

ANSWERING REVIEWERS

Round-1

Reviewer #1

Dear reviewer,

The comments you made on our manuscript were very important to the improvement of its quality and we are grateful for your crucial help. The article underwent a thorough language polishing. Here are the point-to-point changes according your suggestions:

1. We added a broader discussion on the specific immune responses induced by *H. pylori*, highlighted in **green** in the manuscript file.
2. We highlighted the main *H. pylori* virulence factors associated with the development of gastric cancer and added a discussion on another bacterial protein that has been closely associated with this disease, the heat-shock protein 60, as you suggested (highlighted in **pink** in the manuscript file).
3. We included discussions regarding the connection between *H. pylori* infection and gastric cancer and the necessity for further studies on this theme (highlighted in **blue** in the manuscript file).
4. We performed changes in Figure 1 in order to optimize it and make it more accurate, as you suggested.

Best Regards,

Fabício Freire de Melo

Professor, PhD

Reviewer #2

Dear reviewer,

We are grateful for your evaluation on our manuscript. No specific changes were required in your comments.

Best regards,

Fabício Freire de Melo

Professor, PhD

Round-2

Reviewer's code: 05754827

Dear reviewer,

Thank you for the important comments on our manuscript. They were crucial for enhancing the paper quality. We have performed all the changes you suggested, as described below.

1. We added a discussion on other *H pylori* virulence factors that have been closely associated with the development of gastric cancer (highlighted in gray in the manuscript file).
2. We made the Figure 1 available.
3. A thorough language polishing was performed with the elimination of several typos throughout the text (highlighted in yellow in the manuscript file).

Best Regards,

Fabício Freire de Melo

Professor, PhD

Name of Journal: *World Journal of Meta-Analysis*

Manuscript NO: 64742

Manuscript Type: REVIEW

Immune response to *Helicobacter pylori* infection and gastric cancer development

Immunology, H. pylori, and gastric cancer

Breno Bittencourt de Brito, Fabian Fellipec Bueno Lemos, Caroline da Mota Carneiro, Andressa Santos Viana, Nilo Manoel Pereira Vieira Barreto, Gabriela Alves de Souza Assis, Barbara Dicarlo Costa Braga, Maria Luísa Cordeiro Santos, Filipe Antônio França da Silva, Hanna Santos Marques, Natália Oliveira e Silva, Fabício Freire de Melo

Breno Bittencourt de Brito, Fabian Fellipe Bueno Lemos, Caroline da Mota Carneiro, Andressa Santos Viana, Nilo Manoel Pereira Vieira Barreto, Gabriela Alves de Souza Assis, Barbara Dicarlo Costa Braga, Maria Luísa Cordeiro Santos, Filipe Antônio França da Silva, Natália Oliveira e Silva, Fabrício Freire de Melo, Instituto Multidisciplinar em Saúde, Universidade Federal da Bahia, Vitória da Conquista 45029-094, Bahia, Brazil

Hanna Santos Marques, Campus Vitória da Conquista, Universidade Estadual do Sudoeste da Bahia, Vitória da Conquista 45031900, Bahia, Brazil

Author contributions: All authors equally contributed to this paper with conception and design of the study, literature review and analysis, drafting and critical revision and editing, and final approval of the final version.

Corresponding author: Fabrício Freire de Melo, MSc, PhD, Postdoc, Professor, Instituto Multidisciplinar em Saúde, Universidade Federal da Bahia, Rua Hormindo Barros, 58, Quadra 17, Lote 58, Bairro Candeias, 45029-094 Vitória da Conquista, Bahia, Brazil., Vitória da Conquista 45029-094, Bahia, Brazil. freiremelo@yahoo.com.br

Received: February 23, 2021

Revised:

Accepted:

Published online:

Abstract

Gastric adenocarcinoma is a global health concern and has *H. pylori* infection as the main risk factor for its occurrence. Of note, the immune response against the pathogen seems to be a determining factor for gastric oncogenesis, and increasing evidence **has** emphasized several host and bacterium factors that **probably play a role in that scenario**. The development of an inflammatory process against *H. pylori* involves a wide range of mechanisms such as the activation of pattern recognition receptors and intracellular pathways resulting in the production of proinflammatory cytokines by gastric epithelial cells. This process culminates in the establishment of distinct immune response profiles that result from the cytokine-induced differentiation of T naïve cells into specific T helper (Th) cells. Cytokines released from each type of Th cells orchestrate the immune system and interfere in the development of gastric cancer in idiosyncratic ways. Moreover, variants in genes that cytokines such as single-nucleotide polymorphisms have been associated with variable predispositions for the occurrence of gastric malignancy since they influence both in the intensity of gene expression and in the affinity of the resultant molecule with its receptor. In addition, various repercussions related to some *H. pylori* virulence factors seem to **influence host immune response** against the infection and many of them have been associated with gastric tumorigenesis.

Key Words: Gastric cancer; *Helicobacter pylori*; Immune response; Virulence factors; Polymorphisms

de Brito BB, Lemos FFB, Carneiro CDM, Viana AS, Barreto NMPV, Assis GAS, Braga BDC, Santos MLC, Silva FAFD, Marques HS, Silva NOE, de Melo FF. Immune response to *Helicobacter pylori* infection and gastric cancer development. *World J Meta-Anal* 2021; In press

Core Tip: Gastric cancer affects more than 1 million people yearly, and *H. pylori* infection is the main risk factor for that malignancy. Moreover, the immune response against the infection seems to play a pivotal role in gastric carcinogenesis. This article provides a broad and updated

overview on the main aspects regarding *H. pylori* infection, immune response, and gastric cancer development.

INTRODUCTION

About 1 million people are diagnosed with gastric cancer and more than 700,000 individuals die from this neoplasm every year^[1]. That incidence makes gastric adenocarcinoma the 5th most common malignancy, also being the 3rd cause of cancer-related death worldwide^[2]. Among the multiple factors that influence in the development of this disease, *H. pylori* infection stands out. Gastric colonization by this gram-negative, spiral-shaped microorganism is the main risk factor for the occurrence of gastric adenocarcinoma and, worryingly, it infects more than half of the world population^[3]. In that context, studies have emphasized the critical role of the interplays between *H. pylori* and the host immune system in carcinogenesis^[4].

The immune response activation by *H. pylori* infection in the gastric mucosa occurs mainly through the triggering of pattern recognition receptors (PRRs), which leads to the activation of intracellular cascades that culminate in the secretion of proinflammatory cytokines^[5]. The events taking place in the initial phase of infection lead to the recruitment of T cells, and the establishment of specific immune response profiles by T helper (Th) cells is determinant for the development of *H. pylori*-related gastric disorders^[6]. The *H. pylori* has several virulence factors that favor its perpetuation in the gastric hostile environment. Some mechanisms triggered by these bacterial products play pivotal roles in the regulation of the host immune response and seem to influence the genesis of gastric neoplasms^[7]. On the other hand, specific host polymorphisms in genes that encode cytokines also interfere in the risk of developing the disease by altering the expression pattern of these mediators as well as the intensity of the signals that they activate^[8].

Given the background, this article aims to provide a broad and updated review on how the immune response against *H. pylori* infection influences the development of gastric adenocarcinoma, discussing the main bacterial and host variables that interfere in the pathophysiology of the disease.

IMMUNE RESPONSE ACTIVATION BY *H. PYLORI* INFECTION

Complex host immune responses involving innate and adaptive mechanisms are induced by *H. pylori* infection^[9,10,11]. Gastric epithelium plays a pivotal role in the innate immune response to the bacterium since its cells make up the only cell phenotype in direct contact with the pathogen in conditions in which tissue damage is absent yet^[12,13]. The initial contact of the gastric epithelial cells with the pathogen activates **pattern-recognition receptors** including NOD1 and toll-like receptors (TLRs). These innate host defense mechanisms trigger cell **signaling** pathways that induce the activation of nuclear factor kappa B (NF- κ B), activating protein-1 (AP-1), and interferon regulatory factors (IRFs)^[14]. Concerning the TLRs, it has been reported that gastric epithelial cells express TLR1, TLR2, TLR4, TLR5, TLR9, and TLR10, which interact with various *H. pylori* antigens such as lipoteichoic acid, lipoproteins, lipopolysaccharide (LPS), flagellin, HSP-60, neutrophil-activating protein A (NapA), DNA, and RNA^[15- 17]. These receptors are very important to the induction of the expression of pro-inflammatory and antibacterial factors^[16]. For example, the translocation of NF- κ B to the nucleus, aiming at activating the expression of genes associated with the inflammatory process, is directly associated with the engagement of TLRs, particularly TLR2, in a Myeloid differentiation primary response 88 (MyD88)-dependent process^[18]. MyD88 is a key TLR adapter protein used by all TLRs, except TLR3, and transmits signals that result in the induction of inflammatory cytokines^[16]. However, although TLRs are the most studied receptors, *H. pylori* promotes the activation of PRRs other than TLRs. For instance, *H. pylori* peptidoglycan, delivered into host cells by the type IV secretion system (T4SS) or through outer membrane vesicles secreted from the bacterium, is recognized by NOD1^[19-22]. As a result, the interaction between *H. pylori* and PRRs leads to the expression of inflammatory cytokines, antimicrobial peptides, and type 1 interferon (IFN) by gastric epithelial cells^[14]. Subsequently, these cytokines and inflammatory mediators stimulate the recruitment of both polymorphonuclear and mononuclear cells into the gastric mucosa^[23,24]. Lastly, it is important to mention that there are other cell components that also act in the induction of this inflammatory process. Recent reports have led to the conclusion that miRNAs act as modulators of *H. pylori* infection and, concomitantly, have their expression affected by the bacterium^[25].

This inflammatory response is characterized by the chemotaxis of monocytes/macrophages, dendritic cells (DCs), B and T cells, and in particular, neutrophils, whose main **chemoattractant** is the IL-8 secreted by gastric epithelial cells as a result of the

engagement of NOD1, for instance^[26, 27]. Neutrophils are recruited to the lamina propria at the beginning of *H. pylori* infection and several specific *H. pylori* factors are known to interact with these cells and modulate their responses^[28, 29]. One of these factors is a protein produced by *H. pylori* known as neutrophil-activating protein (HP-NAP or NapA). HP-NAP can promote chemotaxis, endothelial adhesion, and the production of reactive oxygen intermediates by neutrophils^[30-32]. Incubation of these cells with HP-NAP results in significant production of cytokines such as IL-12 and IL-23. The same effects of HP-NAP on cytokine secretion were also observed in macrophages and DCs. Therefore, it is possible to conclude that this protein acts on both neutrophils and monocytes, inducing the production of cytokines^[31]. Mononuclear infiltration in the lamina propria is also characteristic of *H. pylori*-induced chronic infection^[13]. Human monocytes and macrophages are important coordinators of the immune response to *H. pylori*-derived products and signals from epithelial cells in direct contact with the bacterium on the surface of the mucosa^[15]. In this infection, both monocytes and macrophages, alongside the DCs, act as activators of adaptive immunity, since they are antigen-presenting cells (APCs), capable of expressing class II MHC molecules that activate CD4⁺ T cells^[33]. Furthermore, monocytes and macrophages also produce factors such as IL-12, responsible to induce a polarized T helper 1 (Th1) immune response, IL-1 β , IL-6, IL-10, and TNF- α , which, except for IL-10, induce the amplification of the inflammatory response^[17]. Moreover, it is important to note that macrophages are also effector cells that are able to produce nitric oxide (NO) derived from the enzyme-inducible NO synthase (iNOS, NOS2) and reactive oxygen species (ROS), both associated with cellular damage^[34]. Although in smaller numbers, DCs are also important in the immune response to *H. pylori* infection, especially because they represent an important bridge between the innate and adaptive immunities^[27]. These cells express a broad spectrum of PRRs, which enables them to capture antigens at the periphery and induce T naive cells to direct T cell differentiation^[35]. This role is played through three main signals: (1) presentation of foreign antigens in form of peptides bound to class II MHC molecules to T cells; (2) co-stimulation of T cell differentiation; and (3) secretion of cytokines, particularly IL-6, IL-8, IL-10, IL-12, IL-1 β , and TNF- α ^[36]. Both aforementioned antigen-presenting cells exhibit remarkable secretion of IL-12, which enables the induction of a Th1-polarized immune response, responsible for the secretion of INF- γ and low amounts of cytokines characteristic of Th2 responses, such as IL-4 and IL-5^[37-41]. Finally, mast cells represent an additional innate cell phenotype that is found within the *H. pylori*-infected gastric

mucosa. These cells can be activated by various *H. pylori* components. For instance, the bacterial virulence factor VacA can induce mast cells to express multiple inflammatory cytokines, including IL-1, TNF, IL-6, IL-23, and IL-10^[42, 43]. Upon the stimulation of epithelial cells, macrophages, and DCs by *H. pylori* bacterial factors, CD4⁺ and CD8⁺ T cells are recruited to the gastric mucosa, with preferential activation of CD4⁺ T cells in detriment of CD8⁺ cells^[44-48].

IMMUNE RESPONSE PROFILES IN *H. PYLORI* INFECTION AND GASTRIC CANCER

As aforementioned, the triggering of an immune response against the *H. pylori* involves the activation of CD4⁺ and CD8⁺ T cells and their migration to the gastric environment^[49]. Among the cytokines expressed in that context, stand out those inducing the differentiation of naïve T cells into Th1 (e.g. IL-12), Th17 (e.g. TGF- β , IL-23, and IL-6), and Treg (e.g. IL-2 and TGF- β) cells^[50].

The establishment of proinflammatory Th1 response in the *H. pylori* infection is associated with the development of corpus gastritis. Depending on the further host and environmental variables, the aforementioned condition can result in gastric atrophy and intestinal metaplasia, which are well-known precancerous lesions^[51,52]. Of note, our group has previously demonstrated that Th1 response varies according to the age among *H. pylori*-positive individuals. In that study, higher gastric concentrations of the Th1-related cytokines IL-2, IL-12p70, and INF- γ were observed in adults than in children. Moreover, the levels of Th1 cytokines were directly correlated with the severity of gastric inflammation^[53].

Regarding Th17 response, although other cytokines such as TGF- β and IL-6 are strongly related to this immune profile, current evidence emphasizes the pivotal role of IL-23 in its induction in the setting of *H. pylori* infection^[54,55]. A study found that chronically infected IL-23(p19)^{-/-} mice had reduced gastric expression of IL-17A as well as milder gastric inflammation and higher levels of *H. pylori* colonization compared to wild-type *H. pylori*-positive mice^[56]. The IL-17A, in its turn, promotes the migration of polymorphonuclear leukocytes to the infection site and is an important component in the control of *H. pylori* gastric infection^[57]. Previous studies using mice have shown that IL-17A^{-/-}, as well as IL-17RA-deficient individuals, have a milder gastric neutrophil infiltration against *H. pylori* infection than wild-type mice. Interestingly, the mice lacking IL-17RA signaling had an enhanced chronic inflammation with intense infiltration of B and CD4⁺ T cells into gastric mucosa^[58].

