$GFR\alpha 3$ is an orphan member of the GDNF/neurturin/persephin receptor family

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Communicated by Stanley J. Korsmeyer, Washington University School of Medicine, St. Louis, MO, March 2, 1998 (received for review February 4, 1998)

ABSTRACT GDNF, neurturin, and persephin are transforming growth factor β -related neurotrophic factors known collectively as the GDNF family (GF). GDNF and neurturin signal through a multicomponent receptor complex containing a signaling component (the Ret receptor tyrosine kinase) and either of two glycosyl-phosphatidylinositol-linked binding components (GDNF family receptor α components 1 and 2, GFR α 1 or GFR α 2), whereas the receptor for persephin is unknown. Herein we describe a third member of the GF coreceptor family called GFR α 3 that is encoded by a gene located on human chromosome 5q31.2-32. GFR α 3 is not expressed in the central nervous system of the developing or adult animal but is highly expressed in several developing and adult sensory and sympathetic ganglia of the peripheral nervous system. GFR α 3 is also expressed at high levels in developing, but not adult, peripheral nerve. GFR α 3 is a glycoprotein that is glycosyl-phosphatidylinositol-linked to the cell surface like GFR α 1 and GFR α 2. Fibroblasts expressing Ret and GFR α 3 do not respond to any of the known members of the GDNF family, suggesting that GFR α 3 interacts with an unknown ligand or requires a different or additional signaling protein to function.

The GDNF family (GF) of neurotrophic factors denotes a subfamily of proteins within the transforming growth factor β superfamily that currently contains three members: glial cell line-derived neurotrophic factor (GDNF), discovered by its ability to maintain the survival of dopaminergic neurons of the embryonic ventral midbrain (1); neurturin (NTN), identified because of its survival-promoting properties on superior cervical ganglion neurons in culture (2); and persephin (PSP), discovered as the result of its homology to GDNF and NTN (3).

The sites of action of the GF ligands are broad and include neurons of the central and peripheral nervous system (CNS and PNS) and the developing kidney. In the CNS, sites of GF ligand action include dopaminergic midbrain neurons (1, 3–7), spinal and facial motor neurons (3, 8–10), Purkinje cells (11), and noradrenergic neurons of the locus ceruleus (12). Many sensory and autonomic ganglia of the PNS are also responsive to GDNF and NTN action, including neurons of dorsal root ganglion (DRG), superior cervical ganglion, trigeminal ganglion, and nodose ganglion (2, 13–15). Developing enteric neurons also respond to GDNF and NTN (R. Heuckeroth, personal communication). Finally, recent investigation has revealed that GF members also can act on ureteric bud branching *in vitro* (3, 16, 17). The intense study of GF ligand activities has revealed two generalized features: (*i*) GDNF and NTN qualitatively share activity on all responsive peripheral and central populations tested (2, 7); (*ii*) PSP shares only a subset of these activities, namely, those in the CNS (dopaminergic and motor neurons) and in the developing kidney (3).

The multicomponent GF receptor system recently characterized accounts for the overlap observed in GDNF and NTN action. The signaling component is the Ret receptor tyrosine kinase, and both GDNF and NTN can activate Ret in transfected cell lines and cultured superior cervical ganglion neurons (18, 19). Activation of Ret by GDNF or NTN requires the presence of a glycosyl-phosphatidylinositol (GPI)-linked ligand binding coreceptor. Two GF receptor α components (GFR α s) have been identified, GFR α 1 and GFR α 2, and both can mediate GDNF or NTN signaling through Ret (20-23), although there is evidence that some specificity exists, with GFR α 1 functioning preferentially as a GDNF receptor and GFR $\alpha 2$ as a preferential NTN receptor (7, 22, 23). The receptor for PSP is unknown; however, neither the Ret-GFR α 1 nor Ret–GFR α 2 receptor complexes are capable of forming functional PSP receptors. This indicates that an additional component is likely to be required for PSP signaling, either another PSP-specific coreceptor or an additional signaling component (3).

Herein we report the identification and characterization of GFR α 3, a member of the GF receptor family. We found that GFR α 3 is a GPI-linked glycoprotein like the related GFR α 1 and GFR α 2; however, GFR α 3 was not capable of mediating Ret signaling for any of the known GF ligands in transfected fibroblasts. GFR α 3 is expressed in developing and adult ganglia of the PNS but was not detected in the CNS, further emphasizing that GFR α 3 likely does not participate in PSP signaling because PSP acts only on neurons in the CNS (3).

