility
Healthy	<i>Vitis riparia</i> Gloire	None
Healthy	Couderc 3309	None
Healthy	Kober 5BB	None

Table 2. Source of serological reagents

Antibody	Source	Reference
GLRaV-1, MAb ^a	BIORBEA AG	8
GLRaV-1, PA ^b	Agritope, Inc.	This report
GLRaV-2, US PA	D. Gonsalves, Cornell University	2
GLRa-2, FR PA	Sanofi Diagnostics Pasteur	26
GLRaV-2b, MAb	P. Gugerli, Nyon Experimental Station, Switzerland	8
RSI-24 kDa, PA	Agritope, Inc.	This report
GLRaV-3, MAb	D. Gonsalves, Cornell University	10
GLRaV-3, PA	D. Gonsalves, Cornell University	25
GLRaV-4, PA	D. Gonsalves, Cornell University	11
Black Spanish, PA	G. Pietersen, Plant Protection Res. Institute, South Africa	21
GCBaV, PA	D. Gonsalves, Cornell University	20
Healthy, PA	Agritope, Inc.	This report

^a MAb, monoclonal antibodies.

^b PA, polyclonal antiserum.

Table 3. Enzyme-linked immunosorbent assay (ELISA) reactivity of virus isolates

Virus isolates	Antibodies ^a			
	GLRaV-1	GLRaV-2	GLRaV-3	GLRaV-4
FC/2	3+ ^{b,c}	+	-	-
LR A	3+	+	-	-
LR B	2+	+	-	-
LR 102	3+	2+	-	-
LR 105	3+	+	4+	-
RSI	-	-	-	-
LR C	-	-	4+	-
LR 101	-	-	+	-
LR 109	-	-	3+	-
LR D	-	-	-	+
LR 106	-	-	-	+
Healthy	- ^d	-	-	-

^a The DAS-ELISA was performed with GLRaV-1 monoclonal antibodies (MAb), GLRaV-2 US polyclonal antiserum (PA), GLRaV-3 PA/MAb, and GLRaV-4 PA.

^b The sample readings were scored in relation to the ratio of the A_{405} of each sample and 2× the reading of the healthy negative control. The following scores, (-), (+), (2+), (3+), (4+) were assigned to each treatment with a ratio of less than 1 (no reactivity), 1 to 2.499, 2.5 to 3.499, 3.5 to 4.499, and above 4.5, respectively.

^c The (+) A_{405} (average of 4-10 wells) of infected extracts were 2.376, 1.711, 2.206, and 0.728 for ELISA using GLRaV-1, -2, -3, and -4 antibodies, respectively.

^d The (-) A_{405} (average of 10 wells) healthy extracts were 0.295, 0.399, 0.251, and 0.271 for ELISA using GLRaV-1, -2, -3, and -4 antibodies, respectively.

