P53 Is Essential for Developmental Neuron Death as Regulated by the TrkA and p75 Neurotrophin Receptors

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Abstract. Naturally occurring sympathetic neuron death is the result of two apoptotic signaling events: one normally suppressed by NGF/TrkA survival signals, and a second activated by the p75 neurotrophin receptor. Here we demonstrate that the p53 tumor suppressor protein, likely as induced by the MEKK-JNK pathway, is an essential component of both of these apoptotic signaling cascades. In cultured neonatal sympathetic neurons, p53 protein levels are elevated in response to both NGF withdrawal and p75NTR activation. NGF withdrawal also results in elevation of a known p53 target, the apoptotic protein Bax. Functional ablation of p53 using the adenovirus E1B55K protein inhibits neuronal apoptosis as induced by either NGF withdrawal or p75 activation. Direct stimulation of the MEKK-JNK pathway using activated MEKK1 has similar effects; p53 and Bax are increased and the subsequent neuronal apoptosis can be rescued by E1B55K. Expression of p53 in sympathetic neurons indicates that p53 functions downstream of JNK and upstream of Bax. Finally, when p53 levels are reduced or absent in p53+/− or p53−/− mice, naturally occurring sympathetic neuron death is inhibited. Thus, p53 is an essential common component of two receptor-mediated signal transduction cascades that converge on the MEKK-JNK pathway to regulate the developmental death of sympathetic neurons.

Key words: neuronal apoptosis • MEKK • nerve growth factor • brain-derived neurotrophic factor • sympathetic neurons

Naturally occurring neuronal death is an essential component of neural development in which <50% of any given neuronal population is lost as a mechanism for establishing appropriate neuron:target cell connections (reviewed in Oppenheim, 1991). Sympathetic neurons of the peripheral nervous system, which require NGF for their survival (reviewed in Levi-Montalcini, 1987), have been a prototype for analysis of the mechanisms regulating this event. During the first two postnatal weeks, sympathetic neurons compete for limiting amounts of target-derived NGF, which binds to and activates TrkA tyrosine kinase receptors (Kaplan et al., 1991a,b; Klein et al., 1991) on neuronal terminals, thereby mediating a retrograde neuronal survival signal (Campenot et al., 1982; Riccio et al., 1997; Senger et al., 1997). Traditionally, it has been thought that the absence of such a TrkA retrograde signal was sufficient to cause rapid neuronal apoptosis during developmental death. However, we have recently shown that the lack of such a NGF/TrkA-mediated survival signal is, of itself, insufficient for appropriate sympathetic neuron death. Instead, a second neurotrophin receptor, p75NTR (Johnson et al., 1986; Radeke et al., 1987), is required for this process; in the absence of p75NTR, sympathetic neuron apoptosis is delayed both in culture and in vivo (Bamji et al., 1998). Moreover, ligand-mediated activation of p75NTR is sufficient to cause sympathetic neuron apoptosis (Bamji et al., 1998). Thus, appropriate developmental sympathetic neuron death occurs when p75NTR is activated coincident with suboptimal NGF/TrkA survival signals (reviewed in Miller and Kaplan, 1998). This convergent regulation of neuronal apoptosis provides a mechanism whereby sympathetic neurons are able to recognize not only whether they are receiving adequate amounts of NGF, but also whether or not they are exposed to “inappropriate” neurotrophins, potentially as a function of late or inappropriate target innervation.

The intracellular mechanisms responsible for transducing these receptor-mediated apoptotic cascades in sympathetic neurons remain ill-defined, although activation of
the JNK pathway occurs after both p75NTR activation (Casaccia-Bonnefil et al., 1996; Bamji et al., 1998) and NGF withdrawal (Estus et al., 1994; Ham et al., 1995) and, in the case of NGF withdrawal, is a necessary early event in the apoptotic pathway. Moreover, Bax is essential for sympathetic neuron apoptosis both in culture and during naturally occurring death in vivo (Deckwerth et al., 1996; Easton et al., 1997). One protein that is known to regulate transcription of Bax (Miyashita and Reed, 1995) and that has been implicated in cellular apoptosis is the p53 tumor suppressor protein (reviewed in Hainaut, 1995; Jacks and Weinberg, 1996; Levine, 1997). However, although p53 has been implicated in neuronal death in response to DNA damage or cellular stress (Li et al., 1994; Sakhi et al., 1994; Wood and Youle, 1995; Morrison et al., 1996; Xiang et al., 1998), this protein has not previously been thought to play a role in naturally occurring cell death nor has it been shown to act downstream of death receptor activation or of the MEKK-JNK pathway.

In this regard, we have previously demonstrated that increased expression of p53 was sufficient to cause sympathetic neuron apoptosis in the presence of NGF (Slack et al., 1996). On the basis of this observation, together with the fact that Bax is essential for naturally occurring sympathetic neuron death (Deckwerth et al., 1996; Easton et al., 1997), we hypothesized that p53 was an essential component of the signaling pathways causing sympathetic neuron apoptosis either in response to NGF withdrawal and/or to p75NTR activation. In this paper, we test this hypothesis, and demonstrate that both p75NTR activation and NGF withdrawal cause increased expression of p53, likely as a consequence of activation of the MEKK-JNK (Derijard et al., 1994; Yan et al., 1994) pathway, and that this convergent regulation of p53 is essential for normal naturally occurring sympathetic neuron death.

Materials and Methods

Primary Neuronal Cultures

Mass cultures of pure sympathetic neurons from the superior cervical ganglion (SCG) of postnatal day 1 rats were prepared as previously described (Bellido et al., 1997; Bamji et al., 1998). Neurons were plated on rat tail collagen coated tissue culture dishes; 6-well plates for biochemistry, and 48-well plates for survival assays. Low density SCG cultures were used for all of the survival assays; for the survival assays in NGF withdrawal and p75 activation experiments, neurons were used at densities of 4,000–6,000 and 2,000–4,000 neurons per well of a 48-well plate, respectively. For biochemistry, ~40,000–50,000 neurons were plated per well of a 6-well dish. For all experiments not involving viral infection, neurons were initially cultured for 5 d in the presence of 50 ng/ml NGF. At the end of this 5 day selection, neurons were washed several times in neurotrophin-free media before addition of the new neurotrophin- or KCl-containing media.

NGF for these experiments was purified from mouse salivary glands (Cedarlane, Hornby, ON). Two sources of recombinant human BDNF (Amgen, Thousand Oaks, CA; Preprotech, Rocky Hill, NJ) were used for these experiments; similar results were obtained with both, as discussed previously (Bamji et al., 1998).

Virus Infections and Survival Assays

Six different recombinant adenoviruses were used for these experiments.

Two, those expressing E1B55K, the E1B55K mutant A262 (Yew et al., 1998), and p53 (Slack et al., 1996) have been previously described and described. We also constructed adenoviruses (Massie et al., 1995) expressing activated MEK1 (Eilers et al., 1998), and Bel-xl (Boisise et al., 1993) in the Ad5 backbone (Bett et al., 1994), which drives expression from the CMV promoter. As a control for these viruses, we used a recombinant adenovirus in the same Ad5 backbone expressing E. coli β-galactosidase, as we have previously described (Slack et al., 1996; gift of Dr. Frank Graham, McMaster University, Hamilton, Ontario).

