

# Modulation of Nrf2-dependent antioxidant response after infection with *Helicobacter pylori* may contribute to gastric cancer

Océane C.B. Martin<sup>1</sup>, Lornella Seeneevassen<sup>1</sup>, Sarah Bacon<sup>1</sup>, Elodie Sifré<sup>1</sup>, Astrid Ducournau<sup>2</sup>, Lucie Bruhl<sup>2</sup>, Philippe Lehours<sup>1,2</sup>, Christine Varon<sup>1</sup>

1-INSERM U1053, Bordeaux Research in Translational Oncology, University of Bordeaux, Bordeaux, France.  
2-CHU de Bordeaux, Bordeaux, France

## Background

- Gastric cancer (GC): 4<sup>th</sup> leading cause of mortality by cancer worldwide
- Major risk factor of GC: chronic infection with *Helicobacter pylori*
- *H. pylori* induces an epithelial-to-mesenchymal transition (EMT), first step of GC
- Nrf2: transcription factor, main regulator of cellular redox homeostasis

## Objectives

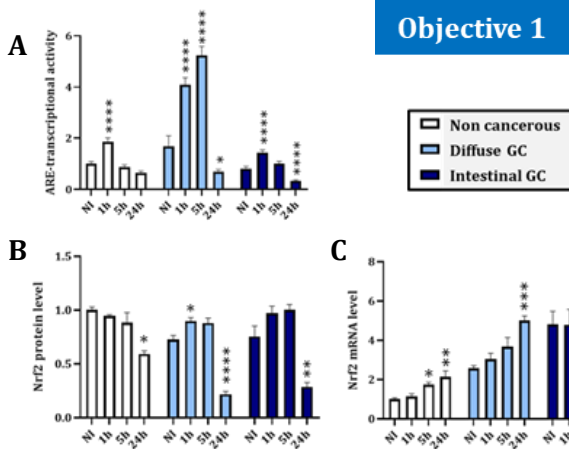
- 1) To assess the modulation of the Nrf2 pathway after infection with *H. pylori*
- 2) To decipher the role of Nrf2 in *H. pylori*-induced EMT

## Material & Methods

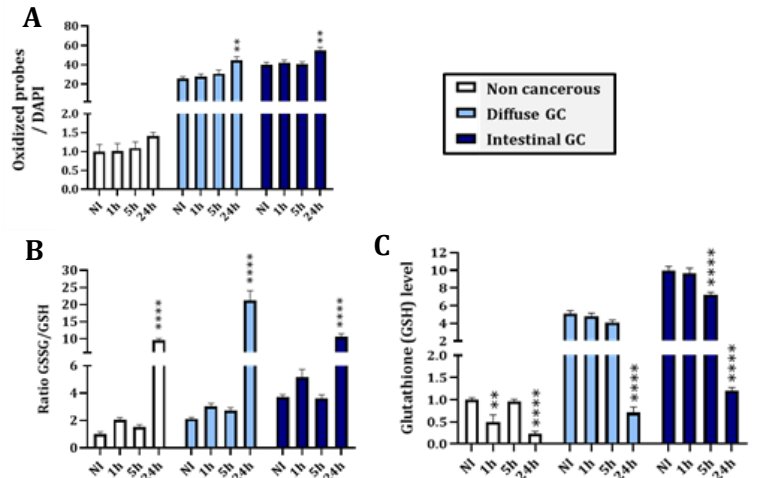
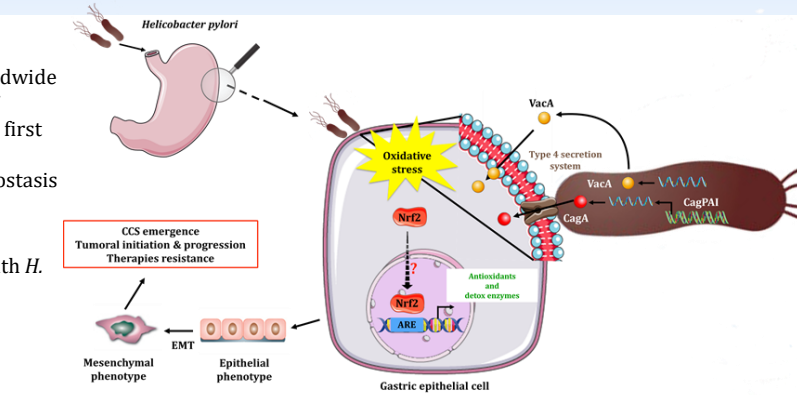
**Objective 1:** 3 gastric epithelial cell lines HFE-145 (non cancerous), AGS (cancerous diffuse subtype), MKN74 (cancerous intestinal subtype) co-cultured with *H. pylori* for 1, 5 or 24h

