

Bacterial molecule may be key to fighting stomach cancer

April 12 2021

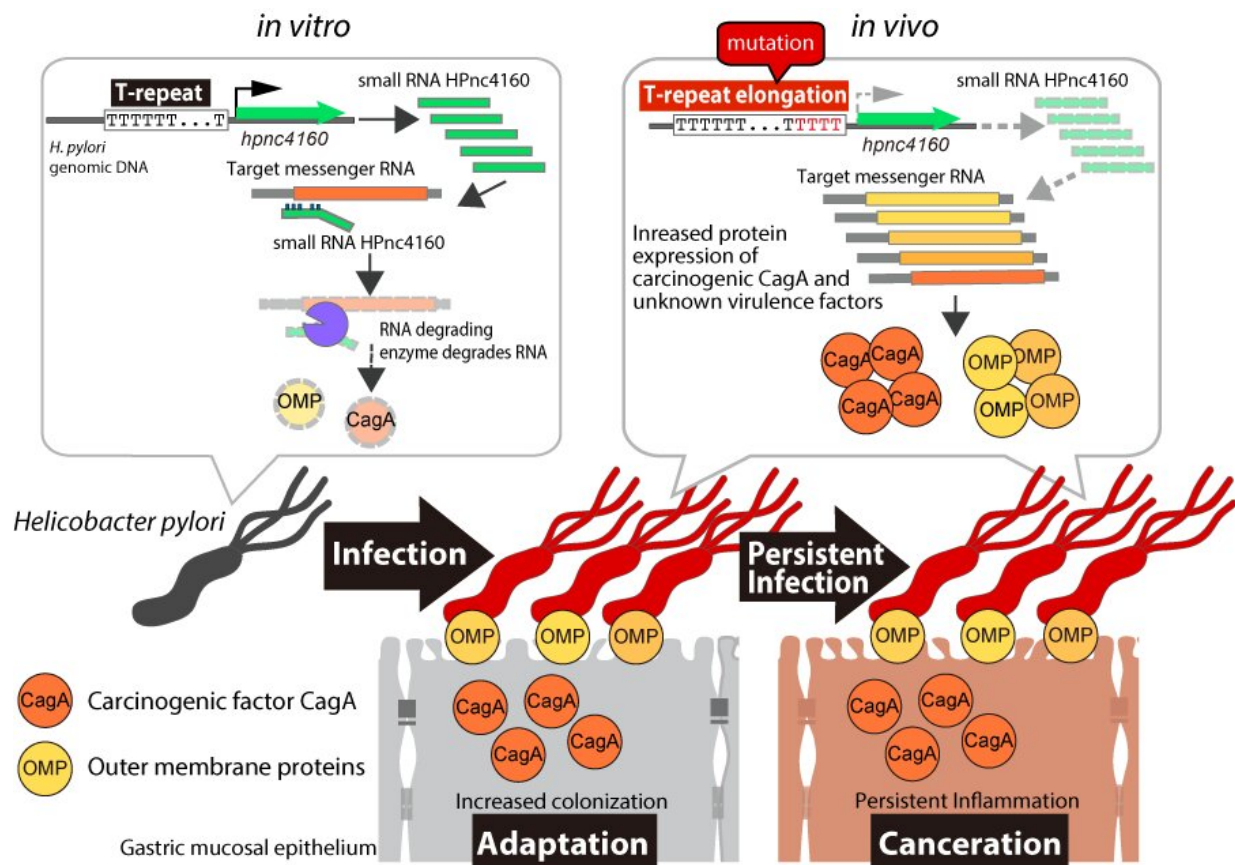


Figure 1. Persistent infection and, potentially, cancer development are regulated by *Helicobacter pylori* (*H. pylori*) small RNA HPnc4160: *H. pylori*'s small RNA HPnc4160 suppresses the expression of messenger RNAs of pathogenic factors. When *H. pylori* infects the gastric mucosa, a mutation is introduced into a continuous thymidine (T) sequence (T-repeat) located upstream of the gene encoding HPnc4160. Because the amount of HPnc4160 subsequently decreases, the number of pathogenic factors increase and *H. pylori* can easily infect the

gastric mucosa for a long period of time. Credit: Osaka University

Humans are exposed to many types of bacteria daily, the majority of which are harmless. However, some bacteria are pathogenic, which means they can cause disease. An extremely common pathogenic bacterial infection is *Helicobacter pylori* (*H. pylori*) in the stomach, where it can lead to chronic inflammation (gastritis), ulcers, and even cancer. A group of researchers from Osaka University have determined a specific molecular mechanism that *H. pylori* uses to adapt to growing in the human stomach for long periods of time.

In a report published in *Nature Communications*, this group found that a small RNA molecule called HPnc4160 plays a key role in how *H. pylori* invades the [stomach](#) and leads to disease.