Dual roles have been attributed to Th17 responses in cancer settings. On one hand, this immune profile seems to be important in the immunosurveillance against malignant cells since it stimulates the migration of leukocytes into tumors and promotes the activation of antitumor CD8⁺ T cells. Intratumoral Th17 cells induce the expression of CCL20, a chemokine that attracts dendritic cells (DCs) to the tumor environment, as shown in a recently published paper by Chen and colleagues^[59]. Subsequently, DCs phagocytose tumor material and migrate to lymph nodes, contributing to the activation of CD8⁺ T cells that migrate to the tumor environment through their chemotaxis to the Th17-induced CXCL9 and CXCL10^[60]. Moreover, studies have shown that Th17 cells can convert into Th1 Lymphocytes *in vivo*, enhancing their antitumor effectiveness^[61,62]. When stimulated by IL-23 and IL-12 in an environment with absent or low TGF- β , Th17 cells are able to express IFN- γ and T-bet, important Th1-related molecules. Interestingly, Th17-derived Th1 cells showed to have a more effective antitumor activity compared to other Th1 Lymphocytes, and this may be due to the prolonged survival and superior functionality of the former compared to the latter^[63].

On the other hand, the Th17 profile **has** shown to be involved in various **protumor** activities. Firstly, IL-17 seems to promote angiogenesis, since elevated intratumoral levels of that cytokine are associated with high expression of vascular endothelial growth factor (VEGF) and increased **tumor** vascular density^[64]. **In addition**, the aforementioned cytokine stimulates cancer cells to release IL-6, which, besides promoting VEGF production, enhances activator of transcription 3 (STAT3) activation, increasing the survival of malignant cells by suppressing apoptosis^[65]. Moreover, studies have described the existence of FOXP3⁺ CD4⁺ Th17 cells, which may play regulatory, **protumor** roles in cancer contexts. This phenomenon seems to occur along with low levels of IL-6, IL-23, and IL-23 as well as with the presence of TGF- β , which activates FOXP3 expression^[66].

A study carried out by Su *et al* showed that IL-17 and ROR γ t (the main IL-17A transcription factor) were highly expressed in both **tumor** microenvironment and peripheral blood mononuclear cells (PBMCs) of gastric cancer patients, mainly among those with metastasis^[65]. This data suggests that the presence of Th17 cells is not only directly associated with the occurrence of gastric cancer but also with the severity of the disease. Indeed, a recently published study embracing stage IV gastric cancer patients from four cohorts **has** reinforced that theory, since it found abnormally high levels of Th17 cell differentiation and activation of IL-17 pathways among patients with

severe disease^[67]. Interestingly, another study evaluating the percentages of Th17 cells in PBMCs among gastric cancer patients before and after tumor resection observed a significant drop in the proportion of Th17 cells after the treatment^[68]. Of note, IL-27 **has** been highlighted as a crucial cytokine that plays dual roles in the regulation of the immune system. As far as this cytokine enhances T-bet expression through IL-27/IL-27R α signaling and subsequent STAT1 phosphorylation leading to Th1 cell differentiation, IL-27 impairs Th17 responses by downregulating ROR γ T^[69,70]. In a recently published study, our group showed that *H. pylori*-infected individuals have higher IL-27 Levels in their serum and gastric mucosa than non-infected individuals do. In contrast, there was a lack of IL-27 in both serum and **gastric mucosa from** gastric cancer patients, who also showed to have a remarkable Th17-polarized inflammatory pattern^[71].

The Treg cells might play pivotal roles in *H. pylori*-induced gastric adenocarcinoma by favoring infection perpetuation and by repressing immune responses against malignant cells through the secretion of regulatory cytokines. Indeed, studies have observed that Treg cells are positively correlated with increased bacterial colonization^[72], **and their levels are enhanced among gastric cancer patients**^[73,74]. Three types of Treg cells have been described by studies: IL-10-secreting Tr1 cells, TGF- β 1-producing Tr3 cells, and FOXP3-expressing CD4⁺CD25^{high} Treg cells^[75]. The latter is a pivotal component in the scenario of *H. pylori* colonization, favoring the pathogen persistence in the gastric environment by suppressing the immune responses. A study demonstrated that FOXP3, TGF- β 1, and IL-10 are highly expressed during *H. pylori* infection and the density of FOXP3⁺ Treg cells was higher in the gastric mucosa of infected individuals than in *H. pylori*-negative people. These cells have been also associated with **an increased bacterial density among individuals with gastritis**^[76].

Advances in the understanding of the interplays between Treg responses and the development of gastric cancer have been achieved. Interestingly, a recent study including gastric cancer patients in various stages of the disease showed that high infiltration of FOXP3⁺ Treg cells was associated with poor outcomes among individuals with advanced disease, but it was a predictor of better prognosis among patients with early-phase **diseases**^[77]. Current evidence emphasizes the role of Wnt/ β -catenin signaling in gastric carcinogenesis, since 70% of gastric cancer patients have dysregulation in pathways associated with this signaling^[78]. The β -catenin induces gastric cancer cells to produce CCL28, which strongly attracts Treg cells to the tumor environment. In

this sense, a recently published study using *H. felis*-colonized mice with gastric cancer found the block of β -catenin-induced CCL28 through anti-CCL28 antibodies leads to the suppression of gastric cancer progression by inhibiting Treg cells infiltration^[79].

A surface glycoprotein known as neuropilin1 (NRP1) seems to be crucial for the immunoregulatory events taking place in the tumor environment. The role of that molecule had already been well described in other malignancies, being related to cell migration, angiogenesis, and invasion^[80]. In a new investigation by Kang *et al*^[81], the expression of NRP1 was associated with increased levels of the regulatory cytokines IL-35, IL-10, and TGF- β 1 as well as with increased infiltration of Treg cells and M2 macrophages in gastric cancer. Moreover, its expression was positively correlated to poorer outcomes, which indicates that the NRP1 has the potential to be used as a prognostic factor in gastric cancer patients.

Besides the aforementioned roles of Treg cells in gastric cancer development, Liu *et al* found that these cells promote the expression of leucine-rich repeat containing G protein-coupled receptor 5 (Lgr5) by tumor cells *via* TGF- β 1 and TGF- β 1 signaling pathway, probably involving the aforementioned Wnt/ β -catenin signaling^[82]. The Lgr5 is a global stem cell marker whose overexpression is observed in gastric cancer and it **has** also been positively correlated with tumor invasion, metastasis, and poor prognosis among individuals with that malignancy^[83]. Figure 1 summarizes the roles played by Th cells in the setting of gastric carcinogenesis.

POLYMORPHISMS IN GENES THAT ENCODE CYTOKINES AND GASTRIC CANCER

IL-1

The Interleukin-1 family has 11 molecules that are able to interact with almost all human cells^[84]. Among these, the pro-inflammatory IL-1 β and the antagonist receptor of IL-1 (IL-Ra) have been associated with an increased risk of developing GC^[85-89]. The genes encoding IL-1 β and IL-1Ra are called *IL1B* and *IL1RN*, respectively^[90]. Single nucleotide polymorphisms (SNPs) found in these genes alter the inflammatory response of these cytokines. The SNPs identified in the coding of IL-1B have a C-T transition base at -511, -31, or +3954 positions. For IL-Ra, the **allele** 2 (IL-RN*2) has been associated with inflammatory responses that increase the risk of **disease** development.^[91]

In the presence of any of the three aforementioned SNPs in the sequences that encode IL-1 β , the production of this cytokine can be enhanced. This overexpression **has** been associated with the development of hypochlorhydria and atrophy of the gastric corpus, in addition to an increased risk for GC, especially on *H. pylori*-positive subjects^[92-102]. The production of IL1-Ra is mediated by a diversity of cytokines such as IL-1 β , and the antagonistic function of the former controls the inflammatory response of the **latter**. In this sense, the SNP IL-1RN*2 **has** been linked to **increased IL-1 β secretion**^[103-106].

IL-8

IL-8, also known as CXCL8, is a proinflammatory chemokine from the alpha subfamily (CXC)^[107]. It can be produced by several cells such as epithelial and endothelial cells, monocytes, macrophages, and tumor cells^[108,109]. Increased expression of IL-8 is promoted by various stimuli, including the initiation, modulation, and maintenance of the host inflammatory response against *H. pylori* infection^[110,111]. This molecule induces the migration and proliferation of endothelial cells, contributing to angiogenesis and tumorigenesis, being related to increased cell migration, invasion, and metastasis^[112].

The *CXCL8* gene is located **in chromosome 4q12-21** and possesses three introns, four exons, and a proximal promoter region^[113]. The genetic polymorphism IL8-251T> A (rs4073) **has** been associated with variations in the expression of IL-8 and increased risk of gastric cancer development mainly in Brazilian, Chinese, and Korean populations^[114]. Curiously, that polymorphism was not significantly associated with a higher risk of gastric cancer in the **Japanese** population, which might be related to specific environmental factors and genetic background^[107].

IL-10

As previously discussed in this review, the IL-10 is an antiinflammatory cytokine; therefore, it inhibits the activity of some defense cells and limits the production of pro-inflammatory cytokines^[115]. Polymorphisms -1082, -592, and, less frequently, -819, can modulate IL-10 transcription, decreasing the expression of this cytokine, which can initiate a hyperinflammatory response that increases the risk of gastric lesions and cancer. Therefore, these polymorphisms have been associated with a possible higher risk of GC, mainly in Asian populations, but also in a study conducted with American subjects.^[116-124]

IL-2

IL-2 is a cytokine that plays proinflammatory and antiinflammatory roles and is encoded by a gene located in chromosome 4q21. Among other repercussions, this molecule contributes to the proliferation of T regulatory cells and regulates the expansion and apoptosis of activated T cells^[26,40]. The *IL2* GG variant genotype -330T> G in *H. pylori*-positive Asians and Brazilians, as well as the SNP *IL-2* + 114T> G and 330g / + 114T haplotype in *H. pylori*-positive Brazilians, have been associated with an increased risk of developing gastric cancer^[125].

IL-4

Similarly to IL-2, IL-4 also plays dual roles in the immune system, being encoded by a gene in the chromosome 5q31.1^[112]. Its function in tumor progression is mainly related to the inhibition of proinflammatory cytokines in the setting of antitumor immune responses, favoring the perpetuation of malignant cells^[111]. Polymorphisms in *IL4* - 590C / T rs2243250 CC, genotype CT + CC, and *IL4* haplotypes have been found to be associated with a higher risk of developing gastric cancer in the Chinese population^[126].

IL-6

IL-6 is a cytokine that plays roles as an proinflammatory immune mediator and as an endocrine regulator^[82]. This protein is encoded by a gene located in the chromosome 7 and have been found to be increased in *H. pylori*-positive individuals^[112]. Polymorphisms in the *IL6* -174C allele and *IL6* -174CC genotype are associated with an enhanced prevalence of diffuse-type gastric cancer, whereas the *IL6* -174CG have been related to intestinal-type gastric cancer^[110]. In addition, the *IL6* SNP -572 (G> C, rs1800796) have been emphasized as a potential genetic biomarker for increased gastric cancer risk in Asian populations^[114].

IL- 22

IL-22 is an antiinflammatory cytokine that belongs to the IL-10 family. It participates in mucosal repair and epithelial immunity processes^[127]. Chinese individuals with the SNP rs1179251 (allele G) encoding IL-22 showed a higher risk of developing gastric cancer associated with *H. pylori*^[128]. Some SNPs of this cytokine have also been found in Chinese patients with increased risk for MALT gastric lymphoma induced by *H. pylori* (alleles C in rs2227485; A in rs4913428; A in rs1026788 and T in rs7314777)^[129].

H. PYLORI VIRULENCE FACTORS, IMMUNE RESPONSE, AND GASTRIC CARCINOGENESIS

CagA

Infection with *cagA*-positive *H. pylori* strains is the main risk factor for the development of gastric cancer^[130-132]. CagA is a multifunctional, pore-forming protein that induces vacuolization, cell necrosis, and cell apoptosis in gastric epithelial cells^[133-138]. Of note, this virulence factor appears to induce an important modulation of the host immune system^[139,140]. A recently published study by He *et al.* using mice revealed that CagA suppresses the expression of proinflammatory cytokines induced by *H. pylori* infection through the inhibition of the MAPK and NF- κ B pathways. In addition, the study **has** shown, for the first time, that this virulence factor downregulates the posttranslational modification of TRAF6, obstructing the transmission of a signal downstream responsible for promoting the release of proinflammatory mediators^[141].

Studies have described that *H. pylori* has **a** molecular mechanism of *cagA* expansion through which its number of copies expands, consequently enhancing its virulence^[142,143]. The analysis of the PMSS1 *H. pylori* strain showed that bacteria that carry more *cagA* copies also produce **higher levels of this toxin**, leading to enhanced cell elongation and IL-8 induction^[144,145]. Yomaoka *et al* **propose** that the levels of IL-8 of the gastric mucosa are related to the presence of CagA and OipA^[146]. Both molecules seem to be involved in the induction of the interferon regulating factor (IRF) and play a role in the complete activation of the IL-8 promoter, using different convergence pathways^[147].

VacA

Vacuolating cytotoxin A (VacA) is a protein encoded by a monocistronic gene known as *vacA*. Secreted VacA molecules have 140 kDa initially, but they are rapidly cleaved into a 10 kDa domain (p10) to produce a mature 88 kDa protein^[148,149]. Generally, they are secreted as soluble proteins in the extracellular space; however, they are found on the bacterial surface as well^[150]. Moreover, this virulence factor is expressed by almost all *H. pylori* strains^[151].

VacA inhibits **the** activation and proliferation of T and B cells, a process that induces the apoptosis of macrophages mainly through the inhibition of INF- β signaling. Moreover, this

virulence factor induces an excessive release of IL-8^[152]. Specifically in T cells, VacA inhibits the production of IL-2, in addition to regulating the surface expression of the IL2- α receptor. This process is possibly due to the ability of VacA to inhibit the activation of the nuclear factor of activated T-cells (NFAT), a global transcription factor that regulates immune response genes for T cell activation. The mechanism by which VacA inhibits activation of NFAT is uncertain; however, it is believed that this virulence factor influences the calcium flow in the extracellular medium, which inhibits the calcineurin-dependent Ca²⁺-calmodulin complex^[153]. Other effects on these cells include the activation of intracellular signaling through MAP kinases, such as MKK3/6 and p38 as well as the Rac/Vav-specific nucleotide exchange factor^[149]. Studies with primary CD4⁺ T cells in humans have demonstrated that VacA inhibits the proliferation of activated T cells through a mechanism that is independent of the effect of VacA on NFAT activation and IL-2 expression^[154,155]. In **antigen-presenting cells**, VacA seems to interfere with the formation of vesicular compartments in macrophages infected with *H. pylori* causing homotypic vacuolar fusion and consequent changes in their physiological properties. It has also been reported that VacA can interfere with the antigen presentation of B lymphocytes by interfering in the MHC II of these cells. Finally, blocking the activation and proliferation of this set of cells helps *H. pylori* to resist the host immune response, **establishing persistent** infection and with worse clinical outcomes^[150].

