



## Review

# p75<sup>NTR</sup> and the concept of cellular dependence: seeing how the other half die

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## Abstract

**Cells depend on specific stimuli, such as trophic factors, for survival and in the absence of such stimuli, undergo apoptosis. How do cells initiate apoptosis in response to the withdrawal of trophic factors or other dependent stimuli? Recent studies of apoptosis induction by neurotrophin withdrawal argue for a novel form of pro-apoptotic signal transduction – ‘negative signal transduction’ – in which the absence of ligand-receptor interaction induces cell death. We have found that the prototype for this form of signaling – the common neurotrophin receptor, p75<sup>NTR</sup> – creates a state of cellular dependence (or addiction) on neurotrophins, and that this effect requires an ‘addiction/dependence domain’ (ADD) in the intracytoplasmic region of p75<sup>NTR</sup>. We have recently found other receptors that include dependence domains, arguing that dependence receptors, and their associated dependence domains, may be involved in a rather general mechanism to create cellular states of dependence on trophic factors, cytokines, adhesion, electrical activity and other dependent stimuli.**

**Keywords:** apoptosis; neurotrophin; receptor; cell death

**Abbreviations:** ADD, addiction/dependence domain; ADD<sub>A</sub>, addiction/dependence domain for androgens; ADD<sub>NT</sub>, addiction/dependence domain for neurotrophins; AR, androgen receptor; BDNF, brain-derived neurotrophic factor; ChAT, choline acetyltransferase; EGFR, epidermal growth factor receptor; IRE, intrinsic receptor effect; NFκB, nuclear factor κB; NGF, nerve growth factor; NT, neurotrophin; NTR, neurotrophin receptor; PC12 cells, pheochromocytoma 12 cells; PTM, post-translational modification

## Apoptosis and negative signal transduction

Cells, especially during development, may depend on specific stimuli for their survival, such as trophic factors, cytokines, adhesion, hormones, or electrical activity. Mature cells may in many cases be more resistant to the loss of these stimuli, but often retain at least some degree of dependence on such stimuli. In the absence of the required stimulus, the dependent cells undergo apoptosis, so the cells are quite literally addicted to the stimulus or stimuli. For example, developing cerebellar granule cells require depolarization for survival, and undergo apoptosis readily in the absence of depolarization (Chang and Wang, 1997; Miller *et al*, 1997).

It has generally been assumed that apoptosis resulting from a lack of such stimuli is simply due to the lack of a positive survival signal, such as that initiated by tyrosine phosphorylation of the Trk receptors and the resultant downstream signaling. While this undoubtedly accounts for part of the effect, results obtained over the past few years argue for a distinct and complementary form of signal transduction (Rabizadeh *et al*, 1993; Bredesen and Rabizadeh, 1997; Rabizadeh *et al*, unpublished observations). Moreover, since apoptosis is an active form of cell death, the signal initiating this active program must be generated or propagated in some way as a result of the lack of binding of growth factor, cytokine, or other dependent stimulus. However, despite the biological rationale for such ‘negative signal transduction’ – i.e., signaling resulting from the absence of binding of a ligand to its receptor – this form of pro-apoptotic signal transduction has not been appreciated previously.

The recognition of this novel form of signal transduction raises a number of questions: How might an unbound receptor participate in signal transduction? How long might a receptor ‘wait’ for ligand binding before initiating or mediating a death signal? Might this form of signal transduction utilize similar pathways to those of classical signal transduction? Our work to date argues that ‘negative signal transduction’ differs from classical signal transduction in critical aspects: for example, in classical signal transduction receptors serve to bind ligand and then initiate a signal; in contrast, in negative signal transduction, molecules that serve as ‘receptors’ for the classical arm of a negative signal transduction pathway may serve as downstream mediators of the signal propagated along the negative signaling arm of a bifid pathway. Feedback loops create a two-state (or multi-state) system that tends toward one stable state over the other(s) by a combination of feedback inhibition of the alternative arm(s) and feedforward enhancement of the designate arm.