All recombinant adenoviruses were purified on CsCl gradients, as we have previously described (Slack et al., 1996). Infectious titer was determined by plaque assay on 293 cells (Graham and Prevec, 1991).

For the experiments involving E1B55 rescue from NGF withdrawal, neurons were plated for 3–4 d in 50 ng/ml NGF, and then were infected overnight with various modified versions of recombinant adenovirus in serum-free media containing 50 ng/ml NGF. The next day, the virus was removed, and the cells were fed with fresh media containing 10 ng/ml NGF. The following day, cells were washed free of NGF with 3-h-long washes in serum-free, NGF-free media, and then were switched to media with or without 10 ng/ml NGF. 2 d later, neuronal survival was assessed. For the p75 activation experiments, the protocol was similar with the exception that, after the washes to remove NGF, neurons were switched to media containing 50 mM KCl with or without 100 ng/ml BDNF for 2 d.

For the experiments with the activated MEK1 or p53 adenoviruses, neurons were plated at a density of 10,000–12,000 neurons/well in 48-well plates (Falcon Plastic, Cockeysville, MD) with 20 ng/ml NGF. On the fifth day of culture, medium was removed, the virus was added in a volume of 200 ul DMEM media (GIBCO-BRL, Gaithersburg, MD) containing 2 mM glutamine, 100 U/ml penicillin, 100 µg/ml streptomycin (all from BioWhittaker, Walkersville, MD), and 10% FBS. After 16–18 h of infection, the virus-containing media was removed and was replaced with UltraTacture media containing 2 mM glutamine, 100 U/ml penicillin, 100 µg/ml streptomycin, and 20 ng/ml NGF. Cultures were maintained for 48 h, washed four times for one hour each with growth factor-free media, and then switched into the same media containing various concentrations of NGF.

Survival assays were performed as previously described (Slack et al., 1996; Bamji et al., 1998) using nonradioactive cell proliferation (MTT) assays (CellTitre 96; Promega Corp., Madison, WI). 50 µl of the MTT reagent was added to each well and left for 90 min followed by the addition of 100 µl of solubilization solution to lyse the cells. Each condition was repeated in triplicate. In each assay, baseline (0% survival) was considered to be 0 ng/ml NGF, and 10 ng/ml NGF was considered to be 100% survival, unless stated otherwise. All other conditions were related to these values.

Western Blot Analysis

For biochemistry, sympathetic neurons were lysed in TBS lysis buffer (Knuel et al., 1994) containing 137 mM NaCl, 20 mM Tris (pH 8.0), 1% (vol/vol) NP-40, 10% (vol/vol) glycerol, 1 mM PMSF, 10 µg/ml aprotinin, 0.02 µg/ml leupeptin, 1.5 mM sodium vanadate, and 0.1% SDS. Cells were collected in cold PBS by gentle scraping to detach them from the collagen substratum, were washed three times with the same buffer, and then were resuspended in 50 to 100 µl of lysis buffer, followed by rocking for 10 min at 4°C. After a 10 minute centrifugation, the lysates were normalized for protein concentration using a BCA Protein Assay Reagent (Pierce Chemical Co., Rockford, IL). Equal amounts of protein (10–50 µg) were then boiled in sample buffer for 5 minutes and separated by SDS-PAGE. After electrophoresis, proteins were transferred to 0.2 µm nitrocellulose for 1.5 h at 0.6 Amps, and the membrane was washed three times with TBS. For all antibodies, the membranes were blocked in 3% nonfat milk in TBST (blotto) for 1.5 h at room temperature. The membranes were then incubated overnight at 4°C with the primary antibodies in blotto: anti-p53 (Oncogene Neurosciences, Manhasset, NY), anti-p53 (Novacstra, Burlingame, CA), anti-p21 (Transduction Laboratories, Lexington, KY), anti-p27 (Transduction Laboratories), anti-Bad (Transduction Laboratories), anti-Bcl-xl (Santa Cruz Biotechnology, Santa Cruz, CA), anti-Bax (Santa Cruz Biotechnology), anti-E1B55K antibody 2A6, anti-human c-myc (PharMingen, San Diego, CA), anti-phospho-p185 (New England Biolabs, Boston, MA; Derijard et al., 1994), anti-phosphoshex473 (New England Biolabs), anti-phospho183 (Protegma Corp.), anti-TrkA (RTA; gift of Dr. L. Reichardt, University of California, San Francisco, CA; Wescamp and Reichardt, 1991), anti-tyrosine hydroxylase (Chemicon Laboratories, Temecula, CA), or anti-tubulin (Oncogene Neurosciences). After incubation with the primary antibodies, membranes were washed four times with...
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Fixed in 4% paraformaldehyde in phosphate buffer (PB) for 1 h to over-
amplify the mutant allele, PCR was conducted for 35 cycles (94
C for 30 s, 55°C for 60 s, 72°C for 90 s) using the following oligonucleotide GTGG-
AGGGGACAAAAGTTCGAGGCC for the 5' primer, and TTTACCG-
GAGCCCTGGCGCTCAGATGT for the 3’ end. This results in a 200 nu-
cleotide fragment in mutant mice and no product in wildtype mice. To
amplify the wild-type allele, the same oligonucleotide was used as the 5'
primer and ATGGGAGGCGTGCCAGTCTAACC was used as the 3’ primer. This results in amplification of a 600 nucleotide fragment in het-
rocyste and wild-type mice, and no product in the p53–/– mice.

For morphometric analysis, the SCG were removed and immersion-
fixed in 4% paraformaldehyde in phosphate buffer (PB) for 1 h to over-
night at 4°C. Ganglia were cryoprotected in graded sucrose solutions, 7-μm-thick sections were serially cut on a cryostat, and every section was
collected and thaw-mounted onto chrom-alum subbed slides. Slides were
stained with cresyl violet and morphometric analyses were performed using
a computer-based image analysis system (Biocor, Paris, France).

Analysis of p53–/– Mice
Mice heterozygous for a targeted mutation in the p53 gene (Donehower et
al., 1992) were obtained from GenPharm International (Mountainview,
CA). p53–/– mice were maintained in a C57Bl/6 background, as homozy-
gotes or heterozygotes. Progeny from p53–/– × p53+/+ or from p53 het-
erozygote crosses were screened for the mutant allele(s) using PCR. To
amplify the mutant allele, PCR was conducted for 35 cycles (94°C for 30 s,
55°C for 60 s, 72°C for 90 s) using the following oligonucleotide GTGG-
AGGGGACAAAAGTTCGAGGCC for the 5' end, and TTTACCG-
GAGCCCTGGCGCTCAGATGT for the 3’ end. This results in a 200 nu-
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collected and thaw-mounted onto chrom-alum subbed slides. Slides were
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a computer-based image analysis system (Biocor, Paris, France).