The various positive VacA-linked bacterial genotypes are associated with a higher prevalence of malignant gastric lesions, in addition **to greater** severity of inflammation induction by the pathogen. VacA is the most studied toxin in *H. pylori* due to its versatility in relation to different receptors in different cell types and functions. VacA is directly involved in the formation of intracellular vacuoles which provide the survival of the bacteria in the gastric environment, even after drug treatment. Therefore, other studies need to be developed with VacA in order to better understand the persistence of the pathogen in the gastric environment^[156].

DupA

The duodenal ulcer promoter A (DupA) protein is a *H. pylori* virulence factor whose gene is located in the plasticity zone of the bacterial genome^[157]. The *dupA* gene contains two overlapping open reading frames (ORF) (jhp0917 and jhp0918) that form a continuous locus^[158].

Of note, only strains that harbor both aforementioned segments are able to produce the DupA protein^[159].

The initial studies on DupA show that its pathogenicity is closely linked to the development of duodenal ulcers^[158]. Based on *in vitro* and *in vivo* studies, such outcome is believed to be due to the role of the DupA gene in the activation of kappa B nuclear transcription factors (NF-κB) and activating protein-1 (AP-1), which enhance the infiltration of neutrophils with the consequent expression of IL-8 in the antrum, which promotes risk of these injuries^[160,161]. These findings have shown that predominant antral gastritis often leads to a reduction in somatostatin, greater gastrin secretion and, consequently, greater release of gastric acid and formation of duodenal ulcers^[162]. In this context, DupA expression is negatively correlated with the risk of gastric atrophy, intestinal metaplasia, and gastric cancer^[163,164]. However, it has to be emphasized that DupA have not been associated with the development duodenal ulcer in western populations^[165,166].

OipA

The 34 kDa external inflammatory protein A (OipA), encoded by the *hopH* gene (hp0638), located approximately 100kd from the Cag Pathogenicity Island (cag PAI), belongs to the family of External Membrane Proteins (OMPs). This protein is associated with gastric inflammation, being one of the main *H. pylori* virulence factors^[167].

The attachment of gastric epithelial cells through OipA occurs with the induction of cellular apoptosis *via* the Bcl-2 pathway, increased levels of Bax, and cleaved caspase 3^[168]. Notably, several studies have shown that positive OipA has been more frequent in individuals with precancerous lesions than those with gastritis alone^[169-172].

The *oipA*-positive *H. pylori* strains are more prone to gastric colonization, and it is also associated with a higher risk of peptic ulcer disease and gastric cancer. This molecule strongly induces inflammation, and the infiltration of neutrophils, as well as the production of IL-8, are significantly higher in *oipA*-positive strains compared to the negative ones^[173]. Some studies indicate that OipA induces the interferon regulatory factor IRF-1, which binds and activates the element similar to the response element stimulated by IFN (ISRE), to induce the genetic transcription of IL-8 and its production^[169]. In addition, nuclear factor kappa B (NF-κB) and

activating protein 1 (AP1) are also involved in the transcription and production of IL-8 by gastric epithelial cells infected with *H. pylori*^[174].

Other proinflammatory cytokines may also be present in *H. pylori* infection caused by the presence of OipA, such as IL-1, IL-6, IL-8, IL-11 IL-17, metalloproteinase 1 matrix (MMP-1), tumor necrosis factor α (TNF- α) or CC chemokine ligand 5^[175]. This is similar to a response linked to the Cag PAI^[8]. However, depending on the OipA states in different strains of *H. pylori*, the secretion of these cytokines may not be observed^[176-178].

The genes that express functional OipA are strong factors of bacterial virulence, and are linked to the genotypes VacA s1, VacA m1, BabA2, and the Cag PAI gene, and can act synergistically with each other to induce worse clinical outcomes of diseases caused by *H. pylori*^[179].

IceA

The Gene Induced by Contact with Epithelium A (*iceA*) is a virulence marker still poorly described. The functions related to this virulence factor remain unclear and it possesses two variants: IceA1 and IceA2. *H. pylori* has only one *iceA* locus from which that protein can be expressed. Therefore, the presence of both aforementioned variations indicates an infection by different strains of the pathogen^[180].

The IceA relationship and the clinical outcomes of gastric diseases are still controversial^[181]. However, studies have emphasized that strains positive for this gene induce the release of the proinflammatory cytokines IL-6 and IL-8 more intensely than negative strains^[182,183]. Dabiri *et al* demonstrated the possible role of IceA1 in the development of gastric cancer, but not in peptic ulcers^[184]. In addition, Yakoob *et al* demonstrated that *iceA2*-positive *H. pylori* strains were more often associated with chronic active inflammation, gastric ulcer, and gastric adenocarcinoma^[185]. Furthermore, studies indicate that IceA has its function preserved regardless of the presence of other *H. pylori* virulence factors^[186-188].

To date, there seems to be a consensus that the global prevalence of IceA1 is higher than IceA2^[181]. However, although there is a greater expression of IceA1 than IceA2, the latter is associated with greater granulocytic and lymphocytic infiltration as well as atrophic gastritis^[189]. Ashwak *et al.* demonstrated in their study that *H. pylori*-infected individuals who express IceA1

or IceA2 alone do not develop gastric carcinoma. On the other hand, 75% of the patients who had both alleles (IceA1 / IceA2) concomitantly developed gastric carcinoma^[190]. Strains that have positive IceA2 tend to stimulate IL-1, resulting in an increased risk of pre-cancerous lesions in the gastric mucosa. This process can become worse if associated with the concomitant effects of other virulence factors, worsening inflammatory processes. Taken all together, the *iceA* gene is an important marker of severe gastric diseases that must be taken into account^[180].

BabA

The mechanisms related to the Blood group antigen-binding adhesin (BabA) pathogenicity are still poorly elucidated. Nonetheless, studies have shown that BabA-dependent *H. Pylori* cell adhesion has great relevance in the initial colonization of the pathogen^[191]. Moreover, BabA works by facilitating the entry of CagA and VacA into host cells^[192]. BabA-negative *H. pylori* strains have been associated with the development of mild gastric lesions and are rarely associated with gastric cancer. This means that BabA positivity might increase the risk of serious gastric lesions and carcinomas^[193].

A study showed a greater expression of IL-33 mRNA in biopsies from patients infected with *H. Pylori* compared to noninfected individuals. Interestingly, a direct relationship was observed between BabA2 and increased gastric levels of that cytokine^[194]. The IL-33 plays an important role in immune regulation, providing protection after damage to epithelial cells^[195]. It also has the potential to reduce colonization in gastrointestinal infections^[196]. In addition, recent studies emphasize its likely role in the development of tumorigenesis^[197].

SabA

Sialic acid A adhesin (SabA) is an *H. pylori* membrane protein whose expression has been explored as a biomarker for increased risk of developing gastric cancer^[198,199]. Yamaoka *et al* demonstrated that SabA is positively associated with gastric cancer, intestinal metaplasia, and body atrophy and is negatively associated with duodenal ulcer^[200]. *H. pylori* uses SabA to recognize the Lewis X antigen from gastric epithelial cells and this virulence factor has been associated with non-opsonic activation of human neutrophils^[201,202]. SabA mediates antigen binding to sialyl-Lewis, which is an established tumor and gastric dysplasia marker^[203]. The available data on this issue highlights how harmful such adhesin can be to the gastric epithelium; however, further studies are needed to better understand the underlying immune system

responses related to this molecule^[204]. Table 1 shows how the *H. pylori* virulence factors interact with the immune system.

Heat-shock protein 60

Heat-shock protein 60 (hsp60) is known to have substantial immunogenic properties. Studies have demonstrated that hsp60 promotes cell signaling upon myeloid and vascular endothelial cells^[205]. *H. pylori*-expressed hsp60 seems to play a **role in** bacterial adhesion to gastric epithelial cells and mucin^[206]. In addition, that virulence factor **has** been shown to effectively inhibit human peripheral blood mononuclear cells (PBMC). A study by Maguire *et al* has demonstrated that the inhibitor effect over human PBMC was more potent with hsp60 from *H. pylori* than hsp60 from *Chlamydia pneumoniae* or human **mitochondrion**^[207]. Evidence **has** shown that hsp60 also promotes immune system responses through the activation of TLRs in human gastric epithelial cells and induces IL-8 expression through TLR-2 and MAPK pathways in human monocytes^[208,209]. Another study evaluating the effects of hsp60 over human monocytes demonstrated that it seems to **promote upregulation** of cytokines such as IL-1a, IL-8, IL-10, IFN- γ , TNF- α , and TGF- β ^[210]. Regarding the oncogenic roles related to this molecule, an enhanced gastric cancer cell and promotion of tube formation by umbilical vein endothelial cells have been positively associated with hsp60, but effects on cell proliferation and cell death prevention have not been attributed to the protein^[211].

HomB

The *homB* gene is the open-reading frame of *jhp0870*, an *H. pylori* outer membrane protein, and its sequence is 90% similar to *homA*^[212]. The former gene has shown to be related to an increased risk of peptic ulcer development among *H. pylori*-infected people^[213]. In addition, *homB* was found to be associated with gastric corpus inflammation and atrophy, which suggests a link between this bacterial gene and gastric cancer development^[214]. Interestingly, a recently published study by Keikha and Karbalaie^[215] demonstrated that *homB*-positive *H. pylori* strains are particularly related to the occurrence of peptic ulcer in western populations and with the onset of gastric cancer in Asian regions.

6HBP

D-glycero- β -D-manno-heptose 1,7-bisphosphate (β HBP) is an *H. pylori* metabolite that was recently identified as a T4SS-dependent NF- κ B activation effector in host cells through its translocation via the ALPK1-TIFA axis^[216]. Moreover, Posselt and colleagues^[217] showed that the β HBP, as well as the protein kinase C, is able to mediate a strong c-Abl threonine 735 phosphorylation (pAbl^{T735}). Their study also demonstrated that pAbl^{T735} interacts with 14-3-3 proteins, leading to c-ABL retention in the cytoplasm, which contributes to cell elongation and migration. In addition, they observed an attenuation of caspase-8 and caspase-9-dependent cell death by pAbl^{T735} nuclear exclusion^[217]. These findings suggest a relationship between β HBP and important *H. pylori*-related pro-oncogenic activities.

CONCLUSION

The knowledge on the relationship between *H. pylori* infection, immune system, and oncogenesis is crucial for the understanding of the mechanisms involved in gastric cancer. Although considerable advances have been achieved in this research field, much has to be done in order to describe underlying mechanisms related to *H. pylori*-related carcinogenesis. A better comprehension on this issue could be useful for the development of tools that may aid in the prevention as well as in the prognostic prediction and treatment of such an important disease. Here, we gathered data showing that *H. pylori* infection promotes multiple immune response activities, such as T helper cell polarization, that are closely related to mechanisms associated with gastric carcinogenesis.

REFERENCES

- 1 **Yang L**, Ying X, Liu S, Lyu G, Xu Z, Zhang X, Li H, Li Q, Wang N, Ji J. Gastric cancer: Epidemiology, risk factors and prevention strategies. *Chin J Cancer Res* 2020; **32**: 695-704 [PMID: 33446993 DOI: 10.21147/j.issn.1000-9604.2020.06.03]
- 2 **Rawla P**, Barsouk A. Epidemiology of gastric cancer: global trends, risk factors and prevention. *Prz Gastroenterol* 2019; **14**: 26-38 [PMID: 30944675 DOI: 10.5114/pg.2018.80001]
- 3 **Wroblewski LE**, Peek RM Jr. Helicobacter pylori, Cancer, and the Gastric Microbiota. *Adv Exp Med Biol* 2016; **908**: 393-408 [PMID: 27573782 DOI: 10.1007/978-3-319-41388-4_19]
- 4 **Camilo V**, Sugiyama T, Touati E. Pathogenesis of Helicobacter pylori infection. *Helicobacter* 2017; **22 Suppl 1** [PMID: 28891130 DOI: 10.1111/hel.12405]

- 5 **Smith SM**. Role of Toll-like receptors in *Helicobacter pylori* infection and immunity. *World J Gastrointest Pathophysiol* 2014; **5**: 133-146 [PMID: 25133016 DOI: 10.4291/wjgp.v5.i3.133.]
- 6 **Wang F**, Meng W, Wang B, Qiao L. *Helicobacter pylori*-induced gastric inflammation and gastric cancer. *Cancer Lett* 2014; **345**: 196-202 [PMID: 23981572 DOI: 10.1016/j.canlet.2013.08.016]
- 7 **Chang WL**, Yeh YC, Sheu BS. The impacts of *H. pylori* virulence factors on the development of gastroduodenal diseases. *J Biomed Sci* 2018; **25**: 68 [PMID: 30205817 DOI: 10.1186/s12929-018-0466-9]
- 8 **de Brito BB**, da Silva FAF, de Melo FF. Role of polymorphisms in genes that encode cytokines and *Helicobacter pylori* virulence factors in gastric carcinogenesis. *World J Clin Oncol* 2018; **9**: 83-89 [PMID: 30254963 DOI: 10.5306/wjco.v9.i5.83]
- 9 **Yoshikawa T**, Naito Y. The role of neutrophils and inflammation in gastric mucosal injury. *Free Radic Res* 2000; **33**: 785-794 [PMID: 11237100 DOI: 10.1080/10715760000301301.]
- 10 **Crabtree JE**, Mahony MJ, Taylor JD, Heatley RV, Littlewood JM, Tompkins DS. Immune responses to *Helicobacter pylori* in children with recurrent abdominal pain. *J Clin Pathol* 1991; **44**: 768-771 [PMID: 1918408 DOI: 10.1136/jcp.44.9.768]
- 11 **Kawai T**, Akira S. The roles of TLRs, RLRs and NLRs in pathogen recognition. *Int Immunol* 2009; **21**: 317-337 [PMID: 19246554 DOI: 10.1093/intimm/dxp017.]
- 12 **Suarez G**, Reyes VE, Beswick EJ. Immune response to *H. pylori*. *World J Gastroenterol* 2006; **12**: 5593-5598 [PMID: 17007009 DOI: 10.3748/wjg.v12.i35.5593.]
- 13 **Wilson KT**, Crabtree JE. Immunology of *Helicobacter pylori*: insights into the failure of the immune response and perspectives on vaccine studies. *Gastroenterology* 2007; **133**: 288-308 [PMID: 17631150 DOI: 10.1053/j.gastro.2007.05.008.]
- 14 **Nagashima H**, Iwatani S, Cruz M, Jiménez Abreu JA, Uchida T, Mahachai V, Vilaichone RK, Graham DY, Yamaoka Y. Toll-like Receptor 10 in *Helicobacter pylori* Infection. *J Infect Dis* 2015; **212**: 1666-1676 [PMID: 25977263 DOI: 10.1093]
- 15 **Meliț LE**, Mărginean CO, Mărginean CD, Mărginean MO. The Relationship between Toll-like Receptors and *Helicobacter pylori*-Related Gastropathies: Still a Controversial Topic. *J Immunol Res* 2019; **2019**: 8197048 [PMID: 30863783 DOI: 10.1155/2019/8197048.]