## p75<sup>NTR</sup>, the prototype addiction/dependence receptor

In 1993, we reported that p75<sup>NTR</sup>, the common neurotrophin receptor, induces apoptosis in the absence of ligand binding, with apoptosis being inhibited by ligand binding. We have reported similar results recently for a related receptor, CD40 (Ruan *et al*, 1997). This novel profile suggested that p75<sup>NTR</sup> creates a state of cellular dependence, or addiction, on neurotrophins (Figure 1). This concept has been supported by data from p75 null ('knockout') mice (Sauer *et al*, 1996; Yeo *et al*, 1997), as well as by *in vivo* antisense experiments (Cheema *et al*, 1996). Sauer *et al* (1996) found that mice deficient in NGF display atrophy and hypoplasia of medial septal cholinergic neurons, but that crossing those NGF hemizygous mice with p75<sup>NTR</sup> null mice restored cell size and number to that of the p75<sup>NTR</sup> null mice, i.e., to supranormal. They concluded that, in the presence of a reduced concentration of NGF, p75<sup>NTR</sup> mediates a reduction in cholinergic neuronal number and size in the medial septal region.

Recent results from Yeo *et al* (1997) are compatible with those of Sauer *et al* (1996): p75 null mice were found to

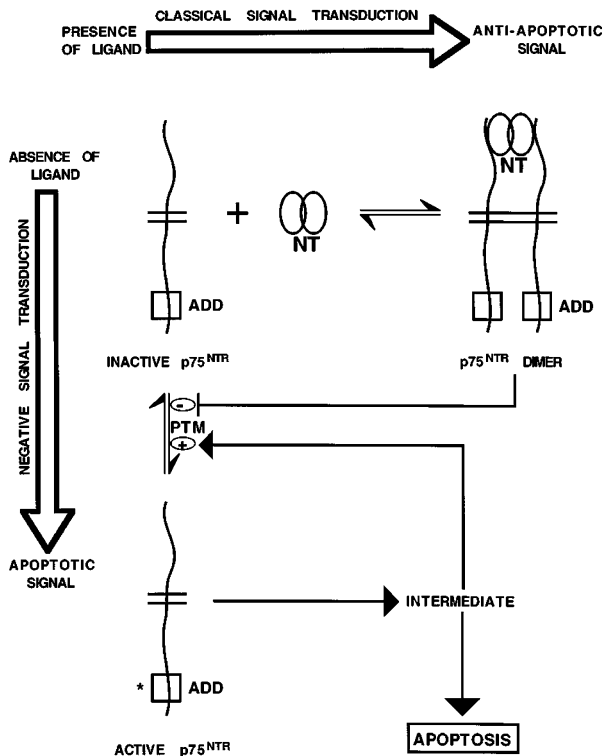
have a 50% increase in cholinergic neuronal number in the diagonal band and medial septal region, with an increase in somal volume of approximately 30%. Perry Bartlett and Graham Barrett, along with their colleagues, found that the use of antisense oligonucleotides to decrease p75 expression also blocks neuronal apoptosis in another paradigm of neurotrophin-withdrawal-induced death, that of axotomy-induced neuronal cell death (Cheema *et al*, 1996).

Not surprisingly, the dependent state created by p75<sup>NTR</sup> is relative rather than absolute. For example, the withdrawal of NGF from PC12 cells leads to apoptosis that can be blocked by downregulation of p75<sup>NTR</sup> (Rabizadeh and Bredesen, 1994; Barrett and Georgiou, 1996) or expression of bcl-2 (Mah *et al*, 1993; Martinou *et al*, 1992). Thus, the net effect is that p75<sup>NTR</sup> expression in the absence of neurotrophin alters the cellular apostat (Bredesen, 1996; Salvesen and Dixit, 1997), enhancing the likelihood of apoptosis. It is of course possible that the same effect will be associated with binding of an antagonistic ligand, rather than the lack of binding of an agonist, but, in the case of p75<sup>NTR</sup>, neurotrophin withdrawal is not currently known to be associated with the binding of a neurotrophin-antagonist ligand; nonetheless, this remains a formal possibility.

