

## The GDNF Family: A Role in Cancer?



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### Abstract

The glial cell line–derived neurotrophic factor (GDNF) family of ligands (GFLs) comprising of GDNF, neurturin, artemin, and persephin plays an important role in the development and maintenance of the central and peripheral nervous system, renal morphogenesis, and spermatogenesis. Here we review our current understanding of GFL biology, and supported by recent progress in the area, we examine their emerging role in endocrine-related and other non–hormone-dependent solid neoplasms. The ability of GFLs to elicit actions that resemble those perturbed in an oncogenic phenotype, alongside mounting evidence of GFL involvement in tumor progression, presents novel opportunities for therapeutic intervention.

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### Introduction

The glial cell line–derived neurotrophic factor (GDNF) family of ligands (GFLs) is comprised of four structurally related factors: GDNF, neurturin (NRTN), artemin (ARTN), and persephin (PSPN) [1–3]. As their name suggests, these transforming growth factor beta–like growth factors have traditionally been implicated in the development and maintenance of central and peripheral neurons [2,4] and consequently have generated therapeutic interest for combating neurodegenerative diseases such as Parkinson's disease [5]. Outside the nervous system, these factors mediate many other processes such as renal morphogenesis in the kidney [6,7].

It is well known that tumors use growth factor pathways to progress an oncogenic phenotype and consequently escape the typical constraints of cellular growth. Growth factor pathways such as the epidermal growth factor (EGF) pathway [8,9] and the vascular endothelial growth factor (VEGF) pathway [10,11] have become the basis for development of anticancer therapeutics. Several lines of evidence suggests that the GFL/RET (rearranged during transfection) receptor tyrosine kinase signaling pathway may also present further potential targets for a multitude of cancers [12,13].

The purpose of this review is to provide a renewed perspective of the emerging role of neurotrophic factors, specifically the GDNF family, in neoplasia.

### GFL Signaling

Each member of the GDNF family is expressed as a pre-pro-precursor protein, which is proteolytically cleaved at a putative furin-like cleavage site (RAAR) by yet unidentified enzymes to generate an active form [14,15]. GFLs act as biologically active homodimers that signal canonically through the transmembrane receptor RET. This is facilitated by each GFL binding to a preferred glycosyl phosphatidylinositol (GPI)–linked GDNF family receptor  $\alpha$  (GFR $\alpha$ ) co-receptor, e.g., GDNF predominantly binds to GFR $\alpha$ 1, NRTN to GFR $\alpha$ 2, ARTN to GFR $\alpha$ 3, and PSPN to GFR $\alpha$ 4. However, this ligand-specific binding to GFR $\alpha$  co-receptors at times is promiscuous with a GFL ligand capable of interacting and functionally signaling

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with one of other nonpreferred GFR $\alpha$  proteins. GDNF can also bind to GFR $\alpha$ 2, NRTN to GFR $\alpha$ 1, and ARTN to GFR $\alpha$ 1 [16,17]. In neuronal populations, transforming growth factor beta is responsible for recruiting GFR $\alpha$ 1 to the plasma membrane to allow GDNF activation [18–20]. The stoichiometry of GFL:GFR $\alpha$ :RET binding interaction (as based on GDNF) is postulated to be one ligand homodimer to two GFR $\alpha$  molecules to two RET receptors, forming a heterohexameric complex [21]. RET homodimerization and subsequent autophosphorylation activate downstream signal transduction. Although not fully understood, all three components appear to be requisite for downstream signaling, as mice with homozygous deletions in either *Ret* or *Gfra1* exhibit a similar phenotype to mutant *Gdnf*<sup>-/-</sup> mice, which die shortly after birth due to kidney defects and a lack of enteric innervation [3]. However, GFR $\alpha$ -independent signaling has also been observed. For example, heparan sulfate proteoglycan syndecan-3 can serve as a co-receptor to transduce GFL signal to the RET with the involvement of Src kinase activation [22].

The RET receptor tyrosine kinase has three functional regions: an extracellular domain containing four cadherin-like domains (CLD1 to 4) followed by a cysteine-rich domain (CRD), a hydrophobic transmembrane region and an intracellular dual tyrosine kinase domain (TK1 and TK2) [23]. RET possesses several glycosylation sites, resulting in a mature protein molecular weight of 170 kDa. RET is present in at least three isoforms, RET51, RET43, and RET9, which contain 51, 43, or 9 amino acids in their unique C-terminal tails, respectively [23]. Isoforms RET9 and RET51 have distinctive signaling properties. The internalization process of RET isoforms was studied using total internal reflection fluorescence microscopy. The RET51 was robustly internalized from the cell surface into endosomal compartments upon the activation by GDNF, while RET9 was considerably slow [24].

GFLs activate several signal transduction pathways, including phosphatidylinositol-3-kinase (PI3K)/protein kinase B (AKT), RAS/mitogen activated protein kinase (MAPK), phospholipase C gamma (PLC $\gamma$ ), and c-Jun N-terminal kinase (JNK) pathways. These pathways propagate the cellular effects of RET activation such as cell survival, proliferation, differentiation, migration, branching morphogenesis, chemotaxis, and potentially oncogenesis (Figure 1).

Four key tyrosine residues, Tyr<sup>905</sup>, Tyr<sup>1015</sup>, Tyr<sup>1062</sup>, and Tyr<sup>1096</sup>, are responsible for initiating the downstream phosphorylation cascade following RET autophosphorylation. Only the long isoform (RET51) possesses Tyr<sup>1062</sup> and Tyr<sup>1096</sup> [25]. Tyr<sup>905</sup>, Tyr<sup>1015</sup>, and Tyr<sup>1096</sup> are binding sites for the adapter proteins GRB7/10, PLC $\gamma$ , and GRB2, respectively. Additionally, Tyr<sup>1062</sup> can bind to at least five additional families of docking proteins: Src homology 2 domain containing (SHC), fibroblast growth factor receptor substrate 2 (FRS2), insulin receptor substrate 1 or 2 (IRS1/2), the DOK (downstream of kinase) family of proteins (DOK1/4/5/6), and Enigma [3,26–28]. Activation of RET via GPI-linked GFR $\alpha$  takes place predominantly within lipid rafts, while signaling outside lipid rafts is mediated by soluble GFR $\alpha$  bound to extracellular GFL [29]. In addition to tyrosine phosphorylation, RET phosphorylation can also occur at Ser<sup>696</sup> in response to increased cAMP levels, resulting in protein kinase A (PKA) activation [30].

A recent development in GFL signaling has been the identification of an essential interaction of heparan-sulfate proteo-glycosaminoglycans (HSPGs), such as syndecans and glypicans, [31,32] with GFLs and their receptors. An interaction between GDNF and HSPGs had always been suspected since the discovery that *ex vivo* kidney development failed

under heparin-sulfate deprivation, in a similar fashion to GDNF or GFR $\alpha$ 1 deletion [33–36]. GDNF, NRTN, and ARTN, but not PSPN, have been shown to bind to syndecan-3 [22]. Likewise, ARTN is also known to bind heparin-sulfates [37]. GDNF promotes migration of cortical neurons via interaction with syndecan-3 [22]. It was demonstrated that the presence of heparin-sulfates is required for GDNF activation of RET tyrosine kinase activation in a MDCK cell line [31]. It was suggested that the low-affinity GDNF:GFR $\alpha$ 1 interaction, identified by [21] in RET-deficient cells, is likely to be due to heparan-sulfate binding. It has been found that HSPGs have a role in facilitating RET activation, acting to increase the local GDNF concentration through low-affinity binding in the vicinity of the GFR $\alpha$  receptors [2], and providing a linking mechanism to activate Src kinases and subsequently Met during RET-independent signaling [38].

