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REVIEW

The role and diagnostic potential of gastrokine 1 in gastric cancer

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Introduction: Gene for *gastrokine 1* (*GKN1*) was identified as one of the most significant in gastric cancer and indicated as a potential therapeutic target.

Aim: The aim was a review of literature reports concerning the role and diagnostic potential of GKN1 in gastric cancer.

Materials and methods: PubMED database was searched for sources using the following keywords: gastrokine 1/GKN1/AMP-18 and gastric cancer, *Helicobacter pylori*, aspirin, nonsteroidal anti-inflammatory drugs. Preference was given to the sources which were published within the past 10 years.

Conclusion: GKN1 is a stomach-specific protein, and its role consists of maintaining mucosal integrity as well as the replenishment of the surface lumen epithelial cells layer. The evaluation of GKN1 expression seems to be a useful indicator of the presence of neoplastic or inflammatory lesions in the gastric mucosa. GKN1 expression is decreased in gastric tumor tissues and derived cell lines and its upregulation in cell lines of gastric cancer induces cells apoptosis. The mechanism by which GKN1 is inactivated in gastric cancer cells is still not fully understood. The future diagnostic capabilities of gastric cancer concern the assessment of serum GKN1 concentration by means of ELISA method. Serum GKN1 concentration is not related to patients' sex. Moreover, the measurement of GKN1 concentration is possible only after the incubation of samples at 70°C for 10 minutes. Nevertheless, the aspect of quantitative serum GKN1 evaluation is new in the context of available literature and requires further studies.

Keywords: gastrokine 1/GKN1, gastric cancer, *Helicobacter pylori*

Introduction

Gastric cancer is one of the most commonly diagnosed malignancies of the digestive tract.[1,2] Literature data show that in the last few decades, both the number of new cases and deaths from gastric cancer have been decreasing.[3,4] Figure 1 shows etiopathological factors of gastric cancer development (Figure 1).[5–11] In about 80% of cases of gastric cancer, the early stage of the disease is asymptomatic, and complaints such as an early feeling of fullness during a meal, nausea or epigastric pain are unspecific and often associated with peptic ulcer disease or other gastrointestinal diseases. Therefore, gastric cancer is diagnosed in advanced stages, which is inevitably combined with ineffective treatment and poor prognosis.[12,13]

The criterion for the diagnosis of gastric cancer is the finding of tumor cells in the histopathological examination of mucosal sections taken during endoscopic examination (gastroscopy). The gastroscopy is supplemented by a radiological examination of the

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upper gastrointestinal tract. Laboratory tests that may suggest a neoplastic process are not very specific (decreased serum total protein and albumin in the proteinogram and a positive stool test for the blood). Iron deficiency anemia usually occurs in the advanced stages of the disease. Cancer markers such as: carbohydrate antigen (CA) 72, carcinoembryonic antigen (CEA), and CA 19-9 are also useful in diagnosis and monitoring of gastric cancer. However, with a diagnostic specificity of 95%, their diagnostic sensitivity for gastric cancer is 35%–50%, 25%–35% and 30%–40%, respectively, which encourages the search for new, more sensitive tests.[14,15]

The literature pays attention to the recently discovered protein – gastrokine 1 (GKN1) with a molecular mass of 18 kDa, which has been isolated from cells of the gastric mucosa of healthy people, as well as other species of mammals (including mice, rats, pigs and cows).[16–19] The gene for GKN1, along with genes for secreted phosphoprotein 1 (SPP1), sulfatase 1 (SULF1), thrombospondin 2 (THBS2), ATPase H+/K+ transporting beta subunit (ATP4B), gastric intrinsic factor (GIF), was identified as the most significant in gastric cancer and, moreover, indicated as potential target for

improving the diagnosis and clinical effects in patients with gastric cancer.[2] Therefore, the aim of the current paper is a review of literature reports concerning the role and diagnostic potential of GKN1 in gastric cancer.

Search strategy

We searched PubMED database for sources using the following keywords: gastrokine 1/GKN1/AMP-18 and gastric cancer, *Helicobacter pylori*, aspirin, nonsteroidal anti-inflammatory drugs. Preference was given to the sources which were published within the past 10 years.

