

Review

TRP channels in gastric cancer: New hopes and clinical perspectives

Andra M. Sterea^a, Emmanuel E. Egom^b, Yassine El Hiani^{a,*}^a Departments of Physiology & Biophysics, Dalhousie University, Halifax, Nova Scotia, Canada^b Egom Clinical & Translational Research Services Ltd, Halifax, Nova Scotia, Canada

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ABSTRACT

Gastric cancer is a multifactorial disease associated with a combination of and environmental factors. Each year, one million new gastric cancer cases are diagnosed worldwide and two-thirds end up losing the battle with this devastating disease. Currently, surgery represents the only effective treatment option for patients with early stage tumors. However, the asymptomatic phenotype of this disease during the early stages poses as a significant limiting factor to diagnosis and often renders treatments ineffective. To address these issues, scientists are focusing on personalized medicine and discovering new ways to treat cancer patients. Emerging therapeutic options include the transient receptor potential (TRP) channels. Since their discovery, TRP channels have been shown to contribute significantly to the pathophysiology of various cancers, including gastric cancer. This review will summarize the current knowledge about gastric cancer and provide a synopsis of recent advancements on the role and involvement of TRP channels in gastric cancer as well as a discussion of the benefits of targeting TRP channel in the clinical management of gastric cancer.

1. Epidemiology

In 2018, the Global Cancer Statistics reported 18.1 million new cancer cases and 9.6 million cancer-related deaths of which gastric cancer represents 5.5% and 8.3%, respectively [1]. Interestingly, the incidence of gastric cancer shows a wide geographical distribution with the highest rates found in Eastern Asia, Eastern Europe, and Southern America, and lowest rates in Northern America, Northern Europe, Australia and New Zealand [2–5]. Obvious clustering of gastric cancer incidence also exists within individual countries [6–9]. For example, in China, the northern populations show higher incidence and mortality rates than the southern populations [10–13] which indicates the presence of an environmental component that contributes, at least in part, to the observed regional variation of gastric cancer occurrence. This is consistent with migration studies that have reported a decrease in the incidence/mortality of gastric cancer among second-generation Chinese and Japanese migrants to the US (the overall gastric cancer incidence of Asian migrants remains higher in the host population) [14,15]. In addition, this variation also exists within genders with 683,754 of the new gastric cancer cases being diagnosed in males and 349,947 in females (equating to 2/3 of all gastric cancer occurring in the male population) [16,17]. This difference was highlighted in an age-standardized study conducted in Korea where the incidence rate of gastric cancer in men

was 32.1 per 100 000 compared to 13.2 per 100 000 in females [1,18,19]. However, despite the unsettling statistics, a steady decline in the incidence of gastric cancer has been observed over the last few decades in both, developed and developing regions. A 2.3% annual decrease in incidence rates is predicted until 2030 which is likely a result of improved hygiene and food conservation methods (for specific risk factor see Etiology section) [20–22].

2. Etiology

Cancer has long been known as a disease of many causes, some that have yet to be uncovered and some that have been established as the drivers of carcinogenesis. Today, there is a consensus that *Helicobacter pylori* (*H. pylori*) is the most important risk factor that contributes to the development of the majority of gastric cancers [23–26]. It is believed that chronic inflammation induced by *H. pylori* facilitates the epigenetic alteration of tumor suppressors genes and paves the way for uncontrolled cell proliferation, apoptosis evasion and cell invasion [27,28]. In addition, *H. pylori* infections can often result in the development of gastritis and polyps [29–32] which can ultimately lead to gastric carcinogenesis [33–36]. Furthermore, the etiology of gastric cancer has also been linked with another pathogen known as the Epstein–Barr virus (EBV) [37–39]. In contrast to *H. pylori* which can be

* Corresponding author at: Dalhousie University, Department of Physiology and Biophysics, 5G, Sir Charles Tupper Building, 5850 College street, Halifax, Nova Scotia, B3H 1 × 5, Canada.

E-mail address: yassine.elhiani@dal.ca (Y. El Hiani).

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found in normal and malignant gastric cells, increasing evidence shows that the EBV pathogen is found only in gastric cancer cells. However, the mechanism of action behind the ability of EBV to cause gastric cancer remains largely unknown [40].