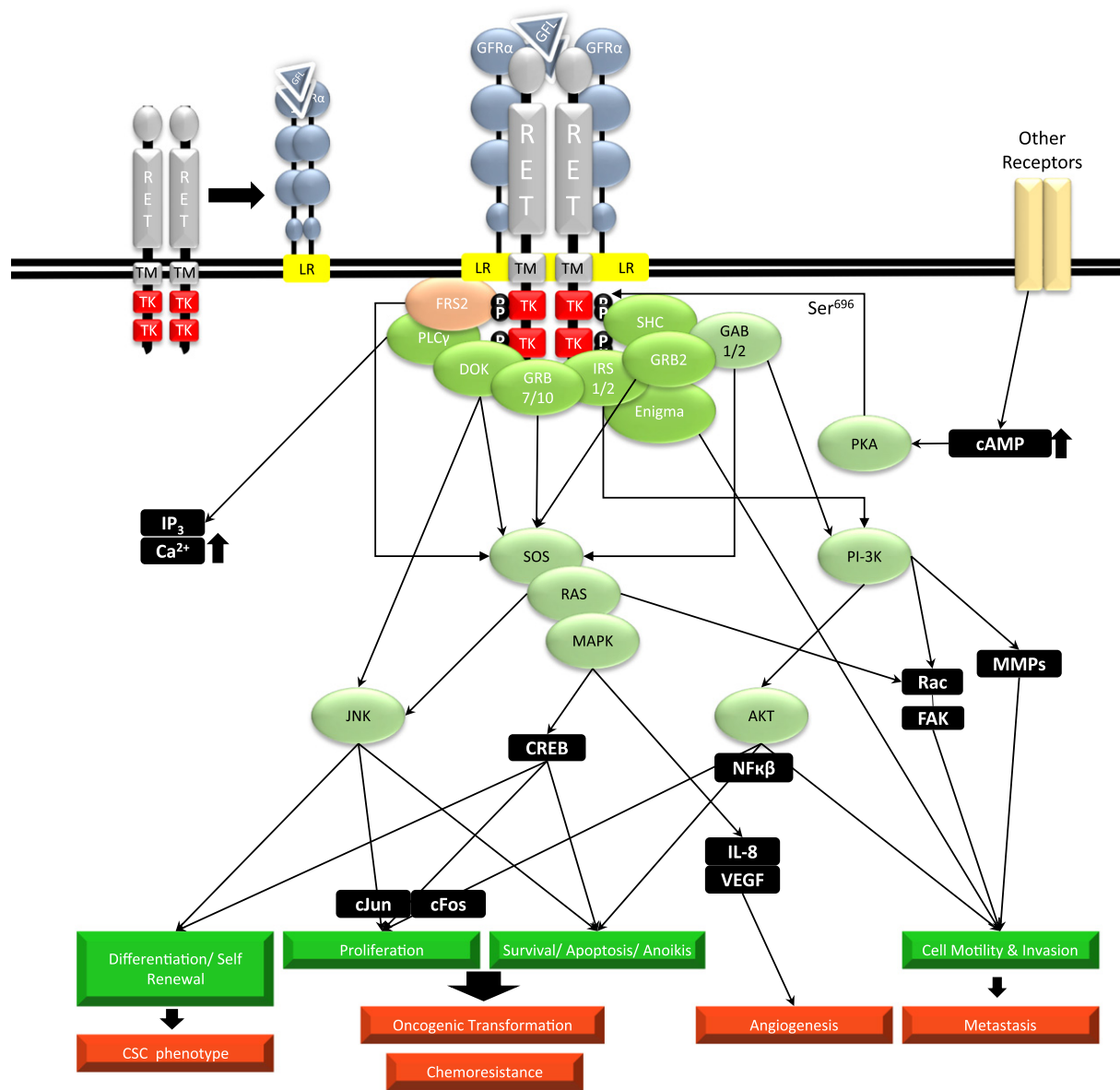
Unlike other receptor tyrosine kinases, binding of different GFLs to their cognate co-receptors GFR $\alpha$  does not appear to result in a differential activation of downstream signal transduction pathways. In fact, different GFLs actually induce coordinated phosphorylation of the same four key RET tyrosine residues (Tyr<sup>905</sup>, Tyr<sup>1015</sup>, Tyr<sup>1062</sup>, and Tyr<sup>1096</sup>) with similar kinetics and eliciting a similar signaling pathway profile [39]. The Tyr<sup>1062</sup> plays a key role in RET signaling during development as revealed by mice with a silencing mutation at Tyr<sup>1062</sup> displaying a similar phenotype as *Ret*<sup>-/-</sup> null mutants [40,41]. The RET Tyr<sup>1062</sup> phosphotyrosine serves as a docking site for multiple intracellular adaptor proteins, which are differentially used by different GFL-GFR $\alpha$  complexes to regulate alternative RET-stimulated cellular events [42]. Therefore, the binding promiscuity of Tyr<sup>1062</sup> may prove to be an avenue of further research in this regard, with yet unidentified adaptor proteins serving as distinct modifiers of biological activity. Furthermore, there is a mounting body of evidence for RET-independent signaling mechanisms.

### RET-Independent Signaling

Recent questions have been raised as to how GFL molecules elicit discrete functions. Adaptor proteins, GFL-GFR $\alpha$ -RET internalization rates, and RET stimulation in *trans* (via soluble GFR $\alpha$ ) can partially explain this; however, RET-independent signaling mechanisms hold more promise. It has been demonstrated that many tissues express GDNF and GFR $\alpha$ 1, but not RET, suggesting the presence of RET-independent pathways [43]. In addition to the aforementioned HSPG-dependent signaling, other mechanisms exist through which GFL members can interact with other growth factor receptors such as neural cell adhesion molecules (NCAMs) or integrins [44] (Figure 2).

In the absence of RET, GDNF is able to signal through lipid raft associated Src family kinases (SFKs). Despite the report of the co-immunoprecipitation of GFR $\alpha$ 1 and Src [38], a direct interaction is not plausible due to their positions on opposite sides of the lipid bilayer [43]. Hence, the existence of a new transmembrane receptor linking Src and GDNF:GFR $\alpha$ 1 has been postulated [2,38]. A potential candidate is the Met tyrosine kinase receptor, as Met phosphorylation in cultured *Ret*-deficient epithelial and neuronal cells is dependent on SFKs [45]. Additionally, there is an overlap between GDNF:Ret and hepatocyte growth factor (HGF):Met signaling pathways, resulting in similar cellular phenotypic consequences [45]. This suggests an indirect GFR $\alpha$ 1-Met association as Met does not immunoprecipitate with GFR $\alpha$ 1 [45].

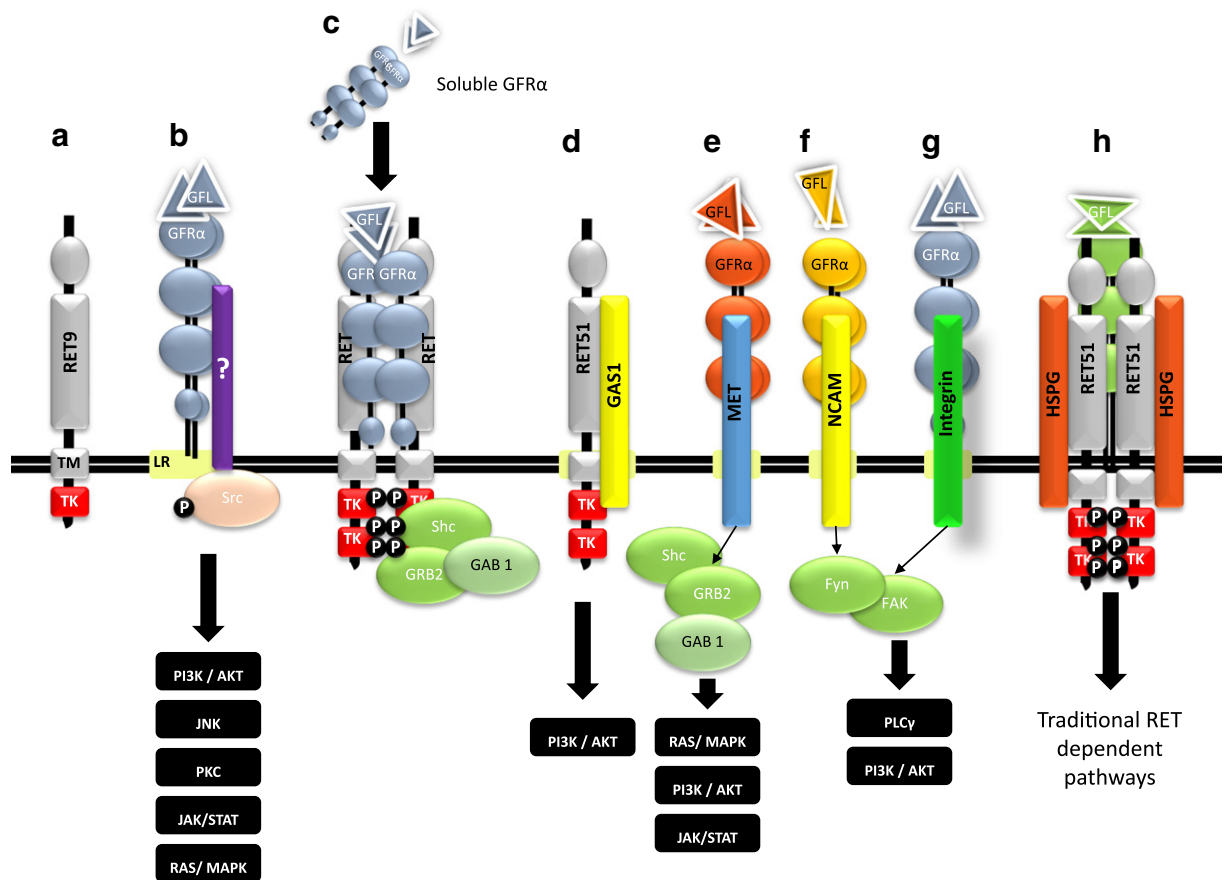
The GDNF:GFR $\alpha$ 1 complex is also able to bind to NCAM with high affinity to activate SFKs and FAK [44]. NCAM may also be



