

## MINI REVIEW

# Interplay between *Helicobacter pylori* and immune cells in immune pathogenesis of gastric inflammation and mucosal pathology

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*Helicobacter pylori* infection is associated with an inflammatory response in the gastric mucosa, leading to chronic gastritis, peptic ulcers, gastric carcinoma and gastric mucosa-associated lymphoid tissue (MALT) lymphomas. Recent studies have shown that apoptosis of gastric epithelial cells is increased during *H. pylori* infection. Apoptosis induced by microbial infections are factors implicated in the pathogenesis of *H. pylori* infection. The enhanced gastric epithelial cell apoptosis in *H. pylori* infection has been suggested to play an important role in the pathogenesis of chronic gastritis and gastric pathology. In addition to directly triggering apoptosis, *H. pylori* induces sensitivity to tumor-necrosis factor (TNF)-related apoptosis-inducing ligand (TRAIL)-mediated apoptosis in gastric epithelial cells via modulation of TRAIL apoptosis signaling. Moreover, *H. pylori* infection induces infiltration of T lymphocytes and triggers inflammation to augment apoptosis. In *H. pylori* infection, there was significantly increased CCR6<sup>+</sup>CD3<sup>+</sup> T-cell infiltration in the gastric mucosa, and the CCR6 ligand, CCL20 chemokine, was selectively expressed in inflamed gastric tissues. These results implicate that the interaction between CCL20 and CCR6 may play a role in recruiting T cells to the sites of inflammation in the gastric mucosa during *Helicobacter* infection. Through these mechanisms, chemokine-mediated T lymphocyte trafficking into inflamed epithelium is initiated and the mucosal injury in *Helicobacter* infection is induced. This article will review the recent novel findings on the interactions of *H. pylori* with diverse host epithelial signaling pathways and events involved in the initiation of gastric pathology, including gastric inflammation, mucosal damage and development of MALT lymphomas.

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## INTRODUCTION

*Helicobacter pylori*, a common human pathogen, which infects about 50% of the world's population, is associated with duodenal and peptic ulcer diseases. The clinical consequences range from asymptomatic gastritis to peptic ulceration and gastric malignancy.<sup>1,2</sup> The outcome of the infection is determined by interactions among *H. pylori* virulence factors, host gastric mucosal factors and the environment. However, the mechanisms by which host factors cause disease remain unclear. The *H. pylori* infection changes the microenvironment of gastric mucosa. The apoptosis of gastric epithelial cells is increased,<sup>3–7</sup> and direct cytotoxicity as well as inflammatory responses occurs in the gastric mucosa cells.<sup>6,8–10</sup> It has been demonstrated that T helper type 1 cells selectively increased during *H. pylori* infection.<sup>11–15</sup> T helper type 1 cytokines, such as gamma interferon (IFN- $\gamma$ ) and tumor-necrosis factor alpha (TNF- $\alpha$ ), can increase the release of proinflammatory cytokines, augmenting apoptosis induced by *H. pylori*.<sup>10</sup> *H. pylori* infection could also induce gastric mucosa damage by increasing expression of Fas in gastric epithelial cells, leading to gastric epithelial cell apoptosis through Fas/FasL interaction with infiltrating T cells.<sup>9,16</sup> These findings suggest a role for immune-mediated apoptosis of

gastric epithelial cells during *H. pylori* infection. Recently, several bacterial pathogens have been found to trigger apoptosis in host cells *in vitro* or *in vivo*, and several types of mechanisms have been elucidated.<sup>17</sup> It was shown that *H. pylori* directly triggers cell death by cytotoxins after interaction with gastric epithelial cells.<sup>18,19</sup> Meanwhile, recent reports have shown that *H. pylori* translocates cytotoxin-associated gene A (CagA) into gastric epithelial cell by type IV secretion, inducing intracellular protein phosphorylation and dysregulate the signal transduction pathways within host cells.<sup>20–23</sup>

