YOLK PROTEIN BIOSYNTHESIS AFTER dsc-4 KNOCKDOWN IN CAENORHABDITIS ELEGANS

Almenara, D.P.¹, Winter, C.E.¹

¹Departamento de Parasitologia, Instituto de Ciências Biomédicas, Universidade de São Paulo, São Paulo, Brazil.

The dsc-4 gene of Caenorhabditis elegans encodes an orthologue of the human Microsomal Triglyceride Transfer Protein (MTP). Mutations in the coding region of this gene affect worm aging processes. MTP is a member of the large lipid transfer protein (LLTP) superfamily, wich includes vitellogenins, insect apolipophorins and human apolipoprotein B (apoB). MTP mutations in man result in severe abetalipoproteinemia. It has been previously demonstrated that MTP promotes the secretion of Xenopus laevis vitellogenin A1 in cell cultures. Here we present an analysis of the dsc-4 gene role in vitellogenin biosynthesis in the nematode worm C. elegans. In C. elegans, double-stranded RNA (dsRNA) provided in the diet can be absorbed from the gut and distributed troughout the body, triggering RNA interference (RNAi) in different tissues. In order to knockdown dsc-4 expression a cDNA fragment corresponding to a partial C. elegans dsc-4 sequence was ligated into the feeding vector L4440, between its two inverted orientation T7 promoters, and transformed into E. coli HT115(DE3). These dsRNA-expressing bacteria were used to fed wild-type worms (strain N2) and transgenic worms (strain DH1033) expressing YP170B::GFP. Living worms, fed on dsRNA-containing bacteria, show YP170B::GFP accumulation in the enterocytes cytoplasm, suggesting that the vitellogenin transport from the gut cells to the pseudocoelom is compromised.

Supported by FAPESP/CNPq.