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GFR α 1 Is an Essential Receptor Component for GDNF in the Developing Nervous System and Kidney

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Summary

Glial cell line–derived neurotrophic factor (GDNF) is a distant member of the TGF β protein family that is essential for neuronal survival and renal morphogenesis. We show that mice who are deficient in the glycosyl-phosphatidyl inositol (GPI)-linked protein GFR α 1 (GDNFR α) display deficits in the kidneys, the enteric nervous system, and spinal motor and sensory neurons that are strikingly similar to those of the GDNF- and Ret-deficient mice. GFR α 1-deficient dopaminergic and nodose sensory ganglia neurons no longer respond to GDNF or to the structurally related protein neurturin (NTN) but can be rescued when exposed to GDNF or neurturin in the presence of soluble GFR α 1. In contrast, GFR α 1-deficient submandibular parasympathetic neurons retain normal response to these two factors. Taken together with the available genetic and biochemical data, these findings support the idea that GFR α 1 and the transmembrane tyrosine kinase Ret are both necessary receptor components for GDNF in the developing kidney and nervous system, and that GDNF and neurturin can mediate some of their activities through a second receptor.

Introduction

Glial cell line–derived neurotrophic factor (GDNF) (Lin et al., 1993), neurturin (NTN) (Kotzbauer et al., 1996), and persephin (PSP) (Milbrandt et al., 1998) constitute a class of secreted proteins that is structurally related to the transforming growth factor protein family. Studies in primary neuronal cultures, as well as in lesioned animal models, have provided evidence that GDNF is a survival factor for embryonic midbrain dopaminergic neurons that degenerate in Parkinson's disease (Lin et al., 1993; Beck et al., 1995; Tomac et al., 1995), spinal motor neurons that degenerate in amyotrophic lateral sclerosis and spinal muscular atrophies (Henderson et al., 1994; Oppenheim et al., 1995; Yan et al., 1995), locus coeruleus noradrenergic neurons (Arenas et al., 1995), and subpopulations of peripheral sensory,

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sympathetic, and parasympathetic neurons (Buj-Bello et al., 1995; Trupp et al., 1995). Likewise, NTN was shown to be effective in attenuating the death of cultured embryonic dopaminergic, motor (Klein et al., 1997), sympathetic, and sensory neurons (Kotzbauer et al., 1996), while PSP was shown to promote the survival of cultured dopaminergic and motor, but not peripheral, neurons (Milbrandt et al., 1998).

The essential physiological role of the GDNF/NTN/PSP protein family is illustrated by the phenotype of mice in whom the *GDNF* gene has been disrupted. These mice display deficits in primary sensory, sympathetic, and motor neurons and also fail to develop metanephric kidneys, ureters, and most of the enteric nervous system (Moore et al., 1996; Pichel et al., 1996; Sánchez et al., 1996). Consequently, although these mice are born, they die shortly after birth, owing to an inability to consume milk and to the lack of renal function.

Despite the physiological and clinical significance of the GDNF protein family, the mechanism by which these growth factors transduce signals is not fully understood. Biochemical and cell culture studies have suggested that GDNF and NTN bind one of several glycosyl-phosphatidyl inositol (GPI)-linked proteins (designated GDNF family receptors GFR α 1–4) and that they also require the presence of the transmembrane tyrosine kinase Ret for signal transduction and neuronal survival (Jing et al., 1996; Treaner et al., 1996; Baloh et al., 1997; Buj-Bello et al., 1997; Klein et al., 1997; Sanicola et al., 1997; Masure et al., 1998; Naveilhan et al., 1998; Thompson et al., 1998; Worby et al., 1998) (data not shown). These studies further revealed that GDNF and NTN facilitate the formation of a physical complex between GFR α and Ret and lead to activation of the Ret tyrosine kinase (Jing et al., 1996; Treaner et al., 1996; Klein et al., 1997). Taken together, the findings supported the proposal that cellular responses to the GDNF protein family are mediated through multicomponent receptor complexes composed of a shared signaling subunit, the orphan tyrosine kinase, Ret, and one of several ~55 kDa ligand binding subunit proteins, which do not have an intracellular domain, and which bind the extracellular membrane leaflet via the GPI lipid modification.

Despite the available information, several major issues concerning the validity of the multicomponent receptor model remained. Most importantly, to date there is no evidence that the GPI-linked proteins are, in fact, functional and necessary receptor subunits for the GDNF protein family *in vivo*. In addition, the ligand specificity of the endogenous GPI-linked proteins remained unclear—this since, in a cell free system, recombinant GDNFR α /GFR α 1 and NTN α /GFR α 2 displayed a high degree of binding specificity for GDNF and NTN, respectively, when tested alone (Klein et al., 1997) but bound GDNF equally well in the presence of Ret (Sanicola et al., 1997). Likewise, GDNF and NTN were shown to be equally effective in activating Ret through GFR α 1, and NTN appeared only 30-fold more efficient than GDNF in activating Ret through GFR α 2 in a fibroblast cell line (Baloh et al., 1997). Nonetheless, in primary neurons, GDNF displayed a preference for GFR α 1, and NTN could promote survival only through GFR α 2 (Buj-Bello et al., 1997).

To examine the physiological significance and ligand specificity of the GFR α proteins, we have generated and analyzed mice who are deficient in GFR α 1 (*GFR α 1^{-/-}*). We show that *GFR α 1^{-/-}* mice display neuronal and renal deficits that are strikingly similar, but not identical to, those of the *GDNF^{-/-}* and *Ret^{-/-}* mice. Moreover, nodose ganglia and midbrain dopaminergic neurons derived from the *GFR α 1^{-/-}* embryos no longer survive in the presence of GDNF and NTN, whereas the response of *GFR α 1^{-/-}* submandibular parasympathetic neurons to these two factors is indistinguishable from that of their wild-type counterpart. The findings verify the physiological importance of the GFR α receptors and validate the multicomponent receptor hypothesis for the GDNF protein family; they further support the idea that, although GDNF and NTN display receptor preferences, they can use multiple GFR α receptors *in vivo*.

Results

Generation of the *GFRα1*^{-/-} Mice

DNA fragments containing the first three of nine exons in the *GFRα1* gene (Eng et al., 1998) were isolated and used to generate a targeting construct in which part of exon 2, encoding for amino acid 14–66, had been deleted (Figure 1A). This targeting construct was electroporated into embryonic stem (ES) cells (Moore et al., 1996), and clones in which the *GFRα1* gene had been disrupted by homologous recombination (Figure 1B) were injected or aggregated into blastocysts to produce *GFRα1* mutant mice (Figure 1C).

Whereas mRNA for *GFRα1* was found in the kidney, gut, and nervous system of wild-type and *GFRα1* heterozygous embryos by in situ hybridization, no *GFRα1* transcripts encoding amino acids 14–66 were detected in null, mutant littermates (Figures 1D and 1E). Heterozygous mice were viable, normal in size, fertile, and did not display any gross morphological or behavioral abnormalities. In contrast, *GFRα1*^{-/-} mice died 1–1.5 days after birth, even though they were initially able to suckle and had normal limb and body movements.