- 16 **Smith SM**. Role of Toll-like receptors in *Helicobacter pylori* infection and immunity. *World J Gastrointest Pathophysiol* 2014; **5**: 133-146 [PMID: 25133016 DOI: 10.4291/wjgp.v5.i3.133.]
- 17 **Cadamuro AC**, Rossi AF, Maniezzo NM, Silva AE. *Helicobacter pylori* infection: host immune response, implications on gene expression and microRNAs. *World J Gastroenterol* 2014; **20**: 1424-1437 [PMID: 24587619 DOI: 10.3748/wjg.v20.i6.1424.]
- 18 **Ding SZ**, Torok AM, Smith MF Jr, Goldberg JB. Toll-like receptor 2-mediated gene expression in epithelial cells during *Helicobacter pylori* infection. *Helicobacter* 2005; **10**: 193-204 [PMID: 15904477 DOI: 10.1111/j.1523-5378.2005.00311.x.]
- 19 **Kawai T**, Akira S. Toll-like receptors and their crosstalk with other innate receptors in infection and immunity. *Immunity*. 2011 May 27;34(5):637-50. [: 21616434. DOI: 10.1016/j.immuni.2011.05.006.]
- 20 **Masumoto J**, Yang K, Varambally S, Hasegawa M, Tomlins SA, Qiu S, Fujimoto Y, Kawasaki A, Foster SJ, Horie Y, Mak TW, Núñez G, Chinnaiyan AM, Fukase K, Inohara N. Nod1 acts as an intracellular receptor to stimulate chemokine production and neutrophil recruitment in vivo. *J Exp Med* 2006; **203**: 203-213 [PMID: 16418393 DOI: 10.1084/jem.20051229.]
- 21 **Fritz JH**, Le Bourhis L, Sellge G, Magalhaes JG, Fsihi H, Kufer TA, Collins C, Viala J, Ferrero RL, Girardin SE, Philpott DJ. Nod1-mediated innate immune recognition of peptidoglycan contributes to the onset of adaptive immunity. *Immunity* 2007; **26**: 445-459 [PMID: 17433730 DOI: 10.1016/j.immuni.2007.03.009.]
- 22 **Kumar Pachathundikandi S**, Brandt S, Madassery J, Backert S. Induction of TLR-2 and TLR-5 expression by *Helicobacter pylori* switches cagPAI-dependent signalling leading to the secretion of IL-8 and TNF- α . *PLoS One* 2011; **6**: e19614 [PMID: 21573018 DOI: 10.1371/journal.pone.0019614.]
- 23 **Jung HC**, Kim JM, Song IS, Kim CY. *Helicobacter pylori* induces an array of pro-inflammatory cytokines in human gastric epithelial cells: quantification of mRNA for interleukin-8, -1 α /beta, granulocyte-macrophage colony-stimulating factor, monocyte chemoattractant protein-1 and tumour necrosis factor- α . *J Gastroenterol Hepatol* 1997; **12**: 473-480 [PMID: 9257236 DOI: 10.1111/j.1440-1746.1997.tb00469.x.]

- 24 **George JT**, Boughan PK, Karageorgiou H, Bajaj-Elliott M. Host anti-microbial response to *Helicobacter pylori* infection. *Mol Immunol* 2003; **40**: 451-456 [PMID: 14568391 DOI: 10.1016/s0161-5890(03)00158-5.]
- 25 **Baltimore D**, Boldin MP, O'Connell RM, Rao DS, Taganov KD. MicroRNAs: new regulators of immune cell development and function. *Nat Immunol* 2008; **9**: 839-845 [PMID: 18645592 DOI: 10.1038/ni.f.209.]
- 26 **Crabtree JE**. Gastric mucosal inflammatory responses to *Helicobacter pylori*. *Aliment Pharmacol Ther* 1996; **10 Suppl 1**: 29-37 [PMID: 8730257 DOI: 10.1046/j.1365-2036.1996.22164003.x.]
- 27 **Andres S**, Schmidt HM, Mitchell H, Rhen M, Maeurer M, Engstrand L. *Helicobacter pylori* defines local immune response through interaction with dendritic cells. *FEMS Immunol Med Microbiol* 2011; **61**: 168-178 [PMID: 21175878 DOI: 10.1111/j.1574-695X.2010.00761.x.]
- 28 **Algood HM**, Cover TL. *Helicobacter pylori* persistence: an overview of interactions between *H. pylori* and host immune defenses. *Clin Microbiol Rev* 2006; **19**: 597-613 [PMID: 17041136 DOI: 10.1128/CMR.00006-06.]
- 29 **Evans DJ Jr**, Evans DG, Takemura T, Nakano H, Lampert HC, Graham DY, Granger DN, Kvietys PR. Characterization of a *Helicobacter pylori* neutrophil-activating protein. *Infect Immun* 1995; **63**: 2213-2220 [PMID: 7768601 DOI: 10.1128/IAI.63.6.2213-2220.1995.]
- 30 **Satin B**, Del Giudice G, Della Bianca V, Dusi S, Laudanna C, Tonello F, Kelleher D, Rappuoli R, Montecucco C, Rossi F. The neutrophil-activating protein (HP-NAP) of *Helicobacter pylori* is a protective antigen and a major virulence factor. *J Exp Med* 2000; **191**: 1467-1476 [PMID: 10790422 DOI: 10.1084/jem.191.9.1467.]
- 31 **Amedei A**, Cappon A, Codolo G, Cabrelle A, Polenghi A, Benagiano M, Tasca E, Azzurri A, D'Elios MM, Del Prete G, de Bernard M. The neutrophil-activating protein of *Helicobacter pylori* promotes Th1 immune responses. *J Clin Invest* 2006; **116**: 1092-1101 [PMID: 16543949 DOI: 10.1172/JCI27177.]
- 32 **Montecucco C**, de Bernard M. Molecular and cellular mechanisms of action of the vacuolating cytotoxin (VacA) and neutrophil-activating protein (HP-NAP) virulence factors of *Helicobacter pylori*. *Microbes Infect* 2003; **5**: 715-721 [PMID: 12814772 DOI: 10.1016/s1286-4579(03)00124-2.]

- 33 **Robinson K**, Argent RH, Atherton JC. The inflammatory and immune response to *Helicobacter pylori* infection. *Best Pract Res Clin Gastroenterol* 2007; **21**: 237-259 [PMID: 17382275 DOI: 10.1016/j.bpg.2007.01.001.]
- 34 **Gobert AP**, Mersey BD, Cheng Y, Blumberg DR, Newton JC, Wilson KT. Cutting edge: urease release by *Helicobacter pylori* stimulates macrophage inducible nitric oxide synthase. *J Immunol* 2002; **168**: 6002-6006 [PMID: 12055207 DOI: 10.4049/jimmunol.168.12.6002.]
- 35 **Muzio M**, Bosisio D, Polentarutti N, D'amico G, Stoppacciaro A, Mancinelli R, van't Veer C, Penton-Rol G, Ruco LP, Allavena P, Mantovani A. Differential expression and regulation of toll-like receptors (TLR) in human leukocytes: selective expression of TLR3 in dendritic cells. *J Immunol* 2000; **164**: 5998-6004 [PMID: 10820283 DOI: 10.4049/jimmunol.164.11.5998.]
- 36 **Rad R**, Brenner L, Krug A, Volland P, Mages J, Lang R, Schwendy S, Reindl W, Dossumbekova A, Ballhorn W, Wagner H, Schmid RM, Bauer S, Prinz C. Toll-like receptor-dependent activation of antigen-presenting cells affects adaptive immunity to *Helicobacter pylori*. *Gastroenterology* 2007; **133**: 150-163.e3 [PMID: 17631139 DOI: 10.1053/j.gastro.2007.04.071]
- 37 **Haeberle HA**, Kubin M, Bamford KB, Garofalo R, Graham DY, El-Zaatari F, Karttunen R, Crowe SE, Reyes VE, Ernst PB. Differential stimulation of interleukin-12 (IL-12) and IL-10 by live and killed *Helicobacter pylori* *in vitro* and association of IL-12 production with gamma interferon-producing T cells in the human gastric mucosa. *Infect Immun* 1997; **65**: 4229-4235 [PMID: 9317031 DOI: 10.1128/IAI.65.10.4229-4235.1997.]
- 38 **Meyer F**, Wilson KT, James SP. Modulation of innate cytokine responses by products of *Helicobacter pylori*. *Infect Immun* 2000; **68**: 6265-6272 [PMID: 11035734 DOI: 10.1128/iai.68.11.6265-6272.2000.]
- 39 **Guiney DG**, Hasegawa P, Cole SP. *Helicobacter pylori* preferentially induces interleukin 12 (IL-12) rather than IL-6 or IL-10 in human dendritic cells. *Infect Immun* 2003; **71**: 4163-4166 [PMID: 12819109 DOI: 10.1128/iai.71.7.4163-4166.2003;]
- 40 **Kranzer K**, Eckhardt A, Aigner M, Knoll G, Deml L, Speth C, Lehn N, Rehli M, Schneider-Brachert W. Induction of maturation and cytokine release of human dendritic cells by *Helicobacter pylori*. *Infect Immun* 2004; **72**: 4416-4423 [PMID: 15271898 DOI: 10.1128/IAI.72.8.4416-4423.2004.]

- 41 **Lindholm C**, Quiding-Järbrink M, Lönroth H, Hamlet A, Svennerholm AM. Local cytokine response in *Helicobacter pylori*-infected subjects. *Infect Immun* 1998; **66**: 5964-5971 [PMID: 9826379 DOI: 10.1128/IAI.66.12.5964-5971.1998.]
- 42 **Nakajima S**, Krishnan B, Ota H, Segura AM, Hattori T, Graham DY, Genta RM. Mast cell involvement in gastritis with or without *Helicobacter pylori* infection. *Gastroenterology* 1997; **113**: 746-754 [PMID: 9287964 DOI: 10.1016/s0016-5085(97)70167-7.]
- 43 **de Bernard M**, Cappon A, Pancotto L, Ruggiero P, Rivera J, Del Giudice G, Montecucco C. The *Helicobacter pylori* VacA cytotoxin activates RBL-2H3 cells by inducing cytosolic calcium oscillations. *Cell Microbiol* 2005; **7**: 191-198 [PMID: 15659063 DOI: 10.1111/j.1462-5822.2004.00446.x.]
- 44 **Crabtree JE**, Shallcross TM, Heatley RV, Wyatt JL. Mucosal tumour necrosis factor alpha and interleukin-6 in patients with *Helicobacter pylori* associated gastritis. *Gut* 1991; **32**: 1473-1477 [PMID: 1773951 DOI: 10.1136/gut.32.12.1473.]
- 45 **Moss SF**, Legon S, Davies J, Calam J. Cytokine gene expression in *Helicobacter pylori* associated antral gastritis. *Gut* 1994; **35**: 1567-1570 [PMID: 7828974 DOI: 10.1136/gut.35.11.1567.]
- 46 **Soehnlein O**, Lindbom L. Phagocyte partnership during the onset and resolution of inflammation. *Nat Rev Immunol* 2010; **10**: 427-439 [PMID: 20498669 DOI: 10.1038/nri2779.]
- 47 **Freire de Melo F**, Rocha AM, Rocha GA, Pedroso SH, de Assis Batista S, Fonseca de Castro LP, Carvalho SD, Bittencourt PF, de Oliveira CA, Corrêa-Oliveira R, Magalhães Queiroz DM. A regulatory instead of an IL-17 T response predominates in *Helicobacter pylori*-associated gastritis in children. *Microbes Infect* 2012; **14**: 341-347 [PMID: 22155622 DOI: 10.1016/j.micinf.2011.11.008.]
- 48 **Lundgren A**, Suri-Payer E, Enarsson K, Svennerholm AM, Lundin BS. *Helicobacter pylori*-specific CD4⁺ CD25^{high} regulatory T cells suppress memory T-cell responses to H. pylori in infected individuals. *Infect Immun* 2003; **71**: 1755-1762 [PMID: 12654789 DOI: 10.1128/iai.71.4.1755-1762.2003.]
- 49 **Lundgren A**, Suri-Payer E, Enarsson K, Svennerholm AM, Lundin BS. *Helicobacter pylori*-specific CD4⁺ CD25^{high} regulatory T cells suppress memory T-cell responses to H. pylori in

infected individuals. *Infect Immun* 2003; **71**: 1755-1762 [PMID: 12654789 DOI: 10.1128/iai.71.4.1755-1762.2003]

50 **Su Z**, Sun Y, Zhu H, Liu Y, Lin X, Shen H, Chen J, Xu W, Xu H. Th17 cell expansion in gastric cancer may contribute to cancer development and metastasis. *Immunol Res* 2014; **58**: 118-124 [PMID: 24402773 DOI: 10.1007/s12026-013-8483-y]

51 **Lindholm C**, Quiding-Järbrink M, Lönroth H, Hamlet A, Svennerholm AM. Local cytokine response in Helicobacter pylori-infected subjects. *Infect Immun* 1998; **66**: 5964-5971 [PMID: 9826379 DOI: 10.1128/IAI.66.12.5964-5971.1998]

52 **El-Omar EM**. The importance of interleukin 1beta in Helicobacter pylori associated disease. *Gut* 2001; **48**: 743-747 [PMID: 11358884 DOI: 10.1136/gut.48.6.743]

53 **Freire de Melo F**, Rocha GA, Rocha AM, Teixeira KN, Pedrosa SH, Pereira Junior JB, Fonseca de Castro LP, Cabral MM, Carvalho SD, Bittencourt PF, de Oliveira CA, Queiroz DM. Th1 immune response to H. pylori infection varies according to the age of the patients and influences the gastric inflammatory patterns. *Int J Med Microbiol* 2014; **304**: 300-306 [PMID: 24373859 DOI: 10.1016/j.ijmm.2013.11.001]

54 **Zhang S**. The role of transforming growth factor β in T helper 17 differentiation. *Immunology* 2018; **155**: 24-35 [PMID: 29682722 DOI: 10.1111/imm.12938]

