

Information pest: Grapevine viruses

Grapevines are affected by many viral diseases; the most harmful and widespread ones are fanleaf degeneration, leafroll, rugose wood, and fleck. The most important strategy to control viral diseases in grapes is preventive and consists of planting virus-free vines during vineyard establishment. The European regulation for grapevine plant material propagation (Directive 2005/43/CE 23/06/2005), included the certification as plant viruses-free for: *Grapevine fanleaf virus* and *Arabis mosaic virus* (infectious degeneration), *Grapevine leaf-roll associated virus-1* and -3, *Grapevine fleck virus* (for rootstock only). In Italy, *Grapevine virus A* has been added for the grapevine certification.

Introduction

The Multiplex End-Point PCR 7 grapevine viruses set has been developed by Qualiplante based on Gambino and Gribaudo, 2006. A verification was performed by Qualiplante (data not published) and the performance characteristics of the set are the same as the original publication. This PCR set allow to detect and to distinguish *Arabis mosaic virus* (ArMV), *Grapevine fanleaf virus* (GFLV), *Grapevine leafroll-associated virus 1, -2 and -3* (GLRaV-1, -2 and -3), *Grapevine fleck virus* (GFKV) and *Grapevine virus A* (GVA).

This method was compared to ELISA method (3 different reagents) during Arnadia Italian project on 122 infected grapevine samples tested by 13 laboratories. The results of the ring-test were presented during the 17th Congress of ICVG, Davis, California, in 2012.

This product should be used only for research purposes.

Intended use

The PCR set is validated for the detection of ArMV, GFLV, GLRaV-1, GLRaV-2, GLRaV-3, GFKV and GVA in Two-Step Multiplex RT-PCR. Suitable tissues are grapevine leaves and bark scrapings from dormant canes.

Set format and content

Two sets are available for 24 and 96 tests.

| Article N° | Product name |
|------------|---|
| 77VV--P2 | PCR ArMV, GFLV, GFKV, GLRaV-1, -2, -3, GVA set 24 |
| 77VV--P9 | PCR ArMV, GFLV, GFKV, GLRaV-1, -2, -3, GVA set 96 |

| Content | set 24 | set 96 |
|--------------------|---------------------------|-----------------------------|
| RT Master Mix | 24 tests 77VV--P2-RTM- | 2x48 tests 77VV--P9-RTM- |
| RT-Enzyme | 24 tests 77VV--P2-RT- | 96 tests 77VV--P9-RT- |
| Direct Master Mix | 24 tests 77VV--P2-DM- | 2x48 tests 77VV--P9-DM- |
| Positive Control 1 | 3 tests 77VV--P2-PC1- | 8 tests 77VV--P9-PC1- |
| Positive Control 2 | 3 tests 77VV--P2-PC2- | 8 tests 77VV--P9-PC2- |
| Negative Control | 3 tests 77VV--P2-NC- | 8 tests 77VV--P9-NC- |

Storage conditions

This set can be shipped at room temperature but upon receipt it should be stored immediately at the recommended storage temperature: **from -30 ° C to -10 ° C**.

Avoid prolonged exposure to light and repeated freeze and thaw cycles.

Shelf life

If the set is correctly stored, at constant-temperature freezer, its performance is guaranteed until the expiration date indicated on the tubes label.

Materials and equipment (not provided)

- RNA extraction tools and reagents
- Nuclease-free filter tips and micropipettes
- Optical grade nuclease-free tubes/plate
- Disposable latex or vinyl gloves
- DNA ladder and loading-dye buffer
- PCR thermal cycler
- Agarose gel reagents and apparatus

Nucleic acids extraction

Extract RNA from samples according to your usual protocol. Qualiplante recommend you to use RNeasy Plant Mini kit from Qiagen (Ref. 74904) according to MacKenzie protocol (MacKenzie D.J., McLean M.A., Mukerji S., Green M., 1997. Improved RNA extraction from woody plants for the detection of viral pathogens by reverse transcription-polymerase chain reaction. Plant disease 81:222-226).

REVERSE TRANSCRIPTION STEP

Preparation of the Master Mix

- Slowly thaw **RT Master Mix** and the **RT Enzyme** by placing it on ice or at 4°C.
- Shake briefly **RT Master Mix** and the **RT Enzyme** and spin down the liquid.
- In a new tube called **RevT master mix**, mix 14,0 µl of **RT Master Mix** and 1,0 µl of **RT-Enzyme** per reaction. This step is not necessary for **Positive Controls** and **Negative Control** due to the fact that they are under DNA form.

| Example: | 1 rxn | 10 rxns |
|----------------------|---------|----------|
| RT Master Mix | 14,0 µl | 140,0 µl |
| RT-enzyme | 1,0 µl | 10,0 µl |

- Store the **RevT master mix** by placing it on ice.

