

## THE OCCURRENCE AND CLINICAL SIGNIFICANCE OF ENTEROHEPATIC *HELICOBACTER* SPECIES IN LABORATORY RODENTS

*Het voorkomen en klinisch belang van enterohepatische Helicobacter-species  
bij laboratoriumknaagdieren*

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### ABSTRACT

To date, 13 enterohepatic *Helicobacter* species have been detected in laboratory rodents. Some of these species, in particular *Helicobacter hepaticus* and *Helicobacter bilis*, cause disease in immunocompromised and immunocompetent laboratory mice, rats and/or hamsters. In addition, subclinical infections with these bacteria may interfere with results obtained from experimental research in these animals and thus may lead to faulty interpretation of the data. Some laboratory rodent-associated *Helicobacter* species may also be considered zoonotic agents. Apart from *Helicobacter hepaticus*, the pathogenesis of *Helicobacter* infections has not been studied extensively, a fact which means there are substantial gaps in the comprehension of the virulence mechanisms of enterohepatic *Helicobacter* species associated with laboratory rodents. For all these reasons, the potential importance of these bacterial pathogens cannot be overlooked and undoubtedly merits further investigation.

### SAMENVATTING

Tot op heden werden 13 enterohepatische *Helicobacter*-species aangetoond bij laboratoriumknaagdieren. Verschillende van deze species, in het bijzonder *Helicobacter hepaticus* en *Helicobacter bilis*, veroorzaken ziekte bij immunodeficiënte en immunocompetente laboratoriummuizen, -ratten en/of -hamsters. Bovendien kunnen subklinische infecties met deze bacteriën interfereren met de resultaten van experimenteel onderzoek bij deze dieren en dus eventueel leiden tot een foute interpretatie van data. Enkele helicobacters die voorkomen bij laboratoriumknaagdieren kunnen ook de mens infecteren en kunnen dus worden beschouwd als zoönotische agentia. Op *Helicobacter hepaticus* na, is de pathogenese van *Helicobacter*-infecties slechts rudimentair bestudeerd waardoor de kennis van de virulentiemechanismen van enterohepatische *Helicobacter*-species bij laboratoriumknaagdieren schaars is. Omwille van al deze redenen kan het potentieel belang van deze bacteriële agentia niet worden ontkend en is er hiernaar duidelijk meer onderzoek nodig.

### INTRODUCTION

The genus *Helicobacter* nowadays includes at least 26 formally named species, with additional novel species in the process of being characterized (Fox, 1997; Whary and Fox, 2004). The genus can roughly be divided into the gastric *Helicobacter* species and the enterohepatic *Helicobacter* species (EHS). All gastric *Helicobacter* species have strong urease activity. They manage to survive gastric acidity by expressing urease at a level higher than that of any other known micro-organism (Dunn et

al., 1997; Sachs et al., 2003). EHS do not normally colonize the gastric mucosa, but do have characteristics of ultrastructure and physiology in common with the gastric *Helicobacter* species. They have been identified in the intestinal tract and/or the liver of humans, mammals and birds (Fox, 1997; Solnick and Schauer, 2001; Inglis et al., 2006). EHS infections are associated mostly with intestinal and hepatobiliary disease in a wide range of animals. In laboratory rodents, 13 species of the enterohepatic *Helicobacter* group have been described so far, of which 11

are recognized and two provisionally named. These bacteria may also interfere with results obtained from experimental research in laboratory animals in which they are highly prevalent, and thus may lead to faulty interpretation of the data (Fox *et al.*, 1994; Ward *et al.*, 1994; Rogers and Fox, 2004; Bohr *et al.*, 2006). Finally, members of the enterohepatic *Helicobacter* group may have zoonotic potential for causing gastroenteritis, hepatitis and other disease signs in humans (Solnick and Schauer, 2001; Ljung and Wadstrom, 2002). For all these reasons, the potential importance of these emerging pathogens cannot be overlooked and undoubtedly merits further investigation.

The discovery of EHS has sparked an interest in exploring the pathogenic potential of these organisms in laboratory rodents. Nonetheless, the knowledge of the pathogenesis of EHS infections in these animals still contains large gaps. Additionally, the actual prevalence and clinical significance of most of the EHS remain to be established (Fox, 1997; Solnick and Schauer, 2001). This paper is intended to give a brief overview of the 11 recognized EHS associated with laboratory rodents used in diverse experimental disciplines, with a focus on bacterium-host interaction and the clinical significance for these animals. The methods used for monitoring laboratory rodents for EHS and for the elimination of EHS from laboratory rodent colonies, as well as the zoonotic potential of these bacterial agents, are also concisely considered.

## OCCURRENCE, CLINICAL SIGNIFICANCE AND BACTERIUM-HOST INTERACTIONS OF EHS IN LABORATORY RODENTS

The different EHS recognized in laboratory rodents and their rodent hosts are presented in Table 1.