Gastrokine 1 (GKN1)

In earlier reports, GKN1 was reported as AMP-18 (antrum mucosal protein-18), CA11, foveolin or TFIZ2.[17,19,20] The *GKN1* coding gene (CA11, accession number: BK0017373) is located on the 2p13 chromosome and consists of six exons. Many studies have shown that GKN1 occurs inside cytoplasmic granules, suggesting that it is rather a secretory protein than an integral membrane protein.[17,19,21,22] Martin et al[17] to localize this protein in the antral part of the stomach, studied
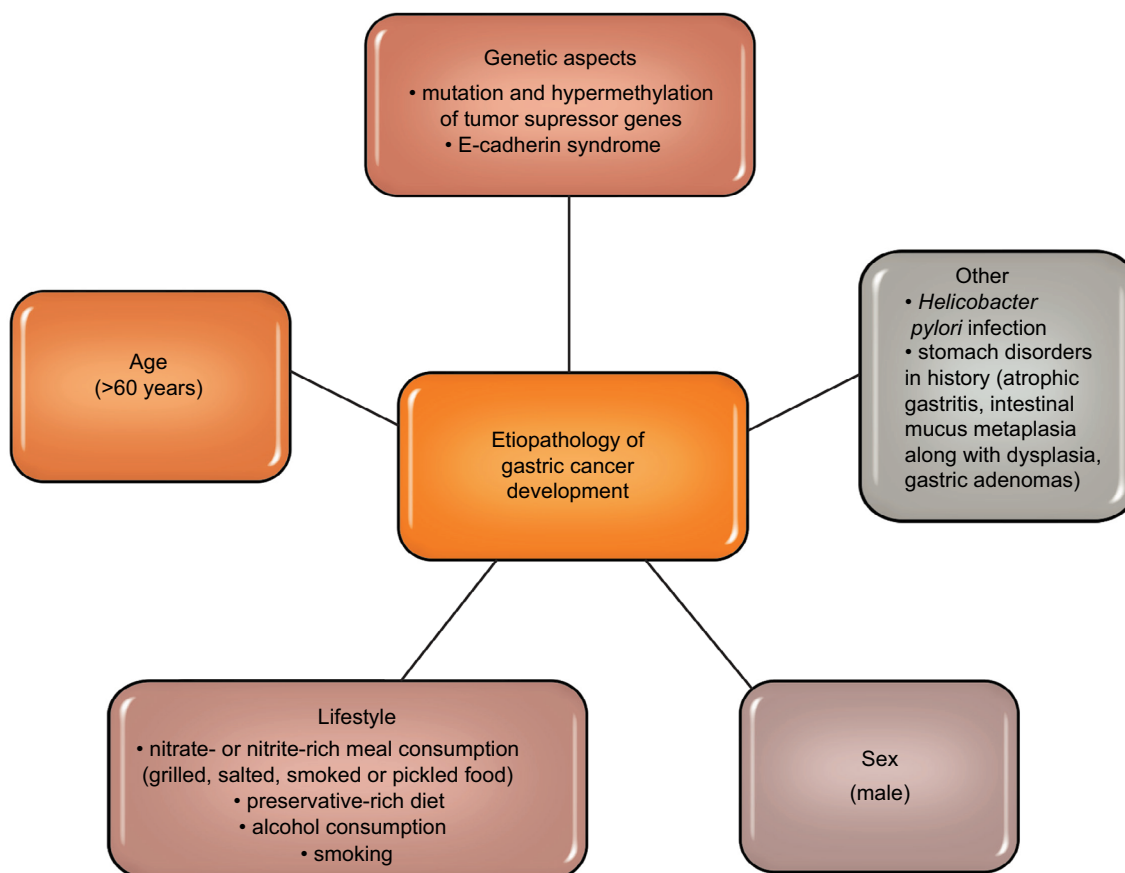

Figure 1 Etiopathology of gastric cancer development.[5–11]

the pigs' gastric mucosal membrane by using the following methods: immunohistochemistry, immunoblotting and northern blot. It was found that the newly discovered protein has tissue specificity, therefore in November 2003 the Human Gene Nomenclature Committee gave it its official name – gastrokine 1.[23,24]

Studies conducted in recent years have shown that gastro-intestinal mucosa cells secrete not only GKN1, but also other gastrokines: 2 (GKN2/GDDR/TFIZ1/blottin) and 3 (GKN3). GKN2 is indicated as a gastric epithelial-secreted protein, however human *GKN3* persists only as a non-expressed pseudogene (Table 1).[5,20–22,25–36] These proteins also have a

Table 1 GKN2 and GKN3 in gastric cancer.