Neuronal Deficits in the *GFRα1* Null Mice

Since GDNF, NTN, and PSP are potent survival factors for embryonic midbrain dopaminergic (Lin et al., 1993; Beck et al., 1995; Tomac et al., 1995; Milbrandt et al., 1998), noradrenergic (Arenas et al., 1995), motor (Henderson et al., 1994; Oppenheim et al., 1995; Yan et al., 1995), sensory, and sympathetic (Buj-Bello et al., 1995; Trupp et al., 1995; Kotzbauer et al., 1996) neurons in culture and/or in animal models in vivo, we first examined whether the *GFRα1*^{-/-} mutant mice displayed any neuronal deficits. As was observed in the *GDNF*^{-/-} mice (Moore et al., 1996; Sánchez et al., 1996), the *GFRα1*^{-/-} embryos exhibited small losses of lumbar spinal (24%) and trigeminal nucleus (22%) motor neurons, but not of facial motor neurons (Table 1). In addition, like their *GDNF*^{-/-} counterparts, the *GFRα1*^{-/-} embryos had a normal complement of tyrosine hydroxylase–positive dopaminergic neurons in the substantia nigra (Table 1; Figures 2A and 2B), did not display a significant reduction in the density of dopaminergic projections in the striatum (Table 1; Figures 2C and 2D), and possessed a normal number of noradrenergic neurons in the locus coeruleus (Table 1).

Surprisingly, however, a comparison of *GDNF*^{-/-} and *GFRα1*^{-/-} embryos showed differences in other neuronal populations. For instance, whereas the *GDNF*^{-/-} embryos possessed a 23% deficit in the L5 dorsal root ganglia sensory neurons, the *GFRα1*^{-/-} embryos had a normal complement of this neuronal population. Similarly, although the *GDNF*^{-/-} mice displayed a decrease of 40% in the number of petrosal-nodose sensory ganglia neurons, the size of this ganglion was reduced by only 15% in the *GFRα1*^{-/-} mice. Finally, even though the *GDNF*^{-/-} and the *Ret*^{-/-} embryos suffered 35% and 100% losses, respectively, in the sympathetic superior cervical ganglion neurons (Durbec et al., 1996; Moore et al., 1996), *GFRα1*^{-/-} mice did not display any significant loss of these neurons (Table 1; Figures 2E–2H; data not shown).

The fact that *GDNF*^{-/-} and the *GFRα1*^{-/-} mice display mild but identical deficits in the number of lumbar spinal and trigeminal motor neurons is consistent with the notion that *GFRα1*^{-/-} serves as an essential receptor for GDNF in these neuronal populations. However, the findings that neither the absence of GDNF (Moore et al., 1996; Pichel et al., 1996; Sánchez et al., 1996), nor the absence of its putative receptor subunit (Table 1; Figure 2), led to a profound degree of neuronal cell losses outside the enteric nervous system suggests that although GDNF can promote the survival of multiple neuronal populations in vitro, it is not a predominant survival factor for central or peripheral neurons in vivo. Since *GDNF*^{-/-} embryos suffer a more severe loss of petrosal-nodose, dorsal root, and superior cervical ganglia neurons, as compared

with the *GFRα1*^{-/-} mice, some of the survival effects of GDNF on these neuronal populations may be mediated by a second receptor.

The finding that the *GFRα1* (no deficit) and *GDNF* (35% deficit) null mice each had a less severe neuronal loss of superior cervical ganglia neurons than their *Ret* null counterpart (100% deficit) (Table 1) is in agreement with the hypothesis that *Ret* is a shared signaling component for the GDNF protein family that can act in conjunction with multiple GFR subunits. Although only a limited survey of neuronal deficiencies has been published for the *Ret*^{-/-} embryos (Schuchardt et al., 1994; Durbec et al., 1996), these findings suggest that the neuronal deficits in the *Ret*^{-/-} mice will be at least as severe as those found in their *GFRα1* and *GDNF* counterparts.

Enteric Nervous System Deficits

As GDNF (Moore et al., 1996; Pichel et al., 1996; Sánchez et al., 1996) and *Ret* (Schuchardt et al., 1994; Durbec et al., 1996) were shown to be essential for the development of the enteric nervous system, we next determined whether ablation of *GFRα1* would lead to deficits in this tissue. In E17 wild-type and *GFRα1*^{+/-} mice, the neural crest-derived enteric neurons belonging to the myenteric (Auerbach) and submucosal (Meissner) plexi were readily visible along the length of the gastrointestinal tract (Figure 3; data not shown). In contrast, these neurons were completely absent in the intestines and colons of age-matched *GFRα1*^{-/-} littermates (Figure 3). In addition, the *GFRα1*^{-/-} animals displayed only a small number of neurons in the stomach (part of the foregut) (data not shown). The absence of myenteric and submucosal neurons in the intestines of the *GFRα1*^{-/-} mice, and the presence of some neurons in the stomachs of these animals, concurs with previous observations in the *GDNF*^{-/-} (Moore et al., 1996; Pichel et al., 1996; Sánchez et al., 1996) and *Ret*^{-/-} (Schuchardt et al., 1994; Durbec et al., 1996) embryos.

Taken together, these findings strengthen the notion that *GFRα1*, *Ret*, and GDNF act in the same signaling pathway and support the idea that a pool of neural crest cells that is derived from the postotic hindbrain and is dependent on the GDNF signal gives rise to most of the enteric nervous system, whereas a distinct lineage that is derived from trunk neural crest and is not completely dependent on the GDNF signal contributes to the enteric nervous system in the foregut (Durbec et al., 1996).

Surprisingly, in the *GFRα1*^{-/-} (Figure 3H) but not in *GDNF*^{-/-} (Figure 3I) embryos, a small number of enteric neuron cell bodies were detected in the descending colon, sigmoid colon, and rectum. Thus, it appears as if a subpopulation of enteric neurons that reside in these derivatives of the hindgut may respond to GDNF in part, through a second receptor.

Renal Deficits

Studies in embryonic kidney cultures have demonstrated that PSP and GDNF can promote the outgrowth and branching of the ureteric bud from the nephric ducts (Milbrandt et al., 1998). In addition, GDNF (Moore et al., 1996; Pichel et al., 1996; Sánchez et al., 1996) and *Ret* (Schuchardt et al., 1994) have been shown to be essential for the development of ureters and kidneys by gene targeting. We therefore further examined the *GFRα1*^{-/-} mice for renal abnormalities.

In agreement with the hypothesis that *GFRα1* is an essential receptor component for GDNF in the developing kidney, most of the *GFRα1*^{-/-} animals (13 of 17) had complete bilateral renal and ureteral deficits. In the remaining *GFRα1*^{-/-} embryos (4 of 17), one rudimentary kidney was detected. Other organs that are derived from the embryonic urogenital intermediate mesoderm, including the pro- and mesonephros, the adrenal glands, and the gonads, as well as the

remaining abdominal viscera and thoracic tissues, appeared normal (Figures 4A and 4B; data not shown). Anatomical examination of the *GFRα1*^{-/-} mice at E12.5, when the metanephric kidney forms, demonstrated the presence of a mesonephric duct and undifferentiated kidney mesoderm but not of a morphologically defined ureter, ureteric bud, or nephrons (Figures 4C and 4D). Moreover, Pax2, a homeo-domain transcription factor that is initially expressed in the early ureteric epithelium and is then induced in the nephrogenic region of the metanephrogenic blastema (Dressler and Douglass, 1992; Rothenpieler and Dressler, 1993), was absent from the kidneys of the E12.5 *GFRα1*^{-/-} embryos (Figures 4E and 4F). In contrast, the Wilm's tumor suppressor gene and putative transcriptional repressor WT1, which is initially expressed in the uninduced kidney mesenchyme (Kreidberg et al., 1993), was found in the kidney region of both the wild-type and *GFRα1*^{-/-} E12.5 embryos (Figures 4G and 4H).