## Modulation of TNF-related apoptosis-inducing ligand (TRAIL)-mediated apoptosis by *H. pylori*

TRAIL (also called Apo2L), a novel TNF superfamily member with strong homology to FasL, is capable of inducing apoptosis in a variety of transformed cell lines *in vitro*,<sup>24,25</sup> but usually not in normal primary cells. It was shown that T cells can kill target cells via TRAIL/TRAIL receptor interaction,<sup>26–31</sup> indicating that TRAIL might serve as a cytotoxic effector molecule in activated T cells *in vivo*. These findings suggest that TRAIL/TRAIL receptor interaction is involved in the interaction between infiltrating T cells and gastric epithelium during

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*H. pylori* gastritis. In recent reports, Wu *et al.* demonstrated that human gastric epithelial cells sensitized to *H. pylori* confer susceptibility to TRAIL-mediated apoptosis, suggesting a role for immune-mediated apoptosis of gastric epithelial cells by infiltrating T cells during *Helicobacter* infection.<sup>7,15</sup> The induction of TRAIL sensitivity by *H. pylori* is independent of expression of *H. pylori* virulent factors vacuolating cytotoxin gene A (VacA) and CagA, and is dependent on viable bacteria and direct contact with cells.<sup>7</sup> The expression of TRAIL receptors did not change after *H. pylori* infection, indicating the *H. pylori*-induced enhanced sensitivity to TRAIL-mediated apoptosis in gastric epithelial cells was not due to upregulation of TRAIL death receptors. *H. pylori*-induced sensitivity to TRAIL-mediated apoptosis in gastric epithelial cells is dependent on activation of caspase 8 downstream pathway to convey the death signal to mitochondria, leading to activation of mitochondrial pathway and breaking the apoptosis resistance. *H. pylori* induces TRAIL-mediated apoptosis via enhancing the assembly of TRAIL death-inducing signaling complex and activation of caspase 8 and its downstream apoptosis signaling pathway. Thus, in addition to directly triggering apoptosis, *H. pylori* induces sensitivity to TRAIL in gastric epithelial cells by modulation of death-receptor signal transduction pathways. Modulation of host cell apoptosis by bacterial interaction adds a new dimension to the immune pathogenesis in *Helicobacter* infection.