55 **Harbour SN**, DiToro DF, Witte SJ, Zindl CL, Gao M, Schoeb TR, Jones GW, Jones SA, Hatton RD, Weaver CT. T_H17 cells require ongoing classic IL-6 receptor signaling to retain transcriptional and functional identity. *Sci Immunol* 2020; **5** [PMID: 32680955 DOI: 10.1126/sciimmunol.aaw2262]

56 **Horvath DJ Jr**, Washington MK, Cope VA, Algood HM. IL-23 Contributes to Control of Chronic Helicobacter Pylori Infection and the Development of T Helper Responses in a Mouse Model. *Front Immunol* 2012; **3**: 56 [PMID: 22566937 DOI: 10.3389/fimmu.2012.00056]

57 **Caruso R**, Pallone F, Monteleone G. Emerging role of IL-23/IL-17 axis in H pylori-associated pathology. *World J Gastroenterol* 2007; **13**: 5547-5551 [PMID: 17948927 DOI: 10.3748/wjg.v13.i42.5547]

- 58 **Algood HM**, Allen SS, Washington MK, Peek RM Jr, Miller GG, Cover TL. Regulation of gastric B cell recruitment is dependent on IL-17 receptor A signaling in a model of chronic bacterial infection. *J Immunol* 2009; **183**: 5837-5846 [PMID: 19812196 DOI: 10.4049/jimmunol.0901206]
- 59 **Chen W**, Qin Y, Liu S. CCL20 Signaling in the Tumor Microenvironment. *Adv Exp Med Biol* 2020; **1231**: 53-65 [PMID: 32060846 DOI: 10.1007/978-3-030-36667-4_6]
- 60 **Bronte V**. Th17 and cancer: friends or foes? *Blood* 2008; **112**: 214 [PMID: 18606882 DOI: 10.1182/blood-2008-04-149260]
- 61 **Bending D**, De la Peña H, Veldhoen M, Phillips JM, Uyttenhove C, Stockinger B, Cooke A. Highly purified Th17 cells from BDC2.5NOD mice convert into Th1-like cells in NOD/SCID recipient mice. *J Clin Invest* 2009; **119**: 565-572 [PMID: 19188681 DOI: 10.1172/JCI37865]
- 62 **Lee YK**, Turner H, Maynard CL, Oliver JR, Chen D, Elson CO, Weaver CT. Late developmental plasticity in the T helper 17 Lineage. *Immunity* 2009; **30**: 92-107 [PMID: 19119024 DOI: 10.1016/j.immuni.2008.11.005]
- 63 **Muranski P**, Borman ZA, Kerkar SP, Klebanoff CA, Ji Y, Sanchez-Perez L, Sukumar M, Reger RN, Yu Z, Kern SJ, Roychoudhuri R, Ferreyra GA, Shen W, Durum SK, Feigenbaum L, Palmer DC, Antony PA, Chan CC, Laurence A, Danner RL, Gattinoni L, Restifo NP. Th17 cells are long lived and retain a stem cell-like molecular signature. *Immunity* 2011; **35**: 972-985 [PMID: 22177921 DOI: 10.1016/j.immuni.2011.09.019]
- 64 **Qian X**, Chen H, Wu X, Hu L, Huang Q, Jin Y. Interleukin-17 acts as double-edged sword in anti-tumor immunity and tumorigenesis. *Cytokine* 2017; **89**: 34-44 [PMID: 26883678 DOI: 10.1016/j.cyto.2015.09.011]
- 65 **Kamran MZ**, Patil P, Gude RP. Role of STAT3 in cancer metastasis and translational advances. *Biomed Res Int* 2013; **2013**: 421821 [PMID: 24199193 DOI: 10.1155/2013/421821]
- 66 **Manel N**, Unutmaz D, Littman DR. The differentiation of human T(H)-17 cells requires transforming growth factor-beta and induction of the nuclear receptor RORgamma. *Nat Immunol* 2008; **9**: 641-649 [PMID: 18454151 DOI: 10.1038/ni.1610]
- 67 **Zhang X**, Yang F, Wang Z. Tumor microenvironment characterization in stage IV gastric cancer. *Biosci Rep* 2021; **41** [PMID: 33416081 DOI: 10.1042/BSR20201248]

68 **Zheng X**, Dong L, Wang K, Zou H, Zhao S, Wang Y, Wang G. MiR-21 Participates in the PD-1/PD-L1 Pathway-Mediated Imbalance of Th17/Treg Cells in Patients After Gastric Cancer Resection. *Ann Surg Oncol* 2019; **26**: 884-893 [PMID: 30565043 DOI: 10.1245/s10434-018-07117-6]

69 **Takeda A**, Hamano S, Yamanaka A, Hanada T, Ishibashi T, Mak TW, Yoshimura A, Yoshida H. Cutting edge: role of IL-27/WSX-1 signaling for induction of T-bet through activation of STAT1 during initial Th1 commitment. *J Immunol* 2003; **170**: 4886-4890 [PMID: 12734330 DOI: 10.4049/jimmunol.170.10.4886]

70 **Diveu C**, McGeachy MJ, Boniface K, Stumhofer JS, Sathe M, Joyce-Shaikh B, Chen Y, Tato CM, McClanahan TK, de Waal Malefyt R, Hunter CA, Cua DJ, Kastelein RA. IL-27 blocks RORc expression to inhibit lineage commitment of Th17 cells. *J Immunol* 2009; **182**: 5748-5756 [PMID: 19380822 DOI: 10.4049/jimmunol.0801162]

71 **Rocha GA**, de Melo FF, Cabral MMDA, de Brito BB, da Silva FAF, Queiroz DMM. Interleukin-27 is abrogated in gastric cancer, but highly expressed in other *Helicobacter pylori*-associated gastroduodenal diseases. *Helicobacter* 2020; **25**: e12667 [PMID: 31702083 DOI: 10.1111/hel.12667]

72 **Kandulski A**, Malfertheiner P, Wex T. Role of regulatory T-cells in *H. pylori*-induced gastritis and gastric cancer. *Anticancer Res* 2010; **30**: 1093-1103 [PMID: 20530414]

73 **Shen LS**, Wang J, Shen DF, Yuan XL, Dong P, Li MX, Xue J, Zhang FM, Ge HL, Xu D. CD4(+)CD25(+)CD127(low/-) regulatory T cells express Foxp3 and suppress effector T cell proliferation and contribute to gastric cancers progression. *Clin Immunol* 2009; **131**: 109-118 [PMID: 19153062 DOI: 10.1016/j.clim.2008.11.010]

74 **Enarsson K**, Lundgren A, Kindlund B, Hermansson M, Roncador G, Banham AH, Lundin BS, Quiding-Järbrink M. Function and recruitment of mucosal regulatory T cells in human chronic *Helicobacter pylori* infection and gastric adenocarcinoma. *Clin Immunol* 2006; **121**: 358-368 [PMID: 16934529 DOI: 10.1016/j.clim.2006.07.002]

75 **Heiber JF**, Geiger TL. Context and location dependence of adaptive Foxp3(+) regulatory T cell formation during immunopathological conditions. *Cell Immunol* 2012; **279**: 60-65 [PMID: 23089195 DOI: 10.1016/j.cellimm.2012.09.009]

- 76 **Gil JH**, Seo JW, Cho MS, Ahn JH, Sung HY. Role of Treg and TH17 cells of the gastric mucosa in children with Helicobacter pylori gastritis. *J Pediatr Gastroenterol Nutr* 2014; **58**: 245-251 [PMID: 24121150 DOI: 10.1097/MPG.0000000000000194]
- 77 **Liu X**, Xu D, Huang C, Guo Y, Wang S, Zhu C, Xu J, Zhang Z, Shen Y, Zhao W, Zhao G. Regulatory T cells and M2 macrophages present diverse prognostic value in gastric cancer patients with different clinicopathologic characteristics and chemotherapy strategies. *J Transl Med* 2019; **17**: 192 [PMID: 31174544 DOI: 10.1186/s12967-019-1929-9]
- 78 **Ooi CH**, Ivanova T, Wu J, Lee M, Tan IB, Tao J, Ward L, Koo JH, Gopalakrishnan V, Zhu Y, Cheng LL, Lee J, Rha SY, Chung HC, Ganesan K, So J, Soo KC, Lim D, Chan WH, Wong WK, Bowtell D, Yeoh KG, Grabsch H, Boussioutas A, Tan P. Oncogenic pathway combinations predict clinical prognosis in gastric cancer. *PLoS Genet* 2009; **5**: e1000676 [PMID: 19798449 DOI: 10.1371/journal.pgen.1000676]
- 79 **Ji L**, Qian W, Gui L, Ji Z, Yin P, Lin GN, Wang Y, Ma B, Gao WQ. Blockade of β -Catenin-Induced CCL28 Suppresses Gastric Cancer Progression *via* Inhibition of Treg Cell Infiltration. *Cancer Res* 2020; **80**: 2004-2016 [PMID: 32156780 DOI: 10.1158/0008-5472.CAN-19-3074]
- 80 **Tse BWC**, Volpert M, Ratther E, Stylianou N, Nouri M, McGowan K, Lehman ML, McPherson SJ, Roshan-Moniri M, Butler MS, Caradec J, Gregory-Evans CY, McGovern J, Das R, Takhar M, Erho N, Alshalafa M, Davicioni E, Schaeffer EM, Jenkins RB, Ross AE, Karnes RJ, Den RB, Fazli L, Gregory PA, Gleave ME, Williams ED, Rennie PS, Buttyan R, Gunter JH, Selth LA, Russell PJ, Nelson CC, Hollier BG. Neuropilin-1 is upregulated in the adaptive response of prostate tumors to androgen-targeted therapies and is prognostic of metastatic progression and patient mortality. *Oncogene* 2017; **36**: 3417-3427 [PMID: 28092670 DOI: 10.1038/onc.2016.482]
- 81 **Kang JY**, Gil M, Kim KE. Neuropilin1 Expression Acts as a Prognostic Marker in Stomach Adenocarcinoma by Predicting the Infiltration of Treg Cells and M2 Macrophages. *J Clin Med* 2020; **9** [PMID: 32408477 DOI: 10.3390/jcm9051430]
- 82 **Liu XS**, Lin XK, Mei Y, Ahmad S, Yan CX, Jin HL, Yu H, Chen C, Lin CZ, Yu JR. Regulatory T Cells Promote Overexpression of Lgr5 on Gastric Cancer Cells *via* TGF-beta1 and Confer Poor Prognosis in Gastric Cancer. *Front Immunol* 2019; **10**: 1741 [PMID: 31417548 DOI: 10.3389/fimmu.2019.01741]

- 83 **Simon E**, Petke D, Böger C, Behrens HM, Warneke V, Ebert M, Röcken C. The spatial distribution of LGR5+ cells correlates with gastric cancer progression. *PLoS One* 2012; **7**: e35486 [PMID: 22530031 DOI: 10.1371/journal.pone.0035486]
- 84 **Schumacher A**, Schwarz R. [Histomorphologic and catamnestic studies of 226 patients with cervical carcinoma in stage Ia in the years 1966 to 1986]. *Zentralbl Allg Pathol* 1989; **135**: 639-648 [PMID: 2588831 DOI: 10.1016/j.jmii.2015.03.0029]
- 85 **Chang YW**, Oh CH, Kim JW, Lee JW, Park MJ, Shim JJ, Lee CK, Jang JY, Dong SH, Kim HJ, Kim SS, Kim BH. Combination of *Helicobacter pylori* infection and the interleukin 8 -251 T > A polymorphism, but not the mannose-binding lectin 2 codon 54 G > A polymorphism, might be a risk factor of gastric cancer. *BMC Cancer* 2017; **17**: 388 [PMID: 28558668 DOI: 10.1186/s12885-017-3378-2]
- 86 **de Brito BB**, da Silva FAF, de Melo FF. Role of polymorphisms in genes that encode cytokines and *Helicobacter pylori* virulence factors in gastric carcinogenesis. *World J Clin Oncol* 2018; **9**: 83-89 [PMID: 30254963 DOI: 10.5306/wjco.v9.i5.83]
- 87 **Garlanda C**, Dinarello CA, Mantovani A. The interleukin-1 family: back to the future. *Immunity* 2013; **39**: 1003-1018 [PMID: 24332029 DOI: 10.1016/j.immuni.2013.11.010]
- 88 **El-Omar EM**, Carrington M, Chow WH, McColl KE, Bream JH, Young HA, Rabkin CS. Interleukin-1 polymorphisms associated with increased risk of gastric cancer. *Nature* 2000; **404**: 398-402. [PMID: 10746728 DOI:10.1038/35006081]
- 89 **Machado JC**, Pharoah P, Sousa S, Carvalho R, Oliveira C, Figueiredo C, Sobrinho-Simões M. Interleukin 1B and interleukin 1RN polymorphisms are associated with increased risk of gastric carcinoma. *Gastroenterology* 2001; **121**: 823-829. [PMID: 11606496 DOI:10.1053/gast.2001.28000]
- 90 **Drici AEM**, Moulessehoul S, Tifrit A, Diaf M, Turki DK, Bachir M, Tou A. Effect of IL-1 β and IL-1RN polymorphisms in carcinogenesis of the gastric mucosa in patients infected with *Helicobacter pylori* in Algeria. *Libyan J Med* 2016; **11**: 1-7. [PMID: 27340011 DOI:10.3402/Ljm.v11.31576]
- 91 **Irtiza S**, Samie AU, Ali S, Siddiqi MA, Naqash SH, Sameer AS. IL-1 β polymorphism and expression associated with decreased risk of gastric carcinoma: a case control study in the ethnic

Kashmiri population, India. *Asian Pac J Cancer Prev* 2015; **16**: 1987-1992 [PMID: 25773799 DOI: 10.7314/apjcp.2015.16.5.1987]

92 **Ruzzo A**, Graziano F, Pizzagalli F, Santini D, Battistelli V, Panunzi S, Canestrari E, Catalano V, Humar B, Ficarelli R, Bearzi I, Cascinu S, Naldi N, Testa E, Magnani M. Interleukin 1B gene (IL-1B) and interleukin 1 receptor antagonist gene (IL-1RN) polymorphisms in Helicobacter pylori-negative gastric cancer of intestinal and diffuse histotype. *Ann Oncol* 2005; **16**: 887-892 [PMID: 15851404 DOI: 10.1093/annonc/mdi184]

93 **Dinarello CA**. Biologic basis for interleukin-1 in disease. *Blood* 1996; **87**: 2095-2147 [PMID: 8630372]

94 **Hollegaard MV**, Bidwell JL. Cytokine gene polymorphism in human disease: on-line databases, Supplement 3. *Genes Immun* 2006; **7**: 269-276 [PMID: 16642032 DOI: 10.1038/sj.gene.6364301]

