

Review Article

Strength and duration of GIPC-dependent signaling networks as determinants in cancer[☆]



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Abstract

GIPC is a PDZ-domain containing adaptor protein that regulates the cell surface expression and endocytic trafficking of numerous transmembrane receptors and signaling complexes. Interactions with over 50 proteins have been reported to date including VEGFR, insulin-like growth factor-1 receptor (IGF-1R), GPCRs, and APPL, many of which have essential roles in neuronal and cardiovascular development. In cancer, a major subset of GIPC-binding receptors and cytoplasmic effectors have been shown to promote tumorigenesis or metastatic progression, while other subsets have demonstrated strong tumor-suppressive effects. Given that these diverse pathways are widespread in normal tissues and human malignancies, precisely how these opposing signals are integrated and regulated within the same tumor setting likely depend on the strength and duration of their interactions with GIPC. This review highlights the major pathways and divergent mechanisms of GIPC signaling in various cancers and provide a rationale for emerging GIPC-targeted cancer therapies.

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Introduction

PDZ domains represent one of the most common protein-protein interaction domains present in all organisms from bacteria to human [1–5]. GAIP interacting protein C-terminus, GIPC/GIPC1, also known as synectin [6], is a key member of this family [1] that functions as an essential trafficking adaptor for membrane receptors, signaling effectors, and protein complexes—altogether comprising more than 50 protein-protein interactions to date [7]. As such, GIPC is considered a highly versatile molecule that controls vastly diverse cellular and pathophysiological processes.

In more recent years, GIPC has gained increasing attention for its roles in cancer. It not only serves as an antigen in breast and ovarian cancer [8–10] but has been strongly implicated in the progression of numerous malignancies including pancreatic [11,12], colon [13,14], skin [15], epidermal cancer stem (ECS) cell [17], glioma [18], lung [19], and gastric cancers [20] by potentiating tumor growth, invasion, metastasis, and cell survival. In contrast, GIPC is downregulated in cervical cancer [21], primary kidney tumor [20], primary colorectal tumor [20], and primary prostate cancer [20] and may promote tumor suppressive effects in these settings. These opposing effects likely arise from varying signaling contexts, the type of PDZ ligand, and the overall strength with which it binds in various cancer cells.

In this review, we discuss the major functions of GIPC in different cancers and their cellular mechanisms by focusing on its ability to differentially interact with membrane receptors, signaling complexes and oncoproteins. We also highlight the emerging roles of GIPC as a potential therapeutic target in cancer therapy.

Intracellular Signaling and Trafficking Functions of GIPC

GIPC was initially identified as an interacting partner of the GTPase-activating protein RGS-GAIP for G-protein coupled receptor subunit G α i [1]. Although the GIPC PDZ domain preferentially interacts with

Abbreviations: ECS, epidermal cancer stem; IGF-1R, insulin-like growth factor-1 receptor; ROS, reactive oxygen species; TGF- β , transforming growth factor- β .

