Role of glycans in *Helicobacter pylori* infection and gastric carcinogenesis

Nuno T. Marcos<sup>1</sup>, Ana Magalhães<sup>1</sup>, Maria J. Oliveira<sup>1</sup>, Ana S. Carvalho<sup>1</sup>, Tim Gilmartin<sup>3</sup>, Steven R. Head<sup>3</sup>, Céu Figueiredo<sup>1,2</sup>, Leonor David<sup>1,2</sup>, <u>Celso A. Reis<sup>1,2</sup></u>.

<sup>1</sup> Institute of Molecular Pathology and Immunology of the University of Porto (IPATIMUP), Portugal;<sup>2</sup> Medical Faculty, University of Porto, Portugal. <sup>3</sup> The Scripps Research Institute, La Jolla CA, USA

Chronic Helicobacter pylori infection is recognized as a cause of gastric diseases. *H. pylori* adhesion to gastric cells has been shown to be mediated by bacterial adhesins such as BabA and SabA. SabA binds the carbohydrate structure Sialyl-Lewis<sup>x</sup>, which is expressed in the gastric epithelium during H. pylori infection. It has been suggested that H. pylori modulates host cell alvcosvlation patterns for enhanced adhesion. Here, we evaluated changes in the glycosylation-related gene expression profile of a human gastric carcinoma cell line following *H. pylori* infection. We observed that *H. pylori* significantly alters expression of 168 of the 1031 human genes tested by microarray, and the extent of these alterations is associated with markers of virulence of the *H. pylori* strain. A highly pathogenic *H. pylori* strain altered expression of several genes involved in glycan biosynthesis, including the  $\beta$ 3GnT5, a GlcNAc-transferase essential for the biosynthesis of Lewis antigens. β3GnT5 induction was specific to infection with *H. pylori* strains carrying the *cag* pathogenicity island. Further,  $\beta$ 3GnT5 overexpression in human gastric carcinoma cell lines led to increased Sialyl-Lewis<sup>x</sup> expression and *H. pylori* adhesion. This study has identified a novel mechanism by which *H. pylori* modulates the biosynthesis of the SabA adhesin ligand in gastric cells, contributing to achieve a successful adhesion.

Supported by FCT POCI/QUI/56393/2004. A.M. supported by FCT (Ref. SFRH/BD/36339/2007). The Gene Microarray Core resources and collaborative efforts provided by CFG were funded by NIGMS - GM62116.