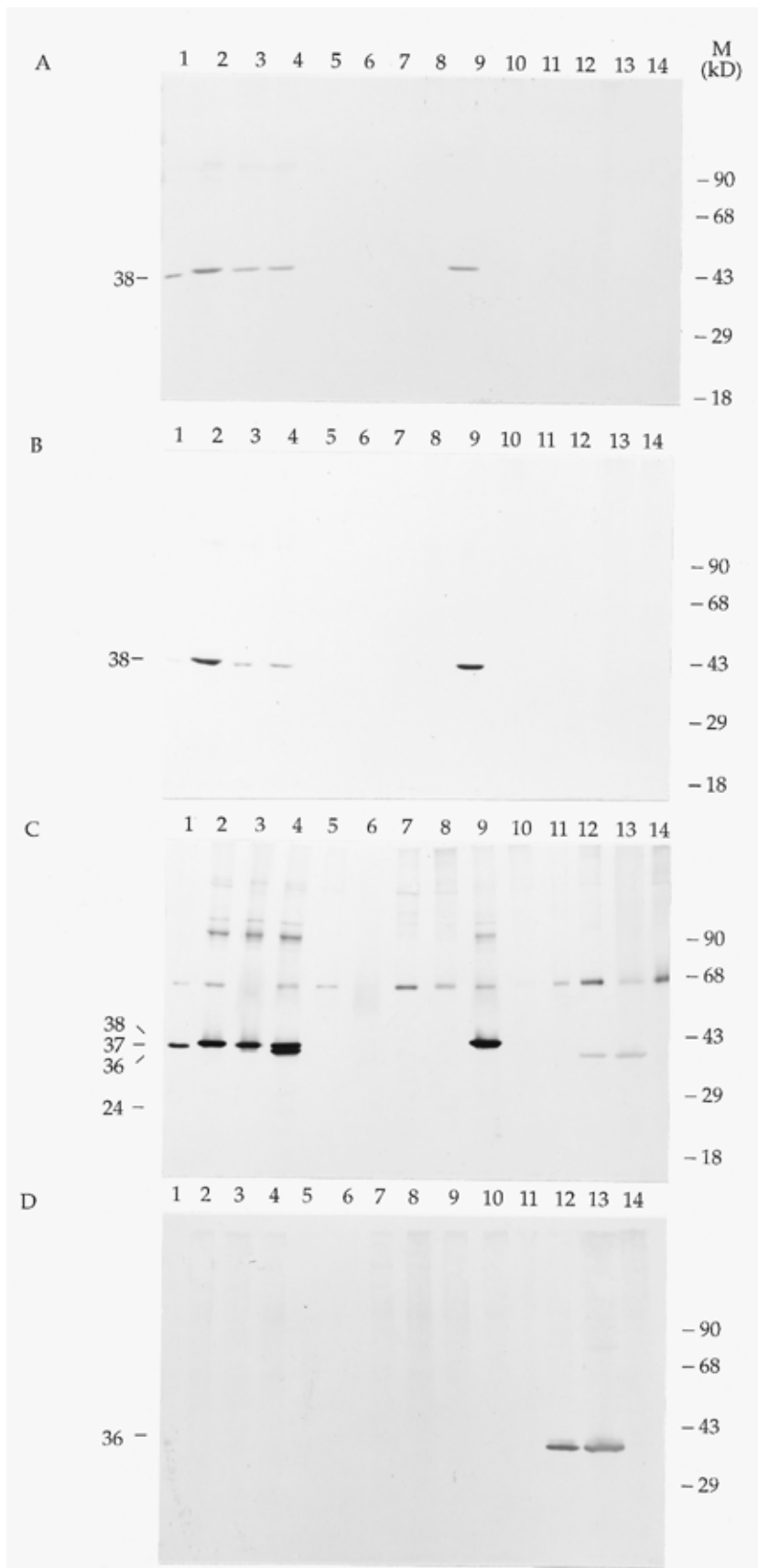


Fig. 1. Western blot assay of protein extracts using the following: GLRaV-1 PA (A); GLRaV-1 MAb (B); GLRaV-2 US PA (C); GLRaV-4 PA (D). The grapevine cultivars or isolates analyzed are: lanes 1, FC/2; lanes 2, LR-A; lanes 3, LR-B; lanes 4, LR 102; lanes 5, RSI; lanes 6, SJ; lanes 7, CB 100; lanes 8, LR 109; lanes 9, LR 105; lanes 10, LR 101; lanes 11, LR-C; lanes 12, LR-D; lanes 13, LR 106; lanes 14, healthy. The molecular weight of size markers and the viral specific proteins are indicated on the sides of each blot.

ELISA. The double antibody sandwich enzyme-linked immunosorbent assay (DAS-ELISA) was performed in duplicate wells for each sample as previously described (18) using GLRaV-1, -2, -3, and -4 antibodies.

Grapevine stem and petioles samples were collected from the bottom of the vine (18) and extracted with a tissue pulverizer (KLECO, Visalia, CA) at 1:5 (wt/vol) with 0.5 M Tris-HCl pH 8.2, 143 mM NaCl, 1% polyethylene glycol (MW 8000), 2% polyvinyl pyrrolidone (MW 40,000), and 0.05% Tween-20. Ground samples were occasionally stored at -20°C prior to ELISA tests. The sample tests were generally incubated for 2 h at room temperature with substrates, but some experiments required overnight incubations at 4°C . The microtiter plates were read at an absorbance of 405 nm.

Initial ELISA tests were performed with the antibodies described above to determine the infection status of each virus isolate. The threshold value for a positive ELISA reading was arbitrarily determined to be two times the average A_{405} value of the healthy control (2 to 6 wells). The sample readings were scored in relation to the ratio of the A_{405} of each sample and two times the A_{405} reading of the healthy negative control. The following scores, (-), (+), (2+), (3+), and (4+), were assigned to each treatment with a ratio of less than 1 (not infected), 1 to 2.499, 2.5 to 3.499, 3.5 to 4.499, and above 4.5, respectively (18).