The effect of p75<sup>NTR</sup> on cellular neurotrophin dependence may underlie aspects of both neural and extraneural cellular behavior: for example, prostate epithelial cells express p75<sup>NTR</sup>, which we have argued (Bredesen, 1994) ties them to a source of neurotrophin, supplied by the stromal cells. Pflug *et al* (1992) found a progressive decrease in p75<sup>NTR</sup> expression associated with the development of prostate neoplasia, with cell lines from metastatic prostate carcinomas failing to express p75<sup>NTR</sup>. We have found that the re-expression of p75<sup>NTR</sup> in PC3 prostate carcinoma cells restores a state of neurotrophin dependence, resulting in apoptosis if NGF is not supplied (Rabizadeh *et al*, unpublished observations). Thus, p75<sup>NTR</sup> may play a role in both neural and extraneural cellular paradigms that share the common thread of neurotrophin dependence.

## An addiction/dependence domain (ADD) in p75<sup>NTR</sup>

What is the mechanism by which p75<sup>NTR</sup> expression creates a state of neurotrophin dependence? Site-directed mutagenesis studies of p75<sup>NTR</sup> have identified a region within the intracellular domain that is required for the creation of cellular neurotrophin dependence (Rabizadeh *et al*, unpublished observations). This region is referred to as an 'addiction domain' or 'dependence domain', since it is required for the induction of apoptosis by p75<sup>NTR</sup> in the absence of NT binding ('addiction' implies that the cells have been exposed to the NT prior to their requirement for it, whereas 'dependence' does not). We have used these non-equivalent terms interchangeably here because p75<sup>NTR</sup> can create a state of cellular NT dependence in cultured cells, but *in vivo*, p75<sup>NTR</sup> expression may follow neurotrophin exposure, and hence in that case the cellular state may more accurately be called an 'addiction' than a 'dependence'. Deletions that include the addiction/



**Figure 1** Creation of a neurotrophin-dependent cellular state by expression of p75<sup>NTR</sup>, the common neurotrophin receptor. The likelihood of undergoing apoptosis (setting of the cellular 'apostat') is increased by p75<sup>NTR</sup> expression, but this effect can be blocked either by binding of neurotrophins to p75<sup>NTR</sup> or by downregulation of p75<sup>NTR</sup>. In the absence of neurotrophin, p75<sup>NTR</sup> is activated, and, in a process that requires the addiction/dependence domain (ADD), triggers apoptosis. Activation of p75 triggers, through an intermediate, both further activation of p75 and apoptosis, creating a positive feedback loop. NT, neurotrophin; PTM, post-translational modification

dependence domain (ADD), and point mutations within this region, both destroy the Trk-independent intrinsic receptor effect (IRE (Bredesen and Rabizadeh, 1997)) on apoptosis. Furthermore, the ADD synthesized as a peptide is a relatively potent pro-apoptotic peptide (Hileman *et al*, 1997; Rabizadeh *et al*, unpublished observations). Thus the ADD is both necessary and sufficient for apoptosis induction. Control peptides, including a scrambled p75<sup>NTR</sup> ADD peptide, mutant p75 peptides, p75-derived peptides from outside the ADD, and control peptides matched for predicted helicity, all failed to induce apoptosis. Mutations that affect the ability of the ADD peptide to induce apoptosis have parallel effects when introduced into the full-length p75<sup>NTR</sup>, further supporting the importance of the ADD in apoptosis induction by p75<sup>NTR</sup> (Rabizadeh *et al*, unpublished observations).

In a cell-free system of neuronal apoptosis (Ellerby *et al*, 1997), the p75<sup>NTR</sup> ADD peptide induces apoptosis, as assessed by caspase activation, mitochondrial cytochrome c release, and other apoptosis parameters (Rabizadeh *et al*, unpublished observations). However, control peptides did not induce the neural cell-free system. This system should allow a determination of the organellar and molecular requirements for apoptosis induction by the p75 neurotrophin addiction domain.