Gastrokine 2 (GKN2) Alternative names: GDDR/TFIZI/Blottin	
Sánchez-Pulido et al[25]	• Proposed GKN2 function in folding and intracellular transport or secretion
Du et al[26]	• *GKN2* mRNA is located in gastric mucosa, and only expressed in gastric tissue • GKN2 encoded protein including a trans-membrane peptide homolog to GKN1
Wesley et al[27]	• GKN2 forms a heterodimer with the trefoil factor family (TFF) protein 1 via an intermolecular disulfide bond between cysteine residues in the carboxy-terminus of TFF1 and in the BRICHOS domain of GKN2
Otto et al[22]	• GKN2 may modulate the integrity of the mucus barrier and its mode of action may be enhanced in a wound situation, perhaps by influencing mucus viscosity
Baus-Loncar et al[21]	• The regulation of GKN2 parallels that of TFF genes, indicating that together they may play a role in the homeostasis of the gastrointestinal tract
Kouznetsova et al[28]	• The TFF1-GKN2 heterodimer was not associated with the mucin fraction, whereas TFF2 did associate with mucins
Moss et al[20]	• GKN2 mRNA was downregulated in distal gastric cancer • GKN2 may be an independent marker of prognosis
Zhang et al[29]	• GKN2 is expressed on the surface area of nontumoral mucosa • GKN2 expression is weak in deeper glands of stomach • In deeper zones of mucosa GKN2 expression coexisted with beta-catenin nuclear accumulation, which may indicate some interactions between them
Chu et al[30]	• GKN2 is expressed in human gastric epithelial cells and significantly downregulated in gastric cancer cells • GKN2 gastric cancer growth inhibition is a TFF1-dependent
Kim et al[31]	• GKN2 may inhibit GKN1 activity • GKN2 expression positively correlated with GKN1 expression
Dai et al[32]	• GKN2 expression was decreased or absent in gastric cancer cell lines, gastric intestinal metaplasia and tumor tissues • GKN2 overexpression inhibited the proliferation, migration, and invasion of gastric cancer cells and arrested the cell cycle at the G1-S transition phase • Overexpression of GKN2 and TFF2 together showed the same inhibitory effect as overexpression of GKN2 alone, which may indicate that TFF2 may not structurally or functionally interact with GKN2
Shi et al[5]	• GKN2 overexpression in gastric cancer cells upregulated Fas expression, but it did not significantly influence the expression of Bcl-2 and Bax, which indicate the extrinsic pathways of apoptosis activation
Menheniott et al[33]	• In vivo GKN2 loss in chronic inflammation induced by *H. pylori* infection or genetically induced hyperactivation of oncogenic gp130/STAT3 signaling plays a role in gastric cancer progression • Gkn2–/– mice lacked spontaneous tumor growth or significant stomach epithelial hyperplasia, which may indicate that isolated GKN2 loss is inadequate for cell-autonomous transformation • Dual overexpression of GKN2 and GKN1 may significantly decrease gastric tumor growth in vivo
Ouyang et al[34]	• GKN2 overexpression significantly inhibits the JAK2/STAT3 signal pathway to further upregulate Bax, and downregulate Bcl-2, Cyclin D1, MMP2 and MMP9, which cause reduced gastric cancer proliferation and invasiveness, increased apoptosis and cell cycle arrest in the G1 phase
Gastrokine 3 (GKN3)	
Menheniott et al[35]	• GKN3 encodes a secreted protein, suggesting an extracellular function • Mouse GKN3 is overexpressed in atrophic gastritis associated with *H. pylori* infection • Human GKN3 persists only as a nonexpressed pseudogene • GKN3 and TFF2 protein are unlikely to form heterodimer
Geahlen et al[36]	• GKN3 is a stomach specific protein expressed in a gastric mucous in mice and pigs • GKN3 protein was not detect in normal gastric tissue of patients, about half of which were heterozygous for the A/G allele • The transcripts for GKN3 gene in humans were not detected, no matter the genotype, which may indicate its pseudogenicity

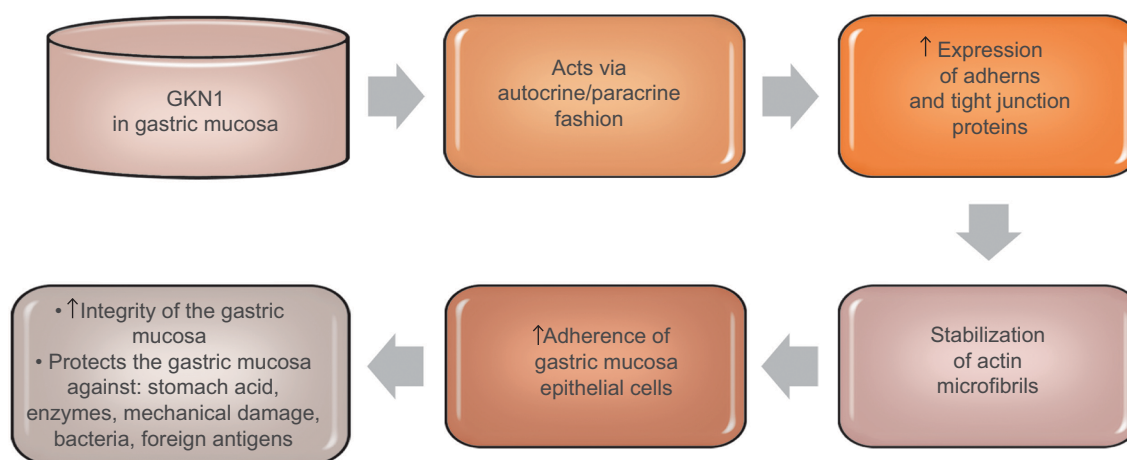**Abbreviation:** *H. pylori*, Helicobacter pylori.