Despite the contribution of extracellular microbes to the pathogenesis of gastric cancer, studies have provided enough evidence to suggest a “sweet and salty” relationship between dietary consumption and the incidence of gastric cancer [41–43]. The latest estimate of the World Cancer Research Fund International indicated that the daily intake of 5 g of dietary salt increased the relative risk of gastric cancer by 12% [44]. The proposed mechanism behind this phenomenon is the potential role that salt plays in mediating *H. pylori* colonization and maintaining the subsequent chronic inflammation [45–47]. On the other hand, studies have also shown that the daily consumption of 100 g of fruit lowered the risk of developing gastric cancer by 5% [22,48,49]. Health conscious dietary choices have always been regarded as the best medicine for many illnesses and in the case of gastric cancer, this theory proves true as the risk for gastric cancer has been linked to obesity [50–54]. Studies investigating the impact of a high body fat index and excess body weight on overall health have identified obese individuals as being at higher risk for gastric cancer [54,55]. This is in part due to the prevalence of *H. pylori* infections in obese patients [53]. Furthermore, other unhealthy habits such as alcohol consumption and tobacco use have also been associated with an increased risk of developing gastric cancer [56–62].

In addition to the aforementioned risk factors and although rare, there is also an inherited component that contributes to gastric cancer [63]. About 10% of reported gastric cancers are found among family members, but less than 3% are truly hereditary [64,65]. Currently, only a few high-penetrance genes have been identified in hereditary gastric cancer and these include *CDH1* and *CTNNA1* [40]. The *CDH1* gene encodes the cadherin 1 (also known as E-cadherin) protein while the *CTNNA1* gene is translated into the catenin a-1 protein (also known as a-E-catenin), both of which facilitate cell-adhesion and gene transcription. Attempts to map the genetic landscape of gastric cancer have also uncovered the presence of mutations in *TP53* (a tumor suppressor gene) and *SKT11* (regulator of autophagic processes), two mutations which are commonly found in many other cancers as well [40,63,66,67]. However, our knowledge of the genetic makeup of gastric cancer is limited and studies are currently ongoing to decipher the genetic predisposition underlying pre-cancerous gastric lesions.

3. Classification

There are several classification systems used for stomach cancer, but the most commonly used are the Lauren and World Health Organization (WHO) systems [40,68]. The Lauren classification distinguishes gastric cancer into two main classes based on histological differences: intestinal and diffuse [69]. The intestinal type tumors often occur in the antrum and incisura portions of the stomach and consist mainly of well differentiated cells that have a slow growth rate and the tendency to form glands. The incidence of this type of gastric cancer is often related to environmental factors and occurs predominantly in men and elderly individuals. In contrast, the diffuse type involves the entire stomach and it is made up of poorly differentiated cells lacking intercellular adhesions that tend to scatter throughout the stomach, which explains the increased incidence of metastasis and poor prognosis associated with this type of gastric cancer. Furthermore, the diffuse type is typically correlated with genetic abnormalities and is more frequently diagnosed in females and young patients [68]. The Lauren classification is complemented by the WHO classification which offers more histological patterns and differentiates gastric cancer into five groups: tubular (the most common type found during the early stages of the disease which consists of small branching tubules), papillary (occurs predominantly in the proximal stomach of older patients), mucinous (characterised by a glandular architecture and extracellular mucinous

pools), poorly cohesive (made up of signet ring cells arranged in clumps), and mixed carcinoma (involves a mix of different types of stomach adenocarcinomas) [70].

4. Diagnosis and treatment

Early stage gastric cancer is often silent or accompanied by non-specific symptoms including heartburn, nausea and poor appetite, all of which can be attributed to other illnesses. The characteristic symptoms (stomach pain with weight loss, and/or blood in stool) that allow for a differential diagnosis of gastric cancer usually emerge during the late stages or once the cancer has metastasized [40,63]. The diagnosis of gastric cancer relies primarily on imaging tests such as endoscopic ultrasounds (and laparoscopy) combined with biopsy [71–74]. The endoscopic ultrasound uses sound waves to create pictures of the inside of the stomach allowing for the identification of abnormal/tumoral structures that can be later confirmed using biopsies and immunohistochemistry (IHC) or fluorescent in situ hybridization (FISH). The IHC test uses antibodies targeting the human epidermal growth factor receptor-2 (HER2) protein which has been demonstrated to be overexpressed in 20% of gastric cancers [75,76]. HER2 exhibits a progressive increase in expression which corresponds to the grade/stage of the gastric tumor (e.g. lower in low grade dysplasia and higher in gastric carcinomas) which establishing HER2 assessment as an accurate testing measure for the diagnosis of gastric cancer patients [75]. The HER2 IHC test generates a score between 0 and 3+ [77]. If the results are 0 or 1+, the cancer is classified as HER2-negative and patients are not treated with HER2-targeted therapies. If the score is 3+, the cancer is considered HER2-positive and patients can be treated with drugs targeting HER2, in particular trastuzumab. However, when the score is 2+, the HER2 status of the tumor is unclear and requires further testing. Often, tumors with a score of 2+ are tested using the FISH method which utilizes fluorescent pieces of DNA that specifically target *HER2* [40]. Once the cancer is detected and has an established HER2 status, laparoscopy is used to exclude the presence of small metastatic tumors and confirm that the tumor remains confined within the boundaries of the stomach [78–80].