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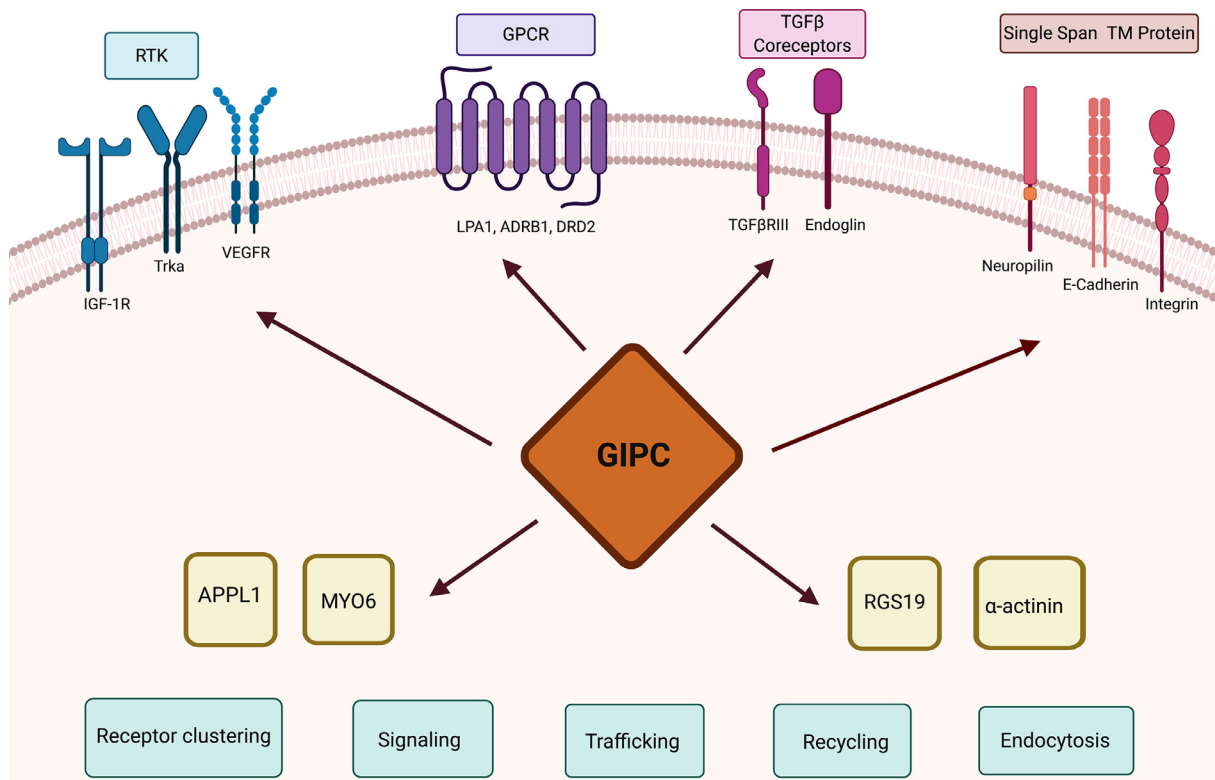


Figure 1. GIPC-dependent signaling complexes. GIPC interacts with various transmembrane receptors to promote their cell surface retention (e.g., T β RIII and endoglin) or internalization for enhanced endosomal signaling (e.g., IGF1R). GIPC also interacts with cytoplasmic signaling and trafficking proteins (e.g., Myo6).

the consensus class I PDZ binding motif, (S/T)X(A/V) at the extreme C-terminus of PDZ ligand [1,22,23], it also sometimes bind with class II and class III PDZ binding motifs, Ψ -X- Ψ -COOH and D/E-X- Ψ -COOH (where X is any amino acid and Ψ is hydrophobic), respectively [24,25]. The GH1 domain within the N-terminal region mediates its dimerization while the GH2 of the C-terminal region is capable of interacting with retrograde protein myosin VI (MYO6) [26,27]. As a result, GIPC functions as an adaptor for loading PDZ-binding proteins as cargo for the MYO6 motor protein to traffic various transmembrane proteins into endocytic vesicles [20]. Indeed, GIPC proves necessary for proper recycling, endosomal signaling and cell surface expression of transmembrane proteins such as RTKs [28,29], GPCR [30,31], TGF β RIII [32], endoglin [33], insulin-like growth factor-1 receptor (IGF-1R) [34], LHCGR [35], and VEGFR [36,37]. GIPC also affects actin bundle stabilization, cell adhesion, cytokinesis and cell migration through interaction with proteins like α -actinin-1 [24], E-cadherin [38] and integrins [5,27]. While the majority of PDZ ligands for GIPC are transmembrane proteins, some are distinctly intracellular proteins like APPL [28] and RGS 19 [1] while others are prominent viral oncoproteins (e.g., Hbc [39], E6 [21], and Tax [23]) (Figure 1). Accordingly, dysregulation of the GIPC activity, either through loss or overexpression, can disrupt crosstalk with multiple growth factor and cell survival signaling networks which ultimately help drive oncogenesis and tumor progression.