Western blot assay. Samples for the WB assay were collected from stems and petioles of actively growing greenhouse plants (Monis and Bestwick, 1996) and prepared essentially as described by Zee et al. (25). The viral proteins were resolved by SDS-PAGE (10% polyacrylamide) (13) and transferred electrophoretically to nitrocellulose membranes (23). The blots were incubated in 0.02 M Tris pH 6.5, with 0.5 M sodium chloride (TBS), containing 5% nonfat dry milk, and 0.03% Tween-20 (blocking buffer) for 1 h at room temperature. Each of the antibodies described in Table 2 were added individually to the blocking buffer and incubated with replica blots for 1 h at room temperature. The membranes were washed (three times for 15 min) with TBS containing 3% nonfat dry milk and 0.03% Tween-20 and incubated for 1 h with alkaline phosphatase conjugated goat anti-rabbit or goat anti-mouse IgG diluted 1:2000 in blocking buffer. The membranes were washed as described above, except that the final wash was performed with TBS. Immuno-reactive proteins were visualized using an NBT/BCIP development solution (Kirkegaard & Perry Laboratories, Inc.)

Excision and culture of apical meristems. For in vitro propagation, 1-cm nodal sections of vegetatively growing stems were surface sterilized and grown in initiation and propagation media (12,19) under

16 h light. For virus elimination, approximately 0.5 mm apical meristems with 1 to 2 leaf primordia were excised and placed in solid medium (12). The regenerated meristems were sub-cultured at least once a month in propagation media (19). The infection status of the regenerated grapevine plantlets was determined using WB. The tissue-culture-grown plantlets were acclimated in the greenhouse and tested again to confirm the lack of virus infection with the WB assay using the RSI-24 kDa, GLRaV-1, and RG PA.

RESULTS

Production of polyclonal antisera to GLRaV-1, RSI-24 kDa, and proteins from healthy plants. The GLRaV-1 PA and RSI-24 kDa PA reacted with polypeptides of the expected molecular weight in FC/2 and RSI-infected material, respectively. The antiserum specific to RG protein extracts reacted with several polypeptides in healthy and virus infected grape extracts in WB assays.

Serological cross-reactivity of virus isolates. ELISA results are summarized in Table 3. Several virus isolates reacted with one or more of the antibodies used in this study. For example, LR-A, LR-B, and LR 102 reacted with GLRaV-1 MAb and GLRaV-2 US PA; LR 105 reacted with GLRaV-1 MAb, GLRaV-2 US PA, and GLRaV-3 PA. Extracts from healthy RG, Kober 5BB, and Couderc 3309 cultivars did not react specifically in ELISA or WB to any of the antibodies described in Table 2.

Western blot immunoassay. Extracts of grapevines were analyzed by WB using the antibodies described in Table 2. GLRaV-1 PA and GLRaV-1 MAb reacted with a single viral-specific polypeptide of circa 38 kDa in extracts from FC/2, LR-A, LR-B, LR 102, and LR 105 isolates (Fig. 1A and B, lanes 1-4, 9).

Using GLRaV-2 US PA, a polypeptide band of circa 38 kDa was detected in extracts from FC/2, LR-A, LR-B, LR 102, and LR 105 viral isolates (Fig. 1C, lanes 1-4, 9); note that in addition to the polypeptide of circa 38 kDa, a faint lower molecular weight band was observed in extracts from LR-B. WB performed with gels that had been run for a longer period of time with LR-B extracts clearly showed a doublet of circa 37 to 38 kDa (not shown). In addition to the doublet of circa 37 and 38 kDa, the GLRaV-2 US PA recognized faintly a polypeptide of circa 24 kDa in extracts from LR 102 isolate (Fig. 1C, lane 4), and one viral specific polypeptide of approximately 36 kDa in LR-D and LR 106 isolates (Fig. 1C, lanes 12 and 13). Crude GLRaV-2 US PA showed stronger reactivity to the 24 kDa polypeptide in LR 102 extracts (not shown). The GLRaV-2 US PA reacted to a number of other non-viral polypeptides (predominantly of circa 68 kDa).

The GLRaV-4 PA (Fig. 1D) reacted with a single viral polypeptide of circa 36 kDa in extracts from LR-D and LR 106 extracts (Fig. 1D, lanes 12-13).