Previous studies showed that HPnc4160 is conserved in different [strains](#) of *H. pylori*, suggesting that it has an important function. Yet, the amount of this molecule produced in the bacteria is variable among strains. Genetic analysis revealed that a specific part of the *H. pylori* gene, called the T-repeat region, varied in length from strain to strain. The researchers became interested in how this region affected HPnc4160 expression. Additionally, they were curious how these fluctuations would allow some strains to colonize the human stomach more efficiently than others.

"Slight differences in the genetic sequence of *H. pylori* can sometimes give certain strains advantages over others that allow it to grow better," says lead author of the study Ryo Kinoshita-Daitoku. "We were interested in the genetic and molecular reasons behind why particular strains are more pathogenic and can live in the stomach for decades, leading to cancer development."

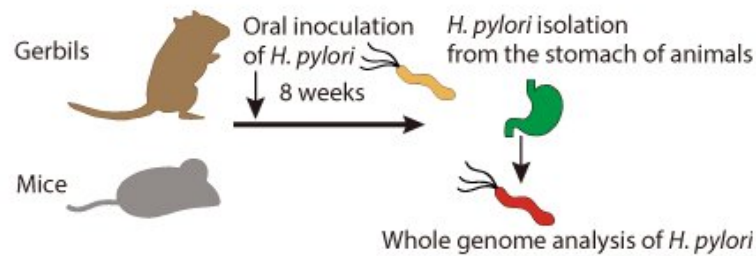


Figure 2. Experimental method: Eight weeks after infecting mice and gerbils with wild type *H. pylori*, we isolated *H. pylori* from each animal's stomach and performed a whole-genome analysis. Credit: Osaka University

To address these questions, the researchers infected gerbils and mice with wild type (normal) *H. pylori* for 8 weeks, then extracted and genetically characterized the mutant strains that appeared. They discovered that strains with a higher number of T-repeats had lower expression of HPnc4160.

"We found that the *H. pylori* strains with low HPnc4160 levels were more infectious," explains Hitomi Mimuro, senior author. "When this RNA was completely absent, the amounts of several bacterial pathogenic factors significantly increased and the strain showed stronger colonization within the rodent stomach."

One of the pathogenic factors was the bacterial protein CagA, which is known as an oncoprotein because it can promote cancer development.

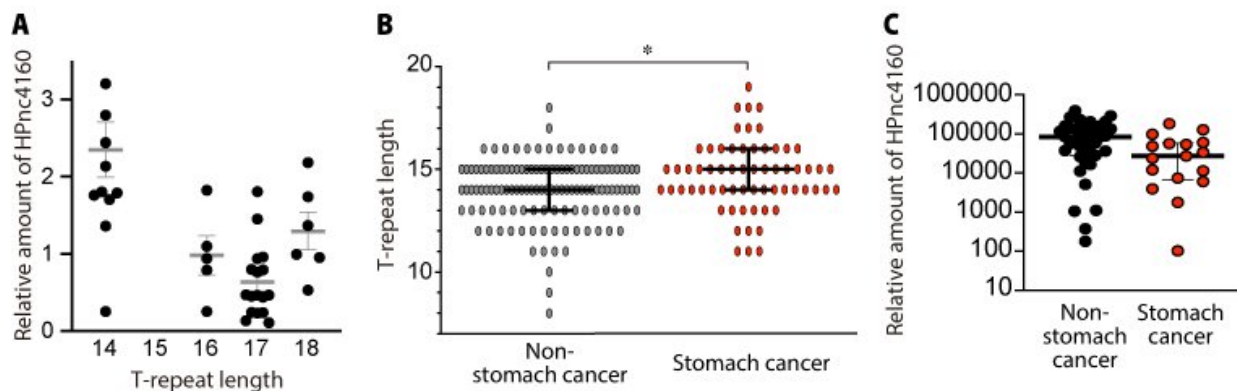


Figure 3. T-repeat length and HPnc4160 expression: (A) *H. pylori* isolated from the stomachs of gerbils contains a mutation that increases the T-repeat length in the region upstream of the small RNA HPnc4160 gene. *H. pylori* strains in the stomachs of patients with gastric cancer have (B) longer T-repeat lengths and (C) lower expression levels of HPnc4160 than *H. pylori* strains in the stomachs of

individuals without cancer. Credit: Osaka University

"Interestingly, we observed longer T-repeat regions, lower levels of HPnc4160, and higher levels of CagA in gastric cancer patients compared with those not suffering from cancer," describes Kinoshita-Daitoku.

These novel findings provide crucial information regarding the molecular and genetic landscape of highly pathogenic *H. pylori* and may assist in the development of innovative therapeutics for the treatment of *H. pylori* related illnesses, including gastric [cancer](#).

More information: Ryo Kinoshita-Daitoku et al. A bacterial small RNA regulates the adaptation of *Helicobacter pylori* to the host environment, *Nature Communications* (2021). [DOI: 10.1038/s41467-021-22317-7](#)

Provided by Osaka University

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