**Figure 1.** Traditional RET-dependent signaling pathways of GFLs with physiological and oncogenic consequences. Upon binding of the GFL dimer to its respective GFR $\alpha$  pair, the complex induces RET dimerization and consequently tyrosine kinase domain (TK) autophosphorylation. A series of SH2 (Src Homology 2) domain adapter proteins (green) including FRS2 (fibroblast growth factor receptor substrate 2), PLC $\gamma$  (phospholipase C gamma), DOK (downstream of kinase) 4/5, GRB (growth factor receptor-bound protein) 7/10, IRS 1/2 (insulin receptor substrate 1 or 2), GRB2 (growth receptor binding protein 2), SHC (Src homology 2 domain containing), and Enigma binds to their respective phosphorylated tyrosine residues, predominantly Tyr<sup>905</sup>, Tyr<sup>1015</sup>, Tyr<sup>1062</sup>, or Tyr<sup>1096</sup>. The signaling cascade activates downstream effector molecules (orange and black), resulting in functional responses such as differentiation, self-renewal, proliferation, survival, apoptosis, and cell motility. When perturbed, these processes become the driving facets of oncogenesis. Arrows are not restricted to only direct interactions. GFL, GDNF family ligand; TK, tyrosine kinase domain; TM, transmembrane domain; LR, lipid raft; P, phosphorylated tyrosine residue; GFR $\alpha$ , GDNF family receptor alpha; PI3K, phosphatidylinositol 3 kinase; AKT, protein kinase B; IP<sub>3</sub>, Inositol triphosphate; SOS, Son of Sevenless; MAPK, mitogen activate protein kinase; FAK, focal adhesion kinase; MMPs, matrix metalloproteinases; cAMP, cyclic adenosine mono phosphate; PKA, protein kinase A; JNK, c-Jun N-terminal Kinase; CREB, cAMP response element binding; NF $\kappa$ B, nuclear factor-kappa beta; IL-8, interleukin 8; GAB1, GRB2 associated binding protein 1; CSC, cancer stem cell.

responsible for facilitating the GFR $\alpha$ -Met interaction alluded to above. GDNF signaling through NCAM is known to occur independently of GFR $\alpha$ 1 through direct interaction with the third immunoglobulin domain of NCAM. However, the introduction of GFR $\alpha$ 1 results in a higher binding affinity to NCAM. In fact, GFR $\alpha$ 1 can also signal through NCAM in the absence of GDNF [44,46]. NCAM is expressed in many human cancers including

mammary and small cell lung carcinoma, and NCAM expression in biliary carcinoma correlates with perineural invasion [47–52]. In the physiological context, GDNF utilizes NCAM signaling pathways to promote axonal growth [44], and may play a role in regulating cell adhesion and synaptic plasticity in the CNS [53]. The effects of GDNF on survival and growth of midbrain dopaminergic neurons can be suppressed by inhibitory anti-NCAM antibodies [54]. GFR $\alpha$ 1



**Figure 2.** Alternative signaling pathways of GFLs. In addition to the traditional GFL:GFR $\alpha$ :RET51 pathway, several other novel pathways have been discovered to modulate GFL signaling [from left to right]. (a) Alternative RET isoforms, e.g., RET9 (ret proto-oncogene isoforms c); (b) activated Src (*v-src* sarcoma viral oncogene homolog) signaling associated with lipid rafts (*LR*) by means of yet an unknown transmembrane protein; (c) soluble GFR $\alpha$  responsible for RET activation outside lipid rafts; (d) GAS1 (growth arrest-specific 1), a recent GFR $\alpha$  alternative receptor; (e) MET (met proto-oncogene); (f) NCAM (neural cell adhesion molecule); (g) integrins; and (h) HSPGs (heparan sulfate proteoglycans) are essential for RET activity. TM, transmembrane domain; TK, tyrosine kinase domain; Fyn, p59fyn kinase; FAK, focal adhesion kinase; GAB1, GRB2 associated binding protein 1; GRB2, growth receptor binding protein 2; PI3K, phosphatidylinositol 3 kinase; AKT, protein kinase B; JAK, Janus kinase; STAT, signal transducer and activator of transcription; SHC, Src homology 2 domain containing.

has two alternative splicing isoforms, GFR $\alpha$ 1a and GFR $\alpha$ 1b, which differ by only five amino acids [55,56]. GFR $\alpha$ 1b, not GFR $\alpha$ 1a, mediates GDNF-stimulated cell migration in C6 glioma cells through GDNF-GFR $\alpha$ 1b-NCAM-RhoA signaling pathway, which is independent of RET [57].

The recently discovered growth arrest specific gene 1 (GAS1), a protein structurally similar to the GFR $\alpha$  family of receptors, has been shown to regulate RET signaling by binding and sequestering the receptor tyrosine kinase to lipid rafts in the plasma membrane in a similar fashion to the “classic” GFL receptors [58]. Like the GFR $\alpha$  molecules, GAS1 is a GPI-linked protein and constitutively localized in the lipid raft compartment of the plasma membrane. GAS1 is a protein with pleiotropic functions depending on its spatiotemporal expression in development. Such functions include growth inhibition, proliferation, and apoptosis [59]. GAS1 has also been shown to be co-expressed with RET in various tissues [60,61]. As a co-receptor, GAS1 modulates RET signaling in a ligand- and GFR $\alpha$ 1-independent fashion, specifically through the reduction of AKT phosphorylation, and without affecting MAPK activation [58].

CD133, also known as AC133 or prominin-1, is a transmembrane glycoprotein expressed on the surface of normal and cancer cells [62]. GDNF signaling can be regulated by CD133 in the absence of RET in neuroblastoma. It was shown that CD133 repressed neuroblastoma cell differentiation in part by downregulating RET transcription [63]. In CD133-expressing cells, GDNF-induced neurite outgrowth can be rescued by RET overexpression, whereas GDNF-induced expression of neuronal cell differentiation markers cannot be recovered by RET in CD133-expressing cells [63]. Therefore, GDNF can also function in a RET-independent manner in neuroblastoma cells.

## The Physiological Role of GFLs

### GDNF

GDNF was originally purified from a rat glial cell line (B49) [64] and is widely expressed throughout the rat and mouse central and peripheral nervous system, as well as the inner ear, corneal keratinocytes, olfactory epithelium, skin, submandibular gland,

bone, seminiferous tubules, cochlea, oocytes, kidney, teeth, gastrointestinal tract, and carotid body [65–67]. GDNF expression is detected in mouse embryonic development during early stages of neurogenesis between E7.5 and E10.5 [68]. During organogenesis, GDNF is found primarily in the mesenchyme and its derived tissues, specifically in tissues where epithelial-mesenchymal interaction occurs, such as in the previously mentioned kidney, submandibular gland, and tooth [66,67]. Similarly, in humans, GDNF is known to be expressed in the nervous system (central and peripheral), retina, kidney, lung [69], pituitary gland [70], skeletal muscle [71], testis, and mammary gland [72].

GDNF was first identified as a promoter of survival and morphological differentiation of rat embryonic midbrain dopaminergic neurons [14]. This observation had dramatic implications for the pursuit of a treatment for Parkinson's disease. But despite early success during human clinical trials [73,74], treatment of patients with GDNF by direct brain infusion failed to improve the symptoms [75].

Recently, it has been reported that GDNF's protection of dopamine neurodegeneration is due to an inhibition of caspase-3 activation and suppression of endoplasmic reticulum stress-related genes [76]. As a survival factor for motor neurons, GDNF is significantly more potent (75-fold) than the neurotrophins brain-derived neurotrophic factor (BDNF), ciliary neurotrophic factor (CNTF), and cholinergic differentiation factor–leukemia inhibitor factor (CDF-LIF); GDNF successfully rescued facial motor neurons from atrophy after deprivation of other survival factors following lesions [77]. Further evidence is provided from studies on *Gdnf* and *Gfra1* knockout mice, which experience a substantial loss of motor neurons due to increased cell death [33,35,78,79]. In these mice, enteric neurons, where GDNF stimulates migration, proliferation, differentiation, and survival of multipotent enteric precursor cells, fail to develop, and the mice die soon after birth [33,34,80,81]. Interestingly, mice lacking other GFLs or co-receptors are viable and fertile [2]. GDNF was shown to induce regeneration of spinal motor neurons after injury [82]. It was observed that GDNF is expressed by skeletal muscle and is localized at the plasma membrane, predominantly near neuromuscular junctions, providing a source of GDNF for nearby motor neurons [83].

Reports of a functional role for GDNF outside the nervous system are accumulating. Most notable is a pivotal role that GDNF plays in renal morphogenesis [7]. GDNF signaling from mesenchymal cells around the ureteric bud tips is required for them to branch out. This happens through the RET receptor kinase and GFR $\alpha$ 1, which then form the renal collecting duct system. It was conferred that the RET/Etv4 signaling promotes directed cell movements in the ureteric bud tips [84]. GDNF also plays a role in spermatogenesis [85,86]. Firstly, GDNF is expressed by the nephrogenic mesenchyme in the developing kidney, located in immediate proximity to the RET-expressing tips of the ureteric bud. GFR $\alpha$ 1 is localized to both the mesenchyme and ureteric bud. This mesenchymal-epithelial interaction induces the initial branching from the Wolffian duct, continuing into morphogenesis of ureteric bud [87]. While mesenchyme-derived GDNF expression is initially quite sparse across the Wolffian duct, upon branching, RET and GDNF become more concentrated around distal tips of the ureteric bud and periphery, respectively. GDNF is essential for kidney development as *Gdnf* knockout mice show defects in this respect and die soon after birth [34]. However, while GDNF is essential, it cannot promote development on its own [88]. It does appear that RET-independent pathways are

utilized as RET-deficient mice exhibit metanephric development to some degree [80]. Moreover, the PI3K/PTEN axis was identified as being critical for chemotaxis and branching morphogenesis [89].