Solution structure of fragments of the intracellular region of p75<sup>NTR</sup> have recently been obtained (Liepinsh *et al*, 1997; Hileman *et al*, 1997). These have disclosed predominantly  $\alpha$ -helical structure in the terminal two-thirds of the p75<sup>NTR</sup> intracellular domain. Liepinsh *et al* (1997) identified six alpha helices,  $\alpha 1 - \alpha 6$ , in this region of p75<sup>NTR</sup>. They pointed out that this region of p75<sup>NTR</sup> is reminiscent of the death domain of Fas, but that there are important structural and functional differences. Whereas Fas and TNFR I have death domains of subtype I, the putative p75<sup>NTR</sup> death domain is of subtype II. Type I death domains display the following characteristics:

1. Binding by ligand induces a pro-apoptotic signal.
2. Self association.
3. Specific arrangement of six alpha helices.

In contrast, the term 'death domain' applied to type II death domains may turn out to be a misnomer. For example, one of the subunits of NF $\kappa$ B displays a type II death domain, but the expression of NF $\kappa$ B inhibits, rather than induces, apoptosis (Van Antwerp *et al*, 1996; Beg and Baltimore, 1996). Furthermore, we have found that mutation of conserved residues in p75<sup>NTR</sup> whose analogous residues are required for apoptosis induction by Fas and TNFR I – not only point mutations but even large deletions – have no effect on apoptosis induction by p75<sup>NTR</sup> following ligand withdrawal (Rabizadeh *et al*, unpublished observations).

An additional distinction between apoptosis induction by Fas and p75<sup>NTR</sup> relates to the requirement for multimerization: for both Fas and TNFR I, multimerization is required for apoptosis induction. However, for p75<sup>NTR</sup>, recent studies utilizing FK binding protein chimeras have shown that the monomeric receptor is required for apoptosis induction, whereas multimerization blocks apoptosis induction (Wang *et al*, unpublished observations;

Rabizadeh *et al*, unpublished observations) – just the reverse of the situation for Fas and TNFR I.

Therefore, we have argued that p75<sup>NTR</sup> has a neurotrophin addiction domain (ADD; Figures 1 and 2) that is distinct from its putative death domain; whether or not it also has a functional death domain remains to be determined, but certainly the studies by Casaccia-Bonnell *et al* (1996) and Frade *et al* (1996) would argue that it does. However, Liepinsh *et al* (1997) found no tendency toward self-aggregation for the putative death domain, which is further confirmation that the putative p75<sup>NTR</sup> death domain is unlike the type I death domains of Fas and TNFR I.

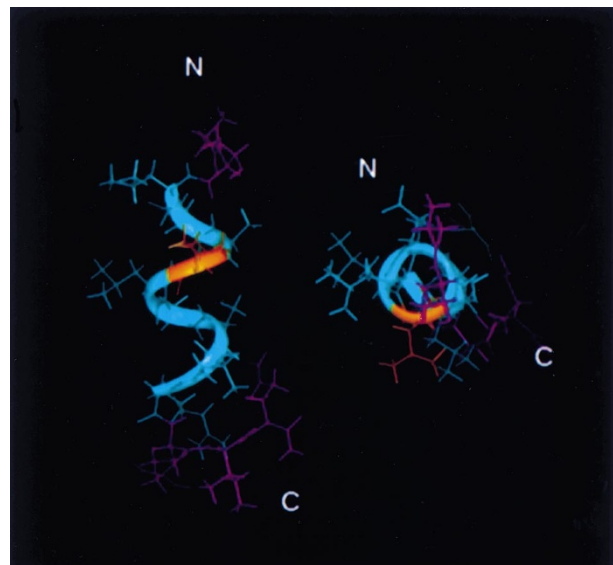
Helix formation may be crucial to the pro-apoptotic effect of p75<sup>NTR</sup>, since disruption of helix formation in the p75<sup>NTR</sup> ADD destroys the pro-apoptotic effect of p75<sup>NTR</sup>. However, additional mutants of the p75<sup>NTR</sup> ADD must be tested to establish this possibility conclusively.

In theory, it should be possible to create states of cellular dependence on other ligands by fusing the p75 ADD to the extracellular and transmembrane domains of other receptors such as the EGFR; such studies are underway.

## Other dependence receptors

p75<sup>NTR</sup> represents a prototype dependence receptor. Identification and analysis of other functionally similar dependence receptors would be advantageous, in part because this may help to shed light on the p75<sup>NTR</sup> dependence mechanism, and in part because it may provide an indication of the breadth of cellular utilization of dependence receptors.