Neuronal numbers were determined by counting all neuronal profiles with
nucleoli on every third section, as per Coggeshall (1984). This sampling
frequency (every 21 μm) ensures that neurons are not double counted,
since the average neuronal diameter does not exceed 21 μm in any of the
groups examined (Bamji et al., 1998). This method does not correct for
split nucleoli. Statistical results were expressed as the mean ± SE of the
mean and were tested for significance by a one-tailed Student’s t test.

For TUNEL analysis, SCGs were dissected from p53+/+ and p53+/+ littermates at postnatal day 7. Ganglia were fixed for 30 min in 4%
paraformaldehyde, cryoprotected in graded sucrose solutions, and 7-μm-thick sections cut on a cryostat. Every third section was collected and thaw-mounted onto chrom-alum subbed slides. Slides were
stained with cresyl violet and morphometric analyses were performed using
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immediately performed using the Boehringer-Mannheim in situ cell death
kit. The number of TUNEL-positive cells on every third section was
counted and these numbers were
determined by fluorescence microscopy, and these numbers were
tested for significance by a one-tailed Student’s t test.

Results

p53 and Bax Are Elevated after NGF Withdrawal from
Sympathetic Neurons
Previously, we have demonstrated that an increase in p53 levels is sufficient to cause sympathetic neuron apoptosis in the presence of NGF (Slack et al., 1996). To determine whether endogenous p53 protein levels were ever similarly
increased during sympathetic neuron apoptosis, we cult-
tured sympathetic neurons from neonatal animals, a time
when developmental death of these neurons is ongoing.
Initially, we examined p53 during sympathetic neuron apoptosis induced by NGF withdrawal; this apoptosis is relatively slow, taking ~48 h, and is transcription-depen-
dent (Deckwerth and Johnson, 1993; reviewed in Johnson
and Deckwerth, 1993). Neurons were cultured for 5 d in the
presence of 50 ng/ml NGF, NGF was withdrawn, and the
then the cellular levels of p53 were quantitated using
Western blots with anti-p53 at various time-point post-
withdrawal (Fig. 1 A). This analysis revealed that p53 lev-
els were elevated approximately threefold by 16 h after
NGF withdrawal (Fig. 1 A). Elevation of p53 protein was
first observed at 12 h (data not shown), and was main-
tained until at least 36 h post NGF-withdrawal (Fig. 1 A).
Interestingly, this timecourse corresponds to the commit-
ment point, after which NGF-withdrawn sympathetic neu-
rons cannot be rescued from apoptotic death (Johnson
and Deckwerth, 1993). Thus, NGF withdrawal leads to an
increase in p53 levels that correlates with the timecourse
of commitment to an apoptotic death.

To determine whether this increase in p53 had func-
tional consequences, we examined the expression of a
well-characterized p53 transcriptional target, the cyclin-
dependent kinase p21 (El-Diery et al., 1993). Western blot
analysis of equal amounts of protein derived from sympathetic neurons either maintained in 10 ng/ml NGF (NGF), or at various time-points from 4 to 48 h after
withdrawal of NGF (−NGF). Western blots were probed with antibodies specific to p53 (A), p21 (B), p27 (C), Bad (D), Bel-2 (E), Bax (F), or Bcl-xl (G). Note that p53, p21, p27, Bad, and Bax
all increase after NGF withdrawal, whereas Bel-2 and Bcl-xl
decrease. (H) Western blot analysis of equal amounts of protein derived from sympathetic neurons either maintained in 20 ng/ml
NGF, or withdrawn from NGF for 12 h (0 NGF). Western blots were probed with antibodies specific to phosphorylated forms of
Akt (P Akt) or ERKS (P Erks). Note that phosphorylation of
these TrkA targets was significantly decreased after NGF withdrawal.

Figure 1. p53 and its transcriptional targets, p21 and Bax, are in-
creased during NGF-withdrawal induced apoptosis of sympa-
thetic neurons. (A–G) Western blot analysis of equal amounts of protein derived from sympathetic neurons either maintained in 20 ng/ml NGF (NGF), or at various time-points from 4 to 48 h after
withdrawal of NGF (−NGF). Western blots were probed with antibodies specific to p53 (A), p21 (B), p27 (C), Bad (D), Bel-2 (E), Bax (F), or Bcl-xl (G). Note that p53, p21, p27, Bad, and Bax
all increase after NGF withdrawal, whereas Bel-2 and Bcl-xl
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NGF, or withdrawn from NGF for 12 h (0 NGF). Western blots were probed with antibodies specific to phosphorylated forms of
Akt (P Akt) or ERKS (P Erks). Note that phosphorylation of
these TrkA targets was significantly decreased after NGF withdrawal.
We next investigated the levels of Bax, another transcriptional target of p53 (Miyaşhita and Reed, 1995) that is known to be essential for naturally occurring sympathetic neuron death (Deckwerth et al., 1996; Easton et al., 1997). Western blot analysis revealed that, like p53, Bax levels increased approximately twofold after NGF withdrawal (Fig. 1 F). This increase was first observed at 16 h, and levels were maximal by 24 h (Fig. 1 F). Interestingly, Western blot analysis revealed that a second proapoptotic member of this family, Bad (Yang et al., 1995), was also increased threefold after NGF withdrawal (Fig. 1 D). Like Bax, this increase in Bad commenced at ~16 h after withdrawal, the commitment point for apoptosis. In contrast to Bad and p21, since levels of p27, a second cyclin-dependent kinase (Toyoshima and Hunter, 1994), were also increased approximately twofold at 24 and 36 h after NGF withdrawal (Fig. 1 C).

values that are significantly different from 50 mM KCl alone, and (***) indicates those values amongst the points that received BDNF that are significantly different from KCL+BDNF (P < 0.005). Note that, as previously reported (Bamji et al., 1998), BDNF reduces neuronal survival in the presence of KCl, and that E1B55K but not A262 is able to completely rescue this BDNF-mediated apoptosis. (C) MEKK1-induced apoptosis. Sympathetic neurons were infected with recombinant adenovirus expressing activated MEKK1 or β-galactosidase (LacZ) at concentrations of 20 to 200 moi, and were maintained in 20 ng/ml NGF for 4 d after infection. As controls, uninfected neurons were maintained for the entire experiment in 20 ng/ml NGF. Cell survival was measured using MTT assays, and results are those obtained in a representative experiment performed in triplicate, and represent the mean ± standard error. Similar results were obtained in 4 separate experiments. **Indicates those values that are significantly different between MEKK1 and the β-galactosidase control at a given moi (P < 0.005). Note that activated MEKK1 decreases sympathetic neuron survival in a concentration-dependent fashion. (D) p53 and MEKK1-induced apoptosis. Sympathetic neurons were infected with 50 moi of MEKK1 plus 500 moi of E1B55K (MEKK + E1B55K), A262 (MEKK + A262) or β-galactosidase (MEKK + LacZ), were maintained in 20 ng/ml NGF for 4 d after infection, and survival was then measured using MTT assays. As further controls, neurons were infected with 50 moi β-galactosidase virus (20 NGF + 50 LacZ) for the same time period, or were withdrawn from NGF for the final two days. Results are those obtained in a representative experiment performed in triplicate, and represent the mean ± standard error. Results are normalized so that 0 NGF in 0% survival, and 20 ng/ml NGF plus 50 moi β-galactosidase (20 NGF + 50 LacZ) is 100% survival. Similar results were obtained in three separate experiments. ** Indicates that E1B55K significantly rescues MEKK1-induced killing of sympathetic neurons (P < 0.005). (E) p53 in JNK-Bax cell death pathway. Sympathetic neurons were infected with 50 moi of the activated MEKK1 adenovirus (50 MEKK) plus or minus 50 moi Bcl-xl (MEKK + Bcl-xl) or 500 moi E1B55K (MEKK + E1B55K) or 500 moi A262 (MEKK + A262) adenoviruses. As a control, neurons were infected with 50 moi β-galactosidase adenovirus (50 LacZ). 4 d after the initial infection, during which time neurons were maintained in 10 ng/ml NGF, survival was measured using MTT assays. Results are those obtained in a representative experiment performed in triplicate, and represent the mean ± standard error. Results are normalized so that 0 NGF is 0%, and 10 ng/ml NGF plus 50 moi β-galactosidase adenovirus is 100% survival. ** Indicates those values that are significantly different from MEKK alone (P < 0.005).
Elevated p53 is Necessary for Sympathetic Neuron Apoptosis in Response to NGF Withdrawal