The GLRaV-2b MAb, RSI-24 kDa PA, and GCBaV PA detected a virus-specific polypeptide of circa 24 kDa in extracts from LR 102, RSI, SJ, CB 100, LR 109, LR 101, and LR-C (Fig. 2A, B, and C; lanes 4-8, 10-11). An additional polypeptide of circa 26 kDa, the significance of which is not known, could be seen in extracts from the LR 109 isolate (Fig. 2B, 8). This 26-kDa polypeptide was not observed

in every virus preparation or WB assay performed using the same virus isolates. A doublet of circa 29 to 30 kDa and other polypeptides of circa 40 kDa and larger than 90 kDa, presumably of plant origin, were detected with GCBaV PA (Fig. 2C, lanes 2, 4-5, 7-8, 10, 12-14). Additional, field selections grown in the Napa Valley in California reacted to the RSI-24 kDa, GLRaV-2b, and GCBaV antibodies, include the Merlot and Petit Shirah cultivars (Table 4). The GLRaV-2 FR PA reacted to a single polypeptide of circa 24 kDa from extracts of LR 102, RSI, SJ, CB 100, LR

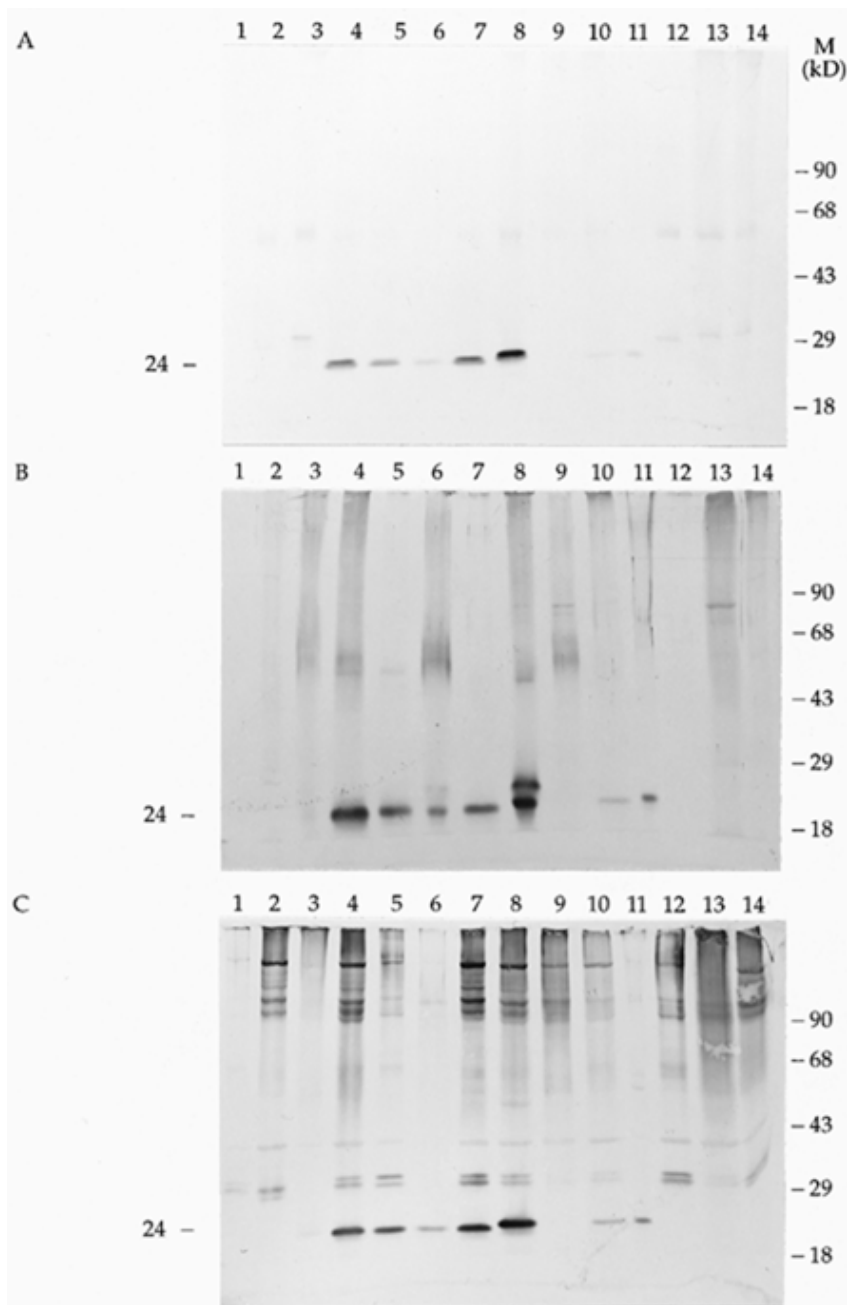


Fig. 2. Western blot assay of protein extracts using the following: GLRaV 2b (A); RSI-24 kDa PA (B); GCBaV PA (C); GLRaV-2 FR PA (D). The grapevine cultivars or isolates analyzed are: lanes 1, FC/2; lanes 2, LR-A; lanes 3, LR-B; lanes 4, LR 102; lanes 5, RSI; lanes 6, SJ; lanes 7, CB 100; lanes 8, LR 109; lanes 9, LR 105; lanes 10, LR 101; lanes 11, LR-C; lanes 12, LR-D; lanes 13, LR 106; lanes 14, healthy. The molecular weight of size markers and the viral specific proteins are indicated on the sides of each blot.