### EHS mainly associated with mice and rats

#### *H. hepaticus*

*H. hepaticus* is a helical, motile bacterium that has been identified in mice and more recently in gerbils (Fox *et al.*, 1994; Ward *et al.*, 1994; Euzéby, 2002). Depending on the mouse strain and detection technique, a moderate to large portion of laboratory mice are found to be infected with *H. hepaticus* (Fox *et al.*, 1998a; Goto *et al.*, 2000; Whary *et al.*, 2000; Goto *et al.*, 2004; Nilsson *et al.*, 2004). Spreading by fecal-oral contact between animals is hypothesized. *H. hepaticus* infection may be spread by dirty bedding (Whary *et al.*, 2000). Livingston *et al.* (1997) demonstrated that *H. hepaticus*-free animals can develop antibodies against this EHS within four weeks after contact with dirty cage bedding from *H. hepaticus*-infected mice. Vertical transmission of *H. hepaticus* has been suggested by Li *et al.* (1998), but may depend on the mouse strain involved. Nonetheless, it has been recommended to foster pups within 24 h of birth to keep them free of *H. hepaticus* (Singletary *et al.*, 2003).

In several mouse strains, *H. hepaticus* infection has been associated with chronic active hepatitis and liver tumors. Susceptibility to liver disease seems to be mouse

**Table 1. Enterohepatic *Helicobacter* species associated with laboratory rodents.**

<i>Helicobacter</i> species	Rodent host species	Type strain
<i>H. hepaticus</i>	mouse, gerbil	ATCC 51448 <sup>T</sup>
<i>H. bilis</i>	mouse, rat, gerbil	ATCC 51630 <sup>T</sup>
<i>H. muridarum</i>	mouse, rat	ATCC 49282 <sup>T</sup>
<i>H. rodentium</i>	mouse, rat	ATCC 700285 <sup>T</sup>
<i>H. trogontum</i>	rat	ATCC 700114 <sup>T</sup>
<i>H. ganmani</i>	mouse	CCUG 43526 <sup>T</sup>
<i>H. typhlonius</i>	mouse, rat	MU96-1 <sup>T</sup>
<i>H. cholecystus</i>	hamster	ATCC 700242 <sup>T</sup>
<i>H. cinaedi</i>	hamster, rat	ATCC 35683 <sup>T</sup>
<i>H. aurati</i>	hamster	ATCC BBA-1 <sup>T</sup>
<i>H. mesocricetorum</i>	hamster	MU97-1514 <sup>T</sup>

strain and sex dependent. While C57BL/6NCr mice appear to be resistant to liver disease caused by *H. hepaticus*, BALB/cNCr, SJL/NCr, A/JCr, SCID/NCr and C3H/HeNCr mice can develop hepatitis which may be worse in males than in females (Fox *et al.*, 1996b; Whary *et al.*, 1998). Table 2 presents the description of lesions and clinical signs which may occur in a number of mouse strains infected with *H. hepaticus*. Finally, Maurer *et al.* (2005) suggested that *H. hepaticus* amongst other EHS might play an important role in the pathophysiology of cholesterol gallstone development in mice and possibly in humans. Forty percent of mice infected with *H. hepaticus* and fed a lithogenic diet developed cholesterol gallstones.

The pathogenesis of *H. hepaticus* infections in mice has been investigated by several research groups with the aim of obtaining a clear view of the relationship of the organism to the host tissues at the molecular level and the development of lesions in the liver and intestines due to *H. hepaticus*. Proteins of *H. hepaticus* have orthologs from both *H. pylori* and *Campylobacter jejuni*, but *H. hepaticus* is deficient in orthologs from most known *H. pylori* virulence factors, including adhesins, VacA cytotoxin, and nearly all *cag* pathogenicity island (PAI) proteins. However, *H. hepaticus* has orthologs of the adhesin PEB1 and the cytolethal distending toxin (CDT) of *C. jejuni*, a 71-kb genomic island (HHGI1) and several genomic islands with a different G+C content than the other sequences of the genome. Interestingly, HHGI1, possessing three basic elements of a type IV secretion system and other virulence protein homologs, constitutes a putative PAI. Within *H. hepaticus*, a large divergence of genome content, including the genomic island HHGI1, is present (Suerbaum *et al.*, 2003). A recent study using male A/JCr mice demonstrated the role of this PAI in the development of hepatitis in these animals (Boutin *et al.*, 2005).

A decade ago, auto-immunity was demonstrated to contribute to hepatocellular damage in *H. hepaticus* infection. *H. hepaticus*-infected mice indeed may build up IgG antibodies against the heat shock protein 70 expressed both by the bacterial agent and the injured liver cells (Ward *et al.*, 1996; Whary *et al.*, 1998).

Additionally, in *H. hepaticus* a toxin activity has been identified that causes vacuole formation in a murine liver cell line, resulting in a granular appearance of the affected cells. The toxin was called granulating cytotoxin (GCT), referring to the induced morphological cell changes (Taylor *et al.*, 1995). Despite the innovative and interesting character of this finding, no further research involving this toxin was performed for almost a decade. Only recently, Young *et al.* (2004) alleged that the cytopathic

effect induced by GCT could be (CDT)-mediated. Indeed, the same research group previously identified three genes encoding CDT and CDT activity in *H. hepaticus* (Young *et al.*, 2000). This toxic factor has the unique ability to stop the proliferation of various cells that are thus blocked before entering into mitosis. CDT-treated cells are enlarged and the nuclei are distended proportionally. Giant cells and multinucleated cells also can be observed (Ceelen *et al.*, 2006).