To determine whether this increase in p53 protein levels was an essential component of the apoptotic cascade that follows NGF withdrawal, we took advantage of the adeno-
viral E1B55K protein, which functionally ablates p53 (Yew et al., 1992). Specifically, neonatal sympathetic neurons were cultured for 3−4 d in 50 ng/ml NGF, and then were infected with one of two recombinant, replication-defective adenoviruses; one of these adenoviruses expressed E1B55K while the second expressed a mutant defective adenovirus expressing E1B55K, which functionally ablates p53 (Yew et al., 1990). As a further con-
trast, p53 protein levels were greatly reduced (Fig. 1 H). Western blot analysis with an antibody specific to the phosphorylated form of the serine/threonine kinase Akt revealed that, 12 h after NGF withdrawal, phosphorylation of Akt was greatly re-
duced (Fig. 1 H). Western blot analysis with an antibody specific to the phosphorylated form of the ERKs revealed a similar decrease in the phosphorylation of these proteins after NGF withdrawal (Fig. 1 H). Thus, NGF withdrawal leads to a decrease in signaling via these TrkA-regulated prosurvival pathways coincident with induction of potentially proapoptotic pathways.

Figure 3. E1B55K reduces p53 levels in sympathetic neurons withdrawn from NGF. (A) Western blot analysis for p53 in sympathetic neurons that were withdrawn from NGF for 36 h (0 NGF), that were maintained in 10 ng/ml NGF (10 NGF), or that were infected with 500 moi of recombinant adenovirus expressing A262 or E1B55K, and 2 d later were withdrawn from NGF for 24 h. Equivalent amounts of protein from lysed neurons were electrophoresed, transferred to nitrocellulose filters, and p53 protein levels were assessed using anti-p53. Note that p53 levels increase after NGF withdrawal, and that expression of E1B55K reverses this increase, while expression of A262 has no effect. (B) The same blot as in panel (A) reprobed for the cytoskeletal protein tubulin. (C) Western blot analysis for E1B55K using anti-
E1B55K in equal amounts of protein derived from sympathetic neurons infected for 36 h with adenoviruses expressing E1B55K or the E1B55K mutant A262. The antibody used for these studies recognizes both the wild-type and mutant proteins. (D) Western blot analysis for p53 in sympathetic neurons that were maintained in 20 ng/ml NGF, and that were infected with 50 moi of recombin-
ant adenovirus expressing activated MEKK1 (MEKK) with or without 200 moi of E1B55K for 36 h. Equivalent amounts of protein from lysed neurons were electrophoresed, transferred to nitrocellulose filters, and p53 protein levels were assessed using anti-p53. (E) The same blot as in D reprobed for TrkA.
Figure 4. p53 levels increase during apoptosis of sympathetic neurons as induced by BDNF-mediated activation of p75NTR. (A) Western blot analysis for p53 in equal amounts of protein derived from sympathetic neurons that were cultured in 50 ng/ml NGF for 4 d, and then were washed free of NGF and switched into 50 mM KCl (KCL), 50 mM KCl plus 100 ng/ml BDNF (KCL + BDNF), or media containing no NGF or KCl (0 NGF) for 24 or 36 h. Note that p53 levels in the neurons treated with KCl and BDNF are similar to those in 0 NGF, and are greater than those maintained in KCl alone. (B) The same blot as in A reprobed for the neurotransmitter enzyme, tyrosine hydroxylase (TH) to demonstrate that equal amounts of protein were present in each of the lanes.

p53 is Downstream of p75NTR and is Essential for p75NTR-mediated Neuronal Apoptosis

We have previously demonstrated that the immediate early protein, c-jun, is hyperphosphorylated in sympathetic neurons after p75NTR activation (Bamji et al., 1998) as it is after NGF withdrawal (Estus et al., 1994; Ham et al., 1995). To determine whether p53 elevation was also a common downstream component of these two apoptotic signaling cascades, we examined p53 levels in sympathetic neurons in conditions where p75NTR activation leads to apoptosis. Specifically, for these experiments, sympathetic neurons were cultured for 5 d in 50 ng/ml NGF and then were switched to KCl, which maintains sympathetic neuron survival in the absence of TrkA activation (Franklin et al., 1995). We then added the neurotrophin BDNF, which selectively binds p75NTR but not Trk receptors on sympathetic neurons (Bamji et al., 1998), and determined cellular p53 levels (Fig. 4A). As seen with NGF withdrawal, p53 levels were increased at 24 and 36 h in neurons maintained in KCl plus BDNF relative to those in KCl alone. Reprobing of the same blot with an antibody to tyrosine hydroxylase revealed that equal amounts of protein were present in all of the samples (Fig. 4B). This increase was already apparent at 12 h, a timepoint when p53 levels were only first starting to increase after NGF withdrawal (data not shown). Thus, p53 elevation is downstream of both p75NTR activation and NGF withdrawal.