Figure 2 GKN1 role in gastric mucosa.

role in maintaining cell integrity, and their expression in the gastric mucosa also decreases in inflammation and gastric cancer.[21,29,35] Gastrokines show similarity in 46% of their amino acid sequences, and the characteristic linking these proteins is the presence in their structure of the BRICHOS domain consisting of ~100 amino acids and associated with the occurrence of gastric cancer.[37–39]

GKN1 and gastric cancer

In vitro studies have shown that both GKN1[18,23,24,40,41] and GKN2[29] occur in the gastric mucosa of healthy individuals. On the other hand, in the gastric mucosa of gastric cancer patients compared to subjects with superficial gastritis, a decreased GKN1 protein and mRNA *GKN1* expression is noted.[42,43] GKN1 was also absent in patients with intestinal metaplasia.[43] Perhaps the reason for reduced expression or absence of *GKN1* in these diseases is damage to the gastric mucosa.

GKN1 acts on cells in an autocrine or paracrine fashion.[18] The studies of Oien et al[24] showed the expression of GKN1 in the gastric mucosa of both the surface epithelium and gastric ulcer. However, no GKN1 expression was found in the deeper parts of the gastric glands. The results of previous studies suggest that GKN1 is a gastric specific mitogenic protein and most likely plays a protective and reparative role in the gastric mucosa.[23,24,44] GKN1 maintains the integrity of the gastric mucosa, protects it against the action of stomach acid and enzymes as well as mechanical damage, bacteria or foreign antigens.[17,19,44] Walsh-Reitz et al[45] conducted studies in mice that were administered dextran sulfate sodium to damage the intestinal mucosa and then injected with GKN1. It was found that under the influence of GKN1 the

immunoreactivity of occludens proteins increased, forming tight connections responsible for the adherence of epithelial cells in the colon mucosa. In addition, during the exposure of tested epithelial cells to monochloramine, it was shown that GKN1 protects them from loss of integrity by increasing the accumulation of intercellular proteins (zonula occludens-1, claudin 5, E-cadherin) and stabilization of actin microfibrils.[45] These findings were further confirmed by studies of Rippa et al,[46] which showed that in normal gastric tissue GKN1 is co-localized with actin and GKN1 overexpression in gastric cancer cells leads to increased expression of adherens and tight junction proteins, which indicates this protein as a main factor helpful in actin stabilization. Figure 2 schematically explains GKN1 role in gastric mucosa.

Toback et al[47] in the in vitro studies demonstrated the possibility of using GKN1 to stimulate and accelerate the regeneration of damaged epithelium. Some authors[19,37,41,48,49] attribute GKN1 as a possible suppressor function in stomach cancer. Yoon et al[19] have shown that GKN1 limits the formation of the MKN-28 gastric cancer cell line colony. GKN1 causes the biological aging of gastric cancer cells leading to their death by activating the p16/Rb and p21 pathways and Ras/Raf/MEK/ERK cell signaling.[18] Xing et al[49] showed that GKN1 inhibits gastric cancer cell invasion via downregulation of MMP2 expression through the NF-κB pathway, which shows that it plays a role in inhibition of cancer metastasis and indicates GKN1 as a novel potential therapeutic target. It is also suggested that GKN1 is able to inhibit migration and invasiveness of gastric cancer cells via downregulation of RhoA protein expression in a miR-185- and miR-34a-dependent manner. Yoon et al[50] found that RhoA GKN1 expression and *GKN1* mRNA were increased in gastric cancer tissues

and related to TNM stage. Also, Yan et al[40] in an in vitro study attempted to explain how GKN1 inhibits the growth of gastric cancer cells. The authors analyzed the effect of GKN1 on the SGC-7901 gastric cancer cell line lacking expression of the tested protein. GKN1 has been shown to inhibit tumor cell growth and reduce the number of cell colonies by stopping the G2/M cell cycle instead of inducing apoptosis.[40] On the other hand, Yoon et al[19] found that in the AGS gastric cancer cell line transfected with *GKN1* both inhibiting the proliferation of these cells and activation of the apoptosis process were induced. In the group of patients with gastric cancer, the lack of or decrease in expression of *GKN1* gene was detected by Western blot analysis. In addition, it was found that the number of DNA copies and transcripts of *GKN1* mRNA were statistically significantly reduced in this study group.[19]