Recently, we have found that the androgen receptor (AR) is a dependence receptor similar to p75<sup>NTR</sup> in functional effect, even though the two receptors are not



**Figure 2** Structure of the p75<sup>NTR</sup> neurotrophin addiction domain (ADD<sub>NT</sub>), based on work by Hileman *et al* (1997) and Rabizadeh *et al* (unpublished observations). The orange region highlights a mutation that, in the context of the peptide, destroys both helix formation and pro-apoptotic effect (Hileman *et al*, 1997)

structurally similar (Ellerby *et al*, unpublished observations). Expression of the AR in cells lacking endogenous AR expression induced the cells to become androgen dependent; apoptosis was induced in the absence of testosterone, but inhibited in the presence of testosterone. Furthermore, the AR has an androgen dependence domain (ADD<sub>A</sub>, to distinguish it from the p75<sup>NTR</sup> ADD<sub>NT</sub>), and, just as for p75<sup>NTR</sup>, a pro-apoptotic peptide sequence is embedded in the AR.

Studies of the AR may shed light on the mechanism by which dependence receptors create cellular states of dependence: we have found that the AR undergoes post-translational modification (PTM), without which apoptotic induction is markedly reduced. Testosterone inhibits both the PTM and apoptosis induction; furthermore, inhibition of AR PTM by mutagenesis of the AR also blocks apoptosis induction (Ellerby *et al*, unpublished observations).

We have identified the same PTM in p75<sup>NTR</sup>, suggesting that the mechanism of apoptosis induction by these two dependence receptors may, at least in principle, be similar.

### Subapoptotic dependence effects of p75<sup>NTR</sup>

The initial studies of p75<sup>NTR</sup> effects on cell death showed that cells overexpressing p75<sup>NTR</sup> in the absence of trophic factor support undergo neurite retraction and atrophy as well as apoptosis (Rabizadeh *et al*, 1993; Rabizadeh *et al*, unpublished results), but it was assumed that the atrophy and neurite retraction simply occurred as the immediate forerunners of apoptosis. Somewhat surprisingly, however, studies in p75<sup>NTR</sup> null mice have confirmed that p75<sup>NTR</sup> expression may indeed lead to neuronal atrophy and synapse loss in surviving, i.e., non-apoptotic (and apparently long-lived) cells, in addition to decreasing neuronal number (Yeo *et al*, 1997). It is tempting to speculate that such changes may represent a stable pre-apoptotic state, i.e., that the cellular biochemical changes resulting from p75<sup>NTR</sup> expression that lead, in the extreme, to apoptosis, may in some cases have subapoptotic manifestations that include neuronal atrophy and alterations in neurite outgrowth. However, it is not yet clear whether the p75 ADD<sub>NT</sub> mediates neuronal atrophy and synapse loss, or whether a different p75<sup>NTR</sup> region is required for those effects.

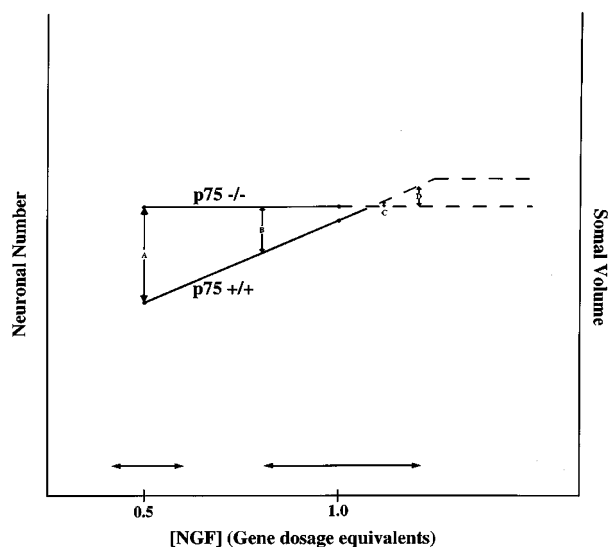
### A role for dependence receptors in input/output matching?