95 **Sicinschi LA**, Lopez-Carrillo L, Camargo MC, Correa P, Sierra RA, Henry RR, Chen J, Zabaleta J, Piazuelo MB, Schneider BG. Gastric cancer risk in a Mexican population: role of Helicobacter pylori CagA positive infection and polymorphisms in interleukin-1 and -10 genes. *Int J Cancer* 2006; **118**: 649-657 [PMID: 16114018 DOI: 10.1002/ijc.21364]

96 **Garza-González E**, Bosques-Padilla FJ, El-Omar E, Hold G, Tijerina-Menchaca R, Maldonado-Garza HJ, Pérez-Pérez GI. Role of the polymorphic IL-1B, IL-1RN and TNF-A genes in distal gastric cancer in Mexico. *Int J Cancer* 2005; **114**: 237-241 [PMID: 15540224 DOI: 10.1002/ijc.20718]

97 **Chang YW**, Jang JY, Kim NH, Lee JW, Lee HJ, Jung WW, Dong SH, Kim HJ, Kim BH, Lee JL, Chang R. Interleukin-1B (IL-1B) polymorphisms and gastric mucosal levels of IL-1beta cytokine in Korean patients with gastric cancer. *Int J Cancer* 2005; **114**: 465-471 [PMID: 15551344 DOI: 10.1002/ijc.20724]

98 **Zeng ZR**, Hu PJ, Hu S, Pang RP, Chen MH, Ng M, Sung JJ. Association of interleukin 1B gene polymorphism and gastric cancers in high and low prevalence regions in China. *Gut* 2003; **52**: 1684-1689 [PMID: 14633943 DOI: 10.1136/gut.52.12.1684]

- 99 **Ismaili A**, Yari K, Moradi MT, Sohrabi M, Kahrizi D, Kazemi E, Souri Z. IL-1B (C+3954T) gene polymorphism and susceptibility to gastric cancer in the Iranian population. *Asian Pac J Cancer Prev* 2015; **16**: 841-844 [PMID: 25684535 DOI: 10.7314/apjcp.2015.16.2.84]
- 100 **Holds JB**, Fogg SG, Anderson RL. Botulinum A toxin injection. Failures in clinical practice and a biomechanical system for the study of toxin-induced paralysis. *Ophthalmic Plast Reconstr Surg* 1990; **6**: 252-259 [PMID: 2271481 DOI: 10.1590/s0100-879x2012007500099]
- 101 **Gehmert S**, Velapatiño B, Herrera P, Balqui J, Santivañez L, Cok J, Vargas G, Combe J, Passaro DJ, Wen S, Meyer F, Berg DE, Gilman RH. Interleukin-1 beta single-nucleotide polymorphism's C allele is associated with elevated risk of gastric cancer in Helicobacter pylori-infected Peruvians. *Am J Trop Med Hyg* 2009; **81**: 804-810 [PMID: 19861615 DOI: 10.4269/ajtmh.2009.08-0494]
- 102 **Hurme M**, Santtila S. IL-1 receptor antagonist (IL-1Ra) plasma levels are co-ordinately regulated by both IL-1Ra and IL-1beta genes. *Eur J Immunol* 1998; **28**: 2598-2602 [PMID: 9710237 DOI: 10.1002/(SICI)1521-4141(199808)28:08<2598::AID-IMMU2598>3.0.CO;2-K]
- 103 **Siech M**, Letko G. Influence of ethanol on survival of acinar cells isolated from rat pancreas. *Res Exp Med (Berl)* 1992; **192**: 57-63 [PMID: 1570415 DOI: 10.1002/ijc.20935.4]
- 104 **Ingerslev J**, Wallevik K. Clinical experience with Hemofil M in a hemophilia patient exhibiting inhibitors. *Ann Hematol* 1991; **63**: 152-154 [PMID: 1932291 DOI: 10.3748/wjg.15.1465.]
- 105 **Abbasian MH**, Abbasi B, Ansarinejad N, Motevalizadeh Ardekani A, Samizadeh E, Gohari Moghaddam K, Hashemi MR. Association of interleukin-1 gene polymorphism with risk of gastric and colorectal cancers in an Iranian population. *Iran J Immunol* 2018; **15**: 321-328 [PMID: 30593746 DOI: 10.22034/IJI.2018.39401]
- 106 **Moghim M**, Dastgheib SA, Heiranizadeh N, Zare M, Sheikhpour E, Neamatzadeh H. ASSOCIATION OF IL-8 -251T>A (RS4073) POLYMORPHISM WITH SUSCEPTIBILITY TO GASTRIC CANCER: A SYSTEMATIC REVIEW AND META-ANALYSIS BASED ON 33 CASE-CONTROL STUDIES. *Arq Gastroenterol* 2020; **57**: 91-99. [DOI: 10.1590/s0004-2803.202000000-16]
- 107 **Felipe AV**, Silva TD, Pimenta CA, Kassab P, Forones NM. Interleukin-8 gene polymorphism and susceptibility to gastric cancer in a brazilian population. *Biol Res* 2012; **45**: 369-374. [PMID: 23558993 DOI:10.4067/S0716-97602012000400007]

- 108 **Zhang Y**, Zeng X, Lu H, Li Y, Ji H. Association between Interleukin-8-251A/T polymorphism and gastric cancer susceptibility: a meta-analysis based on 5286 cases and 8000 controls. *Int J Clin Exp Med* 2015; **8**: 22393-22402 [PMID: 26885219]
- 109 **Ma J**, Wu D, Hu X, Li J, Cao M, Dong W. Associations between cytokine gene polymorphisms and susceptibility to *Helicobacter pylori* infection and *Helicobacter pylori* related gastric cancer, peptic ulcer disease: A meta-analysis. *PLoS One* 2017; **12**: e0176463 [PMID: 28453551 DOI: 10.1371/journal.pone.0176463]
- 110 **Negovan A**, Iancu M, Fülöp E, Bănescu C. *Helicobacter pylori* and cytokine gene variants as predictors of premalignant gastric lesions. *World J Gastroenterol* 2019; **25**: 4105-4124 [PMID: 31435167 DOI: 10.3748/wjg.v25.i30.4105]
- 111 **Yuzhalin A**. The role of interleukin DNA polymorphisms in gastric cancer. *Hum Immunol* 2011; **72**: 1128-1136 [PMID: 21871937 DOI: 10.1016/j.humimm.2011.08.003]
- 112 **Xue H**, Liu J, Lin B, Wang Z, Sun J, Huang G. A meta-analysis of interleukin-8 -251 promoter polymorphism associated with gastric cancer risk. *PLoS One* 2012; **7**: e28083 [PMID: 22279522 DOI: 10.1371/journal.pone.0028083]
- 113 **Cheng D**, Hao Y, Zhou W, Ma Y. Positive association between Interleukin-8 -251A > T polymorphism and susceptibility to gastric carcinogenesis: a meta-analysis. *Cancer Cell Int* 2013; **13**: 100 [PMID: 24143859 DOI: 10.1186/1475-2867-13-100]
- 114 **Wang X**, Yang F, Xu G, Zhong S. The roles of IL-6, IL-8 and IL-10 gene polymorphisms in gastric cancer: A meta-analysis. *Cytokine* 2018; **111**: 230-236 [PMID: 30195978 DOI: 10.1016/j.cyto.2018.08.024]
- 115 **Couper KN**, Blount DG, Riley EM. IL-10: the master regulator of immunity to infection. *J Immunol* 2008; **180**: 5771-5777 [PMID: 18424693 DOI: 10.4049/jimmunol.180.9.5771]
- 116 **Chand-Bhayal A**, Krishnaveni D, Pandu-Ranga-Rao K, Prabhakar B, Vidyasagar A, Murali-Krishna B, Anita P, Jyothy A, Nallari P, Venkateshwari A. Association of interleukin-10 promoter polymorphism (-1082 g/a) and gastric cancer in andhra pradesh population of South India. *Iran J Cancer Prev* 2012; **5**: 117-123 [PMID: 25628830]

- 117 **Liu S**, Liu JW, Sun LP, Gong YH, Xu Q, Jing JJ, Yuan Y. Association of IL10 gene promoter polymorphisms with risks of gastric cancer and atrophic gastritis. *J Int Med Res* 2018; **46**: 5155-5166 [PMID: 30205739 DOI: 10.1177/0300060518792785]
- 118 **Namazi A**, Forat-Yazdi M, Jafari M, Farahnak S, Nasiri R, Foroughi E, Abolbaghaei SM, Neamatzadeh H. ASSOCIATION OF INTERLEUKIN-10 -1082 A/G (RS1800896) POLYMORPHISM WITH SUSCEPTIBILITY TO GASTRIC CANCER: META-ANALYSIS OF 6,101 CASES AND 8,557 CONTROLS. *Arq Gastroenterol* 2018; **55**: 33-40 [PMID: 29561974 DOI: 10.1590/S0004-2803.201800000-18]
- 119 **Yu T**, Lu Q, Ou XL, Cao DZ, Yu Q. Clinical study on gastric cancer susceptibility genes IL-10-1082 and TNF- α . *Genet Mol Res* 2014; **13**: 10909-10912 [PMID: 25526211 DOI: 10.4238/2014.December.19.12]
- 120 **Qi M**, Liu DM, Pan LL, Lin YX. Interleukin-10 gene -592C>A polymorphism and susceptibility to gastric cancer. *Genet Mol Res* 2014; **13**: 8954-8961 [PMID: 25366786 DOI: 10.4238/2014.October.31.10]
- 121 **Li C**, Tong W, Liu B, Zhang A, Li F. The -1082A>G polymorphism in promoter region of interleukin-10 and risk of digestive cancer: a meta-analysis. *Sci Rep* 2014; **4**: 5335 [PMID: 25091209 DOI: 10.1038/srep05335]
- 122 **Lu W**, Pan K, Zhang L, Lin D, Miao X, You W. Genetic polymorphisms of interleukin (IL)-1B, IL-1RN, IL-8, IL-10 and tumor necrosis factor {alpha} and risk of gastric cancer in a Chinese population. *Carcinogenesis* 2005; **26**: 631-636 [PMID: 15579481 DOI: 10.1093/carcin/bgh349]
- 123 **Nishikawa RM**, Yaffe MJ. Effect of various noise sources on the detective quantum efficiency of phosphor screens. *Med Phys* 1990; **17**: 887-893 [PMID: 2233576 DOI: 10.1089/dna.2011.1440]
- 124 **El-Omar EM**, Rabkin CS, Gammon MD, Vaughan TL, Risch HA, Schoenberg JB, Stanford JL, Mayne ST, Goedert J, Blot WJ, Fraumeni JF Jr, Chow WH. Increased risk of noncardia gastric cancer associated with proinflammatory cytokine gene polymorphisms. *Gastroenterology* 2003; **124**: 1193-1201 [PMID: 12730860 DOI: 10.1016/s0016-5085(03)00157-4]
- 125 **Melchiades JL**, Zabaglia LM, Sallas ML, Orcini WA, Chen E, Smith MAC, Payão SLM, Rasmussen LT. Polymorphisms and haplotypes of the interleukin 2 gene are associated with an

increased risk of gastric cancer. The possible involvement of *Helicobacter pylori*. *Cytokine* 2017; **96**: 203-207 [PMID: 28458166 DOI: 10.1016/j.cyto.2017.04.020]

126 **Yun Y**, Dong W, Chen C, Zhang H, Shi N, He M, Chen X. Roles of IL-4 genetic polymorphisms and haplotypes in the risk of gastric cancer and their interaction with environmental factors. *Int J Clin Exp Pathol* 2017; **10**: 8936-8943 [PMID: 31966763]

127 **Vivier E**, Spits H, Cupedo T. Interleukin-22-producing innate immune cells: new players in mucosal immunity and tissue repair? *Nat Rev Immunol* 2009; **9**: 229-234 [PMID: 19319141 DOI: 10.1038/nri2522]

128 **Wang YM**, Li ZX, Tang FB, Zhang Y, Zhou T, Zhang L, Ma JL, You WC, Pan KF. Association of genetic polymorphisms of interleukins with gastric cancer and precancerous gastric lesions in a high-risk Chinese population. *Tumour Biol* 2016; **37**: 2233-2242 [PMID: 26358252 DOI: 10.1007/s13277-015-4022-x]

129 **Okabe H**, Kitamura R, Shibata H, Hase T, Nakanishi Y, Masamune Y. [In vitro replication of plasmid pKYM]. *Yakugaku Zasshi* 1989; **109**: 582-591 [PMID: 2530337 DOI: 10.1038/bcj.2014.70]

130 **Hatakeyama M**. *Helicobacter pylori* CagA and gastric cancer: a paradigm for hit-and-run carcinogenesis. *Cell Host Microbe* 2014; **15**: 306-316 [PMID: 24629337 DOI: 10.1016/j.chom.2014.02.008]

131 . Alipour M. Molecular Mechanism of *Helicobacter Pylori*-Induced Gastric Cancer. *J. Gastrointest. Cancer* 2020;1-8 [DOI: 10.1007/s12029-020-00518-5]

132 **Sukri A**, Hanafiah A, Mohamad Zin N, Kosai NR. Epidemiology and role of *Helicobacter pylori* virulence factors in gastric cancer carcinogenesis. *APMIS* 2020; **128**: 150-161 [PMID: 32352605 DOI: 10.1111/apm.13034]

133 **Hatakeyama M**. Structure and function of *Helicobacter pylori* CagA, the first-identified bacterial protein involved in human cancer. *Proc Jpn Acad Ser B Phys Biol Sci* 2017; **93**: 196-219 [PMID: 28413197 DOI: 10.2183/pjab.93.013]

134 **Lin L**, Wei H, Yi J, Xie B, Chen J, Zhou C, Wang L, Yang Y. Chronic CagA-positive *Helicobacter pylori* infection with MNNG stimulation synergistically induces mesenchymal and

cancer stem cell-like properties in gastric mucosal epithelial cells. *J Cell Biochem* 2019; **120**: 17635-17649 [PMID: 31209915 DOI: 10.1002/jcb.29031]

135 **Ansari S**, Yamaoka Y. *Helicobacter pylori* Virulence Factors Exploiting Gastric Colonization and its Pathogenicity. *Toxins (Basel)* 2019; **11** [PMID: 31752394 DOI: 10.3390/toxins11110677]

136 **McClain MS**, Beckett AC, Cover TL. *Helicobacter pylori* Vacuolating Toxin and Gastric Cancer. *Toxins (Basel)* 2017; **9** [PMID: 29023421 DOI: 10.3390/toxins9100316]

137 **El Khadir M**, Alaoui Boukhris S, Benajah DA, El Rhazi K, Ibrahimi SA, El Abkari M, Harmouch T, Nejari C, Mahmoud M, Benlemlih M, Bennani B. VacA and CagA Status as Biomarker of Two Opposite End Outcomes of *Helicobacter pylori* Infection (Gastric Cancer and Duodenal Ulcer) in a Moroccan Population. *PLoS One* 2017; **12**: e0170616 [PMID: 28125638 DOI: 10.1371/journal.pone.0170616]