It is suggested that inactivation of *GKN1* gene may play an important role in the development of some cases of gastric cancer. GKN1 also regulates protein kinase activity, including inhibition of delta/theta protein kinase activity, or increases ERK 1/2 and JNK 1/2 activity, which has inhibitory effect on the growth of gastric cancer cells. In the studies of Yan et al,[40] it was found that GKN1 affects the activity of 74 proteins, including α-enolase (ENO1) and cathepsin D. ENO1 is an enzyme involved in the metabolism of glycogen, in the presence of magnesium ions, reversibly converts the 2-phosphoglycerate to phosphoenolpyruvate at the phosphotriese stage.[51] In addition, ENO1 exhibits a number of nonenzymatic activities, eg, as a surface protein of epithelial cells, endothelium, fibroblasts, cancer cells or some hematopoietic cells, it acts as a plasminogen receptor, and as a ligand binding cholesterol esters causes the accumulation of cholesterol esters in cells (affects the process of atherosclerosis, multiple sclerosis and encephalopathy). The increase in ENO1 expression has been demonstrated also in small-cell lung carcinoma, breast cancer, head and neck cancer.[40,52] During proliferation, tumor cells have a high demand for ATP, therefore the rate of aerobic glycolysis is elevated and increased expression of enzymes in the glycolysis pathway is observed. ENO1 present in the cytosol of lung cancer cells by association with cytoskeleton proteins facilitated the migration of these cells. Conversely, ENO1 presented on the surface of non-small-lung carcinoma cells caused a local increase in plasminogen density, which accelerated the degradation of fibrin and extracellular matrix proteins, also affecting invasion of cancer cells and metastasis.[51,52] It was shown that in gastric cancer cells the reduction of ENO1 activity and overexpression of GKN1 caused growth and cell cycle arrest. In contrast, overexpression of ENO1 blocked the inhibitory

effect of GKN1 on the growth of tumor cells and the arrest of the cell cycle.[40] Therefore it was suggested that the "down regulation" phenomenon of ENO1 may play an important role in inducing inhibition of gastric cancer cell growth. Rippa et al[38] found that the increased expression of GKN1 in the cell lines (AGS and MKN-28) of gastric cancer stimulated the expression of the Fas receptor which induced programmed death of these cells. The results of research by Rippa et al[38] suggest that GKN1 may be a significant protein modulating apoptotic signal and performs the reparative role of damaged tissues in early stages of neoplastic transformation.

Excessive expression of GKN1 may also inhibit epithelial-mesenchymal transformation of mesothelial cells (epithelial-mesenchymal transition [EMT]) by inactivation of the PI3K/Akt pathway and reduction of ROS.[19] In addition, GKN1 expression is accompanied by increased expression of E-cadherin and reduction of cytoplasmic and nuclear expression of β-catenins, fibronectin, vimentin and cyclin D1. As a result EMT mesothelial cells receive a phenotype of fibroblastic cells expressing α-actin of smooth muscle and the ability to migrate. These data suggest that the *GKN1* gene, in sporadic cases of gastric cancer, can play an important role in its progression by inhibiting EMT and tumor cell migration. Studies by Zhang et al[29] have shown a large accumulation of β-catenin in the nucleus of gastric cancer cells. In addition, nuclear expression of β-catenin occurred in normal cells. The authors suggest that in this case it may be a paraneoplastic element. The conclusion is that the accumulation of β-catenin in the nuclei of normal cells of the mucous membrane may be an alarm signal that there is a probability of neoplastic changes in neighboring cells.[29] Some literature data indicate that during *H. pylori* infection due to the action of cytotoxin-associated gene A (CagA) and proinflammatory cytokines, nuclear accumulation of β-catenin and disturbance of the β-catenin dependent signaling pathway may also occur.[53] These results partially explain why colonization of the gastric mucosa by *H. pylori* increases the risk of cancer development. Figure 3 summarizes suggested GKN1suppressor roles in gastric cancer.