GIPC-mediated Regulation of Signaling Pathways in Cancer

Over the years there has been a steady rise in the number of GIPC targets in different cancer contexts (Figure 2 and Table 1). Here we discuss the 4

major pathways and their signaling components through which GIPC either promotes or suppresses tumorigenesis and disease progression.

GIPC-Neuropilins

Among the many binding targets of GIPC in cancer, arguably the most studied are the neuropilins (NRP1 and NRP2) [15,17,18,40,41], which are VEGF coreceptors [42–44] generally expressed in the neuronal and vascular systems but also found in a number of tumor cells [45,46]. Studies have demonstrated the GIPC-NRP association as a major driver of glioma [18], pancreatic cancer [40,41], skin cancer [15], and ECS cell (ECS) tumor [17].

While NRP1 has been previously shown to promote glioma progression through increased cellular proliferation, invasion and migration, more recent mechanistic evidence demonstrates that GIPC may positively regulate these effects through enhanced clathrin-vesicle trafficking and recycling of this receptor, which drives the oncogenic KRAS-ERK signaling pathway [18]. In ECS, GIPC binds to both NRP1/VEGFA and α 6/ β 4-integrin via a GIPC homodimer. By forming this complex, GIPC facilitates NRP1/VEGFA/ α 6/ β 4-integrin signaling that triggers downstream PI3K/PDK1, LATS1, YAP1, and Δ Np63 α -dependent ECS cell survival and tumor formation [17]. In pancreatic ductal adenocarcinoma, both GIPC and NRP1/NRP2 are frequently overexpressed and their knockdown has been shown to inhibit cell proliferation and migration while enhancing apoptosis to suppress overall tumor growth [40]. However, the underlying mechanism for the GIPC/NRP1-induced tumor promoting effects remain unresolved. Aside from these tumor-intrinsic properties, GIPC also supports the tumor microenvironment by influencing NRP1- α 5 β 1 integrin mediated