The mammalian kidney develops by reciprocal inductive interactions between the ureteric bud, which is an evagination of the mesonephros/Wolffian duct, and the metanephrogenic blastema, a caudal intermediate mesodermal tissue. The metanephrogenic blastema (which makes GDNF) is thought to induce the ureteric bud (which makes GFRα1 and Ret) to form collecting ducts/ureter. The differentiated ureteric bud, in turn, induces the metanephrogenic blastema to form nephrons (Saxen, 1987). The absence of Pax2-positive ureteric bud cells in *GFRα1*^{-/-} mice suggests that the ureteric bud either did not branch from the nephric duct or degenerated shortly after its formation. In the absence of differentiated ureters, the renal parenchyma most likely does not express Pax2 and will not undergo a mesenchymal-to-epithelial conversion. Consequently, no differentiated nephrons will be formed. The fact that WT1 is expressed in the metanephrogenic blastema of the *GFRα1*^{-/-} mice is consistent with the notion that this tissue is dedicated to becoming kidney independent of the ureteric bud and of the GDNF signal.

The Response of *GFRα1* Null Neurons to GDNF

To further elucidate whether the neuronal and renal deficits that were observed in the *GFRα1*^{-/-} mice were indeed caused by a failure in the reception of the GDNF signal, we examined the response of cells derived from the GFRα1-deficient mice to this factor. Embryonic, nodose sensory ganglia neurons were dissected from wild-type, *GFRα1*^{+/-}, and *GFRα1*^{-/-} embryos, and their survival in the presence of GDNF and other neurotrophic factors was examined. The majority of wild-type and *GFRα1*^{-/-} nodose neurons survived in the presence of brain-derived neurotrophic factor, which mediates its signal through the tyrosine kinase receptor TrkB (Klein et al., 1991; Soppet et al., 1991) (Figure 5A). Likewise, GDNF prevented the death of nodose neurons that were derived from wild-type or *GFRα1*^{+/-} neurons (Figure 5A; data not shown). However, consistent with the idea that GFRα1 is an essential GDNF receptor component, GDNF failed to rescue the majority of GFRα1-deficient nodose neurons, even when applied at high concentrations (50 ng/ml) (Figure 5A). Likewise, primary embryonic dopaminergic neurons that were derived from wild-type embryos survived in the presence of GDNF, while similar neurons that originated from *GFRα1*^{-/-} littermates no longer responded to this factor at any of the concentrations tested (Figure 5B). Neuronal survival in the presence of GDNF was, however, restored following the addition of exogenous, soluble, recombinant GFRα1 to the GFRα1-deficient neurons (Figure 5B), supporting the idea that these neurons failed to survive in the presence of GDNF solely owing to the absence of GFRα1, and that they did not degenerate at earlier embryonic stages.

Surprisingly, although the response of both nodose and dopaminergic neurons to GDNF was completely dependent on GFRα1, we found that *GFRα1*^{-/-} parasympathetic, submandibular neurons survived in the presence of GDNF as well as did their wild-type counterparts (Figure 5C). Taken together, these findings support the hypothesis that GFRα1 is an essential GDNF receptor component in many, but not all, populations of neuronal cell types. In addition, it

appears as if distinct classes of neurons and nonneuronal cells may be able to respond to GDNF, possibly with a lower sensitivity, via an alternative receptor.

The Response of *GFRα1*^{-/-} Neurons to NTN

In view of reports that cells expressing *GFRα1* and Ret can respond to the GDNF-related protein NTN (Baloh et al., 1997), we have determined whether the survival of *GFRα1*^{-/-} neurons is still stimulated by this factor.

Examination of wild-type embryonic nodose sensory ganglia neurons revealed that at low concentrations (2 ng/ml), NTN rescued only 25% of the neurons that were rescued by a similar concentration of GDNF. In contrast, at higher concentrations (50 ng/ml), NTN was able to rescue most of the GDNF-responsive neurons (Figure 6A). Nodose ganglia neurons from the *GFRα1*^{-/-} retained their response to the low concentrations of NTN, but no longer responded to high concentrations of this factor (Figure 6A). Thus, it appears as if NTN promotes survival of nodose neurons through two distinct receptors, a high affinity receptor (possibly *GFRα2* [Buj-Bello et al., 1997; Klein et al., 1997]), which might be expressed by a small subpopulation of the embryonic nodose neurons, and *GFRα1*, which appears to be more abundantly expressed, and which functions as a lower affinity receptor for NTN in this neuronal cell type.

Analysis of wild-type embryonic midbrain dopaminergic neurons revealed that this neuronal population survives equally well in the presence of NTN and GDNF (Figure 6B; data not shown). Surprisingly, given previous evidence that NTN could mediate its activities through a distinct receptor (Buj-Bello et al., 1997; Klein et al., 1997), neither GDNF (Figure 5B) nor NTN (Figure 6B) were able to support the survival of dopaminergic neurons that were taken from *GFRα1*^{-/-} embryos. As before, the survival of *GFRα1*^{-/-} dopaminergic neurons in the presence of NTN was restored following addition of exogenous, soluble recombinant *GFRα1* (Figure 6B). Thus, *GFRα1*^{-/-} functions as a receptor for both NTN and GDNF in developing dopaminergic neurons in vitro.

Among the neuronal populations tested, primary embryonic, parasympathetic submandibular neurons were found to be most sensitive to NTN (Figure 6C). Whereas *GFRα1*^{-/-} nodose and dopaminergic neurons no longer respond to NTN, the ability of *GFRα1*^{-/-} parasympathetic submandibular neurons to survive in the presence of this factor, even at very low concentrations (0.08 ng/ml), was indistinguishable from that of wild-type neurons (Figure 6C). Thus, the response of submandibular neurons to NTN does not require *GFRα1* and must be mediated by a distinct receptor that could be *GFRα2* (Buj-Bello et al., 1997; Klein et al., 1997) or *GFRα3* (Masure et al., 1998; Naveilhan et al., 1998; Worby et al., 1998). Interestingly, although *GFRα1*^{-/-} submandibular neurons survive in the presence of GDNF as well as their wild-type counterparts, GDNF was less effective than NTN in promoting the survival of this neuronal population. Thus, in contrast to NTN, GDNF did not elicit significant survival of parasympathetic submandibular neurons at 0.08 ng/ml or 0.4 ng/ml (Figure 5C) and was effective only at concentrations above 2 ng/ml. The fact that GDNF was less potent than NTN in promoting the survival of submandibular neurons is consistent with the idea that the receptors present on these cells are preferentially activated by NTN.

Taken together, these data favor the hypothesis that *GFRα1* is the major high affinity receptor for GDNF on most cell types (Jing et al., 1996; Treanor et al., 1996; Buj-Bello et al., 1997), but that GDNF can also interact with *GFRα2* (Baloh et al., 1997; Sanicola et al., 1997) or other *GFRα* receptors. Conversely, it appears as if NTN has a high affinity receptor (Buj-Bello et al., 1997; Klein et al., 1997) but can mediate signals also through *GFRα1* in vitro (Baloh et al., 1997) and possibly in vivo.