MATERIALS AND METHODS

DNA Cloning, Sequence Analysis, Fluorescence in Situ Hybridization. Sequence analysis and cloning was performed as described (22). Mouse expressed sequence tag (EST) clones obtained from the WashU-HHMI sequencing project were sequenced, and EST AA050083 contained a full-length GFR α 3 cDNA. However, when it was compared with the sequences of other EST clones or mouse or human genomic sequence, it was found to contain a single nucleotide deletion. The mutation was corrected with PCR mutagenesis, and the

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Abbreviations: GDNF, glial cell line-derived neurotrophic factor; NTN, neurturin: PSP, persephin; GF, GDNF family; GFR α , GDNF family receptor α component; GPI, glycosyl-phosphatidylinositol; CNS, central nervous system; PNS, peripheral nervous system; DRG, dorsal root ganglion; EST, expressed sequence tag; HA, hemagglutinin; MRB, master RNA blot; E, embryonic day(s).

Data deposition: The sequences reported in this paper have been deposited in the GenBank database (accession nos. AF051766 for mouse GFR α 3 and AF051767 for human GFR α 3).

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complete cDNA was then cloned into the EcoRV site of pBluescript KS (Stratagene) and then into the HindIII and XbaI sites of pCMV-neo (24). To produce NHA-GFR α 3, the hemagglutinin (HA) epitope was inserted into the cDNA between nucleotides 99 and 100 (amino acids 33 and 34) of the mouse GFR α 3 sequence by PCR mutagenesis, cloned into pCMV-neo, and sequenced. Human and mouse GFRa3 genomic clones were obtained by screening P1 libraries with a PCR assay from the mouse and human GFR α 3 cDNA sequences (Research Genetics, Huntsville, AL). Portions of the human cDNA from 5' rapid amplification of cDNA ends (CLONTECH), genomic sequence, and PCR products from human cDNA libraries were used to obtain the sequence of human GFR α 3. A human genomic P1 clone was labeled with biotin or digoxigenin (Random Primed DNA labeling kit, Boehringer Mannheim) and used for in situ hybridization of human chromosomes derived from methotrexate-synchronized normal peripheral lymphocyte cultures. Conditions of hybridization, detection of hybridization signals, digital image acquisition, processing, and analysis, as well as the procedure for direct visualization of fluorescent signals to banded chromosomes, were performed as described (22, 25) To identify exon-intron boundaries of GFR α 2 and GFR α 3, primers from exon sequence were used to sequence restriction fragments of the mouse P1 genomic clone subcloned into pBluescript. Junctions were identified by comparison of the mouse genomic and cDNA sequences.

Expression Analysis of GFR α **3.** The human master RNA blot (MRB) was probed as described by the manufacturer's instructions (CLONTECH) using a random hexamer ³²P-labeled cDNA probe (nucleotides 266–802 of the human GFR α 3 sequence). The blot was developed by using a PhosphorImager (Molecular Dynamics), and signals were quantified by using the IMAGEQUANT software package. *In situ* hybridization analysis on fresh frozen tissue samples was performed as described (ref. 22 and J.P.G., R.H.B., P. Kotzbauer, P. Lampe, P. Osborne, J.M., and E.M.J., unpublished work). Sense and antisense ³³P-labeled RNA probes were generated from a fragment of the mouse GFR α 3 cDNA (nucleotides 107–291).