109, LR 101, and LR-C isolates (Fig. 3). The reactivity of the GLRaV-2 FR PA was weak even when low dilutions of the antiserum (1:50) were used in the WB assay.

A virus-specific polypeptide of circa 43 kDa was detected by GLRaV-3 MAb (Fig. 4A) and GLRaV-3 PA (not shown) in extracts from LR 109, LR 105, LR 101 and LR-C isolates (Fig. 4A, lanes 8-11). The Black Spanish PA, which reacts with GVA, GLRaV-1, -2 and -3 (21), bound to virus-specific polypeptides extracted from most of the viral isolates tested (Fig. 4B). A polypeptide of circa 38 kDa was detected in extracts from FC/2, LR-A, LR-B, LR 102, and LR 105 (Fig. 4B, lanes 1-4, 9) and another polypeptide of circa 24 kDa can be seen in extracts from LR 102, RSI, SJ, and CB100 (Fig. 4B, lanes 4-8); The same antiserum reacted with polypeptides of circa 43 kDa from LR 109, LR 105, LR

101, and LR-C (Fig. 4B, lanes 8-11). A polypeptide (circa 30 kDa) was also seen in some extracts (Fig. 4B, lanes 1-14); it may be similar to one of the doublet bands seen in Fig. 2C and 4C. The PA prepared with proteins from RG healthy grapevine bound to presumably plant polypeptides from healthy (Fig. 3C, lane 14) and infected grapevine material (Fig. 4C, lanes 1-13). The results of the antibody reactivity of the different viral isolates are summarized in Table 4.

Virus elimination by apical meristem culture. The 38- and 24-kDa polypeptides associated with GLRaV-1 and RSI were not present in explants propagated from regenerated apical meristems excised from the GLRaV-1 infected FC/2 cultivar and RSI when analyzed using WB with GLRaV-1 and RSI-24 kDa PA (not shown). In addition, the RG PA detected the same

polypeptides seen in Figure 4C. The regenerated grapevine meristems have been acclimated and are presently grown in our research greenhouse to be used as negative controls in future experiments. All of the regenerated meristem cultures and greenhouse-grown plants were free of virus-specific polypeptides. These grapevine selections were designated FC/2-vf and Malbec-vf (Table 4).

DISCUSSION

Several leafroll and corky bark associated virus isolates were serologically characterized using ELISA and the WB assay. The ELISA sensitivity of GLRaV-2b MAB (not shown) and GLRaV-4 PA was low, and the GCBaV PA developed unacceptable non-specific ELISA reactivities (not shown). The later is consistent with the number of host bands observed in WB using GCBaV PA. The ELISA and WB results obtained question the utility of performing cross-absorption of PA to decrease their reactivity with host proteins. The low specificity and sensitivity of these antibodies combined with the uneven distribution of grapevine associated closteroviruses might explain the difficulty encountered in detecting these viruses using ELISA (18).

ELISA characterization of virus infections using PA can be complicated by reactions with host proteins as well as uncharacterized mixed virus infections. Our results suggest that the cultivar CA-5 used by Boscia et al. (2) to prepare GLRaV-2 US PA was also infected with other viruses. A doublet protein band of circa 37 to 38 kDa was consistently observed in WB using the GLRaV-2 US PA and the LR 102 viral isolate. The 38-kDa polypeptide from the doublet was identified as the GLRaV-1 capsid protein based on its reactivity with GLRaV-1 MAB and PA. The faster migrating polypeptide did not react with GLRaV-1 MAB or any of the other antisera tested.

In our study, the GLRaV-2 US PA (2) also showed weak reactivity with the 36-kDa polypeptide associated with GLRaV-4 (11) and the 24-kDa polypeptide reported in association with GLRaV-2 (26). GLRaV-2 FR PA (26) showed reactivity with the 24-kDa polypeptide and no reactivity to other viral specific polypeptides (Fig. 2D). These data suggest that the GLRaV-2 US PA reacts predominantly with higher molecular weight proteins, and the 24-kDa polypeptide was not detected in earlier experiments (2) due to the weak reactivity of the GLRaV-2 US PA to this polypeptide.