Only a few studies on the role of CDT in the pathogenesis of EHS infections have been reported thus far. Ge *et al.* (2005) illustrated that *H. hepaticus* CDT is crucial in the persistent colonization of this bacterial agent in the gut of outbred Swiss Webster Mice, especially in males. Young *et al.* (2004) constructed a CDT-negative *H. hepaticus* mutant and challenged C57/BL6-*Il10*<sup>tm</sup> mice with this mutant strain. They noticed that, although the isogenic *H. hepaticus* CDT mutant maintained the capacity to colonize C57/BL6-*Il10*<sup>tm</sup> mice, animals inoculated with the mutant developed markedly less severe disease than mice inoculated with a wild-type *H. hepaticus* strain (Young *et al.*, 2004). Furthermore, the results from a recent study of Pratt *et al.* (2006) suggest that CDT has an important immunomodulatory function allowing the persistence of *H. hepaticus* in IL-10<sup>-/-</sup> mice. In addition, CDT may alter the host immune response, resulting in the development of colitis (Pratt *et al.*, 2006).

Finally, *H. hepaticus* expresses high levels of urease activity comparable to half the activity of *H. pylori* (Sachs *et al.*, 2003). It is not clear why these levels of urease activity would be essential in the lower bowel and liver, both non-acidic environments. Possible roles for this enzyme in *H. hepaticus* include the promoting of better endurance during passage through the stomach and the production of ammonia as a source of nitrogen for protein biosynthesis. Urease activity may also be an important factor in the pathology, given that ammonia harms host cells and urease itself provokes phagocyte chemotaxis, stimulates immune cells and induces cytokine production (Beckwith *et al.*, 2001).

### *H. bilis*

This fusiform bacterial species was first identified in aged inbred mice with chronic hepatitis by Fox *et al.* (1995), who recovered the species from liver, bile and intestines of C57BL/6, CBA/CA, BALB/c and DBA/2 mice strains. *H. bilis* infection seems to be widespread in laboratory rodent colonies: 13%-17.1% in mice and 9.7% in rats (Riley *et al.*, 1996; Whary *et al.*, 2000). Goto *et al.* (2000), however, found a prevalence of only 2.1% in laboratory mice in Japan, and did not detect it at all in

Table 2. Lesions and clinical signs present in *H. hepaticus* infected mouse strains as stated by various authors.

Mouse strain	Lesions and clinical signs		Bacterial exposure timing	Age	Reference	
	males	females				
A/JCr	Generally Liver inflammation and hepatocarcinoma when bacterial exposure at or before 3 weeks of age	Lobular necrogranulomatous and chronic active hepatitis	During conception, 10 days after conception, 3 to 12 months	3 to 12 months	Roger <i>et al.</i> , (2004)	
	Typhlitis, IBD, more severe at older age	NA	3 weeks	Month 1 PI	Myles <i>et al.</i> , (2003)	
	Chronic active hepatitis	Severe perivascular periportal, hepatic parenchymal lesions, lymphohistiocytic and plasmacytic cellular infiltrates	No significant difference between infected and control animals	6 to 8 weeks	Month 6 to 12 PI	Whary <i>et al.</i> , (1998)
		Transmural typhlitis	Mucosal epithelial caecal hyperplasia		Month 3 PI	
A/J	Chronic hepatitis and hepatocarcinoma	Chronic inflammation, cell hyperplasia, hepatomegaly, bile duct proliferation, necrosis, hepatic adenomas, 5-brom-2'-deoxyuridine proliferation	Newborns and weanlings	Month 3 to 18 PI	Fox <i>et al.</i> , (1996a)	
TRC alphabeta mutant	Intestinal epithelial cell hyperplasia and mucosal inflammation	NA	NA	NA	Chin <i>et al.</i> , (2000)	
SCID/NCr	Moderate cecal and colonic lesions	Fibrosis, portal inflammation, including lymphoid nodules, more frequent in older mice	3 to 4 weeks	5 to 7 months	Avenaud <i>et al.</i> , (2003)	
Tac: (SW)f	Hepatitis, proliferate typhlitis, colitis, multifocal to coagulative hepatocyte necrosis	Hepatitis, typhlitis, colitis, colonic epithelial hyperplasia	4 to 6 months	4 to 6 months		
	Chronic hepatitis and enterocolitis	Severe hepatitis, colitis, severe colonic epithelial hyperplasia	9 to 10 months	9 to 10 months	Li <i>et al.</i> (1998)	
		NA	6 to 8 months	6 to 8 months		
Genetically altered mouse lines	IBD, rectal prolapse	NA	Hepatitis, colitis, colonic epithelial hyperplasia	6 to 8 months		
		NA	Multifocal hepatocytic necrosis, lymphocytic and neutrophilic infiltrates	Week 3 PI		
		NA	More severe necrosis	Week 10 to 28 PI		
		NA	Necrosis and multinucleate giant cells present, more prominent inflammation	Week 33 to month 16 PI	Fox <i>et al.</i> , (1996b)	
		Disappearance of necrosis, hepatocellular carcinoma in one mouse, mucosal epithelial caecal hyperplasia, prominent Peyer's patches in colon	Month 16 to 24.5 PI			
		NA	NA	NA	Foltz <i>et al.</i> (1998)	

NA: not available, PI: post inoculation

rats. Although *H. bilis* infections primarily occur in mice and rats (Fox *et al.*, 1995), this organism has also been demonstrated in gerbils, hamsters, dogs, cats and pigs (Eaton *et al.*, 1996; De Groote *et al.*, 2000; Roosendaal *et al.*, 2000; Solnick and Schauer, 2001; Euzéby, 2002).