Studies of p75<sup>NTR</sup> *in vivo*, employing p75 null mice, have produced widely varying results. Lack of p75<sup>NTR</sup> expression may result in an increase in medial septal neuronal number (Peterson *et al*, 1997; Yeo *et al*, 1997), no increase, or a slight decrease (Peterson *et al*, 1997); it may result in an increase in striatal ChAT activity (Van der Zee *et al*, 1996) or no change in striatal ChAT activity (Yeo *et al*, 1997). In the dorsal root ganglia, lack of p75<sup>NTR</sup> may lead to fewer neurons (Lee *et al*, 1992), but may also inhibit apoptosis (Greene and Kaplan, 1995). How can such disparate results be reconciled?

Recent work (Sauer *et al*, 1996) may shed some light on these apparent discrepancies. Sauer *et al* (1996) found that

NGF hemizygotes, with a reduction in NGF concentration, demonstrate both neuronal atrophy and a decrease in medial septal cholinergic neuronal number. This pattern thus represents essentially the inverse of the findings in the p75<sup>NTR</sup> null mice. However, crossing of the NGF hemizygotes with the p75<sup>NTR</sup> null mice produced mice with medial septal cholinergic neuronal sizes and numbers indistinguishable from the p75<sup>NTR</sup> nulls, i.e., with increases in neuronal size and number. The authors concluded that, in the presence of a reduced concentration of NGF, p75<sup>NTR</sup> mediates neuronal atrophy and loss. Interestingly, their work demonstrates that, in the absence of p75<sup>NTR</sup>, medial septal neuronal size and number are *independent* of NGF concentration over the range of concentrations represented by the NGF hemizygotes and homozygotes. If one constructs an idealized graph of these results (Figure 3), it becomes apparent that p75<sup>NTR</sup> is likely to serve a role in input/output matching, in which the input is ligand and the output controls neuronal size, survival, and synaptogenesis. In other words, in the presence of p75<sup>NTR</sup>, neuronal size, and probability of undergoing apoptosis, are matched to the NGF availability; but in the absence of p75<sup>NTR</sup>, this input/output matching does not occur.

The curves shown in Figure 3 may help to explain the disparate results obtained by the different groups: in A, an increase is depicted in neuronal size and number in the p75<sup>NTR</sup> nulls due to the loss of cell number and size mediated by p75<sup>NTR</sup> in the presence of a reduced NGF concentration. This was the result reported by Sauer *et al* (1996). B depicts an increase in neuronal size and number



**Figure 3** Idealized diagram demonstrating the role of p75<sup>NTR</sup> in relating neuronal atrophy and loss to NGF concentration. In the absence of p75<sup>NTR</sup> expression, cholinergic neuronal number and size within the medial septal region are independent of NGF concentration over the range represented by the hemizygotes and homozygotes. In contrast, in p75<sup>NTR</sup> +/+ animals, these same parameters are NGF dependent. A–D represent differences between the p75 nulls and p75 +/+ animals at various NGF concentrations (discussed in the text). The NGF concentration itself may of course vary as a function of time, site, species, and other parameters, and the cellular response to NGF may vary as a function of p75<sup>NTR</sup>: TrkA ratio and other parameters

in the p75 nulls on a homozygous NGF background in which NGF concentration is still limiting. This result is similar to those reported by van der Zee *et al* (1996) and Yeo *et al* (1997). C depicts a lack of difference (insignificant decrease) for the p75<sup>NTR</sup> nulls on a homozygous NGF background, in which a higher concentration of NGF than in B is present. Such a minimal decrease in neuronal number for the p75<sup>NTR</sup> nulls was reported for TrkA<sup>+</sup> septal neurons by Peterson *et al* (1997). D depicts a decrease in neuronal size and number for the p75<sup>NTR</sup> nulls in a setting in which NGF concentration, in a homozygote, is relatively high. This result is reminiscent of that reported by Peterson *et al* (1997) for septal neurons.

### Neurotrophin-induced vs neurotrophin withdrawal-induced apoptosis: is there really a controversy?

The initial reports that p75<sup>NTR</sup> mediates neural apoptosis induced by trophic factor withdrawal appeared in 1993 and 1994 (Rabizadeh *et al*, 1993; Barrett and Bartlett, 1994; Rabizadeh and Bredesen, 1994). Some subsequent reports have described cell death induced by neurotrophins rather than by NT withdrawal (von Bartheld *et al*, 1994; Frade *et al*, 1996; Casaccia-Bonnel *et al*, 1996). Are these reports really at odds?

