In vitro and in vivo models to study biofilm formation by Streptococcus dysgalactiae subsp. dysgalactiae

<u>Cinthia Alves-Barroco</u>¹, Ana Maria Nunes Botelho², Catarina Roma-Rodrigues¹, Márcia Aparecida Guimarães², Marco Antonio Americo², Jayaraman Muthukumaran³, Teresa Santos-Silva³, Euvira Fortunato⁴, Rodrigo Martins⁴, B. Ferreira-Carvalho², Agnes M.S. Figueiredo^{2,*} and Alexandra R. Fernandes^{1,*}

Streptococcus dysgalactiae subsp. dysgalactiae (SDSD) is an important agent of bovine mastitis. This infection causes an inflammatory reaction of the udder tissue, being the most important disease in the dairy industry, due to treatment costs and loss in milk production and quality. Recurrent mastitis is often attributed to biofilm, which allows bacteria to survive and proliferate in hostile environments and resist antimicrobial therapy. Recently, we have described the in vitro biofilm formation by SDSD. Imaging and computer methods were used to characterize the in vitro formation of biofilms. In this work, we demonstrated the capability of SDSE to develop biofilm in vivo, using a murine animal model. For the in vivo assays, bovine SDSD was selected based on the ability to form strong (n=2), moderate (n=2) or weak biofilms (n=1). The expression profile of genes associated with biofilm formation was analyzed. Overall, our results showed an increased ability to accumulate biofilm on catheter surfaces implanted in mice compared with in vitro biofilm formation. In addition, increased biofilm accumulation on glass surfaces was observed after animal passage. An increase of expression of brpA-like (biofilm regulatory protein) and fbpA (fibronectin-binding protein A) genes was observed for all the tested SDSD isolates. However, the mRNA expression of htrA (serine protease) and sagA (Streptolysin S) homologs was more dramatic for SDSD isolates displaying weaker biofilm production. Because the expression of brpA-like correlated with in vitro and in vivo biofilm formation in SDSD, homology modeling was used to predict the 3D structure of BrpA-like protein. Additionally, using high throughput virtual screening and molecular docking, we selected five ligand molecules with strong binding affinity to the hydrophobic cleft of the protein, making them inhibitor candidates of BrpA-like protein from SDSD. These results warrant further investigations for the development of potential novel anti-biofilm drugs for SDSD.

¹UCIBIO, Departamento de Ciências da Vida, Faculdade de Ciências e Tecnologia, Universidade Nova de Lisboa, Caparica, Portugal.

²Instituto de Microbiologia Paulo de Góes, Universidade Federal do Rio de Janeiro, RJ, Brazil.

³ UCIBIO, Departamento de Química, Faculdade de Ciências e Tecnologia, Universidade Nova de Lisboa, Caparica, Portugal

⁴i3N/CENIMAT, Departamento de Ciência dos Materiais, Faculdade de Ciências e Tecnologia, Universidade Nova de Lisboa, Campus de Caparica, Caparica, Portugal

^{*}Corresponding authors - E-mails: ma.fernandes@fct.unl.pt (ARF); agnes@micro.ufrj.br (AMSF)