Stable Cell Line Production and Functional Assays. NIH 3T3 fibroblasts (subclone MG87) expressing Ret, Ret/GFR α 2, Ret/GFR α 3, and Ret/NHA-GFR α 3 were produced as described (19, 22). Briefly, a clonal fibroblast line stably expressing Ret was cotransfected with GFR α 2, GFR α 3, or NHA-GFR α 3 and the SV2-HisD plasmid and selected in 2 mM histidinol (Sigma). Histidinol-resistant clones were isolated and screened for expression by Western (Ret and NHA-GFR α 3) and Northern (GFR α 2, GFR α 3, and NHA-GFR α 3) blotting. For tunicamycin treatment, cells were incubated in tunicamycin at 1 or 5 mg/ml for 24 hr, washed twice with ice-cold PBS, collected in SDS sample buffer, and electrophoresed on 10% polyacrylamide gels. Gels were transferred to nitrocellulose membranes, placed in blocking solution [Trisbuffered saline (TBS)/0.1% Tween 20/5% dry milk] for 1 hr at 25°C, incubated in a 1:100 dilution of anti-HA antibody (mAb 12CA5 hybridoma supernatant) in blocking solution for 1 hr at 25°C and then washed in TBS/0.1% Tween. Blots were incubated in secondary antibody (1:10,000 anti-mousehorseradish peroxidase in blocking solution) 1 hr at 25°C, then washed as above, and developed by using SuperSignal ULTRA substrate (Pierce). For phosphatidylinositol phospholipase C treatment, a confluent 10-cm dish of cells was switched to low serum medium (DMEM/0.5% calf serum) with or without phosphatidylinositol phospholipase C at 1 unit/ml (Oxford Glycosystems, Rosedale, NY) for 1 hr at 37°C. The medium was collected, and cells were washed and collected in SDS sample buffer. Medium and cell samples were subjected to immunoblot analysis as above. For Ret phosphorylation assays, fibroblasts were treated with recombinant GDNF at 100 ng/ml, NTN at 100 ng/ml, or PSP at 100 ng/ml for 10 min, collected, immunoprecipitated with an agarose-conjugated anti-phosphotyrosine antibody (Calbiochem), and immunoblotted (anti-Ret antibody C-19, Santa Cruz Biotechnology) as described (22).

RESULTS

Identification, Sequence Analysis, and Genomic Localization of GFR α 3. The protein sequence of human GFR α 2 was used as a query to the dbEST database by using the BLAST search algorithm (27). Of the EST sequences identified, several did not correspond identically to either GFR α 1 or GFR α 2 but had significant homologies to both sequences (AA049894, AA050083, AA041935, and AA238748). These clones were acquired from the Washington University EST project and sequenced. One of these corresponded to a full-length mouse cDNA that we have called GFR α 3. Sequence information from the mouse cDNA was used to identify human genomic and cDNA clones, and the corresponding human and mouse predicted protein sequences are shown in Fig. 1A. The translations for both human and mouse GFR α 3 indicate an \approx 38.8kDa protein with a putative N-terminal signal sequence (28), three putative N-linked glycosylation sites, and a hydrophobic stretch of residues at the C terminus consistent with a GPI direction sequence (29) like those present in the related proteins GFR α 1 and GFR α 2 (7, 20–22).

GFR α 3 is approximately 34% and 36% identical to GFR α 1 and GFR α 2, respectively, and is therefore more divergent from the two, which are 48% identical to each other. Alignment of the three mouse GFR proteins illustrates the highly conserved cysteine backbone and reveals a relatively large region absent from GFR α 3 at the C-terminal region of the protein just before the GPI direction site and hydrophobic terminus (Fig. 1*B*). Homology between GFR α 1 and GFR α 2 is also low in this region (28% identity), and it is therefore the most variable region between the three family members.

We further identified and sequenced genomic clones for both GFR α 3 and GFR α 2 in mouse and then identified the exon-intron boundaries within the coding region (Fig. 2). These revealed a similar gene structure and confirmed that the region absent in GFR α 3 at the C terminus is not the result of alternative splicing in the cDNAs identified, because there is no putative exon or homologous sequence in intron 7 that corresponds to GFR α 2 exon 8. With a human genomic clone, we performed fluorescence in situ hybridization analysis to determine the chromosomal localization of $GFR\alpha 3$ (Fig. 2C). Hybridization of biotin- or digoxigenin-labeled probes revealed a symmetrical fluorescent signal on 5q31.2-32. Several human disorders are associated with this region of chromosome 5 including Treacher-Collins-Franceschetti syndrome 1, diastrophic dysplasia, progressive low-frequency deafness, and hyperekplexia, an autosomal dominant neurological disorder (30). Also, recurrent balanced and unbalanced rearrangements with breakpoints involving 5q31-32 have been reported in acute lymphoblastic and lymphocytic leukemia, chronic lymphoproliferative and myeloproliferative disorders, non-Hodgkins lymphoma, and adenocarcinoma of the stomach and intestine (31).