Discussion

By creating *GFRα1*^{-/-} mice, we were able to demonstrate that GFRα1 is essential for GDNF signal transduction in the developing kidney and enteric nervous system, and in subpopulations of motor, sensory, and sympathetic neurons. These findings substantiate the hypothesis that the receptors for the GDNF protein family are composed of two subunits; a GPI-linked ligand-binding protein that belongs to the GFRα family and a signaling component that is represented by the transmembrane tyrosine kinase Ret.

Analyses of the *GFRα1*^{-/-} mice further revealed that GFRα1 can function as a receptor for other GDNF-like proteins, such as NTN, and that in some cell types, GDNF can elicit a response through an alternative receptor, possibly another member of the GFR family.

Comparing the *GFRα1*^{-/-}, *Ret*^{-/-}, and *GDNF*^{-/-} Mice

The striking similarities in the phenotypes of the *GFRα1*^{-/-}, *Ret*^{-/-}, and *GDNF*^{-/-} mice, which include deficits in the kidneys and enteric neurons, strongly support the proposal that these three molecules are components of the same signaling cascade, and that Ret and GFRα1 serve as coreceptors for GDNF. Mechanistically, the fact that *GFRα1*^{-/-}, *GDNF*^{-/-}, and *Ret*^{-/-} mice all display a similar loss-of-function phenotype indicates that GFRα1 acts as a coactivator, rather than as a ligand-regulated suppresser of Ret, as suggested for other coreceptors (Stone et al., 1996).

Despite the overall similarities in phenotype, some differences between these mice are notable. First, whereas the *GFRα1*^{-/-} mice have a normal complement of superior cervical ganglion neurons (Table 1), there is a partial (35%) loss of this neuronal population in *GDNF*^{-/-} embryos (Moore et al., 1996), and a complete loss in the *Ret*^{-/-} mice (Durbec et al., 1996). Likewise, while the *GDNF*^{-/-} embryos display deficits of 40% and 23%, respectively, in the number of petrosal-nodose and dorsal root ganglia neurons, the *GFRα1*^{-/-} mice display only a small deficit (15%) in the number of petrosal-nodose ganglia neurons, and they have a normal complement of dorsal root ganglia neurons.

We cannot exclude the possibility that these differences stem in part from the genetic background of these mice. Nevertheless, the severity of the phenotype of the *Ret*^{-/-} mice, when compared with that of the *GFRα1*^{-/-} mice, is consistent with the idea that Ret is an essential, shared signaling component for the GDNF family of receptors (Jing et al., 1996; Treanor et al., 1996; Buj-Bello et al., 1997; Klein et al., 1997). Likewise, the fact that *GDNF*^{-/-} mice display more significant neuronal deficits, as compared with their *GFRα1*^{-/-} counterparts, supports the proposal that GDNF can mediate signals through GFRα2 (Baloh et al., 1997; Sanicola et al., 1997) or through another member of the GFRα receptor family. Further evidence of this hypothesis is provided by our findings that some populations of neurons respond to GDNF in the absence of GFRα1 (Figures 5C and 6C).

Surprisingly, although the comparison between the *GDNF*^{-/-} and *GFRα1*^{-/-} mice suggests that GDNF rescues petrosal-nodose neurons independent of GFRα1 in vivo (Table 1), GDNF was not able to promote the survival of a significant number of *GFRα1*^{-/-} petrosal-nodose neurons in vitro (Table 1; Figure 5A). Thus, it is possible that the effects of GDNF on this neuronal population in vivo are indirect. Alternatively, GDNF may influence the development of a subpopulation of these neurons at a stage before they were cultured.

Ligand Specificity of the GFRα Receptors

Studies in a cell-free system indicated that GFRα1 selectively binds to GDNF, whereas GFRα2 selectively binds NTN (Klein et al., 1997). Surprisingly, although Ret by itself does not bind any known member of the GDNF protein family with a high affinity (Jing et al.,

1996; Treanor et al., 1996; Klein et al., 1997), it can, when coexpressed with the GFR α s, change their ligand-binding specificity, allowing, for example, the binding of GDNF to GFR α 2 (Sanicola et al., 1997). The promiscuousness of the GFR α s, in the presence of Ret, is further illustrated by the findings that cells that express GFR α 2 together with Ret can respond to multiple ligands (Baloh et al., 1997; Buj-Bello et al., 1997).

To examine the ligand specificity of the endogenous GFR α 1 and the receptor specificity of GDNF and NTN, we have analyzed multiple classes of primary neurons derived from the GFR α 1^{-/-} embryos. This analysis revealed that GDNF can, in certain cell types, mediate its response through an alternative receptor, as illustrated by its ability to promote the survival of GFR α 1^{-/-} submandibular neurons. Similarly, examination of GFR α 1^{-/-} neurons for their response to NTN showed that GFR α 1 serves as a receptor for NTN on dopaminergic and nodose sensory ganglia, while submandibular neurons respond to this factor via another receptor, most likely GFR α 2. Although these findings are consistent with the proposal that GDNF and NTN can activate multiple receptors, it is important to note that these receptors are activated with a different potency. Thus, GFR α 1 appears to be activated preferentially by GDNF, whereas the receptor present on submandibular neurons is preferentially activated by NTN (Figures 5C and 6C).

The GFR α Receptors May Display Distinct Functions In Vivo

As revealed by experiments in cultured cells (Baloh et al., 1997) and neurons (Buj-Bello et al., 1997), and through the analysis of the GFR α 1^{-/-} mice, it appears that both NTN and GDNF can interact with multiple receptors (e.g., GFR α 1 and GFR α 2), and that GFR α 1 can mediate the activity of multiple ligands. Surprisingly, however, although such promiscuous receptor-ligand interactions can take place in an experimental setting (Figures 5 and 6) (Baloh et al., 1997), they do not appear to be prevalent in vivo, since most of the major deficits in the GFR α 1^{-/-} mice are recapitulated in the GDNF^{-/-} mice. The limited ability of GFR α 2 to compensate for GFR α 1 in vivo or in vitro is consistent with the distinct tissue distribution of these two receptors. For example, developing dopaminergic neurons, which do not respond to GDNF in vitro, in the absence of GFR α 1, express only GFR α 1, whereas GFR α 2 transcripts are found in high abundance adjacent to, but not on, this neuronal cell type (M. H. and L.-C. W., unpublished data). Likewise, high levels of GFR α 1 mRNA, but not of GFR α 2 mRNA, are found in nodose sensory ganglia neurons (Buj-Bello et al., 1997) and on spinal motor neurons (Klein et al., 1997; Widenfalk et al., 1997), which show deficits in the GFR α 1^{-/-} mice. Finally, GFR α 1, but not GFR α 2, transcripts are expressed in the embryonic day 14 mouse kidney, an organ that fails to develop in the absence of GFR α 1 (Baloh et al., 1997). In contrast, cells such as dorsal root and superior cervical ganglia neurons, which suffer minor or no deficits in GFR α 1^{-/-} embryos as compared with the GDNF^{-/-} embryos, appear to express both GFR α 1 and GFR α 2 during development (Baloh et al., 1997; Nosrat et al., 1997; Widenfalk et al., 1997). Likewise, parasympathetic neurons, which retain responsiveness to GDNF and NTN in the absence of GFR α 1, appear to express multiple GFRs (A. F. and A. D., unpublished data in the chick embryo). The one possible exception to this rule that we observed is enteric neurons. Despite the fact that both GFR α 1 and GFR α 2 are expressed in the developing gut (Jing et al., 1996; Treanor et al., 1996; Baloh et al., 1997; Buj-Bello et al., 1997; Klein et al., 1997), GFR α 2 fails to compensate for GFR α 1 in this tissue. However, the cell type that expresses GFR α 2 in the gut has not been identified, and it is possible that the inability of GFR α 2 to substitute for GFR α 1 is due to its absence from the enteric neurons. This may be analogous to the situation for dopamine neurons, in which both GFR α 1 and GFR α 2 are expressed in the vicinity of dopamine neurons, but only GFR α 1 is expressed at significant levels on the dopamine neurons themselves (M. H. and L. Wang, unpublished data).