Despite technological advancements and improvements in detection strategies, surgical resection remains the only therapeutic option that offers the best prognosis for gastric cancer patients [81]. Different types of surgery can be used to treat gastric cancer patients depending on the extent of the tumor(s) (e.g. endoscopic resection for small tumors, subtotal/total gastrectomy for advanced tumors) [82,83]. Surgery is typically coupled with the appropriate combination of chemotherapy drugs such as platinum, fluoropyrimidines or taxanes [84–87]. However, other treatment options have also been implemented and have shown some success over the years. Prompted by advances in molecular profiling and characterization of gastric cancer, two targeted therapies have been identified: anti-HER2 therapy (Trastuzumab) and anti-VEGF therapy (Ramucirumab) [88]. While these two treatment modalities have paved the way towards precise and personalized medicine, there are still challenges that pose a significant strain on the efficacy of these therapies as well as on the quality of life of the patient. However, recent discoveries have shifted the scientific focus towards a new class of potential therapeutic targets, the transient receptor potential (TRP) channels [89–93]. TRP channels are often regarded as the Jack of all trades due to their extensive involvement in various physiological processes necessary for cell growth and survival, a characteristic which also renders them as suitable candidates for cancer therapy. The features that make TRP channels a favourable anti-cancer target can be summarized into three main points: 1) TRP channels are overexpressed in some cancers; 2) their activation triggers Ca^{2+} signalling cascades that have been shown to contribute to several hallmarks of cancer including uncontrolled proliferation, apoptosis-resistance, and cell migration and invasiveness, 3) most TRP channels are found on the cell surface which makes them more accessible to targeted therapies. The

Current therapies

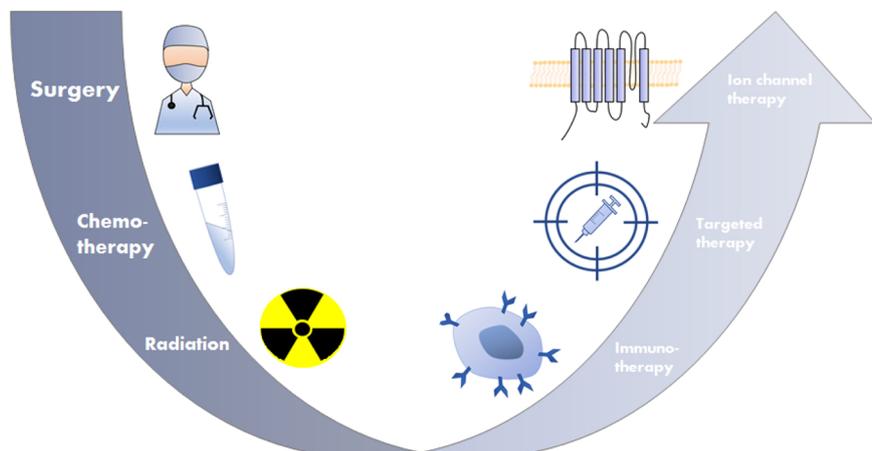


Fig. 1. Schematic depiction of the current and emerging therapies for gastric cancer patients. Surgery has long been the therapy of choice for gastric cancer patients (or any cancer type), in particular those with localized tumors and earlier stages of the disease. However, patients with late stage are often treated with chemotherapy, radiation or a combination of surgery, chemotherapy or radiation. Although successful to a certain extent, these approaches are not always curative and achieve varying clinical outcomes which are largely attributed to tumor heterogeneity. It is these challenges that led to the development of new therapies. Emerging cancer therapies are moving towards personalized medicine where each patient is treated according to their tumor expression profile and genetic makeup. These new therapy options include immunotherapy and targeted therapy which have shown good efficacy at