Expression Analysis of GFR\alpha3 in Developing and Adult Human Tissues. To obtain an extensive survey of tissues in which GFR α 3 is expressed in the developing and adult human, we probed a human MRB array of poly(A)⁺ RNA from 50 tissues by using a fragment of the human GFR α 3 sequence as a probe (Fig. 3). Surprisingly, in contrast to what has been observed for GFR α 1 and GFR α 2, essentially no expression was observed in whole brain or in any of 13 subdissected regions of the adult brain. Moderate expression of GFR α 3 was observed in spinal cord; however, it is likely that this signal

hGFRa3 mGFRa 3 MU hGERg mGER a3 hGER@3 mGFRα3 hGERa mGFRa3 hGFRa3 mGFR mGFRα3 hGER a 3 mGERa hGFR₀ mGFRa3 hGFRα3 Η Q N E N mGFRα3 Q Q N S N

В

GERa GER GFR a 1 sL1 GFRα2 GFRa3 LPLEES KRGMKKEKN GFRa2 SP VPF GFR_α1 GFRa2 GL T GFRa3 A R S GFRAISKGNNCLDAAKACNL GFRa2 A GFRa3 P D S D RRKCHKALROFFDKVPA GFRα1 кШsГ GER a 2 <u>ิ ดิติ</u> ที่ ที่ GFRA3RH LELAQLESFFEKAAESHA CRSR SYEDKE PNCI CRSR GFRa. GER a 3 GFRal DE SRS GFRa2 ASYRT KENYADC ADNYQAC GMIGF GFRα2 A S GFRα3 PM GFRA1 SVAPWC c SG GFRa3 A LS AMTTTAERI RIKNKPLGPAG - - SENEI RVEKTESLEDDLSCSTSL GFRa1 S D V T M W GFRa2 T D V N M S PKGPTFSATOA GFRa.3 СІЗДИРУСКОС GFRa2 E-----SODWADST GED. PESCGLSSLPVMVETALAALLS YSGSCRARLST-ALTALP-LLM PE----RLPILSESILPLILL AETS VSI ISPGSKKVIK GFRa2 GERa3 0 0 0

FIG. 1. Sequence analysis and alignment of GFR α 3. (*A*) Alignment of the amino acid sequences of human and mouse GFR α 3. Identical residues are boxed, the N- and C-terminal pro regions are shaded, and putative N-linked glycosylation sites are marked by dots. (*B*) Alignment of mouse GFR α family members. Identical residues are boxed, and shared cysteines are shaded. Note the divergence of GFR α 3 sequence, particularly the large deletion in the C-terminal region.

corresponds to DRGs that are not removed from the tissue samples before preparation of the RNA (see below).

Expression of GFR α 3 was observed in many nonneuronal tissues, particularly stomach and appendix. The digestive tract showed expression throughout (stomach, small intestine, and colon). GFR α 3 expression was also observed in several tissues of the urogenital system, including the kidney. Several types of glandular tissue showed low expression, and very low or no expression was observed in tissues and cells of the hematopoetic system. A comparison with fetal tissues revealed a similar pattern to adult, with no brain expression evident and a relatively low signal observed in several nonneuronal tissues.

A													
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В					Donor		INTRON		Accej	ptor			
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Exon	4	TTG	TGC	AG	GTGCCCTGCCA	GGGT.	(22 KD)	.TTCCI	TTCTT	CAG	A	TCA	CGC
		Leu	Cys	Ar							g	Ser	Arg
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		Leu	Ile	G			(400 hr	<u>.</u>			ly	Ser	Ala
Exon	6	TGC	CTC	G	GTGTGTATGCT	cccc.	(400 DP	TCTCC	TCTAC	TAG	TG	GAG	GCC
		Cys	Leu	V							al	Glu	Ala
							(195 bp)					
Exon	7	GTC	GTC		GTAGGACTTCT	GCAC.		.ATTTC	TTCCA	CAAG	AAC	AGC	AAC
		Gin	GIN								Asn	Ser	Asn



FIG. 2. Genomic analysis of GFR α 3. (A) Intron-exon junctions in the coding region of GFR α 2 and GFR α 3. Exons are represented by boxes and scaled according to size. Introns (intervening lines) are not to scale. Note that GFR α 3 entirely lacks one exon relative to GFR α 2. (B) Precise intron sites in the mouse GFR α 3 gene. Consensus splice sites are shown in boldface type. (C) Fluorescence *in situ* hybridization analysis of the human GFR α 3 gene location. Digital image of a metaphase chromosome spread derived from methotrexatesynchronized normal human peripheral leukocytes after hybridization with a digoxigenin-labeled GFR α 3 genomic probe and 4',6-diamidino-2-phenylindole counterstaining. Both copies of chromosome 5 have symmetrical rhodamine signals on sister chromatids at region 5q 31.2–32.