Our results are in partial agreement with Boscia et al. (1), who reported that the GLRaV-2 US PA reacted prevalently with GLRaV-1 isolates and weakly with GLRaV-2 isolates. Apparently, the European study (1) using ISEM did not include any American GLRaV isolates. Their inability to detect GLRaV-4 reactivity or to

Table 4. Western blot reactivity of virus isolates

Isolate	Antibodies	Polypeptide size
FC/2	GLRaV-1 ^a , GLRaV-2 US ^b	38 kDa
LR A	GLRaV-1 ^a , GLRaV-2 US ^b	38 kDa
LR B	GLRaV-1 ^a , GLRaV-2 US ^b	38 kDa
	GLRaV-1 ^a , GLRaV-2FR ^b , GLRaV-2 US ^b ,	
LR 102	GLRaV-2b ^c , GCBaV ^b , RSI ^b	38, 37, 24 kDa
LR 105	GLRaV-1 ^a , GLRaV-2 US ^b , GLRaV-3 ^a	38, 43 kDa
RSI	GCBaV ^b , GLRaV-2 FR ^b , GLRaV-2b ^c , RSI ^b	24 kDa
CB100	GCBaV ^b , GLRaV-2 FR ^b , GLRaV-2b ^c , RSI ^b	24 kDa
SJ	GCBaV ^b , GLRaV-2 FR ^b , GLRaV-2b ^c , RSI ^b	24 kDa
MBF	GCBaV ^b , GLRaV-2 FR ^b , GLRaV-2b ^c , RSI ^b	24 kDa
PS	GCBaV ^b , GLRaV-2 FR ^b , GLRaV-2b ^c , RSI ^b	24 kDa
PS3	GCBaV ^b , GLRaV-2 FR ^b , GLRaV-2b ^c , RSI ^b , GLRaV-3 ^a	43, 24 kDa
LR 101	GCBaV ^b , GLRaV-2 FR ^b , GLRaV-2b ^c , RSI ^b , GLRaV-3 ^a	43, 24 kDa
LR C	GCBaV ^b , GLRaV-2 FR ^b , GLRaV-2b ^c , RSI ^b , GLRaV-3 ^a	43, 24 kDa
LR 109	GCBaV ^b , GLRaV-2 FR ^b , GLRaV-2b ^c , RSI ^b , GLRaV-3 ^a	43, 24 kDa
LR D	GLRaV-2 US ^b , GLRaV-4 ^b	36 kDa
LR 106	GLRaV-2 US ^b , GLRaV-4 ^b	36 kDa
Healthy ^d	healthy ^b , GCBaV ^b	No viral specific proteins

^a Monoclonal or polyclonal antibodies.

^b Polyclonal antibodies.

^c Monoclonal antibodies.

^d Couderc 3309, Kober 5BB, Riparia Gloire, Malbec-vf, FC/2-vf.

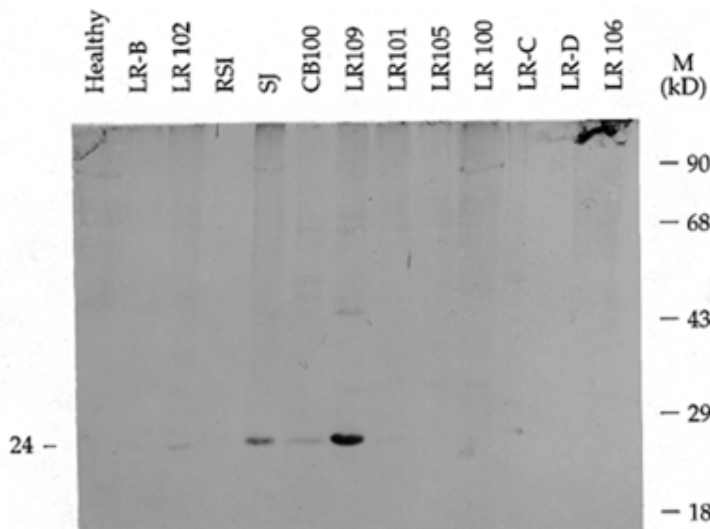


Fig. 3. Western blot assay of protein extracts using GLRaV-2 FR PA. The grapevine cultivars or isolates analyzed are indicated on the top of each lane. The molecular weight of size markers and the viral specific proteins are indicated on the sides of each blot.

discover the 37- to 38-kDa doublet might have been due to the limited amount of virus isolates tested.