Reaction set up

- Shake briefly the **RevT master mix** and spin down the liquid.
- Add 15 µl of **RevT master mix** (without RNA template) to each PCR tubes or wells of an optical-grade PCR plate.
- Add 5 µl of RNA template to the **RevT master mix**.

| Components | Volume/PCR tube or well |
|---------------------------------|-------------------------|
| RNA template | 5 µl |
| RevT master mix | 15 µl |
| Total Volume / PCR tube or well | 20 µl |

In order to confirm the absence of any reagent's contamination, we strongly recommend including a no-template control (e.g. DEPC water) in the assay.

Run and thermal cycling

- Seal carefully the PCR tubes or PCR plate. Centrifuge briefly to collect components at the bottom of the PCR tubes or wells of the plate. Protect from light before thermocycling.
- Load the PCR tubes or plate into the PCR thermal cycler and follow the thermal cycling below:

| Steps | Temp (°C) | Time | Cycle(s) |
|-----------------------|-----------|--------|----------|
| Primer annealing | 25°C | 7 min | 1 |
| Reverse transcription | 50°C | 45 min | 1 |
| Enzyme inactivation | 85°C | 5 min | 1 |
| Storage | 4°C | ∞ | - |

From this step, samples are under cDNA format.

MULTIPLEX END-POINT PCR

Reaction set up

- Slowly thaw **Direct Master Mix** by placing it on ice or at 4°C.
- Shake briefly **Direct Master Mix** and spin down the liquid.
- Add 23 µl of **Direct Master Mix** (without cDNA template) to each PCR tubes or wells of an optical-grade PCR plate.
- Add 2 µl of cDNA template to the **Direct Master Mix**. Do not forget to prepare a PCR tube or well of an optical-grade PCR plate for the **Positive Controls 1 and 2** and the **Negative Control**.

| Components | Volume/PCR tube or well |
|--|-------------------------|
| cDNA template (*) or Positive control 1 or Positive control 2 or Negative control | 2 µl |
| Direct Master Mix | 23 µl |
| Total Volume / PCR tube or well | 25 µl |

(*) See the section "Reverse Transcription Step"

Run and thermal cycling

- Seal carefully the PCR tubes or PCR plate. Centrifuge briefly to collect components at the bottom of the PCR tubes or wells of the plate. Protect from light before thermocycling.
- Load the PCR tubes or plate into the PCR thermal cycler and follow the thermal cycling below:

| Steps | Temp (°C) | Time | Cycle(s) |
|----------------------|-----------|--------------|----------|
| Initial denaturation | 95°C | 15 min | 1 |
| Denturation | 95°C | 30 sec | 35 |
| Annealing | 55°C | 1 min 30 sec | |
| Elongation | 72°C | 1 min 30 sec | |
| Final elongation | 72°C | 10 min | 1 |
| Storage | 4°C | ∞ | - |

Agarose gel electrophoresis

Prepare an agarose gel at **2-2,8% w/v in 0,5X-TAE buffer**.

Gel loading:

- load the DNA ladder (for example 1'000 bp DNA step ladder).
- load 10 µl of PCR products from the previous step adding the loading dye buffer (*not provided in the set*).

Run: run the gel electrophoresis for 40 minutes to 1 hour at 80V.

Results analysis

ANALYSIS VALIDATION

The different DNA fragment size obtained are specific for each virus and are indicated in the table 1.

| Target | Fragment size (bp) | PC 1 | PC 2 | NC |
|---------|--------------------|------|------|----|
| IC | 844 | X | X | X |
| GLRaV-2 | 543 | X | - | - |
| ArMV | 416 | X | - | - |
| GLRaV-3 | 336 | X | X | - |
| GVA | 272 | - | X | - |
| GLRaV-1 | 232 | X | - | - |
| GFkV | 179 | - | X | - |
| GFLV | 118 | X | - | - |