The enteric nervous system (ENS) develops following the migration of neural crest cells (NCC). GDNF is an essential trophic factor for the developing ENS, as knockout mice with deletions in *Gdnf* or *Gfra1* lack enteric neurons [35]. *Ret* knockout mice lack enteric neurons in the small and large intestines but do develop them in the esophagus [90], although the density of neurons in this region is significantly reduced [91]. Nedd4-related E3 ubiquitin ligase-2 (NEDL2) is pivotal in regulating ENS development and GDNF/Ret signaling. The NEDL2-deficient mice die within 2 weeks after birth and have a low body weight. These mice showed a progressive bowel motility defect [92]. GDNF is chemoattractive to NCC, thereby promoting their migration into the gastrointestinal tract [91,93]. Additionally, GDNF supports the survival, proliferation, and differentiation of NCC [94–96].

GDNF is secreted by Sertoli cells, somatic cells of the seminiferous tubule, and ensures the self-renewal of spermatogonial stem cells (SSCs) and prevents terminal differentiation [85,86,97], predominantly acting in the perinatal period [98]. GDNF achieves this through the inhibition of Notch signaling, halting differentiation, and promoting the expansion of the stem cell pool [99], or via the Src family kinase/PI3K/Akt pathway that leads to the upregulation of both c- and N-Myc [100–104]. In addition to this crucial role in the self-renewal and maintenance of SSCs, GDNF may also play a role in the proliferation and differentiation of these cells into  $A_{\text{paired}}$  and  $A_{\text{aligned}}$  spermatogonia via the GDNF-ERK1/2-FOS pathway [105]. Evidence of a role for GDNF in SSC cell fate has been established in mouse *Gdnf* knockout models, which results in the depletion of the SSC population [106]. Conversely, overexpression of GDNF results in a larger, undifferentiated spermatogonia population, leading to the shutdown of spermatogenesis and generation of a nonmetastatic, seminoma-like tumor [106]. To complete this autocrine interaction, both RET and GFR $\alpha$ 1 are located on undifferentiated spermatogonia and are routinely used as markers for spermatogonia isolation [86]. ARTN and NRTN also exhibit the capacity to regulate spermatogonial stem cell proliferation [107]. A study with double-mutant mice model has shown that the NOTCH signaling is activated in the Sertoli cells *in vivo*, and *in vitro*, the NOTCH ligand works through a ligand called JAG1 [108]. However, a negative feedback regulation was indicated in the testicular stem or progenitor cells. The activation of NOTCH signaling in Sertoli cells upregulates the transcriptional repressors HES1 and HEYL, which directly downregulate GDNF expression by binding to the *Gdnf* promoter [108,109].

GDNF is also known to functionally interact with BMP4, a key inhibitory protein of embryonic stem cell differentiation, to enhance neuronal development during kidney development [110,111]. Further evidence of a role for GDNF:RET signaling in stem cell differentiation and biology includes an involvement in the regulation of hematopoietic cell differentiation [112] and the co-localization of GDNF and GFR $\alpha$ 1 with ABCG2 and p63, two known stem cell markers within the human corneal epithelium [113]. It was suggested that GDNF and GFR $\alpha$ 1 may represent a phenotypic property that identifies a population of stem-like precursor cells [113]. GDNF also interacts with integrin  $\beta$ <sub>1</sub> and was observed to be upregulated in mammospheres, neurospheres, and hematopoietic and embryonic stem cells [114]. GDNF is also capable of dimerizing with  $\alpha$ <sub>6</sub> integrin, an adhesion molecule required for cancer stem cell

tumorigenicity [115]. Moreover, integrin  $\beta_1$  could form a complex with GFR $\alpha$ , which is enhanced by GDNF [116]. Furthermore, cancer stem-like cells derived from medullary thyroid carcinoma (MTC) on the basis of CD133 positivity have demonstrated a dependence on RET for their self-renewal and differentiation [117]. Lastly, significant cross talk exists between the Notch and RET:GFR $\alpha$ 1 signaling pathways during nephrogenesis [118].

GDNF also functions in oocyte development [119,120], follicular proliferation in ovaries [121], regulation of hair growth [122], tooth innervation [123], corneal regeneration and wound healing [113,124,125], immune homeostasis and response, and psychoregulation including drug abuse [126]. A recent study observed reduced levels of *GDNF* and *ARTN* mRNA in the peripheral blood cells of patients with major depressive disorders compared to those same individuals in a remissive state [127]. Furthermore, GDNF may promote angiogenesis through increasing production of IL-8, a potent angiogenic factor, in SK-N-MC human primitive neuroectodermal tumor cells [128]. Functionally, GDNF is known to exhibit cross talk with the VEGF:VEGFR and NGF:TrkA pathways [129,130]. GDNF is a key regulator of the endothelial cell network formation. GDNF, by itself and in the presence of adipose-derived stem cells, was associated with enhanced capillary network formation [131].

GDNF expression and activity are regulated by a complex network of factors and mechanisms ranging from epigenetic regulation in peripheral blood cells [132] to regulation by a host of small molecule drugs such as antidepressants [133–136], glucocorticoids [137,138], and follicle-stimulating hormone [139]. This extends to the neurotransmitters dopamine, serotonin, glutamate, and adenosine [140,141]. Additionally, lipopolysaccharides and inflammatory cytokines, such as IL-1 $\beta$ , IL-6, TNF- $\alpha$ , and TNF- $\beta$ , regulate GDNF in C6 glioma cells [142–145]. The regulation of GDNF expression in the nervous system has been extensively reviewed [146].

While the neuroprotective effects of estrogens may be mediated by neurotrophic factors [147], it appears that GDNF expression is not estrogen regulated in certain tissues. Instead, GDNF signaling can be significantly modulated by, or can participate in, cross talk with estrogen. The estrogen receptor (ER) modulator tamoxifen had only a limited effect on GDNF levels in C6 glioma cells [148], and *Esr1* knockout mice demonstrated no alteration of GDNF expression within the murine midbrain [149]. Paradoxically, 17 $\beta$ -estradiol stimulates GDNF expression in developing mice hypothalamic neurons [150]. This effect, however, appears to manifest itself through nonclassical estrogen action since treatment with ER antagonist ICI<sub>182,780</sub> (fulvestrant) does not alter GDNF expression. This nonclassical action is mediated by Ca<sup>2+</sup> and cAMP/PKA signaling [151]. Conversely, in mammary carcinoma, RET and GFR $\alpha$ 1 are highly regulated by estrogen, and significant cross talk exists between the pathways (see below) [152–154]. Genomic profiling of invasive melanoma found that GDNF was one of the genes that were highly amplified in invasive cell lines as compared to noninvasive ones, suggesting an important function in the melanoma metastasis [155].

Signaling pathway, genome-wide chromatin binding, and transcriptome analyses have demonstrated that SPIN1 has a direct impact on the expression of GDNF, which activates RET pathway [156]. It was shown that knockdown of SPIN1 in liposarcoma cells reduced GDNF expression, which was activated in human liposarcoma tissues as compared to normal tissues [156].

## ARTN

ARTN, the most recently discovered member of the GFLs, promotes the survival of peripheral ganglia and dopaminergic neurons *in vitro* [157,158]. Through alternative splicing, human *ARTN* mRNA possesses at least five functional transcript variants, which encode three pre-pro-ARTN isoforms forming identical mature proteins [15]. These splice variants appear to be differentially expressed in different human tissues, with some tissues only expressing nonfunctional variants, with the highest expression levels of total variants observed in nonneural tissues [15]. According to the National Center for Biotechnology Information UniGene database and as reported by our laboratory, human *ARTN* is expressed in a number of tissues, including esophagus, intestine, kidney, larynx, lung, pancreas, parathyroid, placenta, prostate, uterus, colon, trachea, cerebellum, adipose, cartilage, and stomach [159].

*Artn* and *Gfra3* knockout mice are viable and fertile but exhibit ptosis (a characteristic drooping of the eyelids) due to the lack of sympathetic innervation to the superior tarsus muscle [160,161]. In wild-type mice, GFR $\alpha$ 3 is expressed in the ganglia of the peripheral nervous system at a higher level than GFR $\alpha$ 1 or GFR $\alpha$ 2. In particular, GFR $\alpha$ 3-mediated signaling is required for the survival and migration of superior cervical ganglion neurons (SCG), as *Gfra3*<sup>-/-</sup> mice show gross postnatal SCG cell death and impaired rostral migration of SCG neurons to target organs [160,162]. This impaired migration most likely contributes to the death of neurons due to a lack of target-derived growth factors such as nerve growth factor (NGF). Despite expression in other sympathetic ganglia such as the dorsal root ganglion, these ganglia appear to be normal in *Gfra3*<sup>-/-</sup> mice. Specifically, a study using low-density dissociated cultures demonstrated a role for ARTN in the embryonic generation and in the survival and growth of sympathetic neurons, in fact promoting survival to the same degree as NGF [160,162]. ARTN also appears to have a transient effect on the survival of mature SCG neurons as they lose their survival dependence on NGF with age, and concomitant treatment of ARTN and NGF increases neurite growth to a greater extent than NGF alone [162]. The source of this endogenous ARTN, however, is unclear as an inhibitory anti-GFR $\alpha$ 3 antibody did not have any effect on mature sympathetic neurons cultured in the absence of growth factors, indicating the lack of an autocrine loop [160,162]. The parasympathetic ganglia normally innervate some of the same targets as sympathetic ganglia. Mice do not appear to be affected by *Gfra3* deletion. *Gdnf*<sup>-/-</sup> mice have also been shown to have subtle defects in the SCG, suggesting some degree of cross talk between GDNF and GFR $\alpha$ 3 [35]. Unlike *Gdnf* knockout mice, *Artn/Gfra3* knockout mice have normal ENS development [160,161], and ARTN does not induce the migration or neurite outgrowth of NCC [91]. ARTN complexed with GFR $\alpha$ 3 is believed to have a normalizing effect on the pathophysiology of mechanisms of neuropathic pain [163]. In addition to GDNF and NRTN, ARTN was also studied for its effects in the gastrointestinal tract. While GDNF and NRTN promoted the migration of neurite outgrowth in the esophagus, ARTN had no effect [91,164].