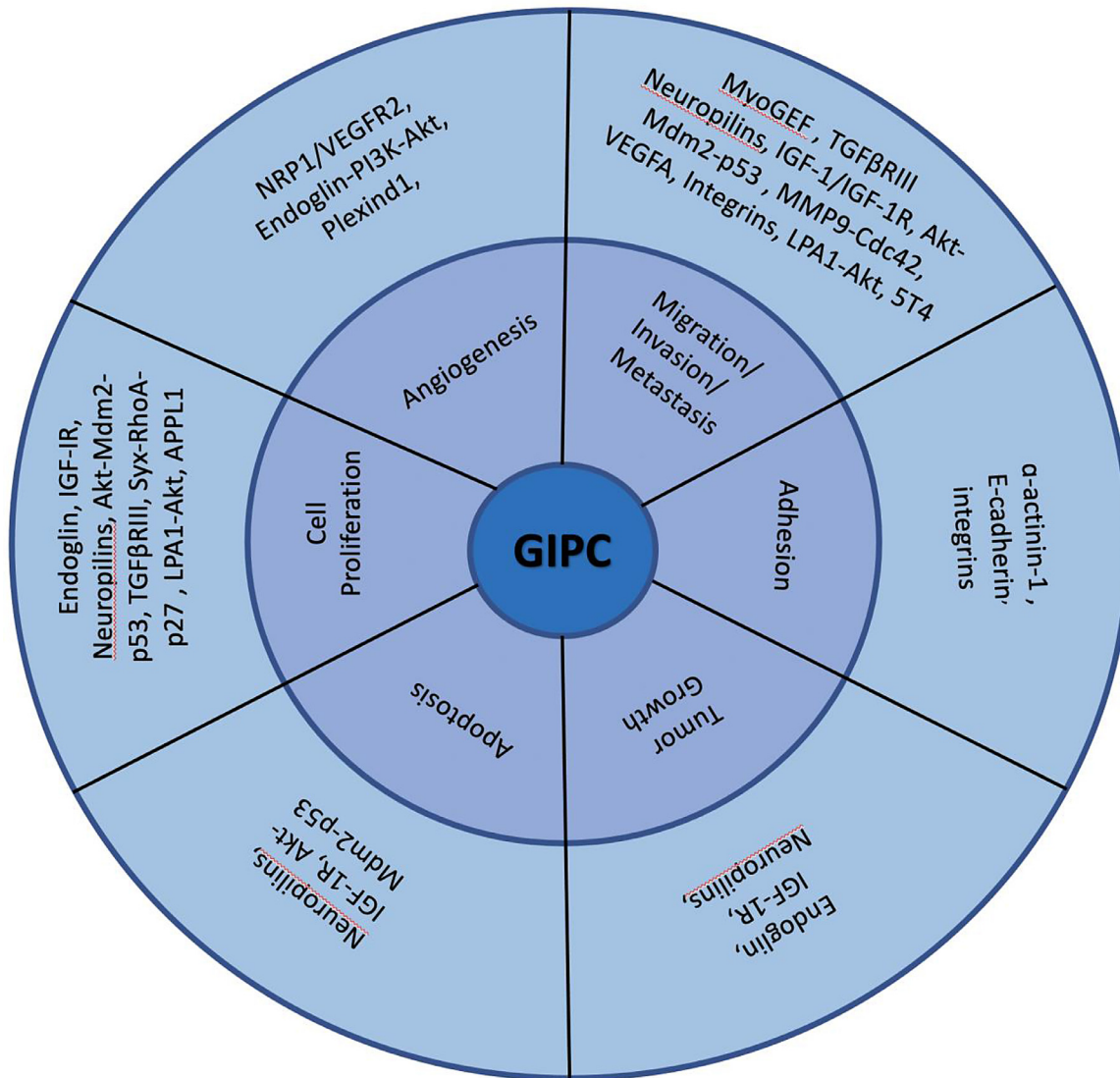


Figure 2. GIPC as a critical node for signaling networks in cancer. GIPC interacts with membrane receptors and intracellular signaling proteins to differentially regulate tumor cell biology and vascularization.

fibronectin fibril assembly in myofibroblasts. Here GIPC scaffolds NRP1 with ABL1, a tyrosine kinase that mediates NRP1 signaling for $\alpha5\beta1$ activation, leading to fibronectin fibril assembly, matrix stiffness, and tumor growth in rodent models [47].

Additionally, the GIPC-NRP interaction strongly supports the tumor microenvironment through neovascularization. Indeed, while both NRPs and GIPC are independently required to promote angiogenesis in developmental and various tumor settings [36,46,48,49], the GIPC/NRP1 interaction is also specifically necessary for efficient VEGFR2 trafficking and recycling to enhance the proangiogenic effects including endothelial migration during sprouting [37,50,51]. Indeed, this vascular function has now been implicated in the growth and metastasis of a vast majority of solid cancers [52–56]. Many preclinical models have shown that targeting NRP1 and/or VEGF exerts antitumor activity by reducing tumor angiogenesis, although the clinical benefits including overall survival in various cancers are still under evaluation [57–63]. Surprisingly, there have been no direct studies on the role of GIPC in tumor angiogenesis or its involvement in NRP1/VEGFR2 signaling during tumor vascularization.

GIPC-IGF-1R

GIPC interaction with IGF-1R is another notable association strongly implicated in the progression of prominent cancers including breast cancer [64], pancreatic cancer [12], and colorectal cancer [13]. IGF-1 signals through the IGF-1 receptor to promote growth and survival, and GIPC has been shown to positively regulate these cellular processes through IGF-1R protein stabilization by inhibiting its proteasomal degradation [11,12,34]. Studies have shown that lipopeptides (e.g., CR1023 and CR1166) which target the GIPC PDZ domain can significantly reduce tumor growth by blocking the GIPC-IGF-1R interaction and thus reducing IGF-1R expression in both breast cancer and pancreatic cancer mouse models [65]. A similar result was observed in *in vitro* studies showing decreased cell proliferation and enhanced apoptosis with reduced IGF-1R expression upon PDZ blocking peptide treatment [11].