The mechanism by which gastrokine 1 is inactivated in gastric cancer cells is still not fully understood and obtained results are controversial.[19,54,55] Yoon et al[19] did not detect *GKN1* gene mutation in gastric cancer, whereas the hypermethylation of *GKN1* promoter was found only in two cases. Conversely, Lu et al[54] revealed that *GKN1* and *GKN2* genes are transcriptionally silenced by methylation of DNA. They also found that the Epstein-Barr nuclear antigen 1 (EBNA1) binds to different promoters of the *GKN1* and *GKN2* genes in

Figure 3 GKN1 suppressor roles in gastric cancer.

gastric cancer cells. The authors suggest that EBNA1 leads to the complex transcriptional and epigenetic deregulation of these tumor suppressor genes in Epstein–Barr virus positive gastric cancer.[54] Altieri et al[55] tested how *GKN1* histonic modification may lead to its downregulation. The authors, for this purpose, immunoprecipitated from human normal and cancerous gastric tissues chromatin for the trimethylation of histone 3 at lysine 9 (H3K9triMe) and its specific histone-lysine N-methyltransferase (SUV39H1). They found that epigenetic mechanisms responsible for *GKN1* silencing in gastric cancer samples were related to high H3K9triMe levels and SUV39H1 recruitment to the *GKN1* promoter. *GKN1* mRNA levels were markedly increased after the inhibition of histone deacetylases with trichostatin A.[55]

GKN1 and *H. pylori* infection

H. pylori is a Gram-negative, spiral shaped bacterium commonly found in the human population. In 1994, the International Agency for Research on Cancer at the WHO

indicated *H. pylori* as a class I carcinogen, whose participation in the formation of gastric cancer has been proven. Colonization of the gastric mucosa by *H. pylori* leads to a local inflammatory reaction with the production of ROS and increases the concentration of nitric oxide, which can affect the DNA of these cells and cause mutations resulting in the development and progression of gastric cancer.[44] It has been observed that *H. pylori* infection leads to the phenomenon of downregulation and lack of expression of GKN1.[24,44,56,57] The number of *GKN1* mRNA copies in patients with chronic gastritis infected with *H. pylori* was statistically significantly lower compared to the group of patients without infection.[44] In addition, the analysis of Western blots of gastric mucosa biopsies in patients with gastric cancer and in patients with premalignant disease did not show gene expression for *GKN1*,[24] and in the reverse transcriptase PCR (RT-PCR), statistically significantly lower *GKN1* mRNA levels were found in patients with *H. pylori* infection.[19] Activation of proinflammatory cytokines such as TNF-α, IL-1β, IL-6 and

IFN-γ occurring in the course of *H. pylori* infection may also inhibit the expression of GKN2 in gastrointestinal mucosa.[21] Moss et al[20] evaluated the expression of *GKN1* and *GKN2* mRNA and the expression of these proteins in biopsies of normal gastric mucosa cells and tumor-altered biopsies. They showed statistically significantly lower expression of GKN1 and GKN2 in gastric adenocarcinoma, especially in the scattered cancer subtype compared to the control group. The loss of GKN1 and GKN2 expression was an unfavorable prognostic marker in surgically treated patients because it correlated with a shorter survival time.[20]

On the molecular level, *H. pylori* CagA directly injected into gastric epithelial cells drives morphology changes of these cells, leads to their apoptosis, increases cell proliferation and motility, and thus promotes gastric carcinogenesis. Additionally, CagA induces activation of NF-κB and PI3K/ Akt signaling pathways and EMT-related proteins.[58] The role of GKN1 relies on inhibition CagA injection into gastric epithelial cells, reduction of ROS via a positive regulatory effect on antioxidant enzymes production, stimulation the negative cell cycle regulators expression (p53, p16 and p21), and induction of miR-185 expression which inhibits genomic DNA epigenetic modification. Moreover, both on human and mice gastric cells and mucosal tissues, it was shown that *H. pylori* CagA decreases gene copy number for *GKN1* and GKN1 expression. Interestingly, increased GKN1 expression decreased the carcinogenic effects of *H. pylori* CagA through binding to CagA, which strongly indicates gastrokine 1 as a potential therapeutic target.[58]

GKN1 and acetylsalicylic acid (ASA)

ASA has anti-inflammatory, analgesic and antipyretic properties, and long-term administration also has anticoagulant activity. ASA is a cyclooxygenase inhibitor which inhibits prostaglandin production and contributes to the reduction of mucus secretion that protects the gastric mucosa. Therefore, about 25% of patients receiving ASA suffer damage to the gastrointestinal mucosa in the form of stomach or intestinal erosions or ulcers.