Like other *Helicobacter* species, this micro-organism colonizes the lower part of the intestinal tract in its hosts, generally without inflammation (Fox *et al.*, 1995). *H. bilis* infection, however, has also been associated with IBD, ulcerative typhlitis, typhlocolitis, proctitis, diarrhea and hepatitis in immunodeficient mice and rats (Shomer *et al.*, 1997; Franklin *et al.*, 1998; Shomer *et al.*, 1998; Fox *et al.*, 2004). Several studies (Eaton *et al.*, 1996; Shomer *et al.*, 1997; Burich *et al.*, 2001; Maggio-Price *et al.*, 2002) illustrate that mice and rats may serve as an animal model for IBD and additionally suggest that *Helicobacter* spp. may act as a valuable tool for studying microbially generated IBD.

Furthermore, Maurer *et al.* demonstrated that *H. bilis* can participate in the development of cholesterol gallstones in mice when the animals are fed a lithogenic diet (Maurer *et al.*, 2005). It should be noted that, despite the occurrence of hepatoenteric lesions in *H. bilis*-infected rodents, it is still unproven that this bacterium actually is the primary cause of the disease.

The pathogenesis of *H. bilis* infection has not yet been totally clarified. Ge *et al.* (2001) recognized outer membrane preparation (OMP) proteins in four *H. bilis* strains derived from different host species. These four strains were similar to each other but revealed a different protein profile than *H. pylori* OMP proteins, thus suggesting that *H. bilis* has a conserved, unique OMP profile. The divergence in the OMP structure of these two helicobacters was also illustrated by the absence of cross-antigenicity between the *H. bilis* OMP and a number of *H. pylori* OMP proteins, except for their flagellins. Another finding in this study was the presence of five heat-modifiable proteins (HMP) in the *H. bilis* OMP. Whether these proteins act as porins *in vivo* still needs to be elucidated (Ge *et al.*, 2001). Additionally, *H. bilis* possesses CDT activity, but to a lesser extent than *H. hepaticus* (Chien *et al.*, 2000). The organism also produces strong urease activity, like most other murine helicobacters. The function of this enzyme in *H. bilis* may be similar to that in *H. hepaticus*.

#### *H. muridarum*

*H. muridarum* has a unique cellular ultrastructure among the EHS: it has nine to 11 periplasmic fibers which emerge as concentric helical edges on the surface of the bacterium (Erlandsen and Chase, 1972; Lee *et al.*,

1992; Zenner, 1999). *H. muridarum* colonizes the intestinal mucus in mice and rats. Phillips and Lee (1983) discovered that the species is present in higher numbers in the ileum than in the large intestine of conventional rodents. However, following experimental inoculation of gnotobiotic animals, the colon and mainly the caecum were colonized with *H. muridarum*, suggesting that in the absence of competing bacterial microbiota, these intestinal parts are predisposed to *H. muridarum* colonization. These animals infected with *H. muridarum* in the ileum, caecum and colon did not develop lesions in a study of Queiroz *et al.* (1992). The results of another experiment, however, showed that SCID mice experimentally infected with *H. muridarum* did develop IBD at a higher rate than conventionally reared mice (Jiang *et al.*, 2002). This bacterium apparently is capable of invading epithelial cells and causing degeneration (Queiroz *et al.*, 1992; Solnick and Schauer, 2001). In aging mice, the species is able to reach the stomach, probably due to the lowered parietal cell mass, and consequently play a part in the etiology of chronic gastritis (Fox, 1997; Euzéby, 2002). When mice are enzootically infected with *H. muridarum* in the lower parts of the intestine and are subsequently inoculated with gastric helicobacters, *H. muridarum* is able to colonize the stomach after displacing these gastric *Helicobacter* species. This EHS may hence take advantage of the less acidic environment (Lee *et al.*, 1993). Additionally, Bury-Mone *et al.* (2003) found *H. muridarum* to be the only EHS harboring the *amiE* and *amiF* genes. These genes encode an amidase and a formamidase, respectively, and are only present in helicobacters capable of surviving in the stomach. Their acquisition might be linked to selective pressure in the acid gastric environment.

#### *H. rodentium*

*H. rodentium* is spiral-shaped and possesses inside circular and intraplasmatic structures containing polyphosphate granules (Shen *et al.*, 1997). *H. rodentium* is present in 5% to 17% of several mouse strains, including C57BL/6, BALB/cA and C3H/HeJ (Sher *et al.*, 2000). Goto *et al.* (2000) detected *H. rodentium* in 38.3% and 30% of mice and rats, respectively. In several colonies, mice were infected with both *H. hepaticus* and *H. rodentium*. *H. rodentium* seems to be a normal inhabitant of the intestinal tract of mice and rats. Like *H. hepaticus* and *H. bilis*, this EHS may be transmitted by dirty bedding (Whary *et al.*, 2000). Its possible pathogenicity is hardly documented. SCID mice co-infected with *H. rodentium* and *H. bilis* suffered from severe diarrhea in one study (Shen *et al.*, 1997; Shomer *et al.*, 1998), while the authors of another study suggest that this bacterial agent is not the

cause of hepatitis or enteritis (Myles *et al.*, 2004). The exact role of *H. rodentium* in the development of diarrhea and other diseases in mice and rats requires further investigation (Shen *et al.*, 1997; Solnick and Schauer, 2001).