To determine whether this increased p53 was essential for p75NTR-mediated apoptosis, we performed rescue experiments using the E1B55K adenovirus. Specifically, neurons were cultured in 50 ng/ml NGF for 3–4 d, were infected overnight with recombinant adenovirus, and 2 d later were switched to media containing 50 mM KCl plus or minus 100 ng/ml BDNF. Neuronal survival was then measured after 48 h using MTT assays (Fig. 2B). As we have previously reported (Bamji et al., 1998), the addition of BDNF to 50 mM KCl caused a decrease in sympathetic neuron survival of ~54% (Fig. 2B). Expression of the mutant E1B55K had no significant effect on this decrease (P = 0.11). In contrast, expression of E1B55K rescued 100% of the p75NTR-driven apoptosis (P = 0.012; Fig. 2B), a rescue similar to that observed for NGF-withdrawal induced apoptosis. Thus, p53 elevation was an essential component of the apoptotic cascades induced in cultured sympathetic neurons by both NGF withdrawal and p75NTR activation.

p53 is Downstream of the MEKK-JNK Pathway, and Is Essential for MEKK-mediated Neuronal Apoptosis

These results indicated that the apoptotic cascades originating from NGF withdrawal and p75NTR activation share two common components; hyperphosphorylation of c-jun and p53 elevation. On the basis of these findings, we hypothesized that p53 was downstream of the MEKK-JNK pathway, which leads to c-jun hyperphosphorylation (Derijard et al., 1994; Yan et al., 1994). To test this hypothesis, we generated a recombinant adenovirus expressing an activated form of MEKK1 which has previously been shown to cause sympathetic neuron apoptosis and c-jun hyperphosphorylation (Eilers et al., 1998). We first confirmed that this virus expressed the recombinant myc epitope-tagged MEKK1 protein, and that it was capable of activating the MEKK1 downstream target, JNK (Yan et al., 1994), in sympathetic neurons. Specifically, sympathetic neurons were grown in 20 ng/ml NGF for 4 d, were infected with 20 moi of recombinant virus expressing activated MEKK1 or β-galactosidase, and then were analyzed 2 d later for expression of the myc-tagged MEKK1 on Western blots with anti-myc (Fig. 5A). Analysis of equal amounts of protein from β-galactosidase versus MEKK1-infected sympathetic neurons revealed the presence of a myc-immunoreactive protein of the appropriate size, 35 kD, in the MEKK1-infected neurons (Fig. 5A). To confirm that virally expressed MEKK1 was capable of activating JNK, we performed similar experiments and then analyzed the lysates for phosphorylation of JNK using a phospho-JNK antibody (Fig. 5B). In this experiment, we compared MEKK1-infected sympathetic neurons to sympathetic neurons withdrawn from NGF, where JNK is known to be activated (Eilers et al., 1998). Western blot analysis revealed that the activated MEKK1 adenovirus caused phosphorylation of JNK to the same level as did NGF withdrawal, relative to neurons maintained in 10 ng/ml NGF alone (Fig. 5B).

We next determined whether adenovirus-mediated expression of activated MEKK1 was sufficient to cause sympathetic neuron apoptosis, as was previously reported with microinjection (Eilers et al., 1998). Specifically, neurons were selected in 20 ng/ml NGF for 5 d, infected with various concentrations of recombinant virus expressing activated MEKK1 or β-galactosidase, maintained in 20 ng/ml NGF for a further 4 d, and then assayed for neuronal survival. Results from four separate experiments revealed that activated MEKK1 led to neuronal apoptosis; at 20, 50, or 100 moi, sympathetic neuron survival was decreased to 52, 54, and 44%, respectively, relative to neurons infected...
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Aloyz et al. uninfected sister cultures that were maintained in 20 ng/ml for the p53 adenovirus and maintained in 20 ng/ml NGF for 48 h, or from protein derived from sympathetic neurons infected with 20 moi MEKK

Western blot analysis for phospho-JNK in equal amounts of protein derived from sympathetic neurons that were infected with 20 moi MEKK1 adenovirus and maintained in 20 ng/ml NGF for 48 h, or from uninfected sister cultures that were infected with 20 moi myc-tagged MEKK1 adenovirus and maintained in 20 ng/ml NGF for 48 h.

Increased Bax protein, but does not affect phosphorylation of JNK.

Western blot analysis for phospho-JNK in equal amounts of protein derived from sympathetic neurons that were infected with 20 moi p53 adenovirus and maintained in 20 ng/ml NGF for 48 h, or from uninfected sister cultures that were maintained in 20 ng/ml for the same time period.

Western blot analysis for phospho-JNK in equal amounts of protein derived from sympathetic neurons infected with 20 moi p53 adenovirus and maintained in 20 ng/ml NGF for 48 h, or from uninfected sister cultures maintained in 20 ng/ml NGF and infected with 20 moi p53 adenovirus or 50 moi activated MEKK1 adenovirus for 30 h. As a control, neurons were withdrawn from NGF (0 NGF) for 30 h.

Figure 5. (A–D) p53 and Bax protein levels increase after activation of the MEKK-JNK pathway in sympathetic neurons. (A) Western blot analysis for c-myc in equal amounts of protein derived from sympathetic neurons infected with 20 moi myc-tagged MEKK1 (20 moi MEKK) or β-galactosidase (control) adenovirus for 48 h. In both cases, neurons were maintained in 20 ng/ml NGF for the entirety of the experiment. (B) Western blot analysis for phospho-JNK in equal amounts of protein derived from sympathetic neurons that were infected with 50 moi MEKK1 adenovirus and maintained for 48 h (50 moi MEKK), or from uninfected sister cultures that were either maintained in 10 ng/ml NGF (10 NGF), or that were withdrawn from NGF for 48 h (0 NGF). Note that the level of phospho-JNK immunoreactivity is similar in neurons withdrawn from NGF or transduced with activated MEKK1. (C) Western blot analysis for p53 in equal amounts of protein derived from sympathetic neurons that were infected with 20 moi MEKK1 adenovirus and maintained in 20 ng/ml NGF for 48 h (20 moi MEKK), or from uninfected sister cultures that were maintained in 10 ng/ml NGF (10 NGF), or that were withdrawn from NGF for 48 h (0 NGF). Note that p53 protein levels are increased by activated MEKK1 as they are by NGF withdrawal. (D) Western blot analysis for Bax in equal amounts of protein derived from sympathetic neurons that were infected with 50 moi MEKK1 adenovirus and maintained in 20 ng/ml NGF for 48 h (50 moi MEKK), or from uninfected sister cultures that were maintained in 10 ng/ml NGF for the same time period. (E and F) Increased expression of p53 in sympathetic neurons causes increased Bax protein, but does not affect phosphorylation of JNK. (E) Western blot analysis for phospho-JNK in equal amounts of protein derived from sympathetic neurons infected with 20 moi p53 adenovirus and maintained in 20 ng/ml NGF for 48 h, or from uninfected sister cultures that were maintained in 20 ng/ml for the same time period. (F) Western blot analysis for Bax in equal amounts of protein derived from sympathetic neurons infected with 20 moi p53 adenovirus and maintained in 20 ng/ml NGF for 48 h, or from uninfected sister cultures maintained in 20 ng/ml NGF and infected with 20 moi p53 adenovirus or 50 moi activated MEKK1 adenovirus for 30 h. As a control, neurons were withdrawn from NGF (0 NGF) for 30 h.

Note that p53 protein levels are increased when sympathetic neurons were infected with 20 or 50 moi of MEKK1 adenovirus, and that the magnitude of this increase was similar to that observed after NGF withdrawal for 48 h (Fig. 5 C). In contrast, no increase in p53 levels was seen upon infection with similar mois of the β-galactosidase adenovirus (data not shown), consistent with the observation that this latter virus had little or no effect on neuronal survival (Fig. 2 C). Thus, activation of the MEKK-JNK pathway led to increased levels of p53 in the presence of NGF.