Immunohistochemistry studies in the normal human brainstem and hippocampus have revealed a distinct temporospatial expression pattern for each of the GFLs and GFR $\alpha$  subtypes at prenatal, perinatal, and adult ages [165–169]. ARTN and its cognate receptor GFR $\alpha$ 3 consistently show a more restricted distribution compared to the other GFLs and associated receptor subtypes [167,168]. In particular, ARTN/GFR $\alpha$ 3 localize to the caudal spinal trigeminal

nucleus, which corroborates with the findings of other studies that suggest a role for ARTN in nociception, touch, and thermal sensory perception [170–172].

Expression of ARTN, GFR $\alpha$ 3, and RET was detected in early murine embryo cultures [173]. They established that endogenous ARTN promoted embryonic development by increasing trophoblast cells, decreasing blastocyst cell apoptosis, and ultimately increasing the percentage of early embryos or zygotes progressing into mature blastocysts [173]. Conversely, depletion of ARTN suppressed embryo development. Similarly ARTN and RET, but not GFR $\alpha$ 3, were detected in early pregnancy oviducts, suggesting that paracrine ARTN from the oviduct ampulla epithelium also facilitates preimplanted embryo development. GDNF has also been reported to exhibit a similar function [174].

ARTN also facilitates the development and maintenance of vascular system innervation. The vascular and nervous systems have a very close relationship, such that vascular-derived neurotrophic factors have long been postulated to regulate vascular innervations [175]. The vascular system largely exerts its effects on the neuronal system by signaling to receptors located on postganglionic sympathetic neurons to direct the innervation of the vasculature [175]. In mice and rats, ARTN is secreted by arterial smooth muscle cells and acts as a chemoattractant to developing axonal processes from SCG at the final stages of innervation [161,176]. Likewise, ARTN expression is present within smooth muscle cells of blood vessels of normal human breast tissue and mammary carcinoma [159]. Vascular-derived ARTN is also a determinant of neurite outgrowth in adult rats [176]. It is alone in this regard, as GDNF- and NRTN-deficient mice do not exhibit abnormal sympathetic nervous systems, and PSPN does not act on the peripheral nervous system [35,177,178]. ARTN-, RET-, and GFR $\alpha$ 3-deficient mice have grossly abnormal development of the sympathetic nervous system [161,179].

### *PSPN and NRTN*

PSPN has several unspliced (nonfunctional) and spliced (functional) transcripts. The unspliced forms are detected in most human tissues, while the spliced transcripts are only detectable at very low levels in the human adrenal gland, cerebellum, spinal cord, and testis [180]. PSPN specifically binds GFR $\alpha$ 4, of which there are several splice variants, one of which is a non-GPI-linked soluble form [180,181]. In newborn and adult mice, the expression pattern of these splice variants is restricted to the thyroid, adrenal medulla, and the pituitary intermediate lobe, with RET co-expression occurring only in the thyroid C-cells and adrenal chromaffin cells [182,183]. In the adult human, expression of GFR $\alpha$ 4 is restricted to the thyroid gland, while RET was found to be widely expressed [180]. PSPN knockdown in oral squamous cell carcinoma cells significantly reduced cell proliferation, and overexpression of PSPN was closely related to tumoral size [184].

NRTN is particularly important in the development, maintenance, and function of the parasympathetic system, both centrally [168,185] and peripherally [186–192]. Mice homozygous for NRTN or GFR $\alpha$ 2 (the co-receptor for NRTN) deletion show defects in parasympathetic ganglia [177,178]. Additionally, a stem/progenitor cell niche was identified within the pituitary (predominantly localized at the marginal zone between the intermediate lobe and adenopituitary) that is characterized by the expression of GFR $\alpha$ 2, Prop-1 (Prophet of Pit-1), and several stem/progenitor cell markers. It has

been suggested that these nonendocrine cells are responsible for maintaining postnatal pituitary homeostasis and expansion as they exhibit the ability to differentiate into hormone-producing cells or neuron-like cells [193].

### **GFLs in Cancers**

As shown in Table 1, GFLs have been implicated in a variety of cancers. In this section, the roles of individual GFL signaling in those neoplasms will be discussed.

#### *RET Germline Mutations and Neuroendocrine Tumors*

Germline activating mutations of the RET proto-oncogene, which is highly expressed by several cell lines of the neural crest lineage [194], cause several forms of neuroendocrine cancer including multiple endocrine neoplasia type 2A and 2B, familial medullary thyroid carcinoma, papillary thyroid carcinoma, pheochromocytomas, and neuroblastomas [195–197]. However, not all cases of MEN2 have mutated RET. In addition, GFR $\alpha$ 4 has also been proposed as a modifying factor for this set of diseases, as it is only detected in MTC, and no other thyroid tumors such as follicular thyroid adenomas, follicular thyroid carcinomas, or papillary thyroid carcinomas [180]. Whether PSPN functions as the GFR $\alpha$ 4/RET ligand in MTC is unclear as its expression pattern is limited. MTC cells secrete calcitonin, which is used as a marker of tumor burden. Exogenously applied PSPN appears to regulate the expression of calcitonin in MTC, a process mediated by the oncogenic forms of RET [198]. Alternatively, one particular mutation in GFR $\alpha$ 4 results in the formation of a truncated, soluble form of the receptor, capable of activating RET independent of its ligand PSPN [199]. However, use of calcitonin as a biomarker in MTC may need to be reassessed in light of the evidence that treatment with a RET inhibitor results in a dose-dependent reduction of calcitonin levels, while tumor volumes remain the same [198].

In normal anterior and neoplastic pituitaries, GDNF and RET are predominantly expressed in almost all somatotrophs but absent in most of the other cell types (corticotrophs and gonadotrophs) in normal pituitaries. Furthermore, GDNF was expressed in all growth hormone-secreting pituitary adenomas and absent in almost all other types of pituitary tumors [70]. These findings suggest a link between the GDNF and growth hormone signaling pathways, although the physiological significance has not yet been ascertained [70]. One role may be the regulation of cell populations because RET was shown to control the somatotroph population numbers through the p53 apoptotic pathway [200]. This is mediated by RET activating Pit-1, a potent transcription factor that also binds and activates the human growth hormone promoter [200,201]. However, upon the application of GDNF, Pit-1 expression and apoptosis are suppressed [200].

In contrast, Hirschsprung disease is a congenital malformation associated with aganglionosis of the gastrointestinal tract resulting from inactivating mutations in RET [25]. These individuals tend to lack certain parts of the enteric nervous system [197]. Polymorphisms in the GFL genes do not appear to be a modifying factor in Hirschsprung disease phenotypes [202]. Beyond these neuroendocrine disorders, GFLs have also been implicated in a series of other cancers of epithelial origin particularly pancreatic, testicular, bile duct, colon, glioma, mammary, endometrium, ovarian, and lung.

#### *Pancreatic Cancer*

GDNF is upregulated following acute induction of pancreatitis in mice [203]. Pancreatic cancer tissues and pancreatic cancer cell lines