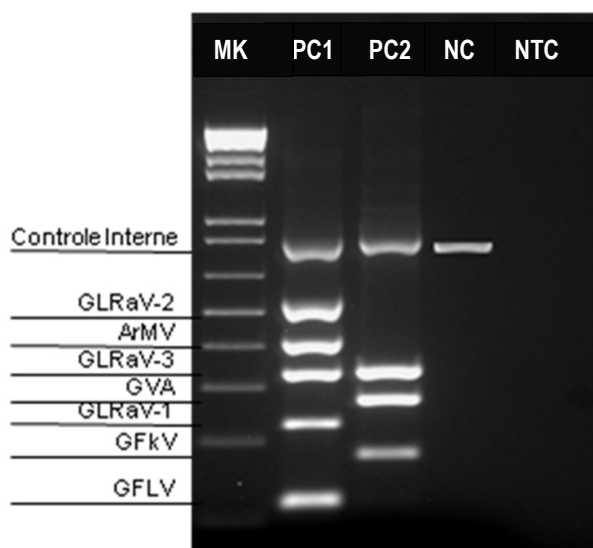
IC: Internal Control / PC: positive control / NC: negative control

A DNA fragment of 350 bp can appear in the **Positive Control 1** but is not applicable with this PCR set.

The analysis is validated when:

- ✓ 7 DNA fragments of 844 bp (Internal Control), 543 bp (GLRaV-2), 416 bp (ArMV), 336 bp (GLRaV-3), 232 bp (GLRaV-1) and 118 bp (GFLV) are visible in the **Positive Control 1** lane.
- ✓ 4 DNA fragments of 844 bp (Internal Control), 336 bp (GLRaV-3), 272 bp (GVA) and 179 bp (GFkV) are visible in the **Positive Control 2** lane.
- ✓ Only one DNA fragment of 844 bp is visible in the **Negative Control** lane corresponding to the Internal Control.
- ✓ All lanes must contain a DNA fragment of 844 bp in order to validate the nucleic acids extraction and the amplification.

The picture below represents a 0,5X-TAE 2,5% agarose gel showing the cDNA grapevine viruses:



MK: DNA MW ladder - **PC1:** **Positive control 1** including amplicons relative to the Internal Control, GLRaV-2, ArMV, GLRaV-3, GLRaV-1 and GFLV - **PC2:** **Positive control 2** including amplicons relative to the Internal Control, GLRaV-3, GVA and GFkV - **NC:** **Negative control** including the Internal Control - **NTC:** no template control (nuclease free water)

RESULTS INTERPRETATION

In order to determinate the infection status of a sample, compare the size of DNA fragment(s) obtained in the sample lane with those corresponding to the **Positive Controls** and **Negative Control**.

A sample is declared as infected when the DNA fragment corresponds to the specific amplicon for one (or more) virus(es).

Special handling instructions

This set was designed to be used by laboratory staff trained to follow the usual molecular biology precautions. Always perform the tests in a nuclease-free work environment. Always wear gloves when handling samples containing DNA/RNA and the components of the set. Do not touch any set components with an ungloved hand. Use appropriate laboratory disposable parts. Use nuclease-free tubes and filter tips to avoid degradation and cross-contamination. Do not use components from sets with different batch numbers in the same test procedure. Do not interchange reagents with other sets. To avoid cross-contamination, use separate rooms for (a) nucleic acids extraction, (b) preparation of the Master Mix and (c) amplification. To avoid cross-contamination and obtain reliable results, it is essential to strictly follow the protocol in this manual. Avoid unnecessary freeze-thaw cycles of the set components. Do not use reagents after their expiration date.

Troubleshooting

Post-PCR data analysis shows no amplification, or amplification plots look grossly abnormal:

| Possible causes | Corrective actions |
|--|---|
| Evaporation of the sample due to inadequate sealing of the plate | Repeat the test using the appropriate tools to seal correctly the plate |
| Consumables are not appropriate for the method | Repeat the test using consumables recommended by the thermal cycler supplier |
| The quality of nucleic acid extracted is low | Repeat the extraction step. Ensure that the method of extraction has been performed correctly. In any doubt, please, contact us |
| Abnormal amplification | Centrifuge the plate briefly to spin down the contents and eliminate any air bubbles |

No amplification reaction is observed in the positive control well, while other samples are positive:

| Possible causes | Corrective actions |
|---|--|
| The positive control provided with the set was not added into the reaction well | Repeat the test. If the problem persists, please, contact us |

An amplification plot is observed in the negative control well:

| Possible causes | Corrective actions |
|---|---|
| Contamination of the negative control or the Master Mix with target-positive nucleic acid | Repeat the test by applying appropriate quality procedures to prevent contamination. Seal the plate correctly |

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