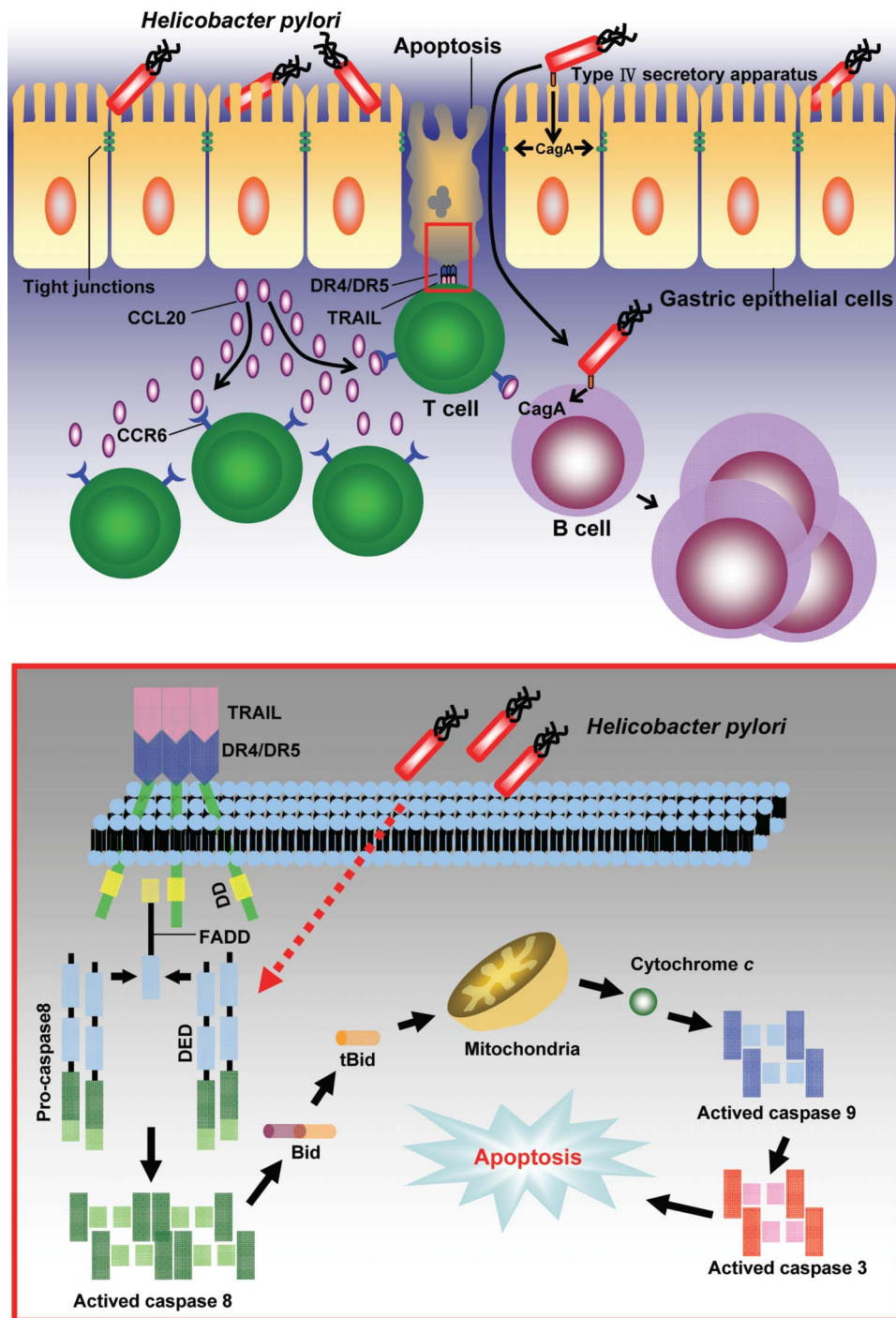
#### Chemokine-mediated lymphocyte trafficking of T lymphocytes in gastric inflammation during *H. pylori* infection

Previous studies have indicated that T lymphocytes play an important role in the pathogenesis of *Helicobacter* gastritis.<sup>11,32</sup> Inflammation of the gastric mucosa develops in response to the host immune reaction against the pathogens. The stimulation of epithelial cells with *H. pylori* contributes to the recruitment of neutrophils and lymphocytes. The activation of macrophages results in the release of cytokines, including IL-1, IL-6, IL-8, IL-12, TNF- $\alpha$  and IFN- $\gamma$ . As described, the features of *H. pylori*-induced inflammatory immune response are orchestrated by sequential elaboration of proinflammatory cytokines including IL-10, IFN- $\gamma$ , TNF- $\alpha$  and IL-1 $\beta$ . Accordingly, factors involved in regulating cytokine responses may confer susceptibility to or protection against *H. pylori*-associated diseases. All these results indicate that immune reaction and inflammation mediators to *H. pylori* play an important role in the pathogenesis of *H. pylori*-associated diseases. Among the T cells in response to *H. pylori* infection, the gastric infiltrating T cells mostly are CD45RO<sup>+</sup>CD69<sup>+</sup>CD4<sup>+</sup> T cells, indicating that there was accumulation of activated memory CD4<sup>+</sup> T cells during *Helicobacter* infection.<sup>15</sup> Recent reports indicated that the T helper type 1 response is induced during infection with *H. pylori*,<sup>11,13,14,33</sup> and the levels of IFN- $\gamma$  and TNF- $\alpha$ , are increased in the gastric mucosa during *H. pylori* infection, augmenting the apoptosis induced by *H. pylori*.<sup>6,7,10</sup> These results suggest a role for immune-mediated apoptosis of gastric epithelial cells by infiltrating T cells during *Helicobacter* infection. Therefore, in addition to bacterial virulence factors, the degree of gastric mucosa damage is also determined by the inflammation response induced during *H. pylori* infection. However, the induction of immune response and the immunopathogenic mechanism(s) of mucosal inflammation in *H. pylori* infections are still not clear, and chemokines are thought to play an important role in this process.<sup>34–37</sup> Chemokines are small, 6–14-kDa heparin binding proteins, which play a role in a variety of biological processes, most notably leukocyte chemotaxis.<sup>38,39</sup> Chemokines are involved in acute and chronic inflammatory processes by attracting neutrophils, monocytes and T cells to the site of inflammation via their corresponding chemokine