138 **Chmiela M**, Karwowska Z, Gonciarz W, Allushi B, Stączek P. Host pathogen interactions in *Helicobacter pylori* related gastric cancer. *World J Gastroenterol* 2017; **23**: 1521-1540 [PMID: 28321154 DOI: 10.3748/wjg.v23.i9.1521]

139 **Kusters JG**, van Vliet AH, Kuipers EJ. Pathogenesis of *Helicobacter pylori* infection. *Clin Microbiol Rev* 2006; **19**: 449-490 [PMID: 16847081 DOI: 10.1128/CMR.00054-05]

140 **Sayehmiri F**, Kiani F, Sayehmiri K, Soroush S, Asadollahi K, Alikhani MY, Delpisheh A, Emaneini M, Bogdanović L, Varzi AM, Zarrilli R, Taherikalani M. Prevalence of cagA and vacA among *Helicobacter pylori*-infected patients in Iran: a systematic review and meta-analysis. *J Infect Dev Ctries* 2015; **9**: 686-696 [PMID: 26230117 DOI: 10.3855/jidc.5970]

141 **He H**, Liu J, Li L, Qian G, Hao D, Li M, Zhang Y, Hong X, Xu J, Yan D. *Helicobacter pylori* CagA Interacts with SHP-1 to Suppress the Immune Response by Targeting TRAF6 for K63-Linked Ubiquitination. *J Immunol* 2021; **206**: 1161-1170 [PMID: 33568397 DOI: 10.4049/jimmunol.2000234]

142 **Wen S**, Moss SF. *Helicobacter pylori* virulence factors in gastric carcinogenesis. *Cancer Lett* 2009; **282**: 1-8 [PMID: 19111390 DOI: 10.1016/j.canlet.2008.11.016]

143 **Ferreira RM**, Machado JC, Figueiredo C. Clinical relevance of *Helicobacter pylori* vacA and cagA genotypes in gastric carcinoma. *Best Pract Res Clin Gastroenterol* 2014; **28**: 1003-1015 [PMID: 25439067 DOI: 10.1016/j.bpg.2014.09.004]

- 144 **Dela Pena-Ponce MG**, Jimenez MT, Hansen LM, Solnick JV, Miller LA. The *Helicobacter pylori* type IV secretion system promotes IL-8 synthesis in a model of pediatric airway epithelium via p38 MAP kinase. *PLoS One* 2017; **12**: e0183324 [PMID: 28813514 DOI: 10.1371/journal.pone.0183324]
- 145 **Jang S**, Su H, Blum FC, Bae S, Choi YH, Kim A, Hong YA, Kim J, Kim JH, Gunawardhana N, Jeon YE, Yoo YJ, Merrell DS, Ge L, Cha JH. Dynamic Expansion and Contraction of *cagA* Copy Number in *Helicobacter pylori* Impact Development of Gastric Disease. *mBio* 2017; **8** [PMID: 28223454 DOI: 10.1128/mbio.01779-16]
- 146 **Yamaoka Y**, Kudo T, Lu H, Casola A, Brasier AR, Graham DY. Role of interferon-stimulated responsive element-like element in interleukin-8 promoter in *Helicobacter pylori* infection. *Gastroenterology* 2004; **126**: 1030-1043 [PMID: 15057743 DOI: 10.1053/j.gastro.2003.12.048]
- 147 **Horridge DN**, Begley AA, Kim J, Aravindan N, Fan K, Forsyth MH. Outer inflammatory protein a (OipA) of *Helicobacter pylori* is regulated by host cell contact and mediates CagA translocation and interleukin-8 response only in the presence of a functional *cag* pathogenicity island type IV secretion system. *Pathog Dis* 2017; **75** [PMID: 29040466 DOI: 10.1093/femspd/ftx113]
- 148 **Nejati S**, Karkhah A, Darvish H, Validi M, Ebrahimpour S, Nouri HR. Influence of *Helicobacter pylori* virulence factors CagA and VacA on pathogenesis of gastrointestinal disorders. *Microb Pathog* 2018; **117**: 43-48 [PMID: 29432909 DOI: 10.1016/j.micpath.2018.02.016]
- 149 **Cover TL**, Blanke SR. *Helicobacter pylori* VacA, a paradigm for toxin multifunctionality. *Nat Rev Microbiol* 2005; **3**: 320-332 [PMID: 15759043 DOI: 10.1038/nrmicro1095]
- 150 **Palframan SL**, Kwok T, Gabriel K. Vacuolating cytotoxin A (VacA), a key toxin for *Helicobacter pylori* pathogenesis. *Front Cell Infect Microbiol* 2012; **2**: 92 [PMID: 22919683 DOI: 10.3389/fcimb.2012.00092]
- 151 **Chauhan N**, Tay ACY, Marshall BJ, Jain U. *Helicobacter pylori* VacA, a distinct toxin exerts diverse functionalities in numerous cells: An overview. *Helicobacter* 2019; **24**: e12544 [PMID: 30324717 DOI: 10.1111/hel.12544]

- 152 **Sukhan DS**, Vernygorodskyi S V., Haidukov N V., Ludkevich HP. Molecular and Genetic Aspects of *Helicobacter pylori* Interaction with Cells of Gastric Mucosa. *Cytol Genet* 2020;54:147–53 [DOI: 10.3103/S0095452720020139]
- 153 **Baj J**, Forma A, Sitarz M, Portincasa P, Garruti G, Krasowska D, Maciejewski R. *Helicobacter pylori* Virulence Factors-Mechanisms of Bacterial Pathogenicity in the Gastric Microenvironment. *Cells* 2020; **10** [PMID: 33375694 DOI: 10.3390/cells10010027]
- 154 **Utsch C**, Haas R. VacA's Induction of VacA-Containing Vacuoles (VCVs) and Their Immunomodulatory Activities on Human T Cells. *Toxins (Basel)* 2016; **8** [PMID: 27322323 DOI: 10.3390/toxins8060190]
- 155 **Sundrud MS**, Torres VJ, Unutmaz D, Cover TL. Inhibition of primary human T cell proliferation by *Helicobacter pylori* vacuolating toxin (VacA) is independent of VacA effects on IL-2 secretion. *Proc Natl Acad Sci U S A* 2004; **101**: 7727-7732 [PMID: 15128946 DOI: 10.1073/pnas.0401528101]
- 156 **Abdullah M**, Greenfield LK, Bronte-Tinkew D, Capurro MI, Rizzuti D, Jones NL. VacA promotes CagA accumulation in gastric epithelial cells during *Helicobacter pylori* infection. *Sci Rep* 2019; **9**: 38 [PMID: 30631092 DOI: 10.1038/s41598-018-37095-4]
- 157 **Alm RA**, Ling LS, Moir DT, King BL, Brown ED, Doig PC, Smith DR, Noonan B, Guild BC, deJonge BL, Carmel G, Tummino PJ, Caruso A, Uria-Nickelsen M, Mills DM, Ives C, Gibson R, Merberg D, Mills SD, Jiang Q, Taylor DE, Vovis GF, Trust TJ. Genomic-sequence comparison of two unrelated isolates of the human gastric pathogen *Helicobacter pylori*. *Nature* 1999; **397**: 176-180 [PMID: 9923682 DOI: 10.1038/16495]
- 158 **Talebi Bezmin Abadi A**, Perez-Perez G. Role of *dupA* in virulence of *Helicobacter pylori*. *World J Gastroenterol* 2016; **22**: 10118-10123 [PMID: 28028359 DOI: 10.3748/wjg.v22.i46.10118]
- 159 **Souod N**, Sarshar M, Dabiri H, Momtaz H, Kargar M, Mohammadzadeh A, Abdi S. The study of the *oipA* and *dupA* genes in *Helicobacter pylori* strains and their relationship with different gastroduodenal diseases. *Gastroenterol Hepatol Bed Bench* 2015; **8**: S47-S53 [PMID: 26171137]
- 160 **Lu H**, Hsu PI, Graham DY, Yamaoka Y. Duodenal ulcer promoting gene of *Helicobacter pylori*. *Gastroenterology* 2005; **128**: 833-848 [PMID: 15825067 DOI: 10.1053/j.gastro.2005.01.009]

- 161 **Youssefi M**, Ghazvini K, Farsiani H, Tafaghodi M, Keikha M. A systematic review and meta-analysis of outcomes of infection with *Helicobacter pylori* dupA+ strains in Iranian patients. *Gene Reports*. 2020;19:100650 [DOI: 10.1016/j.genrep.2020.100650]
- 162 **Miftahussurur M**, Yamaoka Y. *Helicobacter pylori* virulence genes and host genetic polymorphisms as risk factors for peptic ulcer disease. *Expert Rev Gastroenterol Hepatol* 2015; **9**: 1535-1547 [PMID: 26470920 DOI: 10.1586/17474124.2015.1095089]
- 163 **Fatahi G**, Talebi Bezmin Abadi A, Peerayeh SN, Forootan M. Carrying a 112 bp-segment in *Helicobacter pylori* dupA may associate with increased risk of duodenal ulcer. *Infect Genet Evol* 2019; **73**: 21-25 [PMID: 30981881 DOI: 10.1016/j.meegid.2019.04.009]
- 164 **Shiota S**, Nguyen LT, Murakami K, Kuroda A, Mizukami K, Okimoto T, Kodama M, Fujioka T, Yamaoka Y. Association of *Helicobacter pylori* dupA with the failure of primary eradication. *J Clin Gastroenterol* 2012; **46**: 297-301 [PMID: 22298090 DOI: 10.1097/MCG.0b013e318243201c]
- 165 **Argent RH**, Burette A, Miendje Deyi VY, Atherton JC. The presence of dupA in *Helicobacter pylori* is not significantly associated with duodenal ulceration in Belgium, South Africa, China, or North America. *Clin Infect Dis* 2007; **45**: 1204-1206 [PMID: 17918084 DOI: 10.1086/522177]
- 166 **Paredes-Osses E**, Sáez K, Sanhueza E, Hebel S, González C, Briceño C, García Cancino A. Association between cagA, vacAi, and dupA genes of *Helicobacter pylori* and gastroduodenal pathologies in Chilean patients. *Folia Microbiol (Praha)* 2017; **62**: 437-444 [PMID: 28283946 DOI: 10.1007/s12223-017-0514-y]
- 167 **Dossumbekova A**, Prinz C, Mages J, Lang R, Kusters JG, Van Vliet AH, Reindl W, Backert S, Saur D, Schmid RM, Rad R. *Helicobacter pylori* HopH (OipA) and bacterial pathogenicity: genetic and functional genomic analysis of hopH gene polymorphisms. *J Infect Dis* 2006; **194**: 1346-1355 [PMID: 17054063 DOI: 10.1086/508426]
- 168 **Teymournejad O**, Mobarez AM, Hassan ZM, Talebi Bezmin Abadi A. Binding of the *Helicobacter pylori* OipA causes apoptosis of host cells *via* modulation of Bax/Bcl-2 Levels. *Sci Rep* 2017; **7**: 8036 [PMID: 28808292 DOI: 10.1038/s41598-017-08176-7]

- 169 **Baj J**, Forma A, Sitarz M, Portincasa P, Garruti G, Krasowska D, Maciejewski R. *Helicobacter pylori* Virulence Factors-Mechanisms of Bacterial Pathogenicity in the Gastric Microenvironment. *Cells* 2020; **10** [PMID: 33375694 DOI: 10.3390/cells10010027]
- 170 **Braga LLBC**, Batista MHR, de Azevedo OGR, da Silva Costa KC, Gomes AD, Rocha GA, Queiroz DMM. oipA "on" status of *Helicobacter pylori* is associated with gastric cancer in North-Eastern Brazil. *BMC Cancer* 2019; **19**: 48 [PMID: 30630444 DOI: 10.1186/s12885-018-5249-x]
- 171 **Bartpho TS**, Wattanawongdon W, Tongtawee T, Paoin C, Kangwantas K, Dechsukhum C. Precancerous Gastric Lesions with *Helicobacter pylori vacA +/babA2+/oipA +* Genotype Increase the Risk of Gastric Cancer. *Biomed Res Int* 2020; **2020**: 7243029 [PMID: 32149129 DOI: 10.1155/2020/7243029]
- 172 **Sallas ML**, Dos Santos MP, Orcini WA, David ÉB, Peruquetti RL, Payão SLM, Rasmussen LT. Status (on/off) of oipA gene: their associations with gastritis and gastric cancer and geographic origins. *Arch Microbiol* 2019; **201**: 93-97 [PMID: 30255200 DOI: 10.1007/s00203-018-1580-5]
- 173 **Yamaoka Y**. Pathogenesis of *Helicobacter pylori*-Related Gastroduodenal Diseases from Molecular Epidemiological Studies. *Gastroenterol Res Pract* 2012; **2012**: 371503 [PMID: 22829807 DOI: 10.1155/2012/371503]
- 174 **Zhang J**, Qian J, Zhang X, Zou Q. Outer membrane inflammatory protein A, a new virulence factor involved in the pathogenesis of *Helicobacter pylori*. *Mol Biol Rep* 2014; **41**: 7807-7814 [PMID: 25096514 DOI: 10.1007/s11033-014-3673-9]
- 175 **Wroblewski LE**, Peek RM Jr, Wilson KT. *Helicobacter pylori* and gastric cancer: factors that modulate disease risk. *Clin Microbiol Rev* 2010; **23**: 713-739 [PMID: 20930071 DOI: 10.1128/CMR.00011-10]
- 176 **Odenbreit S**, Kavermann H, Püls J, Haas R. CagA tyrosine phosphorylation and interleukin-8 induction by *Helicobacter pylori* are independent from alpAB, HopZ and bab group outer membrane proteins. *Int J Med Microbiol* 2002; **292**: 257-266 [PMID: 12398216 DOI: 10.1078/1438-4221-00205]