We have recently developed MAb to the 37-kDa polypeptide (15,16). The MAb reacted to native and denatured conformations of the 37-kDa polypeptide associated with LR 102. Because the 37-kDa polypeptide did not react with the GLRaV-7 PA developed in Italy (3), which also contains a 37-kDa capsid protein, we have tentatively named our newly uncharacterized grapevine associated virus GLRaV-8 (data not shown). The new MAbs are presently being utilized to determine how widespread this virus strain is present in the United States (J. Monis and R. K. Bestwick, unpublished results).

Although the isolates designated LR-A and FC/2 reacted to GLRaV-1 MAb and GLRaV-2 US PA in ELISA, the WB assay data showed that these isolates are solely infected with GLRaV-1. These results show that when ELISA uses polyclonal antiserum raised against virus mixtures, results must be interpreted with caution.

The WB assay allows virus isolates to be distinguished on the basis of reactivity to different antisera. Because it is possible to correlate the presence of immuno-specific polypeptide bands with specific virus isolates, this assay confirms infection with different GLRaVs and GCBaV. Furthermore, the RG PA allowed the discrimination between viral and non-viral polypeptides. For example, host polypeptide bands were observed in the blots analyzed with RG antibodies as well as the blots analyzed with GCBaV (20) and Black Spanish polyclonal antisera (21). In addition, a host polypeptide of about 68 kDa was observed in healthy and virus-infected material analyzed with the RG PA and GLRaV-2 US PA (2). The same host proteins were observed in extracts from vines from which the virus was eliminated using the apical meristem tissue culture method.

Using the WB assay, the RSI-24 kDa PA, GLRaV-2 FR PA, GLRaV-2b MAb, and GCBaV PA all detected a 24-kDa viral-specific polypeptide in RSI, SJ, CB 100, LR 101, and LR 109 isolates, suggesting that the viruses with 24 kDa molecular weight coat proteins are serologically related. Others have reported the possible serological relationship between GLRaV-2, -2b and GCBaV (8,21). Recently, Boscia et al. (1) suggested that GCBaV and GLRaV-2b are identical to the GLRaV-2 isolate identified in France. Sequence analyses of the nucleic acid of these virus isolates will determine how these virus isolates are related.

A number of field-grown vines infected with viruses with a coat protein of approximately 24 kDa were asymptomatic when grafted onto the AXR-1 and St. George rootstocks. When these virus-infected scions were grafted onto phylloxera-resistant cultivars such as 5C, SO4, Cd

3309, or RG, severe rootstock-scion incompatibility symptoms developed (J. Monis and R. K. Bestwick, unpublished). These observations show the importance of correct virus identification prior to grafting of scion material. In our hands, the only reliable rapid method to detect the 24-kDa polypeptide associated with RSI is the WB assay.

We have shown that mixed infections of grapevine associated closteroviruses are common. Corky bark-, RSI-, and GLRaV-

2b-specific polypeptides were detected in plants infected with GLRaV-2, -3, GCBaV, and vines showing RSI. GLRaV-1 was found in vines infected with GLRaV-3. However, vines with apparent single infections of GLRaV-1 (FC/2, LR-A), GLRaV-3 (CS/8; 18), and GLRaV-4 (LR-D, LR 106) were also identified. Graft-inoculation studies between single and multiple infections will provide information on the interactions between these viruses with respect to symptom expression and disease devel-