**Table 1.** GFL Signaling Involved in Cancers

Cancer Types	GFL Signaling Involved	Other Signaling Pathways Implicated
Neuroendocrine tumors	RET mutations [195–197]; RET [200,201]; GDNF [70]; RET/GFR $\alpha$ 4 [180]; PSPN (?) [198]	p53/Pit-1 [200,201]
Pancreatic cancer	GDNF and RET/GFR $\alpha$ 1 [203–207] ARTN and RET/GFR $\alpha$ 3 [208–211,216,223] NRTN and GFR $\alpha$ 2 [225]	PI3K/AKT and NF $\kappa$ B [213,214]; integrin $\beta$ <sub>1</sub> and MMP-9 [218,219,221,222] MMP-2 and E-cadherin [211]
Glioma	GDNF and RET [227,228]; GFR $\alpha$ 1 [227,229] NRTN and GFR $\alpha$ 1/2 [231,234]	MAPK and JNK [227,228]; GAS1 [232,233]; PCDNA and Ki-67 [235]
Colorectal cancer	GDNF [237]; RET/GFR $\alpha$ 1 [236] NRTN [238]	integrin $\beta$ <sub>1</sub> [236]; VEGF-VEGFR1, p38, PI3K/AKT, and HIF1 $\alpha$ [237]
Breast cancer	RET [255]; GDNF [249,250]; RET/GFR $\alpha$ 1 [72]	TNF $\alpha$ and IL-1 $\beta$ [72]; ER and PR [241]; HER2 [250,252]; p44/42 (ERK1/2)/mTOR and JNK [152,154]; SRC [262]; PAX2 [263,264]. ER [153]; HER2 [159]; TWIST1 [256] [257]; BCL-2 [258]; VEGF-A [259] AKT and CD24 [267]
Endometrial cancer	ARTN [153]; GFR $\alpha$ 1 and GFR $\alpha$ 3 [265] ARTN and RET/GFR $\alpha$ 3 [266]	BCL-2 [268]
Lung cancer	ARTN and RET/GFR $\alpha$ 3 [268]	NCAM/FYN [273]; ERK1/2 and AKT [275]
Ovarian and testicular cancers	GDNF/GFR $\alpha$ 1 [120,273–276]	ERK1/2, c-JUN, MMP-9, c-Kit and p38 [278–280]
Melanoma	GDNF and RET/GFR $\alpha$ 1 [278–282]	NF- $\kappa$ B, MMP-9 and integrin $\beta$ 1 [283]
Oral cancer and salivary adenoid cystic carcinoma	GDNF [283] PSPN and RET [184]	ERK and CDKs [184]
Prostate cancer	GDNF, RET/GFR $\alpha$ 1 [284,285]	ERK and AKT [284]
Liver cancer	ARTN [286] GDNF [131]	AKT/HIF1 $\alpha$ [286]
Bone cancer	GDNF [287,288]	ERK [287,288]
Gastric cancer	GFR $\alpha$ 1 methylation [289]; GFR $\alpha$ 3 methylation [290]	

express GDNF and the RET and GFR $\alpha$ 1 receptors [204]. GDNF promotes pancreatic cancer cell proliferation and invasion through the GFR $\alpha$ -1/RET receptor complex in an autocrine/paracrine manner [205,206]. Endoneurial macrophages are shown to secrete high levels of GDNF, which activates RET and promotes perineural invasion of pancreatic cancer [207]. ARTN may be involved in chronic pancreatitis, a precursor disease to pancreatic cancer characterized by progressive and irreparable damage to the organ, severe fibrosis, and intense abdominal pain leading to functional insufficiency [208,209]. ARTN is closely related to a normalizing effect of neural pain. In chronic pancreatitis where there is increased pain, ARTN and its co-receptor GFR $\alpha$ 3 are upregulated [210]. ARTN, through GFR $\alpha$ 3, was found to promote pancreatic cancer cell motility and invasiveness in MIA PaCa-2 cell lines [211].

Both GDNF and ARTN are strongly expressed in the endocrine cells and intrapancreatic nerves of normal pancreatic tissue, as well as in pancreatic cancer tissue using immunohistochemical methods [212]. Intrapancreatic neural invasion is correlated to GDNF expression and is facilitated through the PI3K/AKT and nuclear factor kappa B (NF $\kappa$ B) pathways [213,214]. GDNF and RET both correlate with survival rate and clinical parameters in pancreatic cancer [215–217]. GDNF enhances the expression of integrins in the pancreas, specifically the  $\beta$ <sub>1</sub> subunit [218,219], which combined with GDNF-induced upregulation of matrix metalloproteinase-9 (MMP-9) production and activation facilitates metastasis [212,218–222]. Another group found that both ARTN and its receptor complex GFR $\alpha$ 3/RET were both increased in expression in cases of pancreatic cancer, with strong expression in both primary cancer cells, in liver metastases, and in surrounding tissues (the strongest of which was in hypertrophic nerves and arterial walls) [216,223]. Most notably, ARTN appeared to increase cell migration and invasion in pancreatic cancer cell lines in a similar manner to GDNF but did not affect proliferation [223]. While ARTN did not affect the expression either of MMP-2 or of MMP-9 mRNA in their study [223], another study showed that ARTN treatment resulted in an increase of MMP-2 and a decrease of E-cadherin expression [211].

Moreover, MMP-9 was found to mediate GDNF-stimulated invasion of pancreatic cancer cells [221,222]. Therefore, the signal transduction pathways involved in ARTN-mediated migration/invasion need to be investigated further. It is possible that the invasion of pancreatic cancer cells into nerves results in injury and inflammation, which in previous research have been shown to result in the upregulation of GFLs in Schwann cells, in an effort to regenerate nerves; GFLs then act in a paracrine/autocrine fashion to perpetuate further intravasation by pancreatic cancer cells [210]. Additionally, certain RET polymorphisms can increase the effect of GDNF-induced pancreatic cell invasion [224].

Pancreatic cancer tissue and pancreatic cancer cells expressed increased amounts of NRTN. NRTN promoted invasiveness and silencing of NRTN reduced proliferation and invasion of pancreatic cancer cells [225]. RET was also upregulated in pancreatic adenocarcinoma, and GDNF depletion in perineurial macrophages, or inhibition of RET with shRNA or a small-molecule inhibitor, reduced perineurial invasion in a mouse model [226].

### Glioma

GDNF expression is abundant in the highly invasive C6 rat glioma cell line but is significantly lower in the noninvasive Hs683 human glioma cell line, suggesting that an autocrine/paracrine mechanism stimulates glioma cell migration [227]. GDNF induces glioma cell migration via MAPK and JNK pathways [227,228]. RET and GFR $\alpha$ 1 are also reported to be highly expressed in glioma cells [227,229]. GDNF appears to confer chemoresistance while promoting mitogenesis in glioblastoma cell lines [230]. It was reported that GDNF additionally permitted neuroblastoma cells to proliferate in the presence of a range of cytotoxic chemotherapeutic agents [231]. GAS1 was shown to inhibit the growth of gliomas by blocking the GDNF-RET signaling pathway [232,233].

There is a contentious role for NRTN and GDNF in neuroblastomas. It was found that NRTN and GDNF potentiated retinoic acid-induced differentiation in neuroblastoma [234]. This effect was much more pronounced in nonaggressive tumors compared



to aggressive ones. Conversely, GDNF and NRTN were found to induce faster growth of neuroblastoma cell lines, conferring protection against chemotherapeutic drug treatment [231]. GDNF has also been shown to stimulate proliferation of glioma cells by upregulating expression of cyclins PCDNA and Ki-67 [235].

### Colorectal Cancer

Colorectal cancer cells express the RET/GFR $\alpha$ 1 receptor complex for GDNF, and their  $\beta$ <sub>1</sub> integrin expression is also significantly enhanced by GDNF, and consequently, the enhancement and associated increase in adhesion and invasive abilities in response to GDNF were inhibited by blocking the GDNF receptor or the integrin  $\beta$ <sub>1</sub> subunit [236]. GDNF is also shown to enhance the migration of colon cancer cells by increasing VEGF-VEGFR interaction, which is mainly regulated by the p38, PI3K/Akt, and HIF1 $\alpha$  signaling pathways [237]. In a case of intestinal ganglion neuromatosis associated with colon adenocarcinoma, GDNF and NRTN were found to be highly expressed in the adenocarcinoma cells, while co-receptors GFR $\alpha$ 1 and RET were expressed in surrounding ganglion and glial cells [238].

Methylation may affect GDNF signaling in colorectal cancer. *GDNF* locus in colonic mucosa of ulcerative colitis patients was found highly methylated and the level of methylation was significantly higher in active inflamed mucosa than in quiescent mucosa [207]. *GDNF* gene is among the 15 genes that have been differentially methylated in colorectal cancer in comparison to adjacent normal mucosae [239].

RET has been shown to be an oncogene in many cancers, but a recent study has shown that RET is a potential tumor suppressor gene in colorectal cancer [240]. *RET* locus was methylated in 27% of colon adenomas and in 63% of colorectal cancers, resulting in a decrease in RET expression, whereas the restoration of RET expression in colorectal cancer cell lines caused apoptosis [240].

### Breast Cancer

Increased expression of GFR $\alpha$ 1 and RET transcripts is observed in mammary carcinoma compared to normal breast tissue [72]. *GFR $\alpha$ 1* mRNA expression was detected in 59.4% of the tumor samples, and was associated with ER receptor expression and lymphovascular invasion/lymph node metastasis at diagnosis. *GFR $\alpha$ 1* mRNA expression was inversely correlated with *p53*, *EGFR*, basal markers, and basal-like tumors but positively correlated with luminal-type tumors [72]. There are many other studies in the cancer microarray database Oncomine that have identified a significant correlation of *GFR $\alpha$ 1* and *RET* receptors with both estrogen and progesterone receptor (PR) positive status [241]. Elevated *GFR $\alpha$ 1* levels in mammary carcinoma have been reported in several studies [242–244]. Of the 212 tumor samples surveyed, *RET* mRNA expression was expressed in 29.7% of the tumors [245]. Of those, only 18.1% co-expressed *RET* and *GFR $\alpha$ 1* compared with 41.9% of the tumors being *GFR $\alpha$ 1* positive/*RET* negative. This suggests that, in mammary carcinoma, GDNF (or other GFL acting through GFR $\alpha$ 1) predominantly acts via a RET-independent signaling mechanism.

We have observed similar correlations for *GDNF* in human normal mammary tissue and mammary carcinoma in an analysis of data extracted from the Oncomine database [159]. However, we did not observe a repeatable correlation between *GDNF* expression and grade, which had been previously demonstrated by other groups

[246–248]. Although additional studies have detected GDNF upregulation in mammary carcinoma [249,250] and GDNF is significantly linked to luminal/apocrine but not basal tumors [251], there is still a lack of evidence that GDNF expression is significantly correlated with disease and outcome. However, it is clear that GDNF correlates consistently with HER2 positive status [250,252] but not ER status [159,250,253].