The aim of the study of Martin et al[23] was to show whether receiving low doses of ASA affects the expression of GKN1 in antral and corpus mucosa. The study group consisted of healthy volunteers without *H. pylori* infection who were taking ASA at a dose of 100 mg/day for 7 days. GKN1 baseline was similarly expressed in both antral and corpus mucosa and it was not changed after 3 days of ASA administration. After 7 days GKN1 expression level was significantly reduced in gastric antrum, whereas in gastric corpus GKN1 transcript

levels were slightly elevated. Mild gastritis and mucosal erosions were limited only to gastric antrum and correlated with GKN1 expression.[23] In an animal model study, intragastric administration of indomethacin showed a decrease in GKN1 concentration.[47] The authors suggest that GKN1 is involved in maintaining the integrity of gastric mucosa cells and stimulates repair mechanisms after its damage. Indeed, exogenous GKN1 applied in vitro to gastrointestinal cells stimulated their reconstruction.[47]

Circulating GKN1 evaluation – diagnostic potential

Currently carbohydrate antigen (CA) 19-9, carcinoembryonic antigen (CEA), and CA 72-4 are used for gastric cancer: diagnosis, monitoring of recurrence, distal metastasis presence, chemotherapy evaluation and prognosis. However, the interpretation of CEA, CA 19-9 and CA 72-4 evaluation should be performed carefully, especially when the tumor biomarker concentration is borderline or only slightly above the reference value. Moreover, the above-mentioned tumor biomarkers are not useful in diagnosis and screening of early gastric cancer. Elevated concentrations of CEA, CA 19-9 and CA 72-4 were observed also in other types of tumors as well as conditions not related to malignancy.[59] It is also highlighted that the combined testing of CEA, CA 19-9 and CA 72-4 is superior compared to single biomarker testing, as it significantly increases their diagnostic usefulness, which is briefly summarized in Table 2.[60–69] Because of these limitations scientists are still looking for an ideal gastric cancer tumor biomarker.

In the available literature there are only two research papers evaluating quantitatively circulating concentration of GKN1 protein or *GKN1* mRNA in individuals with gastric cancer.[70,71] For example Villano et al[70] did not confirm the utility of serum *GKN1* mRNA evaluation in gastric cancer patients compared to serum samples from healthy individuals. However, Yoon et al[71] did indicate the potential clinical application of serum GKN1 protein evaluation by means of ELISA in gastric cancer. The authors showed that GKN1 concentration was significantly lower in gastric cancer subjects compared to healthy individuals, and moreover to the hepatocellular carcinoma as well as colorectal cancer patients.[71]

The relation between the sensitivity and specificity of the test is illustrated using a receiver operating characteristic (ROC) curve, which is generated to calculate the area under the ROC curve (AUC). The ROC curve is a line graph that plots the probability of true positive results – or the sensitivity of the test – against the probability of false positive results

Table 2 Combined testing results of tumor markers in gastric cancer.

Authors	Patients	Tumor markers	Conclusions
Xu et al[60]	GC (n=50) Benign GC (n=50) Healthy control (n=50)	CEA, CA 19-9, CA 72-4, CA 242	The tumor incidence for combined CEA, CA 19-9, CA 72-4, CA 242 testing was higher than that of single tumor marker testing
Ning et al[61]	GC (n=169) Healthy control (n=75)	CEA, CA 19-9, CA 72-4, TK1	The highest AUC (0.895) for single tumor biomarker revealed TK1 The combined TK1, CA 19-9, CA 72-4 and CEA testing showed higher AUC (0.953) than for each tumor marker tested separately
Feng et al[62]	Early GC (n=587)	CEA, CA19-9, AFP, CA 125	The positive rates of CEA, CA 19-9, AFP and CA 125 were relatively low: 4.3%, 4.8%, 1.5% and 1.9%, respectively The positive rate of combined CEA, CA 19-9, AFP and CA 125 testing equaled 10.4%
Chen et al[63]	GC (n=87) Healthy control (n=40)	CEA, CA 19-9, CA 72-4, CA 15-3, CA 125	The combined CEA, CA 72-4, CA 19-9 and CA 125 testing had a higher positive rate (detection rate 60.9%) and diagnostic value for GC than the single tumor markers evaluation
Virgilio et al[64]	GC (n=38) non-GC (n=41)	CEA, CA 19-9, CA 72-4, CA 50	Gastric lavage cytopathology correlated significantly with advancement of tumor invasion depth and the presence of local lymph node metastases Gastric lavage CEA, CA 19-9, CA 72-4, or CA 50 concentration did not achieve a statistically significant correlation with cancer staging
Liang et al[65]	GC (n=2,288) Healthy control (n=1,869)	CEA, CA 19-9, CA 72-4	The sensitivity of CEA, CA 19-9 and CA 72-4 in the GC diagnosis was 20.1%, 21.4% and 27.6%, respectively and increased to 48.2% when tumor markers were used in combination
Rehena et al[66]	Endoscopically suspected GC (71)	CEA, CA 72-4	CA 72-4 sensitivity (48.3%) was higher compared to CEA (31%) in diagnosis of gastric cancer, therefore CA 72-4 is recommended for use in conjunction with other diagnostic tests (eg, endoscopy)
Yu et al[67]	GC (n=216)	CEA, CA 19-9, CA 72-4	The combined positive rate of CEA, CA 19–9 and CA 72–4 was significantly higher compared with the individual CEA, CA 19–9 and CA 72–4 positive rates (44.91% vs 22.69%, 18.98% and 22.69%, respectively)
Yin et al[68]	GC (n=45) GBD (n=40) Healthy control (n=30)	CEA, CA 19-9, CA 72-4, TSGF	The AUCs for single detection of CEA, CA 72-4, CA 19-9 and TSGF in the diagnosis of GC equaled 0.833, 0.805, 0.810 and 0.839, respectively The AUC for combined testing of tumor markers revealed 0.913
Yang et al[69]	GC (n=106) GBD (n=149) Healthy control (n=171)	CEA, CA19-9, CA 72-4, CA 125	The sensitivities of CA72-4, CEA, CA 125 and CA 19-9 at the recommended cut-off level for all patients were 33.0%, 25.5%, 31.1% and 38.7%, respectively When all tumor markers were used in combination the sensitivity increased to 66.0%