receptors.<sup>38,39</sup> Recent reports have shown that there are specific chemokines that mediate the homing of lymphocytes in the intestines,<sup>37,40,41</sup> suggesting that some chemokines may be involved in lymphocyte trafficking in the gut. It has been demonstrated that distinct sets of chemokines and their receptors are responsible for directing lymphocytes to inflammatory sites.<sup>35,36,40,42,43</sup> A set of proinflammatory chemokines has been shown to be involved in *H. pylori* gastritis: Gro- $\alpha$ , IL-8, RANTES, IFN- $\gamma$ -inducible protein-10 (CXCL10), a monokine induced by IFN- $\gamma$  (CXCL11) and CCL20 (MIP-3 $\alpha$ /LARC/exodus).<sup>15,44–46</sup> It has been demonstrated that the gastrointestinal epithelium senses the invading microorganisms and produces cytokines/chemokines that attract lymphocytes and dendritic cells to the site of inflammation.<sup>35,47,48</sup> Recently, it was reported that CCR6 mediates dendritic cell localization, lymphocyte homeostasis and immune responses in mucosal tissue.<sup>49</sup> CCR6, a specific  $\beta$ -chemokine receptor for CCL20, is selectively expressed on dendritic cells and some memory T cells<sup>48,50–52</sup> and may play a role in chemokine-mediated lymphocyte trafficking during gastric inflammation. It has also shown that CCL20, the ligand of CCR6, is abundantly expressed in mouse and human inflammatory enteric mucosa.<sup>53,54</sup> The production of CCL20 was upregulated in response to *H. pylori* in gastric epithelial cells when there was stimulation by the proinflammatory cytokines IL-1 $\beta$  and TNF- $\alpha$ .<sup>15,55–57</sup> These results implicate that the interaction between CCL20 and CCR6 may play a role in recruiting CD45RO<sup>+</sup> memory T cells to the sites of inflammation in the gastric mucosa during *Helicobacter* infection.

#### *H. pylori* CagA protein and the development of mucosa-associated lymphoid tissue (MALT) lymphomas

It has been established that the *cagA* gene product, CagA, can be directly injected into bacterium-attached host gastric epithelial cells via the bacterial type IV secretion system.<sup>20–23</sup> Infection by *cagA*-positive *H. pylori* is associated with gastric carcinomas and gastric MALT lymphomas.<sup>1,58</sup> The development of gastric MALT lymphomas is closely associated with *H. pylori* infection. The pathogenic role of *H. pylori* infection in gastric MALT lymphomas was observed in *in vitro* experiments and clinical evaluations of the effects of eradication on the progression of gastric MALT lymphomas.<sup>59,60</sup> Epidemiological studies further indicated that *cagA*-positive *H. pylori* is present in the gastric mucosa of most patients with gastric MALT lymphomas.<sup>61,62</sup> Clinical observations that eradication of *H. pylori* by antibiotic therapy can lead to the complete remission of MALT lymphomas<sup>60</sup> provide evidence that *cagA*-positive *H. pylori* plays an important role in the development and/or maintenance of MALT lymphomas. The development of gastric MALT lymphomas is dependent on *H. pylori* infection. Bacterial colonization of the gastric mucosa triggers lymphocyte infiltration,<sup>15,63</sup> and the formation of acquired MALTs. Previous studies suggested that MALT lymphoma cells preserve B-cell properties and that their growth may be partially driven by antigenic stimulation. *H. pylori* stimulates lymphoma B cells through tumor-infiltrating T cells, involving CD40 and CD40L costimulatory molecules.<sup>64,65</sup> However, the pathogenesis and how *H. pylori* induces the development of B-cell MALT lymphomas are still not clear. Much attention has been focused on the role of the *cagA* gene product, CagA, in the malignant transformation of cells. CagA was directly injected from bacteria into attached gastric epithelial cells by a type IV secretion system, encoded by the *cag* pathogenicity island,<sup>20,66</sup> and underwent tyrosine phosphorylation.<sup>23,67–69</sup> In human B lymphocytes, overexpression of *cagA* via transfection induces activation of extracellular signal-related kinase and their downstream apoptosis