In summary, the striking similarities between the *GFRα1*^{-/-}, *GDNF*^{-/-}, and *Ret*^{-/-} mice support the hypothesis that GFRα1 is an important receptor component for GDNF and validate the physiological significance of GFRα and the multicomponent receptor hypothesis (Jing et al., 1996; Treanor et al., 1996; Baloh et al., 1997; Buj-Bello et al., 1997; Klein et al., 1997; Sanicola et al., 1997). Although GDNF and NTN display some preferential interactions with different GFRα receptors in vivo, their distinct functions may be controlled, to a large extent, by their unique tissue distribution.

Experimental Procedures

Production of the *GFRα1*^{-/-} Mice

A genomic library, derived from the 129 mouse strain, was screened with synthetic oligonucleotides, and four positive phages encoding for the *GFRα1* gene were identified. A *neomycin* gene under the control of the phosphoglycerate kinase I promoter was then fused to an 890 bp Eco-XhoI fragment, representing a portion of the *GFRα1* gene N-terminal to amino acid 14, and to a 6.8 kbp Bam HI fragment, representing a portion of the *GFRα1* gene, which is C-terminal to amino acid 66, resulting in a deletion of 52 amino acids from the second exon. A thymidine kinase cDNA under the control of the CMV promoter was added to the targeting construct, and the construct was electroporated into R1 ES (Nagy et al., 1993) and ES D3-C12 (Moore et al., 1996) lines. Clones were selected in 400 μg/ml G418 and 2 μM gancyclovir, expanded, and screened by Southern analysis for homologous recombination. The targeted R1 ES was injected into the blastocoel cavity of 3.5-day-old C57BL/6J blastocysts, and six highly chimeric male founders were chosen for further studies. In parallel, clumps of the targeted R1 ES cells were aggregated with diploid embryos from 2.5-day-old C57BL/6J superovulated females (Nagy and Rossant, 1993), and three additional founders were isolated. The resulting embryos were recovered by caesarean section, genotyped, and analyzed.

GFRα1^{-/-} mice were also produced by tetraploid embryo aggregation (Nagy and Rossant, 1993; Nagy et al., 1993). For this, the targeted ES D3-C12 cells were subjected to 2 mg/ml of G418, and clones in which the second *GFRα1* allele was mutated by gene conversion were identified. Clumps of the double mutant ES cells were sandwiched between, or aggregated with, tetraploid embryos that were generated from 2.5-day-old CD-1 mice and implanted. E11–E12 and E14–E16 embryos were recovered by caesarean section, genotyped by polymerase chain reaction and by eye color (the *GFRα1*^{-/-} embryos, which were derived from the 129 mouse, have pigmented eyes, while the wild-type CD-1 embryos have nonpigmented eyes), and serially sectioned for examination. Since the *GFRα1*^{-/-} offspring of the different founders displayed a similar phenotype (data not shown), extensive analysis was performed only on the R1 ES–derived embryos in both the C57BL/6J and CD-1 background, as specified.

Histological Analysis

For histological analysis, embryos and neonate pups were fixed with 10% neutral buffered formalin, embedded in paraffin, serially sectioned, and stained with hematoxylin and eosin or the indicated antibody for microscopic examination. Antibody staining was performed using antiperipherin (1:300; Chemicon), antineurofilament 150 kDa (1:1000; Chemicon), anti-Pax2 (10 μg/ml; Bablo), and anti-wild type 1 (1:500; Santa Cruz Biotechnology) and the peroxidase Vectastain kit (Vector Labs), as previously described (Moore et al., 1996). Intestines from animals perfused with 4% paraformaldehyde were stained in whole mount with antibodies to the neuronal marker PGP9.5 (1:4000; Biogenesis) as previously described (Moore et al., 1996). For neuronal counts, neonate pups were fixed in Carnoy's fixative and embedded in paraffin. Serial sections (7 μm thick) through the head and the lumbar back of homozygous mutants and wild-type littermates were obtained and stained with cresyl violet. Peripheral ganglia and motor nuclei neurons were counted every sixth section (Moore et al., 1996). Some

series were stained with antibodies to tyrosine hydroxylase (1:200; Pel-Freeze), and tyrosine hydroxylase-positive neurons were counted in the substantia nigra and locus coeruleus (Liching and Anderson, 1995; Moore et al., 1996).

Survival Assays

E12 nodose sensory and E17 submandibular parasympathetic ganglia neurons were isolated from wild-type, *GFRα1*^{+/-}, or *GFRα1*^{-/-} embryos and analyzed as described (Davies et al., 1995). The nodose survival assays for each factor were performed on 21 wild-type and 14 *GFRα1*^{-/-} embryos. The submandibular survival assays were done on 5 wild-type, 9 *GFRα1*^{+/-}, and 12 *GFRα1*^{-/-} embryos for each factor. Individual embryos were analyzed. Dopaminergic neurons were isolated from E12 embryos, and 3–5 embryos were analyzed for each condition as described (Poulsen et al., 1994).