It is well established that NGF withdrawal may induce apoptosis in responsive neural populations (Martin *et al*, 1988). This effect is not explained by the finding that NGF may, under certain circumstances, enhance apoptosis. The argument that p75<sup>NTR</sup> is a mediator of NT withdrawal-induced apoptosis is based on the following observations:

1. Overexpression of p75<sup>NTR</sup> in neural cells enhances apoptosis induced by serum withdrawal (Rabizadeh *et al*, 1993).
2. PC12 mutants that do not express p75<sup>NTR</sup> have a reduced apoptotic response to serum withdrawal, whereas those that retain p75<sup>NTR</sup> expression retain the apoptotic response. Return of p75<sup>NTR</sup> expression to the p75-negative mutants returns their apoptotic response (Rabizadeh *et al*, 1993).
3. Cell sorting of PC12 cells into those with high and low expression of p75<sup>NTR</sup> demonstrates a correlation with high and low propensity to undergo apoptosis following serum withdrawal (Rabizadeh and Bredesen, 1994; Barrett and Georgiou, 1996).
4. Antisense inhibition of p75<sup>NTR</sup> expression is associated with inhibition of apoptosis induced by serum withdrawal, both in cell lines and primary cultures of dorsal root ganglion neurons (Barrett and Bartlett, 1994; Rabizadeh and Bredesen, 1994; Barrett and Georgiou, 1996).
5. Expression of p75<sup>NTR</sup> in NIH3T3 cells, which do not express Trk family members, enhances apoptosis following serum withdrawal. This effect is inhibited by NGF, BDNF, and NT-3, but affected only minimally by a mutant NGF (Ibanez *et al*, 1992) that binds Trk but binds p75<sup>NTR</sup> with 100-fold lower affinity than NGF binds p75<sup>NTR</sup> (Rabizadeh and Bredesen, 1994).

6. A peptide that binds p75<sup>NTR</sup> inhibits NGF withdrawal-induced death in cultures of dorsal root ganglion cells (Longo *et al*, 1997). This peptide has no effect on similar cultures from p75-null mice.
7. As noted above, mice with reduced NGF expression due to NGF hemizygosity have a reduced number and size of medial septal cholinergic neurons, but this is rescued by crossing with p75-null mice (Sauer *et al*, 1996).

Thus, there is little argument that NT withdrawal can lead to neuronal apoptosis, or that NTs generally inhibit apoptosis, and, based on the results listed above, it is highly likely that p75<sup>NTR</sup> plays a mediating role. This is not to say that other mediators may not also be discovered.

On the other hand, it is also clear that, under certain circumstances, NGF can enhance cell death by an unknown mechanism in which p75<sup>NTR</sup> participates. The assumption that p75<sup>NTR</sup> cannot mediate both NT withdrawal-induced apoptosis and, under different conditions, NGF-induced apoptosis, ignores both a large volume of data and the complexity of the systems involved. Indeed, some other members of the p75<sup>NTR</sup>/TNFR family, such as the TNF receptors, can induce or inhibit apoptosis following ligand binding (Cheng *et al*, 1994).

How might p75<sup>NTR</sup> mediate both NT withdrawal-induced apoptosis and NGF-induced apoptosis? This is currently a matter of speculation. Based on a comparison of p75<sup>NTR</sup> effects on cells that also express Trk family members with p75<sup>NTR</sup> effects on Trk-negative cells, we have previously suggested that p75<sup>NTR</sup> displays two general types of pro-apoptotic effects:

1. An intrinsic receptor effect (IRE), activated by NT withdrawal, that occurs both in the presence and absence of Trk (although Trk may inhibit this effect, so that it may be less pronounced in the presence of Trk).
2. Trk-dependent effects, which represent the sum of the following:
  - (a) High-affinity binding of NTs by the combination of p75<sup>NTR</sup> and Trk (Hempstead *et al*, 1991). This effect implies that a decrease in p75<sup>NTR</sup> will decrease high-affinity binding of NTs, enhancing apoptosis induction at limiting NT concentrations.
  - (b) Mutual repression of signaling by p75<sup>NTR</sup> and Trk, such that signaling via either alone is inhibited by the other, unless the mutual repression is relieved by binding of the appropriate NT to both p75<sup>NTR</sup> and the co-expressed Trk (Bredesen and Rabizadeh, 1996). Competition from a mismatched NT thus leads to a net pro-apoptotic effect.
  - (c) A decrease in Trk phosphorylation in the presence of p75<sup>NTR</sup> (Kaplan and Miller, 1997).