treating gastric cancer patients, but various limitations still pose a significant strain on their effectiveness. Studies are now being conducted to assess the involvement of ion channels in the pathophysiology of gastric cancer in efforts of improving the efficacy of cancer treatment. Thus far, the TRP ion channel family (shown at the head of the arrow) has been demonstrated to play significant roles in gastric cancer cell survival and are currently under investigation as potential therapeutic targets for gastric cancer patients.

following sections of this review will introduce the TRP channel family and further discuss whether their function could be exploited for targeted cancer therapies (Fig. 1).

5. TRP channels (358 words)

TRP channels were first identified in 1969 by Cosens and Manning, but their structure and function remained a mystery until 1990s when studies revealed that these proteins harboured the defining characteristics of ion channels [94]. This initial discovery was carried forward and TRP channels have since been identified in various animal groups including the mammalian kingdom. In humans, TRP channels consist of transmembrane proteins typically assembled as homo- or hetero-tetramers that form gated pores and function to regulate the ion distribution across cellular compartments. Currently there are 28 members in the TRP superfamily which have been divided into six subgroups: TRPA (ankyrin), TRPC (canonical), TRPM (melastatin), TRPML (mucolipin), TRPP (polycystin) and TRPV (Vanilloid) [92,93]. Structurally, all members of the TRP family share a similar architecture consisting of six transmembrane segments with a pore flanked between the 5th and 6th segments, and cytoplasmic N- and C-termini [95,96]. In contrast to their shared structural similarities, physiologically, each member has a unique ionic signature, mechanism of activation and response to stimuli [97]. Notably, most TRP proteins are non-selective cation channels permeable to Ca²⁺ [98,99] – except for TRPMs 4 and 5 [100]. Ca²⁺ acts as a universal second messenger that interacts with a plethora of intracellular Ca²⁺-binding proteins to regulate a wide range of cellular

functions, from cell survival to cell death [101–103]. It has been suggested that the shape of the Ca²⁺ signals determines the cell's decision to differentiate, divide or die. For instance, local Ca²⁺ oscillations, waves and sparks have been associated with Ca²⁺-mediated cell survival whereas a global sustained cytosolic elevation of Ca²⁺ has been linked to cell death [104]. When open, TRP channels mediate Ca²⁺ entry which activates Ca²⁺ signaling pathways (e.g. Ca²⁺/Calmodulin kinase II (CaMKII), protein kinase C (PKC) and mitogen-activated protein kinase (MAPK)) which, in turn, shape the fate of a myriad of cell processes such as proliferation, apoptosis and migration [99,105–109]. Given the relationship between these processes with cancer cells and their ability to withstand stress and proliferate uncontrollably, it is not surprising that any alteration in the expression/function of TRP channels [89,92,110,111] and the subsequent disruption of Ca²⁺ homeostasis are often associated with the pathophysiology of cancer cells. Functional studies have demonstrated the incidence of altered expression and function of various TRP channels in many types of cancer including gastric cancer [112]. Furthermore, up to 30% of gastric cancer patients develop hypercalcemia and often have a poorer prognosis [113–115]. Over the past two decades, numerous research groups have identified six TRP channels (TRPC6, TRPM2, TRPM5, TRPM7, TRPV4, and TRPV6) that play an important role in the growth and survival of gastric cancer. These channels will be described in detail in the following paragraphs (Table 1).

Table 1
TRP channels involved in gastric cancer and their function.