- 177 **Ando T**, Peek RM Jr, Lee YC, Krishna U, Kusugami K, Blaser MJ. Host cell responses to genotypically similar *Helicobacter pylori* isolates from United States and Japan. *Clin Diagn Lab Immunol* 2002; **9**: 167-175 [PMID: 11777849 DOI: 10.1128/CDLI.9.1.167-175.2002]
- 178 **Teymournejad O**, Mobarez AM, Hassan ZM, Moazzeni SM, Ahmadabad HN. In vitro suppression of dendritic cells by *Helicobacter pylori* OipA. *Helicobacter* 2014; **19**: 136-143 [PMID: 24495278 DOI: 10.1111/hel.12107]
- 179 **Souod N**, Sarshar M, Dabiri H, Momtaz H, Kargar M, Mohammadzadeh A, Abdi S. The study of the oipA and dupA genes in *Helicobacter pylori* strains and their relationship with different gastroduodenal diseases. *Gastroenterol Hepatol Bed Bench* 2015; **8**: S47-S53 [PMID: 26171137]
- 180 **Baj J**, Forma A, Sitarz M, Portincasa P, Garruti G, Krasowska D, Maciejewski R. *Helicobacter pylori* Virulence Factors-Mechanisms of Bacterial Pathogenicity in the Gastric Microenvironment. *Cells* 2020; **10** [PMID: 33375694 DOI: 10.3390/cells10010027]
- 181 **Huang X**, Deng Z, Zhang Q, Li W, Wang B, Li M. Relationship between the iceA gene of *Helicobacter pylori* and clinical outcomes. *Ther Clin Risk Manag* 2016; **12**: 1085-1092 [PMID: 27462162 DOI: 10.2147/TCRM.S107991]
- 182 **Xu Q**, Blaser MJ. Promoters of the CATG-specific methyltransferase gene hpyIM differ between iceA1 and iceA2 *Helicobacter pylori* strains. *J Bacteriol* 2001; **183**: 3875-3884 [PMID: 11395450 DOI: 10.1128/JB.183.13.3875-3884.2001]
- 183 **Dabiri H**, Jafari F, Baghaei K, Shokrzadeh L, Abdi S, Pourhoseingholi MA, Mohammadzadeh A. Prevalence of *Helicobacter pylori* vacA, cagA, cagE, oipA, iceA, babA2 and babB genotypes in Iranian dyspeptic patients. *Microb Pathog* 2017; **105**: 226-230 [PMID: 28215588 DOI: 10.1016/j.micpath.2017.02.018]
- 184 **Feliciano O**, Gutierrez O, Valdés L, Fragoso T, Calderin AM, Valdes AE, Llanes R. Prevalence of *Helicobacter pylori* vacA, cagA, and iceA Genotypes in Cuban Patients with Upper Gastrointestinal Diseases. *Biomed Res Int* 2015; **2015**: 753710 [PMID: 25945344 DOI: 10.1155/2015/753710]

- 185 **Yakoob J**, Abbas Z, Khan R, Salim SA, Abrar A, Awan S, Ahmad Z. Helicobacter pylori: correlation of the virulence marker iceA allele with clinical outcome in a high prevalence area. *Br J Biomed Sci* 2015; **72**: 67-73 [PMID: 26126322 DOI: 10.1080/09674845.2015.11666799]
- 186 **Shiota S**, Watada M, Matsunari O, Iwatani S, Suzuki R, Yamaoka Y. Helicobacter pylori iceA, clinical outcomes, and correlation with cagA: a meta-analysis. *PLoS One* 2012; **7**: e30354 [PMID: 22279585 DOI: 10.1371/journal.pone.0030354]
- 187 **Boyanova L**, Yordanov D, Gergova G, Markovska R, Mitov I. Association of iceA and babA genotypes in Helicobacter pylori strains with patient and strain characteristics. *Antonie Van Leeuwenhoek* 2010; **98**: 343-350 [PMID: 20454856 DOI: 10.1007/s10482-010-9448-y]
- 188 **Ladeira MS**, Rodrigues MA, Salvadori DM, Neto PP, Achilles P, Lerco MM, Rodrigues PA, Gonçalves I Jr, Queiroz DM, Freire-Maia DV. Relationships between cagA, vacA, and iceA genotypes of Helicobacter pylori and DNA damage in the gastric mucosa. *Environ Mol Mutagen* 2004; **44**: 91-98 [PMID: 15278912 DOI: 10.1002/em.20045]
- 189 **Chiurillo MA**, Moran Y, Cañas M, Valderrama E, Alvarez A, Armanie E. Combination of Helicobacter pylori-iceA2 and proinflammatory interleukin-1 polymorphisms is associated with the severity of histological changes in Venezuelan chronic gastritis patients. *FEMS Immunol Med Microbiol* 2010; **59**: 170-176 [PMID: 20482626 DOI: 10.1111/j.1574-695X.2010.00675.x]
- 190 **Abu-Taleb AMF**, Abdelattef RS, Abdel-Hady AA, Omran FH, El-Korashi LA, Abdel-Aziz El-Hady H, El-Gebaly AM. Prevalence of *Helicobacter pylori* cagA and iceA Genes and Their Association with Gastrointestinal Diseases. *Int J Microbiol* 2018; **2018**: 4809093 [PMID: 29849647 DOI: 10.1155/2018/4809093]
- 191 **Yılmaz N**, Koruk Özer M. The Prevalence of Helicobacter Pylori babA, homB, aspA, and sabA Genes and Its Relationship with Clinical Outcomes in Turkey. *Can J Gastroenterol Hepatol* 2019; **2019**: 1271872 [PMID: 31312620 DOI: 10.1155/2019/1271872]
- 192 **Yamaoka Y**. Roles of Helicobacter pylori BabA in gastroduodenal pathogenesis. *World J Gastroenterol* 2008; **14**: 4265-4272 [PMID: 18666312 DOI: 10.3748/WJG.14.4265]

- 193 **Fujimoto S**, Olaniyi Ojo O, Arnqvist A, Wu JY, Odenbreit S, Haas R, Graham DY, Yamaoka Y. Helicobacter pylori BabA expression, gastric mucosal injury, and clinical outcome. *Clin Gastroenterol Hepatol* 2007; **5**: 49-58 [PMID: 17157077 DOI: 10.1016/j.cgh.2006.09.015]
- 194 **Shahi H**, Reisi S, Bahreini R, Bagheri N, Salimzadeh L, Shirzad H. Association Between helicobacter pylori cagA, babA2 Virulence Factors and Gastric Mucosal Interleukin-33 mRNA Expression and Clinical Outcomes in Dyspeptic Patients. *Int J Mol Cell Med*. 2015;
- 195 **Duan L**, Chen J, Zhang H, Yang H, Zhu P, Xiong A, Xia Q, Zheng F, Tan Z, Gong F, Fang M. Interleukin-33 ameliorates experimental colitis through promoting Th2/Foxp3⁺ regulatory T-cell responses in mice. *Mol Med* 2012; **18**: 753-761 [PMID: 22426954 DOI: 10.2119/molmed.2011.00428]
- 196 **Wills-Karp M**, Rani R, Dienger K, Lewkowich I, Fox JG, Perkins C, Lewis L, Finkelman FD, Smith DE, Bryce PJ, Kurt-Jones EA, Wang TC, Sivaprasad U, Hershey GK, Herbert DR. Trefoil factor 2 rapidly induces interleukin 33 to promote type 2 immunity during allergic asthma and hookworm infection. *J Exp Med* 2012; **209**: 607-622 [PMID: 22329990 DOI: 10.1084/jem.20110079]
- 197 **Choi YS**, Choi HJ, Min JK, Pyun BJ, Maeng YS, Park H, Kim J, Kim YM, Kwon YG. Interleukin-33 induces angiogenesis and vascular permeability through ST2/TRAF6-mediated endothelial nitric oxide production. *Blood* 2009; **114**: 3117-3126 [PMID: 19661270 DOI: 10.1182/blood-2009-02-203372]
- 198 **Pakbaz Z**, Shirazi MH, Ranjbar R, Pourm MR, Gholi MK, Aliramezani A *et al* Frequency of sabA gene in helicobacter pylori strains isolated from patients in Tehran, Iran. *Iran Red Crescent Med J*. 2013;15:767-70 [DOI: 10.5812/ircmj.5044]
- 199 **Su YL**, Huang HL, Huang BS, Chen PC, Chen CS, Wang HL, Lin PH, Chieh MS, Wu JJ, Yang JC, Chow LP. Combination of OipA, BabA, and SabA as candidate biomarkers for predicting Helicobacter pylori-related gastric cancer. *Sci Rep* 2016; **6**: 36442 [PMID: 27819260 DOI: 10.1038/srep36442]
- 200 **Yamaoka Y**, Ojo O, Fujimoto S, Odenbreit S, Haas R, Gutierrez O, El-Zimaity HM, Reddy R, Arnqvist A, Graham DY. Helicobacter pylori outer membrane proteins and gastroduodenal disease. *Gut* 2006; **55**: 775-781 [PMID: 16322107 DOI: 10.1136/gut.2005.083014]

201 **Nakasato F**, Shimoyama T, Yoshimura T, Mikami T, Munakata A, Fukuda S. Infection of sabA-positive *H. pylori* does not induce anti-Lewis X antibody in host. *Hepatogastroenterology* 2008; **55**: 1122-1125 [PMID: 18705343]

202 **Sheu BS**, Odenbreit S, Hung KH, Liu CP, Sheu SM, Yang HB, Wu JJ. Interaction between host gastric Sialyl-Lewis X and *H. pylori* SabA enhances *H. pylori* density in patients lacking gastric Lewis B antigen. *Am J Gastroenterol* 2006; **101**: 36-44 [PMID: 16405531 DOI: 10.1111/j.1572-0241.2006.00358.x]

203 **Mahdavi J**, Sondén B, Hurtig M, Olfat FO, Forsberg L, Roche N, Angstrom J, Larsson T, Teneberg S, Karlsson KA, Altraja S, Wadström T, Kersulyte D, Berg DE, Dubois A, Petersson C, Magnusson KE, Norberg T, Lindh F, Lundskog BB, Arnqvist A, Hammarström L, Borén T. *Helicobacter pylori* SabA adhesin in persistent infection and chronic inflammation. *Science* 2002; **297**: 573-578 [PMID: 12142529 DOI: 10.1126/science.1069076]

204 **de Jonge R**, Pot RG, Loffeld RJ, van Vliet AH, Kuipers EJ, Kusters JG. The functional status of the *Helicobacter pylori* sabB adhesin gene as a putative marker for disease outcome. *Helicobacter* 2004; **9**: 158-164 [PMID: 15068418 DOI: 10.1111/j.1083-4389.2004.00213.x]

205 **Bulut Y**, Faure E, Thomas L, Karahashi H, Michelsen KS, Equils O, Morrison SG, Morrison RP, Arditi M. Chlamydial heat shock protein 60 activates macrophages and endothelial cells through Toll-like receptor 4 and MD2 in a MyD88-dependent pathway. *J Immunol* 2002; **168**: 1435-1440 [PMID: 11801686 DOI: 10.4049/jimmunol.168.3.1435]

206 **Huesca M**, Borgia S, Hoffman P, Lingwood CA. Acidic pH changes receptor binding specificity of *Helicobacter pylori*: a binary adhesion model in which surface heat shock (stress) proteins mediate sulfatide recognition in gastric colonization. *Infect Immun* 1996; **64**: 2643-2648 [PMID: 8698490 DOI: 10.1128/IAI.64.7.2643-2648.1996]

207 **Maguire M**, Poole S, Coates AR, Tormay P, Wheeler-Jones C, Henderson B. Comparative cell signalling activity of ultrapure recombinant chaperonin 60 proteins from prokaryotes and eukaryotes. *Immunology* 2005; **115**: 231-238 [PMID: 15885129 DOI: 10.1111/j.1365-2567.2005.02155.x]

208 **Zhao Y**, Yokota K, Ayada K, Yamamoto Y, Okada T, Shen L, Oguma K. *Helicobacter pylori* heat-shock protein 60 induces interleukin-8 *via* a Toll-like receptor (TLR)2 and mitogen-activated

protein (MAP) kinase pathway in human monocytes. *J Med Microbiol* 2007; **56**: 154-164 [PMID: 17244794 DOI: 10.1099/jmm.0.46882-0]

209 **Takenaka R**, Yokota K, Ayada K, Mizuno M, Zhao Y, Fujinami Y, Lin SN, Toyokawa T, Okada H, Shiratori Y, Oguma K. Helicobacter pylori heat-shock protein 60 induces inflammatory responses through the Toll-like receptor-triggered pathway in cultured human gastric epithelial cells. *Microbiology (Reading)* 2004; **150**: 3913-3922 [PMID: 15583145 DOI: 10.1099/mic.0.27527-0]

210 **Lin CY**, Huang YS, Li CH, Hsieh YT, Tsai NM, He PJ, Hsu WT, Yeh YC, Chiang FH, Wu MS, Chang CC, Liao KW. Characterizing the polymeric status of Helicobacter pylori heat shock protein 60. *Biochem Biophys Res Commun* 2009; **388**: 283-289 [PMID: 19664598 DOI: 10.1016/j.bbrc.2009.07.159]

211 **Lin CS**, He PJ, Tsai NM, Li CH, Yang SC, Hsu WT, Wu MS, Wu CJ, Cheng TL, Liao KW. A potential role for Helicobacter pylori heat shock protein 60 in gastric tumorigenesis. *Biochem Biophys Res Commun* 2010; **392**: 183-189 [PMID: 20060384 DOI: 10.1016/j.bbrc.2010.01.010]

212 Oleastro M, Monteiro L, Lehours P, Mégraud F, Ménard A. Identification of markers for Helicobacter pylori strains isolated from children with peptic ulcer disease by suppressive subtractive hybridization. *Infect Immun* 2006; **74(7)**: 4064-4074 [PMID: 16790780 DOI:10.1128/IAI.00123-06]

213 Oleastro M, Cordeiro R, Yamaoka Y, Queiroz D, Mégraud F, Monteiro L, Ménard A. Disease association with two Helicobacter pylori duplicate outer membrane protein genes, homB and homA. *Gut Pathog* 2009; **1(1)**: 12 [PMID: 19545429 DOI: 10.1186/1757-4749-1-12]

214 Jung SW, Sugimoto M, Graham DY, Yamaoka Y. homB status of Helicobacter pylori as a novel marker to distinguish gastric cancer from duodenal ulcer. *J Clin Microbiol* 2009; **47(10)**: 3241-3245 [PMID: 19710266 DOI:10.1128/JCM.00293-09]

215 Keikha M, Karbalaei M. Correlation between the geographical origin of Helicobacter pylori homB-positive strains and their clinical outcomes: a systematic review and meta-analysis. *BMC Gastroenterol* 2021; **21(1)**: 181 [PMID: 33879080 DOI: 10.1186/s12876-021-01764-y]

216 Zimmermann S, Pfannkuch L, Al-Zeer MA, et al. ALPK1- and TIFA-Dependent Innate Immune Response Triggered by the Helicobacter pylori Type IV Secretion System. *Cell Rep* 2017; **20(10)**: 2384-2395 [PMID: 28877472 DOI:10.1016/j.celrep.2017.08.039]

217 Posselt G, Wiesauer M, Chichirau BE, Engler D, Krisch LM, Gadermaier G, Briza P, Schneider S, Boccellato F, Meyer TF, Hauser-Kronberger C, Neureiter D, Müller A, Wessler S. Helicobacter pylori-controlled c-Abl localization promotes cell migration and limits apoptosis. *Cell Commun Signal*. 2019; **17(1)**:10 [PMID: 30704478 DOI: 10.1186/s12964-019-0323-9]