GIPC mediates IGF/IGF-1R-induced tumor growth through reactive oxygen species (ROS) generation, which in turn, regulates phosphatases

Table 1

GIPC mediated key signaling pathways in different cancers.

Cancer Type	GIPC Mediated Signaling Pathways	Effects on Cancer Cell	References
Pancreatic Cancer	GIPC-endoglin	Increases cell proliferation and tumor growth	[82]
	GIPC-IGF-1R	Increases cell proliferation and tumor growth	[11]
	GIPC-NRP1/NRP2	Increases cell proliferation, cell survival, migration and tumor growth	[40]
Breast Cancer	GIPC-IGF-1/IGF-1R-ROS	Increases cell proliferation and migration	[64]
	GIPC/Akt-Mdm2-p53	Increases cell proliferation, cell survival and migration.	[91]
	GIPC-MMP-9-Cdc42	Increases invasion	[91]
Colorectal Cancer	GIPC- TGF β RIII	Inhibits cell migration and invasion	[90]
	GIPC-MyoGEF	Increases cell polarization and invasion	[93]
	GIPC-IGF-1-ROS-ZNF143	ROS generation and tumorigenesis	[13]
Skin Cancer	GIPC-Syx-RhoA-p27	Increases cell proliferation	[15]
ECS Tumor	GIPC-VEGF-A/NRP1/ α 6/ β 4-integrin; LATS1/YAP1/ Δ Np63 α	Increases cell invasion, migration, cell survival and tumor formation	[17]
Glioma Tumors	GIPC/NRP1-APPL1, p130Cas, KRAS-ERK.	Increases cell proliferation, invasion and cell survival	[18]
Cervical cancer (Associated with HPV-18 infection)	GIPC-E6 of HPV 18	GIPC degradation increases cell proliferation	[21]
Melanoma	GIPC-APPL-TRP1	Promotes melanogenesis in melanocytes	[16]

involved in various cancer cell proliferation [64]. In colon cancer, GIPC is believed to play a role in tumorigenesis and survival by inducing zinc finger protein, ZNF143 expression as part of the IGF-1/IGF-1R-ROS cascade [13]. Interestingly, in these cancer cell models, the role of GIPC in IGF/IGF-1R-induced cell proliferation proved to be independent of the canonical MAPK/ERK or PI3K/Akt pathways [13,64]. Therefore, the crucial downstream mechanisms by which GIPC drives ROS generation and tumor cell proliferation through the IGF/IGF-1R signaling system remain to be fully elucidated.

GIPC-LPA1/AKT Pathways

Aside from its above-mentioned roles as a tumor promoter, GIPC can also function as a powerful tumor suppressor often in the same cancer types depending on its ligands. Lysophosphatidic acid 1 (LPA1) is a receptor for LPA that has been shown to promote various carcinomas including breast, prostate, colon, and pancreatic cancer by regulating cell motility, chemotaxis, migration, and proliferation [66–69]. Here, GIPC can oppose these functions by directly binding to the PDZ binding motif of LPA1 to form an endocytic complex with APPL, which promotes LPA1 trafficking to early endosome EEA1 [70]. This in turn results in the downregulation of LPA1-induced Akt signaling from APPL endosome and thus inhibits the cell proliferation and cell migration [70]. Accordingly, GIPC depletion has been shown to delay LPA1 trafficking to EEA1 endosomes and thus maintains the LPA1/APPL association in signaling endosomes to promote Akt signaling towards higher cell proliferation and cell motility [70]. Consistent with the above findings upon GIPC depletion, recent evidence further suggests that disrupting this protein interaction with a point mutation in the LPA1 PDZ-binding motif induces oncogenic transformation by elevating Akt mediated cell proliferation [71].