Channel family	Channel name	Involvement in gastric cancer
TRPC	TRPC6	Upregulated in GC Promotes GC cell growth and expression of EMT markers [120,121,122,123]
TRPM	TRPM2	Increased expression correlates with poor GC patient survival Regulates GC metabolism, enhances invasion and promotes GC survival Regulates gastric inflammation during <i>H. pylori</i> infection [127,128,157]
	TRPM5	Increased expression correlates with poor GC patient survival [129]
	TRPM7	Upregulated in GC Maintains GC cell proliferation and survival [132,133,134]
TRPV	TRPV4	Sustains GC survival Promotes GC cell proliferation [143,144]
	TRPV6	Upregulated in GC Activation induces GC cell death [145,146]

5.1. TRPC channels

The TRPC family includes seven members (TRPC1 through TRPC7), all of which function as mediators of Ca^{2+} -dependent signal transduction in response to G protein-coupled receptor activation and/or endoplasmic reticulum store depletion [116]. Longstanding evidence has demonstrated the expression of TRPC in different types of cancer cells. For instance, the expression of TRPCs 1, 3 and 4 was shown to influence the differentiation of non-small cell lung cancer cells [117]. While TRPC5 and TRPC7 have been demonstrated to contribute to the carcinogenesis of gliomas [118] and neuroblastoma [119], respectively. In gastric cancer, research is now beginning to uncover the central role of TRPC6. Cai et al., provided the first evidence to show that the expression level of TRPC6 is upregulated in human gastric cancer tissues as compared to the surrounding normal gastric tissues. In the same study, the authors demonstrated that TRPC6 is functionally expressed in gastric cancer cell lines and that pharmacological inhibition of this channel significantly reduced cell growth through G2/M cell cycle arrest [120]. These *in vitro* results were complemented with *in vivo* studies showing that inhibition of TRPC6 suppressed gastric cancer tumor growth in nude mice [120]. In addition to its proliferative role, TRPC6 was also found as a key player in gastric cancer epithelial-to-mesenchymal transition (EMT) [121]. EMT is the most common process involved in cancer metastasis and increased expression of EMT markers is often associated with poor prognosis [122]. Ge et al. demonstrated that the loss of TRPC6 function is concomitant with a reduction in the expression levels of many EMT proteins such as E-cadherin, vimentin and alpha smooth muscle actin (α -SMA). Mechanistically, it has been suggested that TRPC6 operates by modulating Ras/Raf/ERK1/2 signaling. Although the studies demonstrating the involvement of TRPC6 in gastric cancer were performed using pharmacological inhibitors (through the use of 2-APB and SKF), a method which lacks in specificity, these reports are still providing an important insight into the molecular mechanisms that mediate the progression of gastric cancer. In addition, these studies also suggest that drugs targeting the function of TRPC6 could be developed and used to improve gastric cancer treatment. For instance, pyrazolo [1,5-a]pyrimidine is a new potent TRPC6 antagonist that has been shown to be effective at inhibiting gastric cancer cell proliferation and migration *in vitro* as well as delaying tumor development in nude mice [123]. This prompts a very important question, could TRPC6 antagonists promote the efficacy of other chemotherapeutic agents such as paclitaxel, doxorubicin and cisplatin? Given that TRPC6 often exists as a hetero-tetramer with TRPCs 1, 3 and 7, and that all of these TRPCs are found in gastric cancer cells, it is likely that the answer to this question is not a straightforward one. To circumvent some of these difficulties, one possible approach could reside in the generation of individual knock out models of all TRPCs that form hetero-tetramers with TRPC6 and evaluate 1) their impact on the functional expression of TRPC6, 2) their role in gastric cancer survival, and 3) their response to TRPC6 antagonists, alone, and in combination other chemotherapeutics.

5.2. TRPM channels

Consisting of eight members (TRPM1 through TRPM8), the TRPM family share the common property of being activated in response to a variety of exogenous and endogenous ligands along with differential permeabilities and structural features. For instance, most TRPM channels are permeable to Ca^{2+} with the exception of TRPM4 and 5 that are Ca^{2+} impermeable. Additionally, three members within the family, TRPM2, 6 and 7 harbor enzymatic domains within their C-terminal regions and are commonly referred to as “chanzymes”. This diversity in structure and selectivity is in part responsible for the involvement of TRPM channels in many physiological processes as well as in various channelopathies [124,125]. The TRPM channels have also been implicated in numerous cancer types [126], but only TRPMs 2, 5 and 7

have been shown to contribute to the pathophysiology of gastric cancer.