To determine whether p53 was an essential component of the MEKK1-induced apoptotic signaling cascade, we asked whether the E1B55K adenovirus could rescue the apoptotic effects of activated MEKK1. To perform these experiments, sympathetic neurons were cultured for 5 d in 20 ng/ml NGF, and were infected with 50 moi of the MEKK1 virus plus or minus 500 moi of the E1B55K virus. Figure 5 D indicates that the E1B55K adenovirus was able to significantly rescue MEKK1-induced neuronal apoptosis (Fig. 2 D) relative to MEKK1 alone, or relative to MEKK1 plus the β-galactosidase or A262 adenovirus (P < 0.005 for E1B55K in all cases; Fig. 2 D). As a baseline for this study, we first determined whether coinfection with E1B55K reduced the p53 levels induced by activated MEKK1 alone (Fig. 3 D). Western blot analysis of sympathetic neurons infected for 2 d revealed that p53 protein levels were reduced in neurons expressing E1B55K plus activated MEKK1 relative to those expressing MEKK1 alone (Fig. 3 D). Reprobing of the same blot with an antibody specific for TrkA confirmed that equal amounts of protein were present in all of the samples (Fig. 3 E).

To determine whether this reduction in p53 levels mediated by E1B55K rescued sympathetic neurons from apoptosis induced by activated MEKK1, we performed similar experiments and, 4 d after infection, MTT assays were performed. As negative controls in this experiment, we used adenoviruses expressing the mutant A262 protein or β-galactosidase. Results of three separate experiments indicated that the E1B55K adenovirus was able to significantly rescue MEKK1-induced neuronal apoptosis (Fig. 2 D) relative to MEKK1 alone, or relative to MEKK1 plus the β-galactosidase or A262 adenovirus (P < 0.005 for E1B55K in all cases; Fig. 2 D). Thus, elevation of p53 pro-
tein levels is necessary for sympathetic neuron apoptosis after activation of the MEKK-JNK pathway.

One potential downstream candidate for the apoptotic effects of p53 in sympathetic neurons after MEKK-JNK pathway activation is the proapoptotic protein Bax. To determine whether Bax levels were increased in response to the MEKK1 adenovirus as they were after NGF withdrawal (Fig. 1F), we selected sympathetic neurons for 4 d in 20 ng/ml NGF, infected them with 50 moi of the MEKK1 adenovirus, switched them into 10 ng/ml NGF and 2 d later performed Western blots with anti-Bax. This analysis revealed that Bax levels were increased approximately twofold in sympathetic neurons expressing activated MEKK1 (Fig. 5D), an increase similar in magnitude to that seen after NGF withdrawal (Fig. 1F).

Previous studies have demonstrated that Bax is necessary for sympathetic neuron apoptosis after NGF withdrawal (Deckwerth et al., 1996). To determine whether proapoptotic proteins like Bax are also necessary for sympathetic neuron apoptosis after activation of the MEKK-JNK pathway, we generated a recombinant adenovirus expressing a prosurvival member of this pathway, Bcl-xl (Boise et al., 1993). Using this adenovirus, we then asked whether altering the balance between prosurvival versus proapoptotic proteins like Bax are also necessary for sympathetic neuron apoptosis after activation of the MEKK-JNK pathway. Specifically, sympathetic neurons were cultured for 5 d, and were infected with 50 moi of the MEKK1 adenovirus plus or minus various concentrations of Bcl-xl virus. For comparison, neurons were infected with 50 moi of MEKK1 adenovirus plus 500 moi E1B55K or of Bcl-xl virus. For comparison, neurons were infected with 50 moi of MEKK1 adenovirus plus 500 moi E1B55K or A262. Neurons were then maintained in 20 ng/ml NGF for four additional days before measuring survival using MTT A262. Neurons were then maintained in 20 ng/ml NGF for four additional days before measuring survival using MTT A262. Neurons were then maintained in 20 ng/ml NGF for four additional days before measuring survival using MTT A262.

\[ p53 \text{ Is Upstream of Bax and Downstream of JNK} \]

These experiments demonstrated that JNK, p53 and Bax are all downstream of activated MEKK, and that elevated p53 is essential for MEKK-induced apoptosis. To determine whether p53 is downstream of JNK and upstream of Bax, as we had hypothesized, we took advantage of a recombinant adenovirus expressing human p53 (previously described in Slack et al., 1996). Specifically, sympathetic neurons were cultured for 4 d in 20 ng/ml NGF, were infected with 20 moi of p53-expressing adenovirus and 48 h later, cell lysates were analyzed by Western blots with anti-p53, anti-phospho-JNK or anti-Bax. This analysis revealed that the p53 adenovirus increased expression of p53 to levels similar to those observed after NGF withdrawal (Fig. 5G), but that this elevated expression of p53 had no effect on JNK phosphorylation in sympathetic neurons maintained in 20 ng/ml NGF (Fig. 5E). In contrast, infection with the p53 virus caused an increase in the levels of Bax protein (Fig. 5F) that was similar in magnitude to that observed after NGF withdrawal (Fig. 1F) or after infection with the activated MEKK1 adenovirus (Fig. 5D). Together with the previous experiments with activated MEKK1, these experiments indicate that MEKK and JNK are upstream of p53, while Bax is downstream (Fig. 7).
p53 is Essential for Naturally Occurring Sympathetic Neuron Death In Vivo

Our previous work indicates that naturally occurring sympathetic neuron death is a result both of suboptimal activation of the TrkA receptor and of coincident activation of p75NTR (Bamji et al., 1998). Since, in culture, both of these types of neuronal apoptosis require p53, we hypothesized that p53 would also be essential for sympathetic neuron death in vivo. To test this hypothesis, we examined the SCG of transgenic mice in which the p53 gene was deleted by homologous recombination (Donehower et al., 1992). In control mice, the SCG contains ~20,000–25,000 neurons at birth, depending on the genetic background. Over the ensuing two weeks approximately half of these neurons are lost so that by P15, control SCGs contain ~13,000 neurons (Bamji et al., 1998). We therefore chose to analyze the SCG from p53+/+, p53+/− and p53−/− mice at two timepoints; postnatal days 1 and 15, timepoints immediately preceding and after the normal period of naturally occurring cell death (Fig. 6, A and D).

Analysis of the SCG at postnatal day 1 revealed that sympathetic neuron numbers before the period of naturally occurring death were similar regardless of the presence or absence of p53 (Fig. 6 D). The SCG of p53+/+ animals contained 23,976 ± 2764 neurons (n = 3), a number similar to mice of other genetic backgrounds (Bamji et al., 1998), while those from p53+/− and p53−/− animals contained 20,537 ± 2514 (n = 5) and 19016 ± 2675 (n = 3), respectively (Fig. 6 D). All of these numbers were statistically similar (P > 0.1 for all comparisons). However, analysis of the SCG from animals of these same genotypes at P15 (Fig. 6, A and D), after the period of naturally occurring death, revealed significant differences. In control p53+/+ animals the P15 SCG contained 13,163 ± 875 neurons (n = 5; Fig. 6 D). In contrast, the SCG of p53+/− animals contained 20,352 ± 944 neurons (n = 6), and that of p53−/− mice contained 20,600 ± 1709 neurons (n = 3), statistically significant increases of 55 and 56%, respectively (P < 0.005 in both cases). A comparison within the same genotype from P1 to P15 revealed that while the p53+/+ ganglia lost ~45% of its sympathetic neurons over this timeframe (P < 0.05; Fig. 6 D), there was no significant loss of sympathetic neurons in either the p53+/− or p53−/− SCG over the same period (Fig. 6 D; P > 0.3).