It is noteworthy that the reports of NGF-induced apoptosis have occurred in systems in which mismatched Trk members were expressed (TrkB or TrkC), or in which TrkB and TrkC expression were not evaluated. Thus, it will be of interest to determine whether or not NGF-induced apoptosis requires the expression of a mismatched Trk family member.

## Models of apoptosis induction by p75<sup>NTR</sup>

Two models for NGF-induced apoptosis have been proposed recently. Both have been proven incorrect. The first model (Kaplan and Miller, 1997) featured NGF-induced apoptosis mediated by p75<sup>NTR</sup> when the p75<sup>NTR</sup>:Trk ratio is high, including the limiting case in which it is infinite (i.e., Trk is absent). Published results have shown just the opposite: Tagliabatella *et al* (1996) found that BDNF inhibited, rather than induced, apoptosis in PC12 cells, but that this occurred only with a high p75<sup>NTR</sup>:TrkA ratio (approximately 70:1), which in the reported experiments was achieved through the use of antisense oligonucleotides. Similarly, Cortazzo *et al* (1996) found that neuroblastoma cells with a high p75<sup>NTR</sup>:Trk ratio (approximately 100:1) were inhibited from undergoing apoptosis, rather than induced to undergo apoptosis, by binding p75<sup>NTR</sup> to the exclusion of TrkA. In neuroblastoma cells lacking Trk expression, Bunone *et al* (1997) found that p75<sup>NTR</sup> induced apoptosis, and NGF inhibited apoptosis. Thus Tagliabatella *et al* (1996), Cortazzo *et al* (1996), and Bunone *et al* (1997) all reported the inverse of the model's predictions.

The second model (Dechant and Barde, 1997) proposed that signaling through Trk is anti-apoptotic, whereas signaling through p75<sup>NTR</sup> is pro-apoptotic. Again, published results clearly exclude this model. In addition to the studies of Tagliabatella *et al* (1996), Cortazzo *et al* (1996), and Bunone *et al* (1997) quoted above, Longo *et al* (1997) developed a peptide that binds p75<sup>NTR</sup> but not Trk. This peptide was shown to block the apoptosis of dorsal root ganglion neurons in primary culture, demonstrating that binding to p75<sup>NTR</sup> may initiate an anti-apoptotic signal. Interestingly, the peptide was only effective as a cyclic dimer, with the monomer being inactive; this finding argues that p75<sup>NTR</sup> dimerization is crucial to the initiation of an anti-apoptotic signal.

The FK binding protein results quoted above argue that, at least in some paradigms, monomeric p75<sup>NTR</sup> is required for apoptosis induction, and that dimerization blocks the pro-apoptotic effect. However, whether or not this will be the case in neurons *in vivo* remains to be seen.

## Conclusions

p75<sup>NTR</sup>, the common neurotrophin receptor, is a prototypic dependence receptor. The characteristics of receptors with this newly recognized effect include the following:

1. Expression in the absence of the dependent ligand shifts the cellular apoptat toward a greater probability of apoptosis.
2. Ligand binding blocks the pro-apoptotic effect.
3. These receptors include novel domains referred to as addiction domains or dependence domains (ADDs), which are required for apoptosis induction by the receptors in the absence of ligand. These ADDs are distinct from death domains.
4. Peptides representing ADDs are pro-apoptotic.

p75<sup>NTR</sup> may also mediate subapoptotic events such as somal size specification and synaptogenesis (the latter perhaps via an effect on neurite retraction).

In some cases, NGF may induce, rather than inhibit, apoptosis. Whether this effect will turn out to require the coexpression of a mismatched Trk (TrkB or TrkC) with p75<sup>NTR</sup> remains to be determined.

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