In summary, this initial survey of fetal and adult human tissue expression revealed that in contrast to its two known family members, $GFR\alpha 3$ does not appear to be expressed in the fetal or adult brain but is expressed in the digestive tract and urogenital system.

Expression of GFR α 3 in the Developing Mouse Embryo by in Situ Hybridization. To analyze the location of GFR α 3 expression within developing mouse tissues, we performed in situ hybridization analysis of embryonic day 14 (E14) mice (Fig. 4). From the human MRB data above, we expected low-level expression in several mouse tissues and no expression in brain. Indeed, no expression was observed in mouse brain



FIG. 3. Survey of GFR α 3 expression in human tissues. A human MRB (CLONTECH) was probed with a fragment of the human GFR α 3 cDNA. The hybridization signals are shown next to the signal intensity quantified by using a PhosphorImager. GFR α 3 was low or absent in structures of the CNS and moderate in some nonneuronal tissues, particularly those of the digestive system, urogenital system, and several glandular structures. Note that the spinal cord sample contained DRG material, which is likely the source of GFR α 3 expression (see text and Fig. 4).

at E14 in any region (Fig. 4A and data not shown). Highest expression was observed in developing peripheral nerve and ganglia. DRGs and the preaortic abdominal sympathetic ganglia (of Zuckerkandl) showed high-level expression that appeared diffusely throughout the ganglia (Fig. 4 B and C). The embryonic and adult spinal cord itself showed no staining in any region, in contrast to the results of the MRB (Fig. 4B and data not shown). As mentioned above, because DRGs are a component of the RNA preparations on the human MRB (CLONTECH), it is likely that the GFR α 3 expression observed in the human spinal cord sample was because of its presence in DRGs. High-level expression was also observed in cranial ganglia, including the trigeminal and superior glossopharyngeal ganglia (Fig. 4A) and also in the sympathetic chain ganglia and stellate ganglion (Fig. 4 B and C and data not shown). Only low-level diffuse staining was observed in nonneuronal tissues at E14. Finally, peripheral nerve showed high-level expression of GFR α 3, including trigeminal nerve, spinal nerve roots, and nerves of both the brachial and lumbosacral plexi (Fig. 4A-C and data not shown). Therefore, in situ hybridization analysis of GFR α 3 in the developing mouse indicates that it is highly expressed in peripheral nerve and ganglia, including sensory (DRG, trigeminal, and glossopharyngeal), and sympathetic ganglia (stellate, chain, and preaortic ganglia), and not expressed in the central nervous system.



FIG. 4. In situ hybridization analysis of GFR α 3 expression in mouse. (A) Saggital section of E14 mouse head. GFR α 3 is observed in trigeminal ganglion (tg) and nerve (tn) and in the glossopharyngeal ganglion (IX) but not in brain (br). (B) Transverse section of E14 mouse showing GFR α 3 expression in DRG but not in the spinal cord (sp.c). Around the abdominal aorta (aa), staining is observed in the sympathetic chain ganglia (sc) and the preaortic ganglia of Zuckerkandl (pag). (C) Saggital section of E14 mouse spinal column showing DRGs (drg), sympathetic chain (sc), and labeled nerve roots (r). (D) Dark-field photomicrograph of adult trigeminal ganglion at higher power, showing GFR α 3 is localized to neurons (arrows).

Expression of GFR α 3 at E14 was diffuse in all of these ganglia and was present in peripheral nerve, much like the expression of both GFR α 1 and GFR α 2 (22), suggesting that GFR α 3 is expressed in Schwann cells in nerve and ganglia in the absence of Ret. However, examination of the adult trigeminal nucleus and DRGs revealed punctate cellular staining that is clearly in a subpopulation of neurons when examined with bright-field microscopy (Fig. 4 D and E and data not shown). Furthermore, GFR α 3 was undetectable in adult trigeminal nerve by *in situ* hybridization (data not shown). Therefore, as demonstrated in the trigeminal system, initial diffuse GFR α 3 expression during development that is both neuronal and nonneuronal becomes restricted to neuronal expression in adulthood.

