

november 16 to 18, 2010

Abstracts

Book





Topic: Transcriptomics and Proteomics PI: TP03

MOLECULAR, BEHAVIORAL AND ANATOMICAL SOPHISTICATION IN SPIDER WEBS: INSIGHTS FROM SPINNING GLAND RNA-SEQ EXPERIMENTS IN PRIMITIVE AND MODERN SPIDERS

Prosdocimi F¹, Bittencourt D³, Silva F R D², Motta P C⁴, Rech E²

¹Universidade Católica de Brasília

²EMBRAPA-CENARGEN

³EMBRAPA-Amazônia Ocidental

⁴Universidade de Brasília

RNA-seq experiments conducted in 454 sequencers were carried out to produce 87.000 short-reads representing the transcriptome of two spiders'spinning glands. We produced sequences from (i) Actinopus sp., a spider from the Mygalomorphae clade, and (ii) Gasteracantha cancriformis, an Orbicularia spider. Mygalomorphae spiders are known to retain a number of primitive morphological and behavioral characters. They use mixtures of a primitive web, soil, and plants only to cover a burrow they make on the ground for shelter and predation. On the other hand, Orbicularia spiders show a number of derivative spider's characters and they are capable to build different and complex silks used in a variety of situations. It is interesting to note that the complexity of web production, usage and behavior in these spiders is reflected both by (i) the variety of the repertoire of protein molecules (spidroins) they use to make their webs and (ii) the complexity of their anatomical spinning gland apparatus used to produce silk. Here we have first conducted a broad analysis of the spinning gland transcriptome in both spiders producing unigenes and categorizing annotated genes in biological functions. Then we started to analyze the number and variety of spider silk proteins and families found in different spider clades. We have shown that spiders using web only for a limited number of situations present a less sophisticated morphological spinning apparatus and produce a small repertoire of spidroin molecules. Phylogenetic analyses were conducted in the 3'region of spidroins and we try to relate (i) the evolution of silk protein families, (ii) the evolutionary complexification of silk production behavior and web usage, and (iii) the appearance of new specialized spinning glands along the evolution of specimens and clades in the Araneae order.

Supported by: CNPq, FAP-DF.