To confirm that this difference in neuron number in the SCG during the period of naturally occurring cell death reflected a deficit in apoptosis in the p53+/− and p53−/− mice, and was not due to an increase in the proliferation of neuronal progenitors that occurs for a short period postnatally in the SCG (Hendry, 1977), we analyzed the total number of apoptotic cells in the SCG at postnatal day 7, when sympathetic neuron apoptosis is ongoing. To perform this analysis, SCG from p53+/− and p53+/+ littermates were sectioned, and every third section was analyzed by in situ TUNEL-labeling. This analysis revealed that TUNEL-positive cells were detected in the ganglia of both p53+/− and p53+/+ animals (Fig. 6 B), but that the total number of apoptotic nuclei was significantly decreased in the p53+/− ganglia (p53+/− SCG, mean = 661 ± 19, n = 5; p53+/+ SCG, mean = 1025 ± 155, n = 4; P = 0.016; Fig. 6 C). Thus, the magnitude of sympathetic neuron apoptosis is decreased when p53 levels are decreased in vivo.

Discussion

In this study, we have examined the role of the p53 tumor suppressor in naturally occurring sympathetic neuron death. Our findings support an essential role for p53 in this process and, more specifically, support the following conclusions. First, after NGF withdrawal of neonatal sympathetic neurons, p53 levels are increased with a timecourse that is consistent with it playing a role in the “commitment” to apoptosis. The magnitude of this increase is similar to that required to induce sympathetic neuron apoptosis using adenovirus-mediated transduction of p53 (Slack et al., 1996). Second, levels of two p53 transcriptional targets, p21 and Bax, are also increased; the latter of these two, Bax, has already been shown to be essential for sympathetic neuron apoptosis after NGF withdrawal (Deckwerth et al., 1996). Third, p53 levels are also increased when sympathetic neuron apoptosis is induced by p75NTR activation, with a similar timecourse to that observed after NGF withdrawal. Fourth, a decrease in levels of p53 mediated by the E1B55K protein is sufficient to inhibit sympathetic neuron apoptosis induced either by NGF withdrawal or by p75NTR activation, indicating that increased p53 levels are both necessary and sufficient (Slack et al., 1996) for sympathetic neuron apoptosis. Fifth, NGF withdrawal and p75NTR activation may induce apoptosis via a pathway involving MEKK-JNK-p53-Bax (Fig. 7) since (a) expression of activated MEKK1, which is sufficient to cause sympathetic neuron apoptosis, leads to elevated levels of p53 and Bax, (b) expression of p53 leads to elevated Bax levels, but does not affect JNK activation, and (c) elevated p53 is essential for MEKK-induced apoptosis. Finally, the physiological relevance of these observations is indicated by our findings that, when p53 is reduced or absent, naturally occurring sympathetic neuron death is inhibited in vivo. Thus, our data indicate that p53 is a common, essential target during developmental sympathetic neuron death that is downstream of the MEKK-JNK pathway and that may well integrate apoptotic signals deriving both from p75NTR activation and from a lack of NGF/TrkA receptor activation.

The p53 tumor suppressor protein encodes a transcriptional regulator that functions to control cell proliferation and apoptosis in a cell context-dependent fashion (reviewed in Levine, 1997; Jacks and Weinberg, 1998). The precise mechanism by which p53 mediates apoptosis is not well understood, but it is believed to proceed by a number of mechanisms including direct transactivation, transcriptional repression, and direct involvement in DNA cleavage (Lane, 1993; Caelles et al., 1994; Elledge and Lee, 1995; Miyashita and Reed, 1995). Within the nervous system, p53 is upregulated in neurons after a number of traumatic insults, including kainic acid and ischemia (Li et al., 1994; Sakhi et al., 1994; Wood and Youle, 1995), and is necessary for excitotoxicity-induced apoptosis of hippocampal and cortical neurons (Morrison et al., 1996; Xiang et al., 1998). However, p53 has not been thought to be involved in naturally occurring neuronal cell death, nor has it been thought to be an essential downstream effector of
death receptor activation. In particular, a number of previous studies have failed to implicate p53 in sympathetic neuron death. In one study, Martinou et al. (1995) demonstrated that microinjection of the adenoviral protein E1B19K, but not E1B55K, rescued sympathetic neurons from apoptosis due to NGF withdrawal. The inability to rescue sympathetic neuron apoptosis with microinjected E1B55K as opposed to adenovirally expressed E1B55K is likely due to the lower levels of expression obtained using microinjection. In this regard, we have ourselves compared the rescue of sympathetic neurons using adenovirally expressed E1B19K versus E1B55K, and E1B19K was able to rescue sympathetic neurons at titers ~10-fold lower than those needed for E1B55K (Pacquet, L., F. Miller, and D. Kaplan, unpublished data). In a second set of studies, cultured p53−/− sympathetic neurons died after NGF withdrawal, leading to the conclusion that p53 played no role in developmental sympathetic neuron death (Davies and Rosenthal, 1994; Sadoul et al., 1996). The studies presented here demonstrate that neurons with lowered p53 levels still die in vivo, although the magnitude of this death is significantly lower than in controls; these results do not necessarily contradict the finding that p53−/− neurons die in culture after NGF withdrawal. Moreover, results might also differ in response to acute inhibition of p53 (as mediated via E1B55K) versus a chronic loss of p53 (as in p53−/− mice). For example, the developmental absence of p53 may well cause compensatory upregulation of parallel apoptotic pathways. Precedent for such compensation derives from our previous studies with another tumor suppressor protein, pRb. In Rb−/− mice, cortical neurons are unable to differentiate appropriately and, as a consequence, undergo neuronal apoptosis in vivo (Slack et al., 1998). However, when the same Rb−/− cortical progenitors are cultured and undergo the transition to postmitotic neurons in vitro, they differentiate normally and do not undergo apoptosis (Slack et al., 1998); this difference is explained by a selective upregulation of p107 and p130, two other Rb family members, in culture but not in vivo (Ruth Slack, personal communication). A similar explanation could be invoked for cultured p53−/− neurons; sympathetic neurons express p73 (Pozniak, C., F. Miller, unpublished data), an apoptotic p53 family member (Jost et al., 1997; Kaghad et al., 1997) that might well compensate for the lack of p53 in a mutant p53 background.