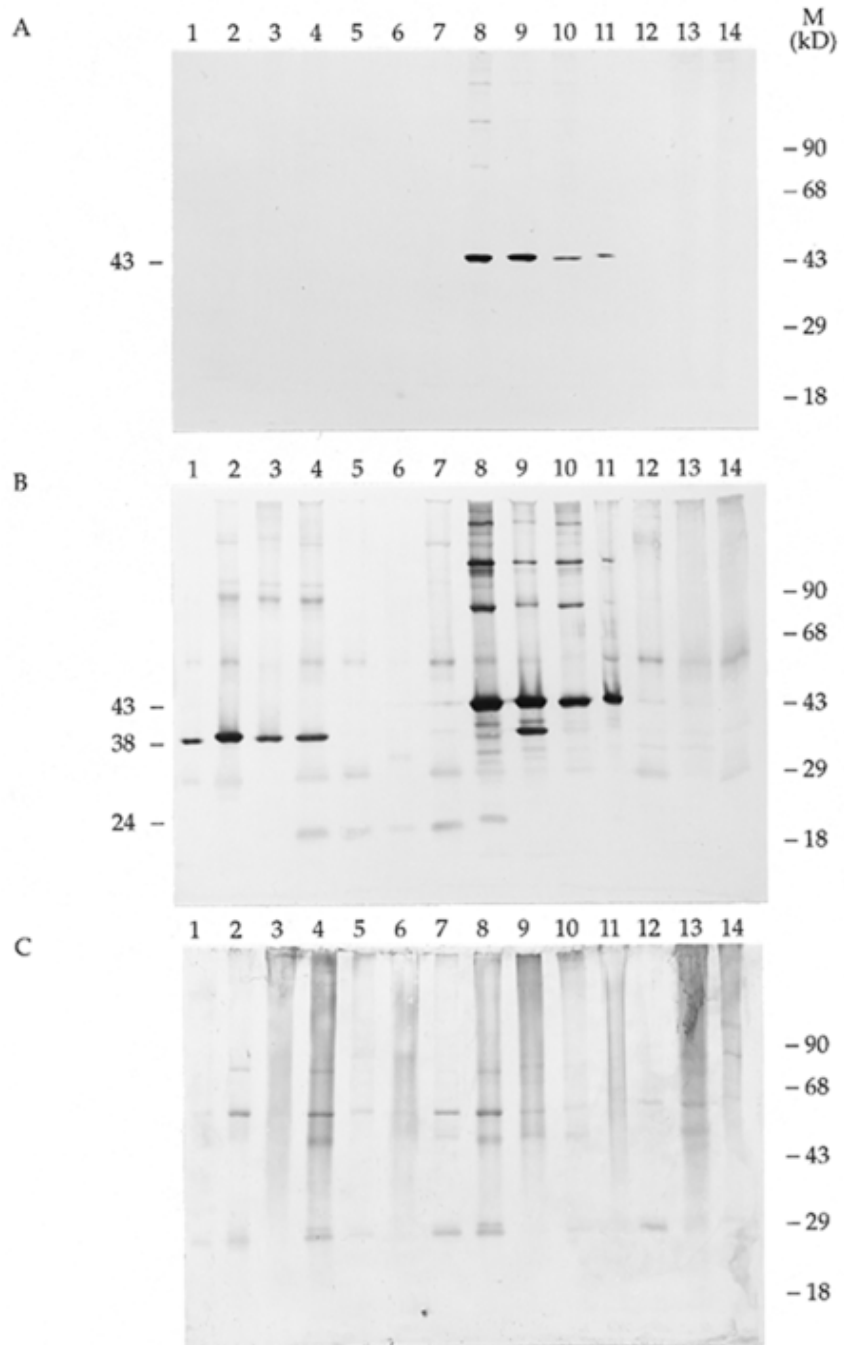


Fig. 4. Western blot assay of protein extracts using the following: GLRaV-3 MAb (A); Black Spanish PA (B); healthy grape PA (C). The grapevine cultivars or isolates analyzed are: lanes 1, FC/2; lanes 2, LR-A; lanes 3, LR-B; lanes 4, LR 102; lanes 5, RSI; lanes 6, SJ; lanes 7, CB 100; lanes 8, LR 109; lanes 9, LR 105; lanes 10, LR 101; lanes 11, LR-C; lanes 12, LR-D; lanes 13, LR 106; lanes 14, healthy. The molecular weight of size markers and the viral specific proteins are indicated on the sides of each blot.

opment. It is possible that certain virus combinations might intensify symptom and disease development.

Although the WB assay is not a new technique, it has not been used to diagnose grapevine viruses. The WB assay is routinely used in our laboratory to confirm questionable ELISA results and to detect viruses associated with RSI. The disadvantages of the WB assay are the requirements of preparing concentrated virus extracts and using characterized viral isolates and antibodies. We believe that these disadvantages are outweighed by the reliable detection of viruses that otherwise would remain undetected until an infected scion is grafted onto an incompatible rootstock cultivar.

Future research will focus on the molecular and serological characterization of the described viruses. Sequence analyses of the viral nucleic acid and reactivity to panels of monoclonal antibodies will further elucidate the relatedness of these grapevine associated closteroviruses. For example, we will be able to determine if closterovirus-like particles with coat protein subunits in the range of 24 to 43 kDa should be grouped within the closterovirus group or with other groups, based on refined molecular and serological characteristics. The information obtained from molecular studies should also allow the rationalization of the nomenclature of the viral isolates and antisera used for the diagnosis of grapevine rootstock scion incompatibility, leafroll, and corky bark diseases.

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