**Objective 2:** 4 gastric epithelial cell lines (AGS and MKN74) WT or Nrf2-KO by CRISPR-Cas9 co-cultured with *H. pylori* for 24h

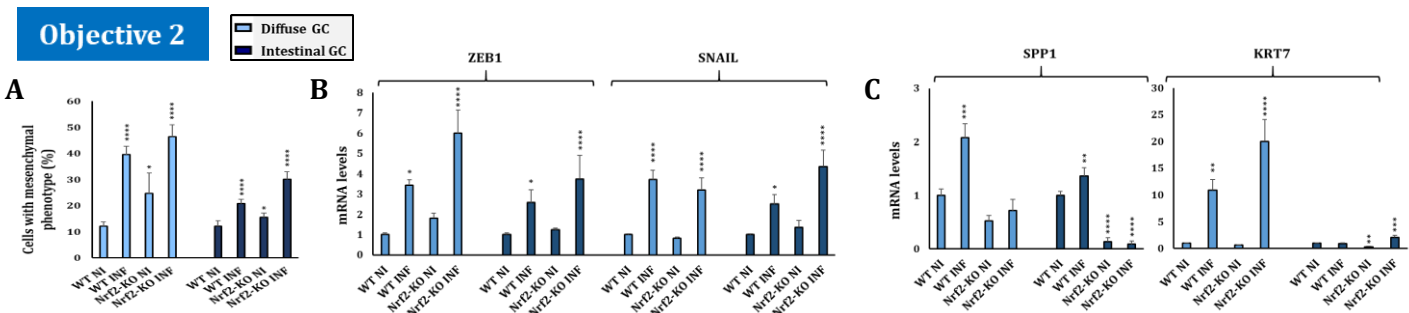
## Results



**Figure 1:** Non cancerous (white), cancerous diffuse subtype GC (light blue) and cancerous intestinal subtype GC (dark blue) cell lines were colcultured with *H. pylori* for 1, 5 and 24h. ARE-transcriptional activity (A) and Nrf2 protein (B) and mRNA (C) levels were assessed.



**Figure 2:** Non cancerous (white), cancerous diffuse subtype GC (light blue) and cancerous intestinal subtype GC (dark blue) cell lines were colcultured with *H. pylori* for 1, 5 and 24h. Intracellular level of ROS (A), ratio between oxidized glutathione (GSSG) and total glutathione (GSH) (B) and level of total glutathione (C) were assessed.



**Figure 3:** Cancerous diffuse subtype GC (light blue) and cancerous intestinal subtype GC (dark blue) cell lines either WT or Nrf2-KO were colcultured with *H. pylori* for 24h. Percentage of cells with a mesenchymal phenotype (A) and mRNA levels of mesenchymal (B) and epithelial (C) markers were assessed.

## Conclusions

**Objective 1:** Upon infection with *H. pylori*, Nrf2 pathway is activated in the early phase while inhibited after 24h, the late inhibition is associated with an increased oxidative stress and a depletion of glutathione.

**Objective 2:** Nrf2 seems to be a guardian of the cellular epithelial phenotype integrity and *H. pylori*-induced EMT is increased when cells are deleted for Nrf2

**Nrf2 could play a key role in *Helicobacter pylori*-induced gastric carcinogenesis**