What is the signal transduction pathway that derives either from p75NTR activation or from a lack of activation of TrkA and that leads to increased p53? Our data implicate the MEKK-JNK pathway (Derijard et al., 1994; Yan et al., 1994; Fig. 7). This pathway is activated in sympathetic neurons in response to either NGF withdrawal (Ham et al., 1995; Eilers et al., 1998) or p75NTR activation (Bamji et al., 1998), and the resultant hyperphosphorylation of c-jun is necessary for NGF-withdrawal induced sympathetic neuron death (Estus et al., 1994; Ham et al., 1995). The data presented here indicate that activation of the MEKK-JNK pathway causes sympathetic neuron death via a p53-dependent mechanism, thereby providing a potential direct link from p75 to p53. Moreover, our finding that the stress-induced (Herdegen et al., 1997) MEKK-JNK pathway acts via p53 in neurons may explain why excitotoxicity-induced death of CNS neurons can be inhibited by deletion of either the JNK3 (Yang et al., 1997) or p53 genes (Morrison et al., 1996; Xiang et al., 1998).

Are there other pathways that cause elevated p53 after NGF withdrawal or p75NTR activation? A second candidate upstream pathway involves deregulation of the cell cycle; in cycling cells, p53 is viewed as a "fail-safe" mechanism that causes cellular apoptosis when proliferative mechanisms are deregulated (reviewed in Levine, 1997; Jacks and Weinberg, 1996). Such cell cycle deregulation has been proposed to occur in sympathetic neurons after NGF withdrawal (Rubin et al., 1993; Freeman et al., 1994; Park et al., 1997). Therefore, it is possible that NGF withdrawal causes two different signals, MEKK-JNK activation and cell cycle deregulation, that together converge on p53. Thus, p53 may function as a sensor to induce neuronal apoptosis when the sum of these upstream signals reaches a critical level.

What are the downstream events that are responsible for p53-mediated neuronal apoptosis? One potential downstream mechanism involves the protein Bax, which is known to be essential for sympathetic neuron apoptosis (Deckwerth et al., 1996), and for p53-dependent death of cortical neurons (Xiang et al., 1998). Data presented here indicate that, after the commitment of sympathetic neurons to apoptosis, the balance of proapoptotic versus pro-survival members of the bcl-2 family is considerably shifted to the proapoptotic, and that one of the changes contributing to this shift is an increase in Bax protein levels. Moreover, we demonstrate that increased expression of p53 alone is sufficient to cause increased Bax protein levels in sympathetic neurons. Since Bax is a direct tran-
scriptional target of p53 (Miyashita and Reed, 1995), we propose that elevated p53 protein levels cause increased p53-dependent transcription of Bax, and that this is sufficient to tip the balance towards neuronal apoptosis. In support of this hypothesis, exogenous p53 is unable to mediate apoptosis of Bax−/− cortical neurons (Xiang et al., 1998), Bax−/− sympathetic neurons do not undergo apoptosis in response to NGF withdrawal (Deckwerth et al., 1996), and increased expression of Bel2 or Bel-xI (Garcia et al., 1992; Gonzalez-Garcia et al., 1995) in sympathetic neurons rescues NGF withdrawal induced cell death.

The in vivo data presented here support a number of additional conclusions. First, our studies demonstrate that the postnatal period of naturally occurring sympathetic neuron death is inhibited, but not completely eliminated when p53 levels are lowered, indicating that p53 is essential for normal apoptosis in the SCG, but also suggesting that other, potentially parallel death pathways are important. Such pathways may involve other p53 family members such as p73 (Jost et al., 1997; Kaghad et al., 1997) or p51 (Osada et al., 1998; Trink et al., 1998). Second, our data showing decreased apoptosis in the p53+/− SCG support the idea that it is not the presence or absence of p53 that determines apoptosis, but it is instead the levels of p53 that are important. Precedent for such a gene dosage effect has been documented with regard to p53 and tumorigenesis (Donehower et al., 1992; Harvey et al., 1993). Third, our in vivo data demonstrate that, although p53 may be essential embryonically for regulating the proliferation and survival of sympathetic neuroblasts (Vogel and Parada, 1998), that by birth any resultant deficits are not obvious, since neuron numbers in the newborn p53+/+ versus p53−/− SCG are not significantly different.

What is the biological rationale for p53 acting as a “threshold” to neuronal apoptosis? During the period of naturally occurring death, sympathetic neurons compete for limiting amounts of trophic support to ultimately ensure an appropriate target innervation density. The net result of this competition is a loss of ~40–50% of sympathetic neurons over the first 2 wk of postnatal life. Our previous data indicate that both TrkA and p75NTR are required for this process (Bamji et al., 1998). Moreover, these same studies demonstrate that robust activation of TrkA antagonizes p75NTR-derived apoptotic signals and that, conversely, p75NTR can override TrkA-derived survival signals. On the basis of the current study, we propose that p53 is a convergent downstream target that “sums” opposing signals deriving from TrkA versus p75NTR activation, and that apoptosis is triggered when the balance of “death” signals deriving from p75NTR is greater than the “survival” signals deriving from TrkA. In cellular terms, this ensures that all neurons that are incapable of sequencing sufficient target-derived NGF are rapidly and efficiently eliminated from competition.

One final issue derives from the emerging similarities between sympathetic neuron apoptosis as induced by NGF withdrawal and p75NTR activation. In both cases, the MEKK-JNK pathway is induced, and in both cases, p53 is required for apoptosis. In the case of p75, while this receptor has been demonstrated to signal via a number of pathways, including ceramide production (Dobrowsky et al., 1994), JNK activation (Casaccia-Bonnefil et al., 1996), c-jun hyperphosphorylation (Bamji et al., 1998) and NFkB activation (Carter et al., 1996), the results presented here are the first to identify a required component, p53, of the p75NTR apoptotic pathway. Given the similarities between NGF withdrawal and p75NTR activation, we propose that NGF-withdrawal-induced apoptosis may be, to a large extent, a p75NTR-mediated process. In particular, our previous work (Bamji et al., 1998) indicates that NGF-withdrawal induced apoptosis of sympathetic neurons is greatly delayed in the absence of p75NTR. We propose that, during naturally occurring sympathetic neuron death, p75NTR may well provide the basis for a constitutive apoptotic signal in sympathetic neurons via one of two different mechanisms. First, there may be an autocrine p75NTR-dependent apoptotic loop; sympathetic neurons synthesize BDNF (Causing et al., 1997) that they process and secrete (Causing, C.G., R. Aloyz, and F.D. Miller, unpublished observations). Since BDNF can bind to p75NTR to cause sympathetic neuron apoptosis (Bamji et al., 1998), and since BDNF is required in vivo for a portion of appropriate naturally occurring sympathetic neuron death (Bamji et al., 1998), this may indicate the presence of an ongoing autocrine BDNF:p75NTR death signal. A second possibility is that p75NTR functions to mediate a constitutive death signal in the absence of ligand, as previously suggested (Rabizadeh et al., 1993). However, it is clear that addition of exogenous BDNF is essential for a maximal p75NTR-dependent apoptotic signal (Bamji et al., 1998), indicating that even if a constitutive death mechanism is in place, activation of p75NTR by exogenous, potentially target-derived, neurotrophins is likely to play a key role. In either case, exposure to an appropriate neurotrophin, in this case NGF, would lead to TrkA activation, thereby providing a mechanism for overriding p75 and p53-mediated apoptosis.

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