Similar to LPA1, GIPC depletion has been reported to delay TrkA trafficking from APPL signaling endosome to early endosome [28], although their functional outcomes can differ significantly. The delay in TrkA trafficking has been shown to reduce Akt and Erk signaling [28], whereas the delay in LPA1 trafficking enhances the Akt signaling for cellular proliferation [70]. This indicates that the GIPC-APPL/Akt signaling pathway can yield highly divergent effects depending on its target proteins even within the

same cancer cell type, suggesting that the overall strength and duration of the interaction between GIPC and its ligands serve as fundamentally crucial yet elusive determinants of cancer progression. Further highlighting the complexity of these GIPC-dependent signaling networks, GIPC has also been reported to bind directly to APPL to form a GIPC-APPL/Akt complex [16]. Hence, precisely how GIPC preferentially interacts with these trafficking components to modulate the oncogenic properties of Akt signaling remains to be fully elucidated.

GIPC Regulation of the TGF-beta pathways

GIPC-Endoglin

Endoglin is a specialized TGF β co-receptor expressed in proliferating endothelial cells of normal and tumor vessels [72–74]. GIPC binds to this coreceptor through the C-terminal PDZ-binding motif SMA, which promotes endoglin cell surface retention and potentiates canonical Smad1/5/8 signaling downstream to promote angiogenesis [33]. While endoglin has received considerable attention as a vascular target in many solid cancers [73,75–77], more recent findings in cell lines and patient samples demonstrate that endoglin is also expressed in a subset of human breast [78], pancreatic [79], colorectal [80], and prostate cancer cells [81], which raises the question of the precise role of GIPC in regulating the tumor-intrinsic versus vascular endoglin in the surrounding tumor vessels. Notably, in pancreatic cancer, Pal et al., has shown that the GIPC-endoglin interaction is necessary for cell proliferation as blocking this interaction via a peptide-based inhibitor AP1032 inhibited *in vivo* tumor growth, and induced differentiation while also sensitizing pancreatic cancer cells to the frontline chemotherapeutic drug gemcitabine [82]. Furthermore, similar to the vascular system, inhibiting the GIPC-endoglin interaction also abrogated Smad 1/5/8 activation in pancreatic tumor cells [82].

Interestingly, the enhanced Smad 1/5/8 activation mediated by the GIPC-endoglin complex serves to inhibit endothelial cell migration [33], whereas GIPC stimulates migration when it interacts with NRP1 [51]. This dynamic again exemplifies the versatility of GIPC in differentially regulating cellular functions depending on the PDZ ligand, although precisely how GIPC favors one PDZ ligand over another, such as the case in endoglin versus NRP1,

remains to be characterized. In any event, it is also notable that the GIPC-endoglin interaction also regulates noncanonical signaling functions at least in the vascular system. In one study, GIPC has been shown to scaffold PI3K to endoglin at the plasma membrane to promote PI3K/Akt activation which acts to stabilize endothelial capillary sprouts during developmental angiogenesis [83]. While it is unclear whether GIPC functions in a similar manner towards endoglin signaling in pancreatic cancer cells, targeting the endoglin/GIPC complex may be an effective dual strategy to counter the tumorigenic and angiogenic properties of endoglin in certain tumor types.

GIPC-TGF β RIII

TGF β RIII is another coreceptor of the transforming growth factor- β (TGF- β) superfamily that shares structural homology with endoglin but is much more widely expressed [84]. Previous studies have established numerous roles for this receptor protein including tumor suppressive effects in certain cancers in part by inhibiting tumor motility and invasion *in vitro* [85–89] and angiogenesis, invasion and metastasis *in vivo* [86]. As was observed for the GIPC-endoglin interaction in endothelial cells, GIPC also directly interacts with TGF β RIII through the PDZ-binding motif to facilitate TGF β RIII cell surface stability, expression and TGF β responsiveness for Smad2/3 activation [32], which potentiates the suppressive effects of TGF β RIII on cell proliferation, migration and invasion in breast cancer [90].