### *H. trogontum*

Mendes *et al.* (1996) isolated this rod with pointed ends from the colonic mucosa of Wistar and Holtzman rats. *H. trogontum* is genetically most closely related (97%) to *H. hepaticus*, but has the ability to grow at 42°C, in contrast to *H. hepaticus* (Mendes *et al.*, 1996). The prime colonization site in rats probably is the colon. It also has been cultured from the caecum, colon and stomach in experimentally *H. trogontum*-inoculated axenic outbred mice. An organism ultrastructurally indistinguishable from *H. trogontum* was noticed in bile ducts in rats experimentally infected with the liver fluke. However, this bacterium was not cultured and thus not conclusively characterized as *H. trogontum* (Mendes *et al.*, 1996; Solnick and Schauer, 2001). Whether *H. trogontum* induces gastrointestinal lesions in rats has not been studied (Moura *et al.*, 1999). In mice, *H. trogontum* may be associated with gastrointestinal and perhaps hepatic lesions (Moura *et al.*, 1999). Because of the presence of urease activity in this EHS, we can infer that *H. trogontum* has the capacity to colonize the stomach as well, similar to other urease-positive intestinal microbes such as *H. muridarum* in mice (Queiroz *et al.*, 1992; Mendes *et al.*, 1996).

### *H. ganmani*

*H. ganmani* has been detected in the caecum and liver of both specific pathogen free and conventional mice in Australia (Robertson *et al.*, 2001). A prevalence of 33% and 90% in C3H/HeJ mice and C57BL/6 mice, respectively, has been documented (Nilsson *et al.*, 2004). Remarkably, *H. ganmani* grows anaerobically at 37°C, but not microaerobically. No clear pathogenic significance has yet been attributed to *H. ganmani* (Euzéby, 2002). However, Zhang *et al.* (2005) suggested an association between natural *H. ganmani* infection in *I110* knock-out mice and the development of IBD.

### *H. typhlonius*

*H. typhlonius* is a spiral organism that has been isolated from *I110* knock-out mice suffering from IBD resulting in diarrhea and rectal prolapse and from the intestinal content of BALB/c mice with typhlocolitis. It has also been detected in B6sJl, SCID, BALB/cA, C57BL/6 and C3H/HeJ mouse strains (Fox *et al.*, 1999; Franklin *et al.*, 2001). No conclusive proof of liver colonization by this

micro-organism has yet been established. Infected immunodeficient mice only revealed mild portal inflammation (Franklin *et al.*, 1999). Recently, the presence of *H. typhlonius* in sex organs has been reported in three different mouse strains, but no vertical transmission was found (Scavizzi and Raspa, 2006).

Fox *et al.* (1999) demonstrated that *H. typhlonius* was as prevalent (4.88%) as *H. bilis* (4.33%) in laboratory rodents. In a more recent study, *H. typhlonius* was found in not less than 94% of examined SPF-SCID mice by means of pyrosequencing (Nilsson *et al.*, 2004).

## **EHS mainly associated with hamsters**

### *H. cholecystus*

*H. cholecystus* is the most important EHS in Syrian hamsters. *H. cholecystus* is similar to *H. bilis*, being fusiform in shape. It lacks urease activity (Franklin *et al.*, 1996; Euzéby, 2000; Solnick and Schauer, 2001). The intestinal tract of Syrian hamsters is possibly the normal habitat of *H. cholecystus* and animals older than five weeks seem to be more colonized with this agent than younger ones (Franklin *et al.*, 1996). *H. cholecystus* also may be present in ferrets (Garcia *et al.*, 2002).

In epidemiological studies, isolation of *H. cholecystus* has been strongly associated with cholangiofibrosis and centrilobular pancreatitis in Syrian hamsters. In fact, this micro-organism was first isolated from the gall bladders of hamsters suffering from these diseases (Franklin *et al.*, 1989, 1996). Liver lesions consist of portal neutrophilic and lymphoplasmatic infiltration with hyperplasia of the bile ducts. Formation of lymphoid follicles may also be present. More severe lesions are characterized by massive bile duct hyperplasia and fibrosis with lymphoplasmacytic infiltration. Pancreatic lesions range from mild periductular neutrophilic and lymphoplasmatic infiltration to severe inflammation resulting in interstitial pancreatitis (Brunnert and Altman, 1991; Franklin *et al.*, 1996). Affected hamsters suffering from cholangiofibrosis or centrilobular pancreatitis do not always show overt clinical signs (Franklin *et al.*, 1996). Hamsters have a common duct joining the bile and pancreatic ducts before entering the intestine. Consequently, one may assume that lesions in liver, gall bladder or pancreas may arise from an ascending infection in the common duct (Chen *et al.*, 2003). It is however also possible that *H. cholecystus* is not the primary cause of these lesions, but that already damaged liver, gall bladder or pancreatic tissue simply represents a favorable environment for growth of this micro-organism (Phillips and Lee, 1983; Franklin *et al.*, 1996).

*H. cinaedi*

The morphology of *H. cinaedi* resembles that of *H. hepaticus*. The agent was first detected in homosexual men, both asymptomatic individuals and men suffering from proctitis, proctocolitis or enteritis. The name 'cinaedi' means homosexual in Latin (Fennel *et al.*, 1984). Since then, many reports regarding individuals infected with *H. cinaedi* have been documented (Cimolai *et al.*, 1987; Ng *et al.*, 1987; Sacks *et al.*, 1991; Orlicek *et al.*, 1993; Burman *et al.*, 1995; Mammen *et al.*, 1995; Van der Ven *et al.*, 1996; Sullivan *et al.*, 1997; Peña *et al.*, 2002; Simons *et al.*, 2004; Uckay *et al.*, 2006).