Functional Analysis of GFR α 3 as a Component of GF Signaling *in Vitro*. Because the two closely related proteins GFR α 1 and GFR α 2 are GPI-linked proteins that can both mediate GDNF or NTN signaling through Ret (22, 23), we analyzed the properties of the GFR α 3 protein and its ability to function as a GPI-linked component of a GDNF, NTN, or PSP receptor with Ret. Fibroblasts were generated that stably express both Ret and an HA-tagged form of GFR α 3 (NHA-GFR α 3) to assess the production of GFR α 3 protein. Immunoblot analysis of NHA-GFR α 3-containing cells with an anti-HA antibody revealed a doublet at approximately 47 and 51 kDa that was not observed in the control parental line (Fig. 5*A*). Because both species migrated more slowly than the analogous protein generated *in vitro* (data not shown), we examined whether GFR α 3 was glycosylated at any of the Neurobiology: Baloh et al.



FIG. 5. Analysis of NHA-GFR α 3 protein in transfected fibroblasts. (A) Anti-HA immunoblot of fibroblasts stably transfected with HAtagged GFR α 3 and Ret (Ret/GFR α 3) show a doublet at approximately 47 and 51 kDa (open arrowheads) that was not present in the parental line (Cn). Tunicamycin (Tunica) treatment of the cells for 24 hr to block N-linked glycosylation resulted in loss of the upper band at 1 mg/ml, and appearance of a lower molecular mass band. At a tunicamycin concentration of 5 mg/ml, only the lower band running at the predicted molecular mass for GFR α 3 was visible (solid arrowhead). (B) Phosphatidylinositol-specific phospholipase C (PIPLC) treatment of NHA-GFR α 3 expressing fibroblasts to specifically cleave GPI-linked proteins resulted in depletion of the NHA-GFR α 3 from the cells and induced the presence of a band in the medium corresponding to NHA-GFR α 3. Reprobing of the blot with an anti-ERK p42/44 antibody is shown below to indicate equal loading of cell lysates.

putative N-linked glycosylation sites present in the sequence (see Fig. 1*A*). Tunicamycin treatment of the cells, which blocks N-linked glycosylation, depleted both forms of the protein, and at high levels of tunicamycin, only a single band was observed at ≈ 39 kDa, the predicted molecular mass of GFR α 3. Treatment of NHA-GFR α 3-containing cells with the enzyme phosphatidylinositol phospholipase C, which specifically cleaves GPI-linked proteins from the cell surface, depleted the level of NHA-GFR α 3 from the cells and induced the presence of a similarly sized band in the cell medium (Fig. 5*B*). These results indicate that GFR α 3 is a multiply N-glycosylated protein that is GPI-linked to the cell surface like the related proteins GFR α 1 and GFR α 2.

GDNF and NTN can signal through the Ret receptor tyrosine kinase and are capable of using both either GFR α 1 or $GFR\alpha^2$ as a coreceptor (22). Recently, it was shown that PSP does not use either of these receptor complexes (3) and likely uses a different receptor complex. Therefore, we investigated whether stimulation of fibroblasts expressing both GFR α 3 and Ret with any of the three GF ligands induced activation of the Ret receptor tyrosine kinase. Although stimulation of Ret/ GFR α 2-containing fibroblasts with NTN induced strong Ret phosphorylation as expected, fibroblasts that express Ret and NHA-GFR α 3 did not respond via Ret phosphorylation to stimulation with any of the three known GF ligands (Fig. 6). Fibroblasts expressing a non-HA-tagged form of GFR α 3 and Ret also did not respond to ligand stimulation (data not shown). Therefore, although GFR α 3 is similar in structure to GFR α 1 and GFR α 2, it does not form a signaling receptor complex with Ret for any of the known ligands and may



FIG. 6. Functional analysis of the fibroblasts expressing Ret and GFR α 3. Fibroblasts that stably express Ret and GFR α 2 (Ret/GFR α 2) or Ret and NHA-GFR α 3 (Ret/GFR α 3) were left untreated (-) or stimulated with the indicated GF ligand (100 ng/ml) for 10 min, immunoprecipitated (IP) with anti-phosphotyrosine antibodies, separated by SDS/PAGE, and immunoblotted with an anti-Ret antibody. Fractions of the total lysates before immunoprecipitation are shown below to demonstrate presence of Ret (arrowheads) and equal loading (Total). As expected, stimulation of Ret/GFR α 2 containing fibroblasts with NTN shows strong Ret phosphorylation, whereas stimulation of Ret/GFR α 3 is not able to form a functional receptor with Ret for any of these GF ligands.

therefore have an as yet unknown ligand or require the presence of additional proteins to take part in GF signaling.