TRPM2 was first demonstrated to play an active role in gastric cancer in 2018 by Almasi et al. The authors used two shRNA directed against TRPM2 to downregulate its expression and function in the AGS and MKN45 gastric cancer cell lines [127]. Results demonstrated that TRPM2 is functionally expressed as a plasma membrane ion channel permeable to Ca^{2+} in gastric cancer cells and that its inhibition reduced cell bioenergetics, suppressed cell invasion, and decreased cell survival. These results were further confirmed *in vivo* using a SCID mouse model where the loss of TRPM2 led to a decrease in tumor growth. The authors identified that TRPM2 operates via a JNK-dependent and mTOR-independent autophagy pathway to mediate gastric cancer survival [127,128]. Almasi et al. later confirmed that TRPM2 inhibition also promotes the effectiveness of chemotherapy drugs, specifically paclitaxel and doxorubicin, suggesting that TRPM2 inhibition in conjunction with known chemotherapeutics could represent an efficacious strategy for the treatment of gastric cancer [128]. Moreover, the authors reported that TRPM2 expression levels correlate with poor patient survival, in particular in those with late/advanced stages; thus, showcasing the possibility of a potential role for TRPM2 as a prognostic marker for late stage gastric cancer [128]. Similarly, other studies have found that high expression of TRPM5 was associated with shorter survival in gastric cancer patients [129]. However, further studies are needed to document the importance of TRPM2 and TRPM5 in gastric cancer patient survival and its clinical applicability.

TRPM7 is another member of the TRPM family that has captured the attention of certain research groups [130,131]. Results from Kim et al. showed that TRPM7 is upregulated in many gastric cancer cell lines including AGS, one of the most common human gastric adenocarcinoma cell lines [132]. In addition, using pharmacological tools and small interfering RNA as well as four gastric cancer cell lines (AGS, MKN45, SNU1 and SNU484), the authors demonstrated that TRPM7 inhibition caused a dramatic reduction in cell proliferation and a significant increase in apoptosis. These findings made TRPM7 a promising target for gastric cancer therapy. Accordingly, a variety of potent compounds against TRPM7 have been developed and their role in gastric cancer cell has been documented [133,134]. For instance, pharmacological studies showed that 5-lipoxygenase (LOX) inhibitors [135], nordihydroguaiaretic acid (NDGA), 3-[(p-chlorobenzyl)-5-(isopropyl)-3-tert-butylthioindol-2-yl]-2,2-dimethylpropanoic acid (MK886), 2,3,5-trimethyl-6-(12-hydroxy-5,10-dodecadiynyl)-1,4-benzoquinone (AA861) and waixenicin A [136] significantly decreased the TRPM7 current and caused gastric cancer cell death. Some fruit and vegetable extracts have also been proposed as promising TRPM7 antagonists (e.g. quercetin [137] and ginseng [131] have been shown to decrease TRPM7 activity and cause gastric cancer cell death). However, despite promising results following TRPM7 blockade, the development of anti-TRPM7 drugs faces its own challenges stemming from the ability of TRPM7 to exist as a hetero-tetramer with TRPM6. More studies are required to investigate whether the currently available inhibitors are specifically targeting TRPM7 or the TRPM7/6 hetero-tetramers. In addition, the function of TRPM7 in regard to some of the other major hallmarks of cancer is currently unknown (migration, invasion, and metastasis) and should be investigated in efforts to expand our knowledge of the underlying mechanisms of cancer cells. Although many questions remain open about the involvement of the TRPM family in gastric cancer, the quest for answers has already begun.

5.3. TRPV channels

Similar to the other TRP subgroups, the TRPV family consists of six members (TRPV1 through TRPV6) that function as Ca^{2+} entry mediators, either directly or indirectly following GPCR activation. Based on their function, two TRPV groups have been proposed: the non-selective group consisting of TRPVs 1–4, and the Ca^{2+} -selective group including TRPVs 5 and 6 [138]. The first group is characterised by its temperature