**Figure 1** Immune pathogenesis of gastric mucosa damage in *H. pylori* infection. In the absence of *H. pylori* infection, there are very few T cells infiltrating into the gastric mucosa, which do not induce apoptosis in gastric epithelial cells. In contrast, in the presence of *H. pylori* infection, *H. pylori* induce inflammation and production of chemokine CCL20 to recruit CCR6 expressing activated CD4<sup>+</sup> T cells infiltrated to the sites of inflammation in the gastric mucosa. The TRAIL expressing T cells subsequently induce apoptosis in the *H. pylori*-infected gastric epithelia cells. Meanwhile, immune cells constituting MALTs migrate to and infiltrate the site of *H. pylori* infection in the gastric mucosa, and in such circumstances, CagA may be injected into lymphocytes as well as gastric epithelial cells. When CagA is translocated into B lymphocytes, it may induce activation of B lymphocytes to proliferate. The molecular mechanism of *H. pylori*-induced susceptibility to TRAIL-mediated apoptosis in gastric epithelia cells is shown in the lower inlet of the figure. In the absence of *H. pylori* infection, TRAIL engagement with death receptors on gastric epithelial cells induces only weak activation of caspase 8, and is not able to activate the caspase 8 downstream signals to trigger cell death. In contrast, in the presence of *H. pylori* infection, *H. pylori* enhance the assembly of TRAIL death-inducing signaling complex (DISC) after TRAIL engagement, to augment the activation of caspase 8 and to convey the death signal to mitochondria via cleavage of Bid, leading to activation of mitochondrial pathway and breaking the apoptosis resistance. CagA, cytotoxin-associated gene A; FADD, Fas-associated protein with death domain; MALT, mucosa-associated lymphoid tissue; TRAIL, tumor-necrosis factor-related apoptosis-inducing ligand.

regulators, indicating that CagA has effects on the growth and survival of B lymphocytes and may play a role in the development of MALT lymphomas.<sup>70,71</sup> *H. pylori* infection stimulates immune lymphocytes in the gastric mucosa and induces the formation of MALTs, from which MALT lymphomas of B-cell origin develop. Immune cells constituting MALTs migrate to and infiltrate the site of *H. pylori* infection in the stomach. In such circumstances, CagA may be injected into lymphocytes as well as gastric epithelial cells. Recent results in our laboratory have demonstrated that CagA could be directly translocated into human B cells from *H. pylori*. This implies the direct role and importance of CagA in the development of *H. pylori*-associated MALT lymphomas.

## SUMMARY

Human gastric epithelial cells sensitized to *H. pylori* conferred susceptibility to TRAIL-mediated apoptosis. Although the induction of TRAIL sensitivity by *H. pylori* in gastric epithelial cells was independent of *H. pylori* virulent factors CagA and VacA, the degree of apoptosis was linked to the presence of *H. pylori* and the associated inflammatory response. Therefore, the degree of mucosal damage was also determined by the inflammatory response induced by *H. pylori* within gastric epithelium. These results suggest a role for immune-mediated apoptosis and mucosa damage by infiltrating T cells during *Helicobacter* infection (Figure 1). In conclusion, *H. pylori* enhances susceptibility of gastric epithelial cells to TRAIL-mediated apoptosis. The induction of TRAIL sensitivity by *H. pylori* is dependent upon direct contact of viable bacteria with gastric epithelial cells. Modulation of host cell apoptosis by bacterial interaction adds a new dimension to the immune pathogenesis in chronic *Helicobacter* infection. The interplay between *H. pylori* and immune cells may induce activation of B lymphocytes via direct interaction or indirect immune stimulation leading to the development of *H. pylori*-associated MALT lymphomas.

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