

## PROTEINASE ACTIVITY OF Tetranychus evansi AND Tetranychus urticae: EFFECTS OF INTERACTIONS AND ADAPTATIONS TO TOMATO-PLANT DEFENSES ATIVIDADE DAS PROTEASES DE Tetranychus evansi E Tetranychus urticae: EFEITOS DAS INTERAÇÕES E ADAPTAÇÕES À DEFESA DO TOMATEIRO

## <u>M.A. Solís-Vargas</u><sup>1,4</sup>, F.R. Ribeiro<sup>2,4</sup>, B. Amorim<sup>3,4</sup>, A.M.G. Bernardo<sup>1</sup>, M.A.G. Oliveira<sup>2,4</sup> & A. Pallini<sup>1</sup>

<sup>1</sup>Department of Entomology/Laboratory of Acarology, UFV; <sup>2</sup>Departament of Biochemistry and Molecular Biology, UFV; <sup>3</sup> Undergraduate Agronomy Course, UFV; <sup>4</sup>Laboratory of Enzymology Proteins and Peptides Biochemistry of the Biotechnology Institute Applied to Agricultural (BIOAGRO), UFV, Viçosa, MG, Brazil.

Tetranychus evansi is a spider mite pest that successfully suppresses salicylic and jasmonate acid pathway of tomato induced defenses, which made it a novel model of herbivore. By the other hand, Tetranychus urticae shows a normal model of inducing plant defenses. However, when T. evansi interacts with T. urticae, in the same plant, proteinase inhibitors (PI's) are induced. This work aimed to describe digestive proteinases activity related to these two species and PI's effects. Clean tomato plants where infested with a) only T. evansi or b) T. urticae and c) both species simultaneously, control consisted on non-infected plants. Proteinase inhibitors from plant samples and total proteinase, serine and cysteine proteinase activity from herbivores were determined by spectrophotometry. Oviposition rates over disc leaves from treatments were obtained to determine T. evansi performance. Only plants attacked by T. evansi did not increase levels of PI's. Meanwhile, T. urticae induced plant defenses. The total proteinase activity of T. evansi has a two fold increase when shared plants with T. urticae and a high PI concentration were induced, which can be associated to direct effects of PI's concentration on T. evansi performance. T. urticae presented the higher total proteinase activity compared with T. evansi and induced a high PI concentration. The total proteinase activity patterns are correlated to T. evansi and T. urticae trypsin-like activity. Adversely, quimotrypsin-like activity showed an opposite pattern to thus of trypsin-like proteases, suggesting a low quimotrypsin-like production when the trypsin-like production increased. It seems that the action of PI's in the digestive system stimulates trypsin-like proteases in detriment of quimotrypsin-like as response to toxic PI's effects. Quimotrypsinlike proteases are more active under low PI's concentrations and trypsin-like proteases are produced as a response to few available aminoacids and PI's antimetabolic effects. T. evansi PI's suppression could become an advantage for other herbivore species that can be reflected in proteinase activity. However, sharing food resources decreased T. evansi performance due to PI's antimetabolic effects.

Keywords: inhibitors, performance, sharing resources Financial support: CAPES, FAPEMIG, CNPq, OEA-GCUB, INCT-IPP