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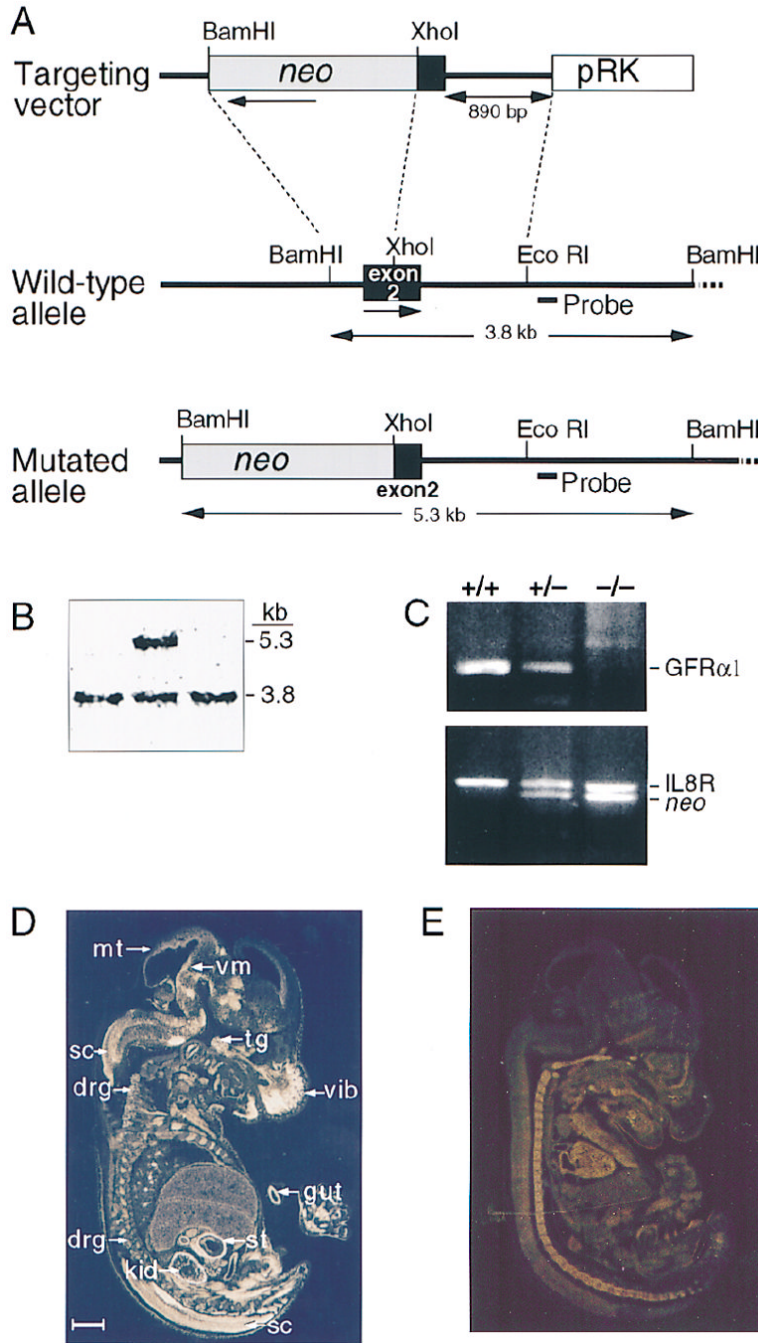


Figure 1. Disruption of the *GFRα1* Gene

(A) Targeting vector, wild-type *GFRα1* allele and the disrupted allele. Amino acids 14–66 are missing from the disrupted gene. The location of the probe used in Southern blot is indicated (Probe). The directions of gene transcription are marked by horizontal arrows.

(B) Detection of homologous recombination event in an ES clone by a Southern blot.

(C) Genotype analysis of wild-type (*+/+*), heterozygous mutant (*+/-*), and homozygous mutant (*-/-*) animals by polymerase chain reaction. The band in the upper panel (*GFRα1*) is specific for the wild-type *GFRα1* gene. The upper band in the lower panel (*IL8R*) represents a control fragment from the *IL8* receptor gene. The lower band in this panel (*neo*) is specific for the *neo* gene.

(D and E) In situ hybridization of wild-type (D) and *GFRα1*^{-/-} (E) E15 mouse embryos with exon 2 *GFRα1* probe. Abbreviations: drg, dorsal root ganglia; gut, gut; kid, kidney; sc, spinal cord; st, stomach; tg, trigeminal ganglion; vib, vibrissa; and vm, ventral midbrain. Scale bar, 1 mm.

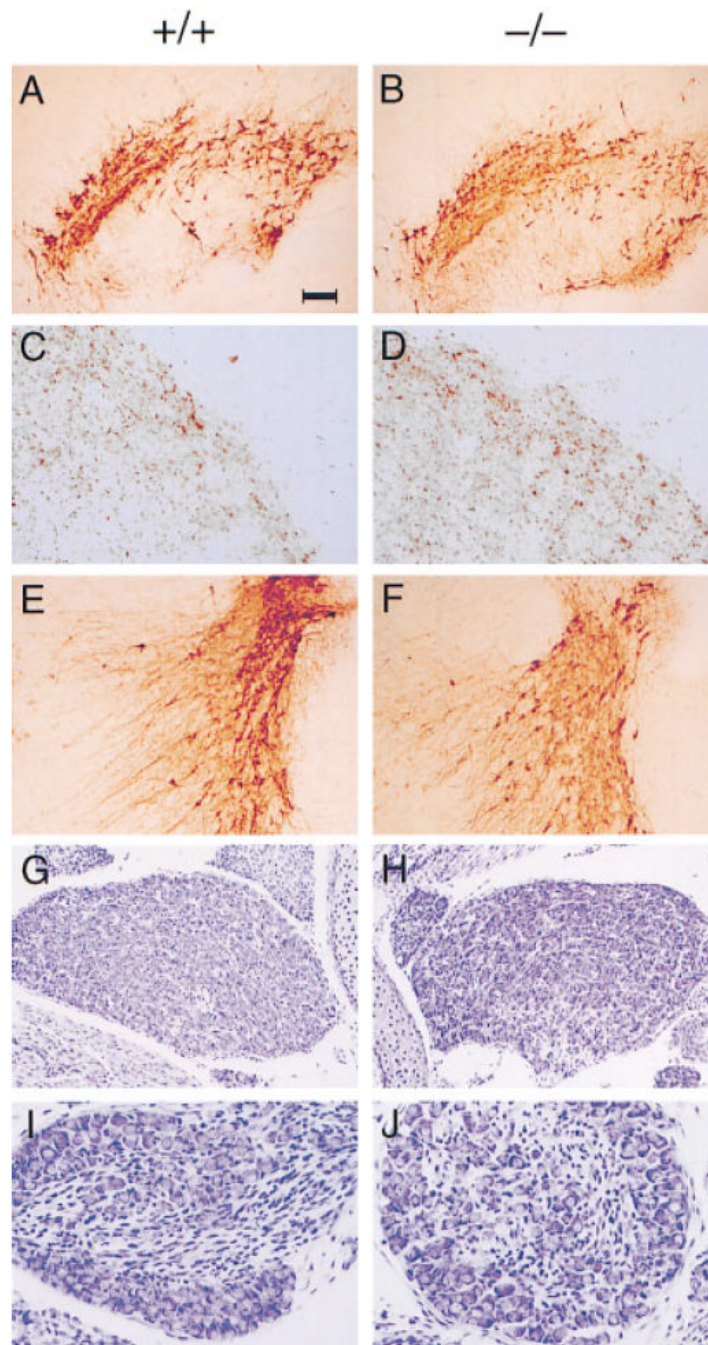


Figure 2. Neuronal Populations in P0 wild-type (+/+) and *GFRα1*^{-/-} (-/-) Mice

Tyrosine hydroxylase staining of substantia nigra (A and B), striatum (C and D), and locus coeruleus (E and F). Tyrosine hydroxylase is the rate-limiting enzyme in dopamine and noradrenaline synthesis. Cresyl violet staining of superior cervical ganglia (G and H) and petrosal nodose ganglia (I and J) neurons from 129 × CD-1, F2 mice. No deficits or abnormalities in neuronal number, morphology, or innervation pattern are detected in the *GFRα1*^{-/-} mice. Scale bar: ~100 μm in (A), (B), (E), and (F), 30 μm in (C), (D), (I), and (J), and 50 μm in (G) and (H).

