

Introduction

Helicobacter pylori is a species of epsilon proteobacteria which colonizes the harsh environment of the human stomach (**Chalmers et al., 2004**). Its name refers to both its spiral shape (*Helicobacter*) and the area of the lower stomach which it habitually colonizes: the gateway (pylorus) between the stomach and small intestine (**Meyers, 2007**). This bacterium is thought to be present within up to 50% of the human population and has been linked to the development of a number of different medical conditions (**Chalmers et al., 2004**). This treehouse will provide information about the discovery of *H. pylori* as well as its classification, morphology, physiology and its effects on its human hosts.

Since its first isolation 20 years ago, *Helicobacter pylori* (*H. pylori*) infection continues to generate considerable interest in the medical and scientific community. Today, *H. pylori* is considered one of the most common pathogenic infections of mankind, infecting nearly one half of the population.

Support for a pathogenic role for *H. pylori* in humans initially came from independent studies by Marshall and Morris who established that ingestion of a large inoculum of *H. pylori* results in infection (**Marshall et al., 2009 and Morris and Nicholson, 2009**).

Data to support the early acquisition of *H. pylori* infection comes from retrospective studies which estimate the incidence of new *H. pylori* infections to be 0.37% per patient year (**Valle et al., 2010**).

At the present time, we know that *H. pylori* infection clusters within families and that the acquisition of this infection is strongly linked to conditions associated with lower socioeconomic status during childhood, such as residential crowding (**Malaty et al., 2010**).