It has been demonstrated that in MCF-7 cells (RET<sup>+</sup>/GFR $\alpha$ 1<sup>+</sup>), treatment with exogenous GDNF resulted in increased S-phase entry, enhanced cell survival as measured by cell attachment after serum starvation, and increased cell scattering [72]. GDNF treatment also resulted in the loss of cortical actin organization and formation of actin stress fibers [72]. In a wild-type MCF-7 xenograft mouse model, the authors also detected strong GDNF-expressing, infiltrating fibroblasts around the tumor and a low level of GDNF expression by the tumor itself, with localization of GDNF on the invasive margin of the tumor [72]. Untreated MCF-7 cells showed very little endogenous GDNF mRNA, but treatment with inflammatory cytokines TNF $\alpha$  and IL-1  $\beta$  resulted in a substantial increase [72]. Together, these results suggest that mammary carcinoma cells not only secrete GDNF in response to inducing cytokines, but also respond to paracrine- and autocrine-derived GDNF sources. Additionally, MCF-7 cells have been found to have robust expression of both GFR $\alpha$ 1 and RET proteins [72,159,254].

*RET* gene has previously been reported to contain an estrogen response element located in its promoter region [255]. Both *RET* and *GFR $\alpha$ 1* mRNA expressions were strongly increased following estradiol treatment with similar kinetics to that of early response genes such as the trefoil family factor TFF1 and c-Myc. It appears that, rather than altering GDNF levels, estrogens contribute to an enhanced level of GDNF signaling, albeit in a cell line-dependent manner [152]. In fact, GDNF stimulates anchorage-independent proliferation through RET, an effect mediated by p44/42 (ERK1/2) MAPK and JNK [152].

ARTN expression is regulated by estrogen. Overexpression of ARTN is associated with resistance to antiestrogen drugs like tamoxifen in patients with ER-positive mammary carcinoma, and inhibition of ARTN restores the tamoxifen sensitivity [153]. ARTN is also expressed in ER-negative breast cancers. ARTN synergizes with TWIST1 to promote metastasis and poor survival outcome in patients with ER-negative tumors [256]. TWIST1 is a transcription factor. High TWIST1 expression is associated with breast cancer invasion and metastasis [257]. ARTN has also been reported to stimulate radio- and chemoresistance by promoting TWIST1-BCL-2-dependent cancer stem cell-like behavior in mammary carcinoma cells [258] and to promote *de novo* angiogenesis in ER-negative mammary carcinoma through activation of TWIST1-VEGF-A signaling [259]. Enhanced GDNF/RET signaling in ER positive breast cancers promotes resistance to aromatase inhibitors in postmenopausal patients with ER breast cancers [260]. The RET kinase inhibitor NVP-AST487, superior to the aromatase inhibitor letrozole, inhibited the GDNF-induced motility and tumor spheroid growth in ER-positive breast cancer cells *in vitro* and demonstrated similar efficacy in impairing tumor growth *in vivo* [261].

GDNF was able to neutralize trastuzumab-induced apoptosis in HER2+ breast cancer cells *in vitro* and induce *in vivo* growth in xenograft tumors [262]. Interestingly, the SRC kinase inhibitor saracatinib effectively blocked GDNF-stimulated growth of

trastuzumab-sensitive cells but did not inhibit GDNF-promoted growth of trastuzumab-resistant cells, indicating that SRC mediates GDNF pro-survival functions by bridging RET-HER2 cross talk in trastuzumab-responsive breast cancer tumors, whereas GDNF is also linked to trastuzumab resistance by acting independently from SRC in trastuzumab-resistant tumors [262].

Therefore, targeting the GDNF- and ARTN-induced RET signaling may be an effective way to overcome endocrine resistance in ER-positive breast cancers [153,154,261] as well as trastuzumab resistance in HER2-positive breast cancers [262].

A previous study demonstrated that PAX2 (paired box 2 gene), a transcriptional activator, coordinated GDNF expression by binding to the 5'UTR of exon 1 within the gene, driving kidney development [263]. This is interesting not only because GDNF may be under similar transcriptional control within the breast but also because PAX2 has been implicated in mediation of tamoxifen resistance through an interaction with ER and the *HER2* promoter [264].

Unlike GDNF, ARTN appears to play more of an autocrine role within mammary carcinomas. ARTN protein expression was detected in 65% of human mammary carcinoma samples and correlated to HER2 positivity and higher tumor stage, and increased ARTN expression is linked to decreased overall survival of stage III and HER2-negative patients [159].

Forced expression of ARTN in MCF-7 mammary carcinoma cells increased cell survival, promoted anchorage-independent growth and enhanced cell migration and invasion, and promoted a more aggressive cellular morphology with formation of proliferative and disorganized colonies in a three-dimensional basement membrane culture model [159]. Like GDNF [152], ARTN had little effect on anchorage-dependent proliferation under serum-replete conditions but displayed a stimulatory effect under serum-depleted conditions [159].

ARTN significantly increased the expression of the antiapoptotic BCL-2 protein and several genes involved in invasion and metastasis, such as SERPINE1, MMP1, and PLAU [159]. Forced expression of ARTN significantly promoted tumor formation and tumor progression *in vivo*, with a greater than two-fold increase in size after 6 weeks compared to control tumors [159]. Additionally, ARTN is correlated with ER-positive status in mammary carcinoma, and its expression is correlated with poorer distant metastasis-free survival [153]. ARTN both increases ER transcriptional activity and function, as well as mediates resistance to antiestrogen therapies. Resistance appears to be driven by ARTN-induced BCL-2 expression [153]. Concordantly, targeting of the RET receptor sensitizes mammary carcinoma cells to antiestrogen therapies, and recurrent invasive tumors are twice as likely to exhibit RET positivity following adjuvant tamoxifen treatment [154]. GDNF stimulation of RET was also shown to increase ER $\alpha$  phosphorylation and estrogen-independent activation of ER $\alpha$  transcriptional activity by a mammalian target of rapamycin (mTOR)-dependent mechanism [154]. We have demonstrated that ARTN co-receptors GFR $\alpha$ 1 and GFR $\alpha$ 3, but not Syndecan-3, are significantly upregulated in breast cancer and that their expression is significantly associated with survival outcome of breast cancer patients, especially in ER-negative or HER2-negative mammary carcinoma [265].

It has been demonstrated that ARTN can induce epithelial to epithelial-to-mesenchymal transition (EMT) in ER-negative mammary carcinoma cells through the upregulation of TWIST1 [256]. *In vivo*, this resulted in enhanced local invasion and distant metastasis.

Interestingly, low expression of both ARTN and TWIST1 was sufficient to predict 100% relapse-free and overall survival in patients with ER-negative breast cancer [256]. On the other hand, high expression of both ARTN and TWIST1 was correlated with a poor survival.

### Endometrial Cancer

ARTN exhibits similar oncogenic effects in endometrial carcinoma to mammary carcinoma [266]. Expression of ARTN was significantly associated with higher-grade tumors and myometrial invasion. ARTN stimulated the *in vitro* proliferation and cell survival of RL95-2 and AN3 cells, regulating such key genes as CDC25A, CDK2, Bcl2, p53, Bad, Bax, and TERT. Furthermore, anchorage-independent growth was enhanced, while ARTN overexpression also induced a mesenchymal phenotype, which was correlated to deregulation of vimentin, Met, MMP1, MMP9, PLAUR, SERPINE1, and SERPINE5. Mechanistically, AKT phosphorylation was indicated as pivotal to the survival and invasive response demonstrated by ARTN. *In vivo*, xenografts yielded almost two-fold larger tumors that were poorly differentiated and significantly more invasive, infiltrating surrounding supporting tissues. Enhanced proliferation and survival were also evident *in vivo*. We have also demonstrated that ARTN confers endometrial carcinoma chemoresistance through transcriptional activation of CD24 [267]. Specifically, ARTN alleviates the G2/M arrest initiated by doxorubicin and the apoptosis induced via microtubule disruption by paclitaxel treatment. Additionally, ARTN inhibition by functional antibody or small-interference RNA (siRNA) delivery enhanced the efficacy of the respective chemotherapeutic drug treatments [267]. Forced expression of ARTN in endometrial carcinoma cells was shown to decrease sensitivity to chemodrugs doxorubicin and paclitaxel, and ARTN-stimulated resistance can be abrogated through inhibition of CD24 expression [266].