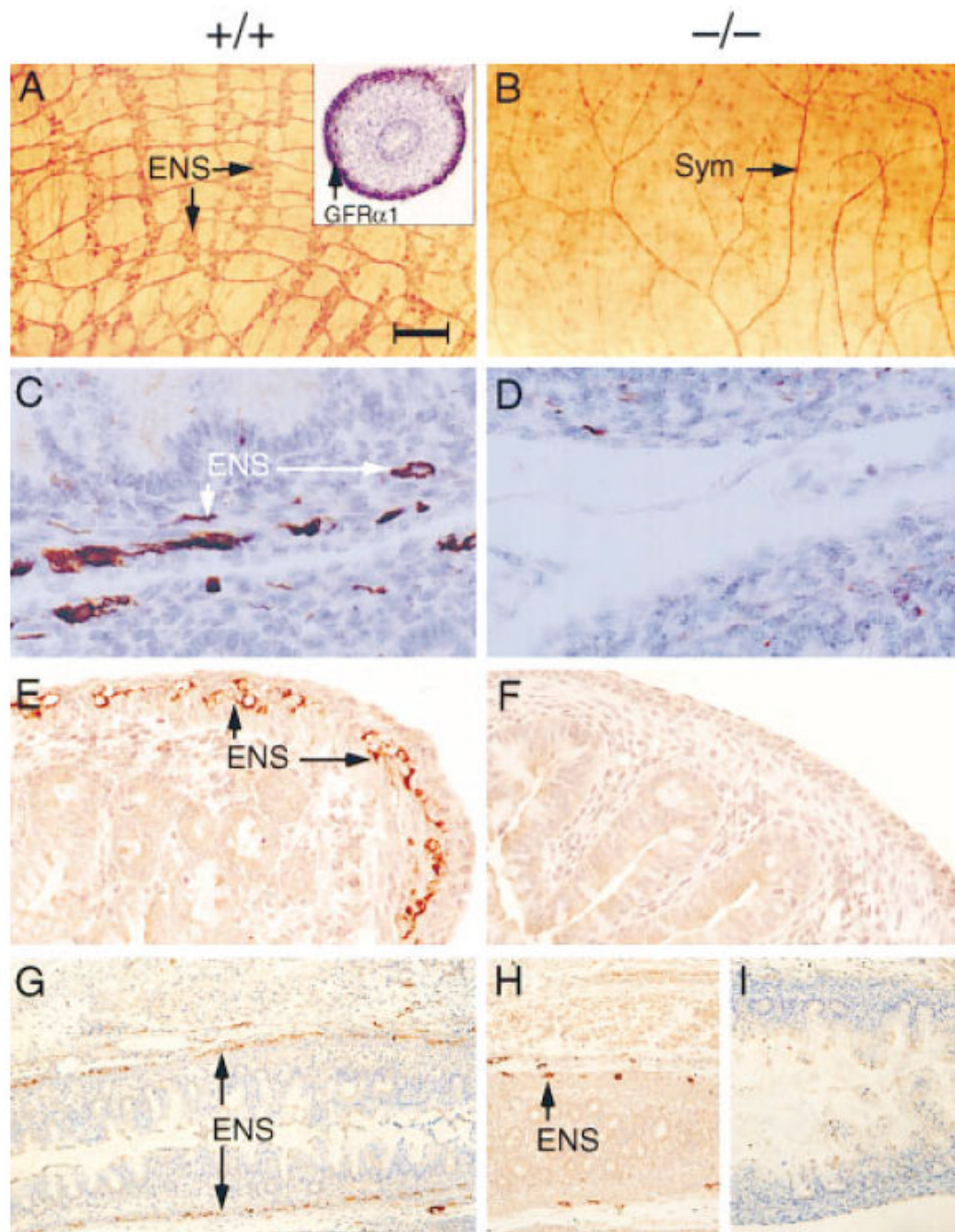


Figure 3. Enteric Nervous System in Wild-Type (+/+) and *GFRα1*^{-/-} (-/-) Mice

(A and B) Whole mounts of small intestine from E18 mice stained with the general neuronal antibody PGP9.5. Inset in (A) depicts the expression of *GFRα1* mRNA in E18 wild-type mouse gut as detected by in situ hybridization.

(C–I) Section through the small intestine (C and D), colon (E and F), and rectum (G–I) of E17 embryos stained with neurofilament (C and D) or peripherin (E–I). Enteric neurons (ENS) were not found in the intestine and are very rarely found in the stomach and colon of the *GFRα1*^{-/-} 129 × CD-1, F2 mice. No ENS neurons were detected in the colon of the *GDNF*^{-/-} mice.

(I) Sym represents afferent fibers probably derived from sympathetic innervation to the gut. Scale bar: ~100 μm in (A) and (B), 30 μm in (C) and (D), 50 μm in (E) and (F), and 300 μm in (G) through (J).

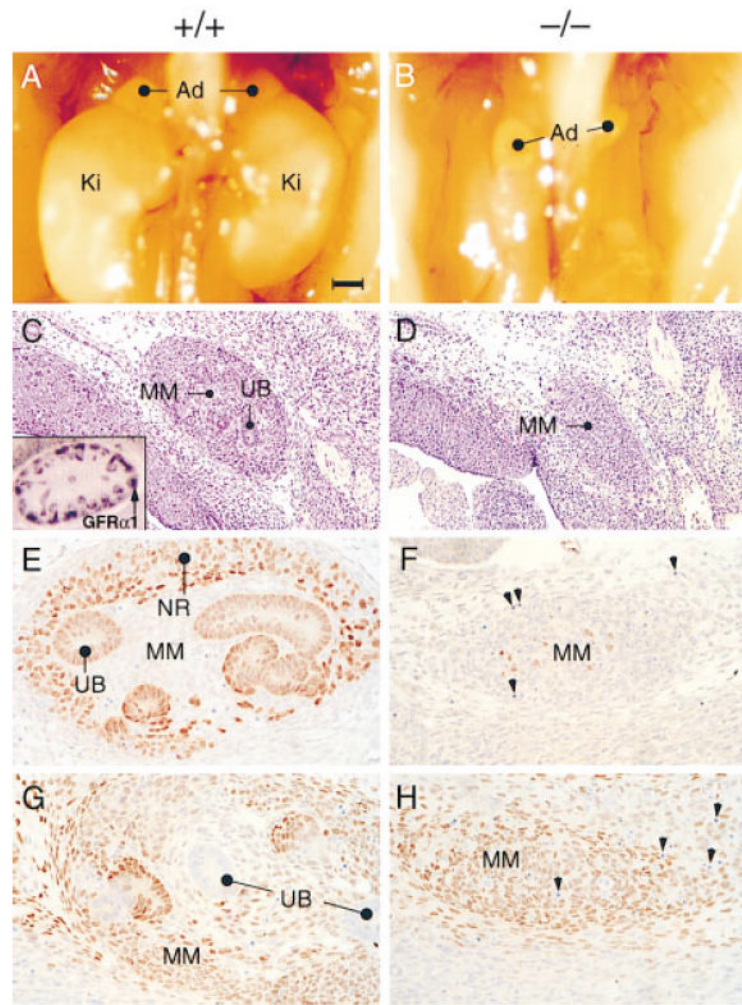


Figure 4. Kidneys in Wild-Type and *GFRα1*^{-/-} Mice

(A and B) Photographs of the abdomen in E17 wild type (A) and *GDNF*^{-/-} (B) 129 × CD-1, F2 mice. Note the position of the kidneys (Ki) subadjacent to the adrenals (Ad) in (A) and their absence in the mutant (B).