Loss of GIPC expression, as evidenced in HPV-18-associated cervical cancer, appears to dampen the tumor suppressive effects of certain PDZ ligands including TGF β RIII. Studies have shown that GIPC interacts with E6 oncoprotein of HPV-18 which induces GIPC polyubiquitination for proteasomal degradation [21]. As GIPC enhances TGF β RIII protein stability and responsiveness to TGF β , E6-mediated GIPC degradation results in higher proliferation due to decreased antiproliferative effects of TGF β signaling [21]. This mechanism may apply similarly to many other cancers including primary kidney tumors, primary prostate cancer and primary colorectal tumor where GIPC gene expression is downregulated [20] and therefore GIPC may exhibit as a tumor suppressor role in such cancers.

Targeting GIPC in cancer

As GIPC is highly expressed in a number of human malignancies, inhibiting its activity can be an effective strategy for cancer therapy. Indeed, multiple studies have already demonstrated *in vivo* and *in vitro* antitumor activity by targeting GIPC, both through RNAi-mediated knockdown and by peptide-based competitive inhibition of GIPC-PDZ specific interactions [11,14,15,40,82,91,92]. GIPC depletion was shown to promote apoptosis [14,40], G2 cell-cycle arrest [14,40], and autophagy [92] while impairing cell proliferation [14,40,91], motility [14], invasion [40,91,93] and tumor growth [11,40,91], in breast cancer [14,91,93], colorectal cancer [14], and pancreatic cancer [11,40,92]. In some cases, GIPC silencing [92] and inhibition of the GIPC-endoglin interaction [82] also influences therapeutic efficacy as evidenced by the sensitization of pancreatic cancer cell lines to chemotherapeutic drug gemcitabine. In particular, data shows that GIPC depletion promotes this effect by causing exosome exocytosis of drug resistance gene ATP-binding cassette sub-family G member 2 (ABCG2), rendering it nonfunctional and thus sensitizing the cells to gemcitabine [92]. In another study, Patra et al., developed a series of cell-permeable lipopeptides that selectively block the PDZ domain of GIPC to elicit inhibitory effects against pancreatic and breast cancers in both cellular and animal models [65]. By mimicking the unique C-terminal PDZ-binding motif of GAIP, especially the prevalent SEA motif, these lipopeptides appear to competitively block the protein-protein interactions occurring between GIPC and PDZ ligand proteins [65]. Among the peptides, CR1166 was found most efficient in binding to the GIPC PDZ domain to disrupt the GIPC/IGF-1R association,

which consequently inhibits IGF-1R activity, cell proliferation and tumor growth [65]. Likewise, in skin cancer competitive inhibition of the GIPC/Syx interaction with a PDZ blocking peptide prevented RhoA activation which potently inhibited cell proliferation [15].

RNAi-mediated knockdown is another potential approach of targeting GIPC function. Although siRNA-mediated drugs have their own caveats like poor stability and tissue penetration [94], polymeric nanoparticle mediated siRNA delivery is considered a remedy to overcome the aforementioned obstacles [95–97]. As demonstrated by Borchardt et al., nanoparticle based GIPC-siRNA treatment can significantly reduce tumor volume in animal models with minimal adverse effects [40].

However, like the vast majority of targeted therapies in human clinical trials, both peptide-based and siRNA approaches must fully consider the unanticipated off-target effects of abrogating the essential homeostatic actions of GIPC. Conversely, because GIPC can act as a tumor suppressor and is downregulated in some cancers, alternative approaches should include identifying and targeting the negative regulators of GIPC. Taken together, these findings suggest that GIPC can serve as a promising therapeutic target in various human cancers but its expression pattern and dominant functional roles in each tumor type must be carefully considered.

Conclusion

GIPC is a versatile adaptor protein that regulates the functional trafficking of RTKs, GPCRs, TGF β receptors, and integrins. The dynamic actions of GIPC is highlighted by its vastly diverse signaling network of molecular partners, their interplay and biological consequences. Increasing evidence of its dichotomous roles in cancer clearly warrants further investigation into not only more selective GIPC-targeted therapies but also understanding the quantitative aspects of the strength and duration with which these ligands bind to GIPC. A clear grasp of these molecular mechanisms will be crucial to deciphering how these dynamic processes are altered in various cancers.

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