This micro-organism seems to be a normal inhabitant of the intestinal tract of Syrian hamsters, and it causes neither histological lesions nor clinical signs in these animals. Since 75% of the hamsters harbor this bacterium, these animals may, however, serve as a reservoir of *H. cinaedi* infections for humans (Fennel *et al.*, 1984; Kiehlbauch *et al.*, 1995; Euzéby, 2001; Fox, 2002). Other animals where the species has been found are cats, dogs, foxes, rats, pigtailed macaques and rhesus macaques, in some cases displaying clinical symptoms (Kiehlbauch *et al.*, 1995; Flores *et al.*, 1990; Vandamme *et al.*, 2000; Fernandez *et al.*, 2002).

*H. aurati*

*H. aurati* is a fusiform *Helicobacter* species that was first isolated from inflamed stomachs and caeca of Syrian hamsters (Patterson *et al.*, 2000a). The presence of urease in *H. aurati* distinguishes it from the other three helicobacters found in these animals. The preferential colonization site of *H. aurati* in hamsters is the intestinal tract, particularly the caecum. However, subsequent spreading of this bacterial agent to the stomach may occur as well. The micro-organism colonizes the stomach following coprophagia or by retrograde ascending from the intestine in a similar way to *H. muridarum* (Queiroz *et al.*, 1992; Patterson *et al.*, 2000a). Urease activity allows the bacteria to survive in the acid gastric environment.

The role of *H. aurati* in gastric disease of hamsters is still unknown. Nonetheless, *H. aurati* has been identified in hamsters suffering from gastritis, accompanied by lesions similar to those in *H. muridarum*-infected mice. Hamsters naturally infected with helicobacters suffering from gastritis provide a model for the study of the development of gastric diseases in humans (Patterson *et al.*, 2000a,b; Nambiar *et al.*, 2005, 2006).

*H. mesocricetorum*

*H. mesocricetorum* is a urease-negative, spirally curved *Helicobacter* species. It was originally recovered from fecal pellets obtained from clinically healthy hamsters without enteric or hepatic lesions (Simmons *et al.*, 2000). It is phylogenetically most closely related to *H. rodentium*. The habitat and pathogenic potential of *H. mesocricetorum* are not clear. Up to now, the species has not been associated with any lesions or clinical signs in hamsters. *H. mesocricetorum* should possibly be considered a commensal organism of the intestinal tract of hamsters (Simmons *et al.*, 2000; Solnick and Schauer, 2001).

## ZOO NOTIC POTENTIAL OF EHS ASSOCIATED WITH LABORATORY RODENTS

During the last two decades, *Helicobacter* colonization of the gastrointestinal tract of humans has been the subject of intensive research (Solnick and Schauer, 2001; Fox, 2002; Ljungh and Wadstrom, 2002). The recovery of different helicobacters from both immunocompromised and immunocompetent human patients suffering from enteric and hepatobiliary disease has raised the question about the origin and impact of these infections.

Up to the current time, the causal role of EHS in human hepatoenteric disease has been mostly presumptive (Fox, 1997; Fox *et al.*, 1998b; Nilsson *et al.*, 1999; 2000a,b; 2001; Solnick and Schauer, 2001; Fox, 2002; Rocha *et al.*, 2005).

The rodent EHS *H. hepaticus*, *H. bilis* and *H. cinaedi* are generally considered zoonotic. *H. hepaticus* may play a role in liver carcinogenesis, IBD, irritable bowel syndrome (IBS) and chronic pancreatitis in humans (Nilsson *et al.*, 2000b; Apostolov *et al.*, 2005; Nilsson *et al.*, 2006; Zhang *et al.*, 2006). *H. bilis* is the only murine EHS that has actually been isolated from human gallbladder (Andersen, 2001) and its DNA has been demonstrated in patients suffering from chronic cholecystitis (Vorobjova *et al.*, 2006), cholecystolithiasis (Matsukura *et al.*, 2002), choledochocystolithiasis (Murata *et al.*, 2004, Kobayashi *et al.*, 2005), biliary tract and gall bladder cancer (Matsukura *et al.*, 2002; Murata *et al.*, 2004; Apostolov *et al.*, 2005; Kobayashi *et al.*, 2005).

*H. cinaedi* was first isolated from homosexual men, both asymptomatic individuals and men suffering from proctitis, proctocolitis and enteritis. Next, many reports regarding individuals infected with *H. cinaedi* have been documented. The agent is mainly found in immunocompromised persons, where it often causes a non-lethal disease with a large possibility for recurrence (Uckay *et al.*,

2006). *H. cinaedi* infection has been associated with septicemia and meningitis in a neonate (Orlicek *et al.*, 1993), bacteremia in an afebrile patient with X-linked agammaglobulinemia (Simons *et al.*, 2004), acute diarrhea (Tee *et al.*, 1987), bacteremia in immunosuppressed persons due to AIDS or cancer (Cimolai *et al.*, 1987; Ng *et al.*, 1987; Sacks *et al.*, 1991; Mammen *et al.*, 1995; Sullivan *et al.*, 1997; Uckay *et al.*, 2006) and multifocal cellulitis and monoarticular arthritis (Burman *et al.*, 1995; Sullivan *et al.*, 1997). Van der Ven *et al.* (1996) reported a case of an HIV-seropositive man who was suffering from a *H. cinaedi* bacteremia with involvement of the soft tissue in the right lower leg causing a localized pain in this area. It was illustrated that endovascular infection was present and could thus be a feature of *H. cinaedi* bacteremia. Despite the association of this species with extragastric infections, Peña *et al.* (2002) detected *H. cinaedi* DNA in antral gastric biopsies obtained from patients. One patient was diagnosed with erosive gastritis. Another patient had a history of colitis. Very recently, *H. cinaedi* DNA was detected in patients with pancreatic exocrine cancer (Nilsson *et al.*, 2006).

