

# Molecular Detection of *Rupestris stem pitting-associated virus* in Grapevines in Brazil

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## RESUMO

### Detecção por métodos moleculares do *Rupestris stem pitting-associated virus* em videiras no Brasil

RT-PCR foi utilizada para amplificar parte do gene da replicase do *Rupestris stem pitting-associated virus* (RSPaV) a partir de

videiras (*Vitis* spp.) no Brasil. O segmento amplificado foi clonado e seqüenciado e, por comparação da seqüência de nucleotídeos e dos aminoácidos deduzidos, verificou-se que correspondiam, respectivamente, em 88% e 94% com as de isolados do RSPaV de outros países.

*Rupestris stem pitting* (RSP), a component of the rugose wood complex, is one of the most widespread graft-transmissible grapevines (*Vitis spp.*) virus. It was detected in Brazil late in the 60's (Kuniyuki. Fitopatol. Bras. 5:137, 1972). The RSP is characterized by the presence of small pits in the woody cylinder below the point of inoculation by chip budding on *Vitis rupestris* Scheele cv. St. George ('du Lot') (Goheen. Compendium of Grape Diseases. 1988. p.53). A virus named *Rupestris stem pitting-associated virus* (RSPaV), *Foveavirus* genus, has been associated with the disease (Zhang *et al.* Phytopathology 88:1231, 1998; Meng *et al.* Eur. J. Plant Pathol. 105:191, 1999). This work reports the molecular detection of RSPaV in grapevines by RT/PCR, nucleotide sequence analysis and a non-isotopic cDNA probe specific to RSPaV. The RNA was extracted from petioles and used in the RT-PCR reaction with RSPaV-specific primers (Zhang *et al.* Phytopathology 88:1231, 1998). The amplified fragment was cloned and sequenced. The nucleotide (88%) and deduced amino acid (94%) sequences of the gene fragment (831 pb; Figure 1A) showed high homologies with those of two other RSPaV isolates (GenBank AF6278; AF057136). A digoxigenin-labeled cDNA probe was generated by PCR and used to positively detect RSPaV in RNA extracted from varieties of *V. vinifera* L. (Figure 1B). These results confirm the presence of RSPaV in grapevines that had indexed positive for RSP on *V. rupestris* 'St. George' in Brazil. The RT-PCR technique together with the cDNA probe will be a useful procedure for rapidly detecting the RSP disease in grapevines.

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**A**

	10	20	30	40	50	60
001	AGCATGCTCTTTGGCAACTGCAGTGATCCCCCAGAAGTGTGGCTCTCCTGAAAGT					
	S M L L A T A V I P P E V L V G S P E S					020
061	CTAAATCCTTGGGCTACCAGTACAGAATTAGTGATAATCACTGCTCTTCGCACCTGAT					
	L N P W A Y Q Y R I S D N Q L L F A P D					040
121	GGTAATTGGAGTGAATGATTCACAGCCTTTGTGCATGCAGATACCTACTTAAGGCTAGA					
	G N W S E M Y S Q P L S C R Y L L K A R					060
181	TCCGTTGTTTGCCTGATGGTTCAGCTATTGACATCATCTTCAAAATTTAGC					
	S V V L P D G S R Y S V D I I H S K F S					080
241	CACCACTTGCTTAGCTTACCCCCATGGGCAATCTTTAGCTCAAACATGAGGTGCTTC					
	H H L L S F T P M G N L L A S N M R C F					100
301	TCTGGCTTTGATGCAATAGGCATAAAGGATCTGAACCTCTAAGCCGTGGCATGCACAGT					
	S G F D A I G I K D L E P L S R G M H S					120
361	TGTTTTCCAGTGATCATGATGTTGTACCAAATATATCTTACTTGAGGACCCCTCAA					
	C F P V H H D V V T K I Y L Y L R T L K					140
421	AAGCCAGACAAGGAGTCTGCAGAGGCAAAGCTTCGCAACTCATTGATAAGCCACAGGG					
	K P D K E S A E A K L R Q L I D K P T G					160
481	AGGGAGATAAAATTCATTGAAGATTTTCTCACTAGTTATAAGTTGTGGAAGGAGTGGT					
	R E I K F I E D F S S L V I S C G R S G					180
541	TCTTGTCTTATGCCCAACATTCTAAGTTGGTAAATCATTCTTTCFCCGAATGATGCCA					
	S L L M P N I S K L V I S F F C R M M P					200
601	AATGCACTTGCTAGGCTTTCTCCAATTTTCGGAGTGTCTCACTGGATTCATTTGTGTAT					
	N A L A R L S S N F R E C S L D S F V Y					220
661	TCACCTTGAGCCTTCAATTTTCAATTAATTTGGTGGATATCACTCCCGATTTCTTTGAG					
	S L E P F N F S I N L V D I T P D F F E					240
721	CATTATTTCTTTCTCTGTCTCAATGAGTTAATCGAGGAGATGTTGAAGAGGTCATG					
	H L F L F S C L N E L I E E N V E E V M					260
781	GACAACCTTTGGTTTGGACTCGGGATTTGCAATTCATCCGACAGAGGG					
	D N S W F G L G D L Q F N R Q R					276



**FIG. 1** - (A) Partial nucleotide (above) and deduced amino acid (below) sequences of *Rupestris stem pitting-associated virus* (RSPaV). (B) Dot-blot hybridization with RSPaV-specific probe. RNA extracted from infected plants of *Vitis vinifera* cultivars: (1) Itália (Biritiba Mirim, SP); (2) Itália (São Miguel Arcanjo, SP); and (3) Benitaka (Londrina, PR). RNA extracted from healthy plants of 'Itália' (4) and 'Benitaka' (5).

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