(C–H) Sagittal sections through the kidney region of E12.5 wild-type (+/+) and *GFRα1*^{-/-} (-/-) embryos stained with hematoxylin and eosin (C and D), Pax2 antibodies (E and F), or WT1 antibodies (G and H). Inset in (C) represents in situ hybridization of *GFRα1* cDNA probe to developing nephrons and ureteric bud in wild-type kidney. Abbreviations: UB, ureteric bud; MM, metanephric (condensing) mesenchyme; and NR, nephrogenic region (the region that undergoes mesenchymal-to-epithelial conversion and differentiated nephrons). Scale bar: ~500 μm in (A) and (B), 100 μm in (C) and (D), and 20 μm in (E) through (H).

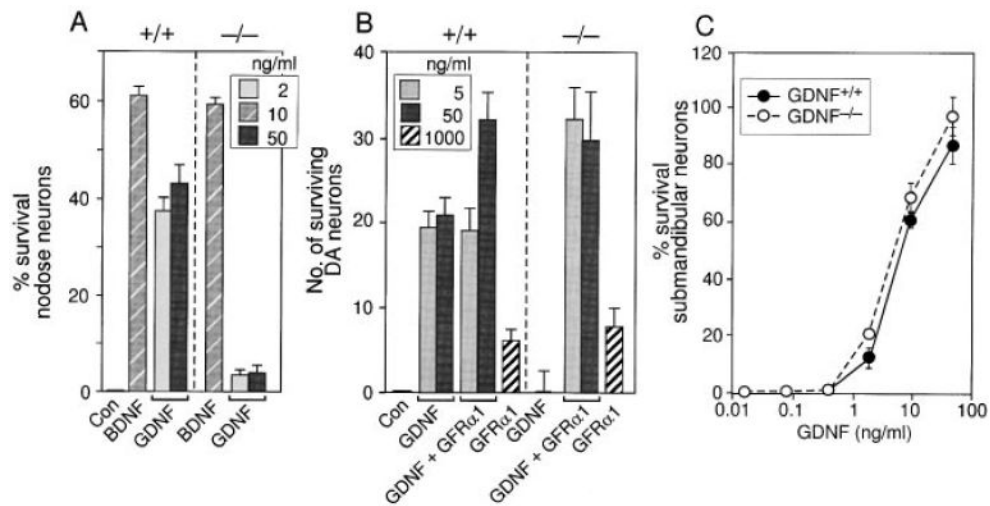


Figure 5. Survival of Primary Embryonic Neurons from Wild-Type and *GFR α 1*^{-/-} Mice in the Presence of GDNF

The response of primary embryonic wild-type (+/+) and *GFR α 1*^{-/-} (-/-) nodose (A), dopaminergic (B), and submandibular (C) neurons from 129 \times CD-1, F2 mice to GDNF. Neuronal survival is presented as percent or absolute number over control.

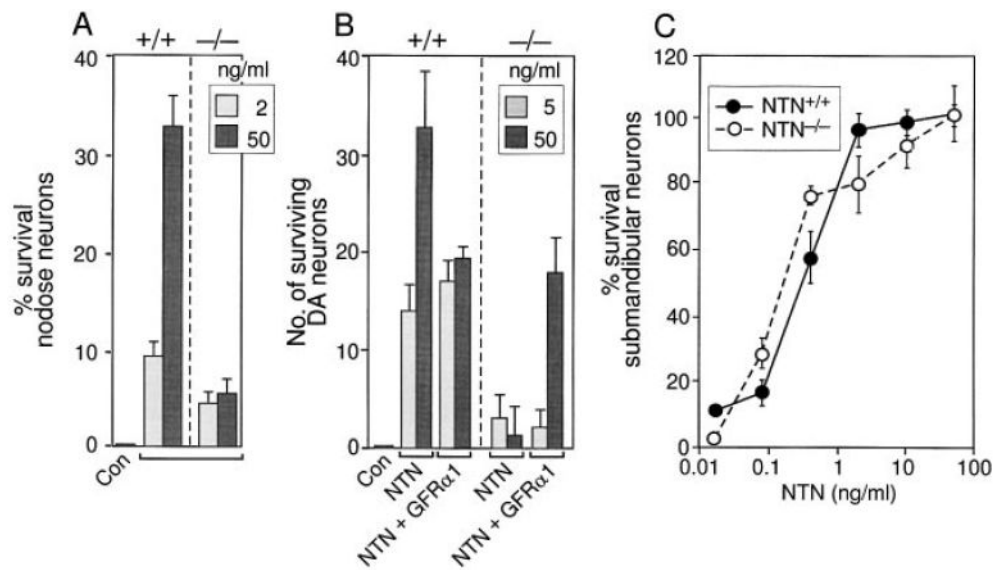


Figure 6. Survival of Primary Embryonic Neurons from Wild-Type and $GFR\alpha 1^{-/-}$ Mice in the Presence of NTN

The response of primary embryonic wild-type (+/+) and $GFR\alpha 1^{-/-}$ (-/-) nodose (A), dopaminergic (B), and submandibular (C) neurons from 129 \times CD-1, F2 mice to NTN. Neuronal survival is presented as percent or absolute number over control.

Table 1

Neuronal Counts

<u>CNS catecholaminergic neurons</u>	+/+	-/-	<i>GRFα</i> ^{-/-} % Deficit
Dopaminergic (SN)	4822 ± 336 (3)	5060 ± 405 (3)	n.s. (n.s.)
Noradrenergic (LC)	1912 ± 82 (3)	2198 ± 130 (3)	n.s. (n.s.)
<u>Sensory ganglia</u>			
Trigeminal	42700 ± 996 (3)	39886 ± 2048 (3)	n.s. (n.s.)
Vestibular	3804 ± 339 (3)	4060 ± 134 (5)	n.s. (n.s.)
Petrosal-Nodose	7884 ± 403 (3)	6714 ± 127 (5)	15 ^{**} (40%)
L5 dorsal root	7428 ± 493 (3)	7486 ± 925 (3)	n.s. (23%)
<u>Sympathetic ganglia</u>			
Superior cervical	21790 ± 616 (3)	21364 ± 938 (3)	n.s. (35%) [100%]
<u>Motor nuclei</u>			
Facial (VII)	4586 ± 251 (4)	4236 ± 148 (6)	n.s. (n.s.)
Trigeminal (V)	1047 ± 27 (4)	816 ± 20 (6)	22 ^{***} (19%)
Spinal lumbar (L1-L6)	3272 ± 92 (3)	2503 ± 320 (4)	24 [*] (22%)

Cell counts are expressed as the mean number of neurons ± SEM. The number of F2 129 × CD-1 animals that were analyzed is shown in parentheses. The percent values depicted in parentheses represent the observed deficits in the *GDNF*^{-/-} embryos. The percent values depicted in brackets represent the published deficits in the *Ret*^{-/-} embryos.

One-tailed Student's t test: n.s., not significant,

* p < 0.05,

** p < 0.01,

*** p < 0.001.