At least one other EHS present in rodents that may be transmitted to human beings is *H. ganmani*. This bacterial species has been reported in pediatric patients with liver disorders (Tolia *et al.*, 2004).

An association has been seen between *Helicobacter* species DNA in the liver and hepatitis C cirrhosis, with or without hepatocellular carcinoma (HCC) (Rocha *et al.*, 2005). *Helicobacter* species DNA has also been detected in livers from patients with cholangio- (71%) and hepatocellular (75%) carcinoma, but not in patients with hepatic metastases from colorectal carcinoma (Nilsson *et al.*, 2001). Moreover, neither *H. hepaticus* nor *H. bilis* DNA could be demonstrated in liver samples from human patients with primary sclerosing cholangitis (PSC) or primary biliary cirrhosis (PBC) (Nilsson *et al.*, 2000a), or in patients with gallstone formation (Maurer *et al.*, 2005).

Results from a study of Nilsson *et al.* (2001) demonstrated that 39% of patients with chronic liver disease and 20% of patients with PSC showed augmented antibody levels in an *H. hepaticus* enzyme immuno-assay (EIA). Patient serum samples retested by the *H. hepaticus* EIA after absorption with sonicated *H. pylori* cells remained positive in 12 of 37 serum samples. Distinctive antibody reactivity to 55–65 kDa proteins was noticed in *H. hepaticus* immunoblot (IB), after the absorption step, and was supposed to be unique for *H. hepaticus*. These authors suggest that antibodies to *H. hepaticus*, frequently cross reacting with *H. pylori*, occur regularly in persons with chronic liver diseases (Nilsson *et al.*, 2000b; Vorobjova

*et al.*, 2006). There may, however, be contradictions between results obtained with molecular techniques, on the one hand, and serology, on the other hand. While Apostolov *et al.* (2005) did not detect *H. bilis* DNA in gallbladder and liver biopsy specimens, 9% of the examined patients did reveal IgG antibodies to *H. bilis* using IB. The same research group also detected anti-*H. hepaticus* antibodies in the sera of patients with chronic cholecystitis using IB. Using the polymerase chain reaction (PCR)-denaturing density gradient gel electrophoresis (DDGE) and DNA sequencing as molecular methods or IHC, no positive results were obtained. The positive serologic reactions might be due to cross-reactivity to antigens of other (not yet identified) helicobacters or antigenically related bacteria. Alternatively, they may also result from an extrahepatic or extra-biliary infection (Apostolov *et al.*, 2005).

Altogether, the data from studies on biliary and hepatic diseases, as well as on pancreatic disorders, suggest that bile-tolerant *Helicobacter* species may induce a chronic infection with possible malignant transformation. Whether they truly participate in the genesis of biliary disease, however, requires additional investigation. At least there is evidence that both gastric and intestinal helicobacters can circulate in human bile (Queiroz *et al.*, 2003; Kobayashi *et al.*, 2005).

#### SCREENING OF LABORATORY RODENTS FOR THE PRESENCE OF EHS

Since EHS have been associated with gastrointestinal disease in laboratory rodents, it is important to screen accurately for EHS in laboratory facilities. These infections also may interfere with *in vivo* experiments and thus may lead to faulty interpretation of the data (Roger and Fox, 2004; Jacobsen *et al.*, 2005). Rodent EHS can be demonstrated in fecal samples or intestinal, gallbladder and liver tissue by culture, molecular methods, and histologic examination. Alternatively, sera can be examined for the presence of antibodies to these micro-organisms by ELISA (Fox *et al.*, 1994; Shames *et al.*, 1995; Fox *et al.*, 1996a,b; Riley *et al.*, 1996; Livingston *et al.*, 1997).

Columbia, Trypticase Soy and Brucella agar supplemented with 5% sheep or horse blood and occasionally with TVP (trimethoprim, vancomycin, polymyxin) are mainly used as cultivation media. EHS associated with laboratory rodents grow best in a microaerobic environment at 37°C, but not at 25°C. Several species may also grow at 42°C. *H. ganmani* is unusual in that this species grows anaerobically at 37°C, but cannot be cultivated under microaerobic conditions. Isolation can take place using nylon acetate filters with a pore size of 0.45 µm or



0.65  $\mu\text{m}$ , which may reduce contamination by other bacteria. Agar-grown EHS may present as swarming or single pointed colonies. Brucella broth supplemented with 5% fetal calf serum can also be adopted (Fox *et al.*, 1994; Fox *et al.*, 1995; Fox *et al.*, 1996a,b; Franklin *et al.*, 1996; Mendes *et al.*, 1996; Livingston *et al.*, 1997; Foltz *et al.*, 1998; Whary *et al.*, 1998; Chien *et al.*, 2000; Euzéby, 2000; Franklin *et al.*, 2001; Robertson *et al.*, 2001; Euzéby, 2002; Garcia *et al.*, 2002). Satisfactory culture of EHS may be hampered by the fastidious growth requirements of EHS. Additionally, the phenotypic similarity between member species of the genera *Helicobacter* and *Campylobacter* may result in misidentification (On *et al.*, 1996). Moreover, culture may require prolonged periods of time (one to three weeks) (Shames *et al.*, 1995). Because of these shortcomings, different alternative detection methods are appreciated (On *et al.*, 2002).