DISCUSSION

Herein we reported the identification and initial characterization of GFR α 3, the third member of the GF receptor family. As predicted by the sequence, GFR α 3 is GPI-linked to the cell surface like its relatives GFR α 1 and GFR α 2. However, the protein sequence of GFR α 3 is the most divergent of the GFR family and is missing a relatively large C-terminal region that contains two conserved cysteines in GFR α 1 and GFR α 2. Analysis of the GFR α 3 gene structure excludes the possibility that this deletion is the result of alternative splicing and, therefore, GFR α 3 is entirely missing one exon relative to GFR α 1 and GFR α 2. This divergence in the structure of GFR α 3 suggests a more divergent function, which is supported by its inability to function as a receptor for any of the known GF ligands with Ret.

Expression analysis of GFR α 3 also indicates a divergence from the other members of the GFR family, namely, the lack of expression observed in CNS structures. GFR α 1 and GFR α 2 are both highly expressed in several CNS structures including embryonic motor neurons and dopaminergic midbrain neurons and likely mediate the effects of GDNF and NTN on these neuronal populations (7, 21, 22). GFR α 3 is expressed highly in developing peripheral and cranial ganglia, including DRGs and trigeminal neurons, which express Ret and are responsive to GDNF and NTN (2, 13-15). Although we did not observe any GFR α 3 involvement in GF ligand signaling through Ret, GFR α 3 may play a more subtle modulatory role in GF signaling in peripheral ganglia that was not apparent in our analysis. Lastly, the inverse relationship of sites of $GFR\alpha 3$ expression (PNS only) and sites of PSP action (CNS only) strongly suggests that GFR α 3 is not involved in PSP signaling.

With the demonstration that GFR α 3 is expressed in developing peripheral nerve, all three GFR α s are developmentally expressed in the supportive cells of nerve and ganglia in the absence of Ret (21, 22, 26). Because both peripheral neurons and Schwann cells arise from neural crest precursors, the induction of Ret in a subset of coreceptor expressing neural crest precursors may yield responsiveness to GF ligands that can then function in lineage determination or as neuronal survival and/or maturation factors. The expression of GFR α s in nonneuronal cells during embryogenesis may be important in the development of peripheral nerve and ganglia or may be vestigial. However, the increased expression of GFR α 1 after nerve injury indicates it plays a functional role in nerve regeneration and suggests a similar role may occur during development (22, 26).

Our analysis of transfected fibroblasts indicates that GFR α 3 does not form a functional receptor complex with Ret for any of the known GF ligands. There are several possibilities to explain this result, all of which suggest the presence of additional receptor system components. We cannot exclude the possibility that known GF ligands interact with GFR α 3 in the presence of another Ret-like signaling protein. The existence of another Ret-like signaling molecule has also been proposed to explain the expression of GFR α 1 and GFR α 2 in several structures without Ret (refs. 22 and 26; J.P.G., R.H.B., P. Kotzbauer, P. Lampe, P. Osborne, J.M., and E.M.J., unpublished work). Alternatively, GFR α 3 may mediate Ret signaling for an as yet unknown ligand of the GDNF family. Finally, the lack of PSP interaction with any known receptor complex suggests either an additional coreceptor or signaling component is required to mediate PSP signaling. We have analyzed neuroblastoma cell lines that express all three coreceptor proteins and Ret but do not respond to PSP, indicating that combinations of known signaling components are also not sufficient for PSP signaling (M.G.T. and R.H.B., unpublished data). Furthermore, PSP does not support neuronal survival in the DRGs and superior cervical ganglia where GFR α 3 is expressed (3), providing further evidence that PSP does not interact with GFR α 3, even in biological systems where additional GFR α 3-associating signaling proteins are likely present.

In summary, we have reported the identification and initial characterization of GFR α 3. Functional and expression analyses of all known components of the GDNF ligand and receptor families suggest that additional members of both the ligand and receptor families are likely to exist.

We thank T. Fahrner, K. Simburger, and S. Audrain for their outstanding technical support and R. O. Heuckeroth for critical analysis of the manuscript. This work was supported by National Institutes of Health Grants R01 AG13729 and R01 AG13730 and Genentech. Washington University, E.M.J., and J.M. may receive income based on a license to Genentech.

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