Abbreviations: AFP, alpha fetoprotein; AUC, area under receiver operating characteristics curve; CA 125, cancer antigen 125; CA 15-3, cancer antigen 15-3; CA 19-9, cancer antigen 19-9; CA 242, cancer antigen 242; CA 50, cancer antigen 50; CA 72-4, cancer antigen 72-4; CEA, carcinoembryonic antigen; GBD, benign gastric diseases; GC, gastric cancer; TK1, an enzyme involved in the regulation of the mammalian cell cycle; TSGF, tumor-specific growth factor.

for a range of different cut-off points.[72] In the study of Yoon et al[71] AUCs in differentiating gastric cancer patients from healthy subjects, hepatocellular carcinoma and colorectal cancer patients were: 1.00, 0.99 and 0.99, respectively. Moreover, Yoon et al[71] found that GKN1 serum concentration was also useful for the assessment of gastric cancer cell invasion, as protein concentration was significantly higher in early gastric cancer (EGC) compared to advanced gastric cancer. EGC patients, based on the GKN1 concentration in the serum, additionally could be distinguished from: healthy subjects (AUC =1.00), atrophic gastritis patients (AUC =1.00), and individuals with atrophy and intestinal metaplasia (AUC =0.98). Interestingly, authors did not find differences between males and females for serum GKN1 concentration. It should be noted that the measurement of

GKN1 concentration was possible only after the incubation of samples at 70°C for 10 minutes; under nonheating conditions Yoon et al[71] did not detect GKN1 in human sera. Nevertheless the aspect of utility of GKN1 serum quantitative evaluation requires further studies.

Conclusion

Gastric cancer is one of the most common lethal cancers worldwide. GKN1 is a protein specific for stomach, and its role consists of maintaining mucosal integrity as well as the replenishment of the surface lumen epithelial cells layer. The evaluation of GKN1 expression seems to be a useful indicator of the presence of neoplastic or inflammatory lesions in the gastric mucosa. Because GKN1 expression is decreased in gastric tumor tissues and derived cell lines and

its upregulation in cell lines of gastric cancer induces apoptosis, a suppressor role of this protein is suggested in tumor progression. However, the mechanism by which GKN1 is inactivated in gastric cancer cells is still not fully understood and obtained results are controversial. In the available literature few studies evaluate quantitatively circulating concentration of GKN1 protein or *GKN1* mRNA in gastric cancer individuals. The future diagnostic capabilities of gastric cancer concern the assessment of serum GKN1 concentration by means of ELISA. Serum GKN1 concentration did not differ depending on patients' sex. Interestingly, the measurement of GKN1 concentration is possible only after the incubation of samples at 70°C for 10 minutes. Nevertheless, the aspect of quantitative serum GKN1 evaluation is new in the context of available literature and requires further studies.

Acknowledgments

Author contributions

All authors contributed toward data analysis, drafting and revising the paper, gave final approval of the version to be published and agree to be accountable for all aspects of the work.

Disclosure

The authors report no conflicts of interest in this work.

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