A frequently applied method is PCR. PCR now makes it possible to detect and identify different and novel species of the *Helicobacter* group. PCR also allows rapid and sensitive detection of the pathogen compared with bacterial culture, electron microscopy, histology and serology (Shames *et al.*, 1995; Livingston *et al.*, 1997). For each EHS in laboratory rodents, specific simplex, nested or reverse transcriptase (RT)-nested PCR methods have been described (Fox *et al.*, 1994; Battles *et al.*, 1995; Foltz *et al.*, 1995; Livingston *et al.*, 1997; Fox *et al.*, 1999; Moura *et al.*, 1999; Goto *et al.*, 2000; Franklin *et al.*, 2001; Nilsson *et al.*, 2004; Jacobsen *et al.*, 2005). However, a PCR assay with fecal samples is often hampered by inhibitory agents. Therefore, Shames *et al.* (1995) developed a PCR method using polyvinylpyrrolidone and Chelex 100 for demonstrating *H. hepaticus*. A *Tth* polymerase supplemented with an enhancer was applied for the DNA amplification. This PCR assay can be employed as a specific, non-invasive way of rapidly screening mice for *H. hepaticus*.

For demonstration of antibodies to *H. hepaticus*, *H. bilis* and *H. rodentium*, a species-specific ELISA technique can be used (Livingston *et al.*, 1997; Whary *et al.*, 2000; Euzéby, 2002). However, cross-reaction with some unidentifiable *Helicobacter* spp. may be noticed. False positive results can thus occur and confirmation by means of PCR is therefore required (Livingston *et al.*, 1997).

EHS can be visualized in tissue sections using the routinely adopted hematoxylin and eosin (H&E) stain. A Steiner modification of the Warthin-Starry stain, a Silver stain, may be more adequate and specific (Ward *et al.*, 1994a; Zenner, 1999; Rogers *et al.*, 2004).

## ELIMINATION OF EHS FROM LABORATORY RODENT COLONIES

Up to the present, only limited information about the effective treatment of EHS infections in laboratory animals has been available. Studies about this subject have mainly been performed in *H. hepaticus*- and *H. bilis*-infected laboratory rodents. Since EHS infections may cause not only gastrointestinal disease in these animals, but may also interfere with *in vivo* experiments, thus leading to the misunderstanding of data, it is of course important to eliminate EHS from laboratory rodent colonies (Rogers and Fox, 2004; Jacobsen *et al.*, 2005).

The best option for obtaining *Helicobacter*-free rodent colonies is probably rederivation by means of embryo transfer. Embryo transfer to rederive infected mouse strains from *H. hepaticus* (and others) has been well established (Van Keuren en Saunders, 2004; Watson *et al.*, 2005). Another possibility for getting rid of *H. hepaticus* infection may be caesarean section (Bergin *et al.*, 2005). Since vertical transmission of this bacterium has been implied (Li *et al.*, 1998), caesarean section may not be appropriate. On the contrary, the presence of *H. typhlonius* in sex organs of mice without vertical transmission to their offspring was documented very recently (Scavizzi and Raspa, 2006).

Antibiotic treatment of EHS-infected mice and rats might be an alternative (Bergin *et al.*, 2005; Jury *et al.*, 2005). Russell *et al.* (1995) claimed that amoxicillin administered orally for two weeks eliminates or prevents *H. hepaticus* infection in weanlings, but not in older mice with established enteric colonization. A triple therapy of amoxicillin, metronidazole and bismuth administered orally appears to be effective for the eradication of *H. hepaticus*, but not for *H. bilis* or *H. rodentium* infections (Foltz *et al.*, 1995; 1996; Shomer *et al.*, 1998). Recently, an amoxicillin-based triple therapy in combination with cross-fostering onto *Helicobacter*-free foster mothers showed some promise for the eradication of *Helicobacter* infections in several mouse strains. However, a cross-fostering control without amoxicillin triple therapy was not carried out, so no clear-cut conclusions can be drawn about the antibiotic treatment in this study (Kerton and Warden, 2006).

## CONCLUSIONS

The number of rodent helicobacters is rapidly growing and will probably continue to increase in the future. Despite the high prevalence of EHS in laboratory animals, little is currently known about the pathogenicity of these bacteria. Only *H. hepaticus* is firmly recognized as

a pathogen in mice, although its pathogenicity is still not totally understood. The pathogenicity of other rodent helicobacters has yet to be established. Some of these may constitute harmless inhabitants of the intestinal tract. In addition, very little information regarding the epidemiology of EHS in rodents is available. The evidence for the zoonotic potential of most EHS and their interference with experimental research is still circumstantial and requires more clarification. This certainly implies the necessity of prospective studies to better understand the pathogenicity of these bacterial organisms.

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