### Lung Cancer

ARTN has also been implicated in non-small cell lung carcinoma (NSCLC) [268]. *ARTN*, *RET*, and *GFR $\alpha$ 3* have been demonstrated to be unregulated in primary neoplasms relative to their normal counterparts, while high *ARTN* expression also correlated with lymph node metastasis and increasing grade of NSCLC, according to an examination of the OncoPrint database [268]. Transcriptional activation of BCL-2 by ARTN enhances *in vitro* survival in monolayer and anchorage-independent conditions. Furthermore, ARTN significantly enhanced invasive capability of H1299 and H1975 NSCLC cell lines. The former, when injected into immunodeficient mice, yielded 63% larger tumors at day 30 compared to control. Tumors exhibited increased S-phase entry and reduced apoptotic levels. Conversely, these oncogenic effects abated following siRNA or functional antibody inhibition. GDNF and GFR $\alpha$ 1 on the other hand lacked any significant correlation between normal and cancerous lung tissue [268]. GFR $\alpha$ 2 in the majority of cases exhibited a downregulation in the neoplastic setting (squamous cell carcinoma and adenocarcinoma) [268]. In terms of RET mutations causing SCLC, several novel somatic RET mutations were reported in squamous small cell lung carcinoma cell lines and tumor samples [269]. However, another study was unable to detect the same RET (or GDNF) mutations in 54 SCLC cell lines [270]. Moreover, there was no consistent pattern for RET, GDNF, or GFR $\alpha$ 1 expression in 21 SCLC cell lines assayed [270]. Therefore, RET mutations may not be an important step in the tumorigenesis of

SCLC. Nevertheless, RET fusion genes occur at a rate of approximately 1% (84/6899) in NSCLC patients, and female patients (or those less than 60 years old of age) usually have higher frequencies than male patients (or those aged 60 years and older), particularly in patients from Asian [271]. For example, RET fusion genes occur at 1.9% (3 of 156) of NSCLC patients in Koreans [272].

### Ovarian and Testicular Cancers

GDNF and GFR $\alpha$ 1 are localized to follicles at various stages of development, being actively secreted by oocytes and acting in an autocrine/paracrine fashion [120]. GDNF has also been implicated in ovarian tumorigenesis, being influenced by androgens [273,274]. Interestingly, the expression of GDNF in the ovary is distinct to that in the testes, where it is the somatic Sertoli cells which express GDNF. Testicular tumors develop regularly in older GDNF-overexpressing mice [275], and GDNF promotes invasive behavior in testicular seminoma cells [276]. Because spermatocytic seminoma cells share many phenotypic markers with SSCs, whose self-renewal is triggered by GDNF, it is possible that GDNF contributes to spermatocytic seminoma by promoting the formation of SSCs [277].

### Melanoma

Activated RET signaling is found to be correlated with the development of malignant melanoma in a mouse model [278–280]. Several human melanoma cell lines express RET and GFR $\alpha$ 1, and GDNF stimulation significantly enhances the proliferation of human melanoma cells [281,282]. The expression level of intrinsic *Ret*, *Gdnf*, and *Gfra1* transcripts in malignant melanomas from RET-transgenic mice was significantly upregulated compared with those in benign melanocytic tumors, and GDNF treatment activated RET via phosphorylated Tyr<sup>905</sup> in human malignant melanoma cells [282]. Therefore, GDNF-mediated RET kinase activation is associated with the pathogenesis of malignant melanoma.

### Oral cancer and salivary adenoid cystic carcinoma

*PSPN* mRNA and protein were significantly upregulated in oral squamous cell carcinoma (OSCC)–derived cells compared with human normal oral keratinocytes. *Pspn* knockdown significantly decreased cell proliferation and reduced receptor tyrosine kinase signaling and cell cycle arrest at the G1 phase. Primary OSCCs have significantly higher PSPN protein expression than normal counterparts, and overexpression of PSPN is closely related to tumoral size, suggesting that PSPN is a regulator of OSCC progression and may be a diagnostic marker for OSCC [184].

GDNF protein was strongly expressed in salivary adenoid cystic carcinoma and adjacent nerve fibers and positively correlated to the expression of NF- $\kappa$ B, MMP-9, and integrin  $\beta$ 1 [283]. This observation suggests that GDNF may increase the matrix degrading and cell adhesion in the process of perineural invasion of salivary adenoid cystic carcinoma.

### Prostate Cancer

A recent study found that RET was expressed in all prostate cancer cell lines tested, but GFR $\alpha$ 1 was only expressed in 22Rv1 cells; as a result, 22Rv1 cells responded to exogenous GDNF, while all cell lines responded to combined GDNF and GFR $\alpha$ 1 treatment [284]. RET knockdown inhibits tumor growth *in vivo*, and mechanically, RET activates ERK or AKT signaling to promote transformation-associated

phenotypes, including perineural invasion via activation of p70S6 kinase [284]. Immunohistochemical analysis of tumor tissues revealed that GDNF expression was significantly stronger in higher-stage prostate tumors [285].

Our group has unpublished evidence examining the role of ARTN in prostate carcinoma. Immunohistochemistry indicated a significant increase in the expression of ARTN in 50 cases of human prostate carcinoma in comparison to 23 cases of human benign prostate hyperplasia. ARTN does not appear to regulate cell proliferation or survival, as ARTN overexpression, knockdown by siRNA, and inhibitory antibody treatment had negligible effects (unpublished observations). This was also the case for chemotherapeutic resistance to doxorubicin and paclitaxel. However, ARTN overexpression increased the metastatic potential of cells, registering significant changes in monolayer and 3D Matrigel culture cell morphology, lamellipodia formation, cell migration and invasion, and characteristic deregulation of mRNA species involved in EMT (unpublished observations). ARTN also promoted anchorage-independent growth, an effect that is inexplicably not abrogated by ARTN inhibitory antibody or chemodrugs.

### Liver Cancer

ARTN has been shown to be a hypoxia-responsive factor essential for hypoxia-induced expansion of cancer stem cells in hepatocellular carcinoma (HCC). In addition, increased ARTN expression was associated with larger tumor size and worse clinical outcome of HCC patients. Similar to the role of ARTN in breast cancer, forced ARTN expression reduced apoptosis; increased proliferation; and enhanced EMT and motility, tumorsphere formation, and the tumor-initiating capacity of HCC cells. ARTN was also shown to dramatically increase xenograft tumor size and metastasis *in vivo* [286]. GDNF is secreted by adipose-derived stem cells and HCC and contributes to pathological neovascularization [131]. Therefore, targeting ARTN and GDNF signaling may be an effective approach to treating HCC.

### Bone Cancer

Metastasis of any cancers to the bone causes the bone cancer pain in cancer patients. GDNF was shown to be involved in bone cancer pain in an animal model. Lentivirus-mediated GDNF RNAi significantly attenuated mechanical and thermal hyperalgesia and downregulated the ratio of pERK/ERK, where its activation is crucial for pain signaling [287,288]. Therefore, it is possible to target GDNF signaling as a therapeutic treatment for bone cancer pain.

### Gastric Cancer

Genome-wide DNA methylation profiling of metastatic and nonmetastatic gastric carcinomas and their surgical margins has identified that *GFR1* gene is 1 of the 15 genes that were significantly differentially methylated in gastric carcinoma compared with the surgical margins, and methylation changes of *GFR1*, together with *SRF* and *ZNF382*, may be potential biomarkers for gastric carcinoma metastasis prediction [289]. Similarly, *GFR3* promoter region was shown to be markedly hypermethylated in almost all gastric tumors [290].

### Concluding Remarks

The GDNF family represents a group of four structurally related ligands that have traditionally taken up developmental roles within the neuronal system. However, more recently, they have been ascribed additional developmental and maintenance functions within

extraneuronal tissues, including the kidney, testis, eyes, ovary, hematopoietic system, vascular system, pituitary, and endometrium. Owing to their ability to regulate numerous physiological actions in these tissues such as cell survival, motility, and proliferation, their deregulation has always been suspected to yield oncogenic effects. Central to this has been their indirect implication in neuroendocrine tumors, and from the aforementioned studies, this has been recently expanded to include cancers of epithelial origin in endocrine-responsive tissues—most notably breast, endometrial, and pancreatic cancers. Here we have reviewed GFL biology and discussed the evidence and rationale behind whether components of the GFL signaling complex, particularly the ligands themselves, present novel opportunities for therapeutic intervention in cancer.

Taken together, the literature suggests that GFLs have a very complex and tissue-dependent role in both homeostatic physiology and oncogenesis. Indicative of this is the multitude of heterogeneous phenotypes and mechanisms that cell populations can exhibit with respect to combinations of GFLs and GFR $\alpha$ s: both membrane-bound and -soluble forms, RET, nontraditional receptors (HSPG, NCAM, integrins, Met), GFL paracrine versus autocrine actions, and GFL extracellular concentration and release. This inherent complexity ensures that challenges remain and highlights the importance of further research that links *in vitro* biology with clinical significance.

All GFLs have a pronounced role in early development and postnatal repair. Oncogenesis sees their reactivation and/or deregulation of homeostatic levels. With respect to GDNF, it appears that it functions primarily in a paracrine nature whether it is from the surrounding microenvironment—more specifically, support or stromal cells (adipose, fibroblasts, immune, epithelial, and muscle)—or the neuronal architecture supporting the tissue. Endogenously sourced GDNF seems to act in a supporting role as opposed to being a key driver of nonneuronal function. The lack of clinical and primary tumor correlation is indicative of this. Conversely, ARTN exhibits a predominantly autocrine role within tumors of various origins. Currently, there is a lack of sufficient data to attribute an oncogenic role to NRTN and PSPN.

### Disclosure of Potential Conflicts of Interest

D. X. L. and P. E. L. are inventors on PCT application PCT/NZ2008/00152, US provisional application 61/234902, China ZL200880104674.9, EPO 2164870, and US 14/482512. P. E. L. is an inventor on US provisional application 61/252513 and US 2015/0079106A1. The other authors declare that they have no conflict of interest.

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