**TITLE:** SHIGA TOXIN-PRODUCING *Escherichia coli* ISOLATED FROM *Musca domestica* COLLECTED IN DAIRY FARM

AUTHORS: ALVES, T.S.<sup>1</sup>; LARA, G.H.B.<sup>2</sup>; RIBEIRO, M.G.<sup>2</sup>; LEITE, D.S.<sup>1</sup>

**INSTITUTION:** <sup>1</sup>INSTITUTO DE BIOLOGIA, UNIVERSIDADE ESTADUAL DE CAMPINAS -UNICAMP, CAMPINAS, SP (CAIXA POSTAL 6109, CEP 13083-862). <sup>2</sup>FACULDADE DE MEDICINA VETERINÁRIA E ZOOTECNIA, UNIVERSIDADE ESTADUAL PAULISTA "JÚLIO DE MESQUITA FILHO" – UNESP, BOTUCATU, SP (CAIXA POSTAL 560, CEP 18618-681).

## ABSTRACT:

Musca domestica species (Linnaeus, 1758) are known for their pronounced synanthropic behavior, being able to transit between contaminated (feces and garbage) and uncontaminated (food) sources, acting as vectors of several pathogens in the environment. Since healthy cattle is the main STEC reservoir, this study sought to investigate the presence of such strains in flies developing close to calves. Escherichia coli strains were isolated from the external surface of flies collected nearby the calves from a dairy farm in the city region of Botucatu, SP, Brazil, using a sweep net. The virulence factors stx1, stx2, eae, hlyA, K99 (F5), LT2 and STa were searched by PCR assay and the stx1 and/or stx2 positive strains were selected for the in vitro cytotoxicity assay in Vero cells. The antimicrobial susceptibility was determined by disk diffusion. Of 57 flies, 35.1% (20/57) were identified as Musca domestica and 64.9% (37/57) as other species of the Muscidae, Fanniidae, Calliphoridae, Chloropidae and Acalyptratae family. In total, 135 E. coli strains isolated, with 7/135 (5.2%) of strains from M. domestica showing a positive genotype for stx1 and cytotoxic effect in Vero cells but not the eae gene. The relative frequencies of the other virulence factors tested were LT2 (14.1%), hlyA (8.1%), K99 and STa (3.7%) and eae (2.9%). None isolate was positive for stx2. One STEC strain showed multidrug resistance ( $\beta$ -lactam, sulphonamide + folate pathway inhibitor, and tetracycline). Considering that *M. domestica* exhibits a transiting behavior between contaminated and uncontaminated environments, as well as the importance of STEC infection in humans, usually linked to contaminated food, the results indicate that *M. domestica* can carry strains of STEC in the environment, probably from cattle, and contribute to STEC transmission.

**Keywords:** flies, STEC, mechanical vectors, calves **Development Agency:** FAPESP (2015/15425-2).