sensitivity and low Ca^{2+} conductance while the second group is not temperature sensitive and is highly selective for Ca^{2+} [139]. Many members of the TRPV family have been linked to various cancer types including breast cancer [140,141] and non-small cell lung carcinoma [142], but only TRPV4 and TRPV5 have been shown to play a prominent role in gastric cancer. Functional studies indicated that TRPV4 is upregulated in gastric cancer cells and its activation evoked a large outwardly rectifying current leading to a marked elevation in cytosolic Ca^{2+} [143]. Additional studies have also uncovered that the function of TRPV4 in gastric cancer cells is facilitated through the activation of GPCRs. For example, TRPV4 contributes significantly to the influx of Ca^{2+} through the calcium sensing receptor (CaSR)- and vasoactive intestinal polypeptide receptor 1 (VPAC1), both of which are GPCRs [143,144]. Furthermore, pharmacological inhibition and/or genetic depletion of TRPV4 expression markedly abolished CaSR and VPAC1-mediated gastric cancer cell proliferation and invasion, as well as tumor growth and metastasis. Interestingly, the mechanism behind these phenomena varies depending on the function of the GPCR receptor. CaSR/TRPV4 promotes gastric cancer survival through the Ca^{2+} /AKT/ β -catenin pathway while VPAC1/TRPV4 sustains survival via the Ca^{2+} /ERK1/2/JNK/ β -catenin pathway [143,144]. Although these studies have provided convincing evidence on the role of TRPV4 in gastric cancer, further research is needed to determine whether TRPV4 alone has a biological role in gastric cancer and if targeting its function, alone or in combination with GPCR inhibitors represents a viable option for the treatment of this disease.

The first indication of the potential involvement of TRPV6 in gastric cancer was provided by the Hediger group that showed an upregulation in TRPV6 expression in glandular gastric cancer cells [145,146]. Later, Chow et al. discovered the importance of TRPV6 when examining the role and selectivity of capsaicin in gastric cancer cell death [147]. Remarkably, capsaicin-mediated gastric cancer cell death was not mediated through TRPV1, a traditionally known capsaicin target [148], but rather through TRPV6. In the same study, the authors found that TRPV6 expression was upregulated in gastric cancer cells and its activation increased cytosolic Ca^{2+} levels. This discovery paved the way to uncovering that the mechanism behind Capsaicin/TRPV6-mediated gastric cancer cell death was mediated by the Ca^{2+} /p53/JNK pathway. While these findings indicate a therapeutic potential for TRPV6 in gastric cancer, they also raise several questions. Does capsaicin target TRPV6 directly to facilitate Ca^{2+} entry and gastric cancer cell death or does it activate TRPV6 indirectly through a complex signaling network? Is TRPV6 important in gastric cancer by itself or as a part of cascade signaling? Answering these questions will validate the clinical potential of TRPV6 and improve our understanding of the complex etiology of gastric cancer.

6. Conclusions and perspectives

Over the last two decades, a plethora of studies have demonstrated the involvement of TRP channels in various cancers, so much so that they are often regarded as *oncochannels* [104]. However, their contribution to the pathophysiology of cancer remains poorly understood and only a few TRP channels (TRPC6; TRPMs 2, 5 and 6; TRPVs 4 and 6) have been studied in the context of a gastric cancer model. It is of interest to examine the expression and function of TRP channels in correlation with oncogenes such as c-myc which is upregulated in more than 40% of gastric cancers [149–151] and it is often associated with *H. pylori* infections [152–156]. Some studies have demonstrated the involvement of TRPM2 [157] and TRPV4 [158] in *H. pylori* infections; therefore, it is important to establish the role of these channels in *H. pylori*-dependent tumours, especially when this pathogen remains the primary risk factor for gastric cancer.

Finally, despite marked progress in drug development, there are currently no pharmacological agents targeting any of the TRP channels that have successfully completed a clinical trial. Nonetheless, regardless

of previous shortcomings, SOR-C13 (NCT03784677), a small, synthetic TRPV6 antagonist is currently undergoing a phase I clinical trial to validate its safety/tolerance and patient response in several solid tumors [159,160]. Upon binding, SOR-C13 disrupts the function of TRPV6 and inhibits Ca^{2+} entry [160]. As shown previously, TRPV6 is upregulated in different cancers including breast, ovarian and gastric cancer cells where it facilitates the entry of Ca^{2+} . Manipulating TRPV6-mediated Ca^{2+} transport using SOR-C13 may enable the inhibition of Ca^{2+} -dependent functions in cancer cell survival and growth. Thus far, pre-clinical studies have demonstrated that SOR-C13 is effective at inhibiting the growth of breast and ovarian tumors in animal models and it is currently classified as an orphan drug by the FDA for the treatment of ovarian and pancreatic cancer [160–162]. Although this molecule is still undergoing testing and validation, it does provide a foundation for its potential use in gastric cancer patients. TRP channels represent a relatively new field of research with most studies still in their infancy, but these channels hold tremendous potential that has yet to be uncovered in the hopes of achieving major clinical breakthroughs in the treatment of gastric cancer.

Declaration of Competing Interest

No conflicts.

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