

Modulation of host gene expression during the initial steps of tomato infection  
by a potyvirus

Poliane Alfenas-Zerbini<sup>1\*</sup>, Francisco Murilo Zerbini<sup>1\*</sup>, Riani Moreira Neto<sup>1</sup>, Ivan  
G. Maia<sup>2</sup> e Júlio C.M. Cascardo<sup>3</sup>.

<sup>1</sup>Dep. de Fitopatologia/BIOAGRO, UF Viçosa, Viçosa, MG, 36570-000. <sup>2</sup>Dep.  
de Genética, UNESP, Botucatu, SP; <sup>3</sup>Dep. de Biologia e Genética Molecular,  
UESC, Ilhéus, BA. \*E-mail: [palfenas@ufv.br](mailto:palfenas@ufv.br); [zerbini@ufv.br](mailto:zerbini@ufv.br)

Plant defense responses against pathogens cause up- and downward shifts in gene expression. To identify differentially expressed genes in a plant-virus interaction, a subtractive library was constructed from tomato plants inoculated with *Pepper yellow mosaic virus* (PepYMV). Several genes were up- or down-regulated, including transcriptional regulators (e.g., SCARECROW and WRKY transcription factors), signaling proteins (e.g., CAX-interacting and SNF1 kinases), proteins involved in stress responses (e.g., Hsp90, DNA-J, TCTP) and ubiquitins. Differential expression of 42 genes was validated by macroarray analysis. Four genes (CRC9, CXIP4, SNF1 and TCTP) were validated by qRT-PCR. The genes encoding SNF1 and TCTP were selected for additional analyses. The kinetics of differential expression was analyzed by qRT-PCR. Expression of both TCTP and SNF1 was induced at 48 hpi, and remained high until 96 hpi. To verify whether the induction of TCTP and SNF1 is part of a general response to biotic stresses, tomatoes were inoculated with *Pseudomonas syringae* pv. *tomato*, *Alternaria solani*, *Meloidogyne incognita*, TMV, CMV and ToYSV, besides PepYMV. Leaves were collected at 0 and 72 hpi and expression levels of TCTP and SNF1 were analyzed by qRT-PCR. TCTP was induced only upon infection by PepYMV, while SNF1 was induced upon infection by PepYMV and ToYSV. Functional analysis of the role of TCTP, SNF1 and Sub $\alpha$  (alpha subunit of the 26S proteasome) is being investigated by virus-induced gene silencing (VIGS). Preliminary results suggest that silencing of all three genes renders the plants resistant to infection.

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