



# v-Crk, an effector of the nerve growth factor signaling pathway, delays apoptotic cell death in neurotrophin-deprived PC12 cells

Robert H. Glassman<sup>1,2</sup>, Barbara L. Hempstead<sup>1</sup>,  
Lisa Staiano-Coico<sup>3</sup>, Melissa G. Steiner<sup>3</sup>, Hidesaburo  
Hanafusa<sup>2</sup> and Raymond B. Birge<sup>2,4</sup>

<sup>1</sup> Division of Hematology-Oncology, Department of Medicine, The New York Hospital-Cornell University Medical College, 1300 York Avenue, New York, NY 10021, USA

<sup>2</sup> Laboratory of Molecular Oncology, The Rockefeller University, 1230 York Avenue, New York, NY 10021, USA

<sup>3</sup> Division of Otorhinolaryngology, Department of Surgery, The New York Hospital-Cornell University Medical College, 1300 York Avenue, New York, NY 10021, USA

<sup>4</sup> corresponding author: Laboratory of Molecular Oncology, The Rockefeller University, 1230 York Avenue, New York, NY 10021. e-mail: birger@rockvax.rockefeller.edu

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## Abstract

v-Crk is a member of a class of SH2 and SH3-containing adaptor proteins that have been implicated in regulating the TrkA receptor tyrosine kinase and potentiating Nerve Growth Factor (NGF)-mediated neurite outgrowth in pheochromocytoma (PC12) cells (Hempstead *et al*, Mol. Cell Biol. 14: 1964–1971). Given the fact that NGF induces both differentiation and survival by binding to TrkA, we examined the rate of apoptotic cell death elicited by NGF-withdrawal in native, v-Crk, and TrkA-expressing PC12 cells. While more than 50% of native PC12 cells underwent apoptosis within 48 h of NGF withdrawal, the v-Crk and TrkA-expressing cells were much more resistant to apoptosis under these conditions, whereby approximately 70 and 95%, respectively, of the cells were alive. The ability of v-Crk to delay apoptosis required prior NGF-dependent differentiation, since naive undifferentiated v-Crk expressing PC12 cells or cells that express v-Crk mutants that are defective in NGF signaling were not protected from apoptosis during growth factor withdrawal. Moreover, addition of 50 ng/ml EGF to serum and NGF deprived v-Crk expressing cells, which also causes neurite outgrowth, promoted complete and long-term survival, although such EGF replacement had no neurotrophic effect on wild-type PC12 cells or PC12 cells overexpressing Human Bcl-2. These experiments suggest that v-Crk potentiation of a receptor tyrosine kinase under conditions of growth factor deprivation is essential for preventing apoptosis. However, unlike native PC12 cells, neither v-Crk or TrkA-expressing PC12 cells exhibited a G1 arrest when incubated for 2 weeks in NGF. Thus, v-Crk and TrkA may protect NGF deprived PC12 by preventing cell cycle arrest and hence an aborted entry into a defective

cell cycle. Moreover, during NGF-withdrawal, v-CrkPC12 cells exhibited down regulation in MAP kinase and JNK activities while in native cells, these activities increased within 6–8 h after NGF deprivation. Thus, unlike v-Crk-mediated augmentation of differentiation, sustained activation of MAP kinase may not be required for v-Crk-induced cell survival.

**Keywords:** SH2 and SH3-containing adaptor proteins, TrkA receptor tyrosine kinase, nerve growth factor, Bcl-2, MAP kinase

**Abbreviations:** NGF: nerve growth factor; EGF: epidermal growth factor; PC12: pheochromocytoma cells; SH2: Src Homology 2; PLC- $\gamma$ : phospholipase C- $\gamma$ ; PI-3 kinase: phosphatidylinositol-3-kinase

## Introduction

Programmed cell death is an essential feature during development of the nervous system, to eliminate redundant cells and to achieve a precise balance between target cells and neurons. The survival and differentiation of immature neuroblasts in the peripheral nervous system is regulated by the limiting supply of neurotrophic factors (Barde *et al*, 1989), the best characterized of which is nerve growth factor (NGF) (Thoenen and Barde, 1980). The neural crest-derived rat pheochromocytoma (PC12) cell line has served as a model system to study the role of NGF during neuronal differentiation and survival (Greene and Tischler, 1976, 1982). Upon exposure to NGF, PC12 cells stop dividing, develop neurites, and increase neurotransmitter synthesis so as to acquire a phenotype similar to mature sympathetic neurons (Greene and Tischler, 1982; Rudkin *et al*, 1989). NGF initiates these effects by binding to its receptor, p140 Trk A, and activating its intrinsic tyrosine kinase activity (Kaplan *et al*, 1991; Klein *et al*, 1991). A number of cytoplasmic signaling proteins, including phospholipase C- $\gamma$  (PLC- $\gamma$ ), phosphatidylinositol-3-kinase (PI-3' kinase), and Shc, which contain either Src Homology 2 (SH2) or Phosphotyrosine binding domains (PTB) (Geer and Pawson, 1995) bind to specific sequences within the auto-phosphorylated TrkA receptor (reviewed in Kaplan and Stephens, 1994). Mutagenesis of the Shc (Y490) and PLC- $\gamma$  (Y785) binding sites on TrkA block NGF-mediated neurite outgrowth demonstrating the importance of receptor phosphorylation in mediating neural differentiation (Obermeyer *et al*, 1994; Stephens *et al*, 1994). While the signaling pathways that lead to differentiation following NGF addition are reasonably well understood, the molecular mechanisms regulating apoptosis during NGF withdrawal are particularly lacking.

An emerging hypothesis that may explain mechanisms by which mature neurons activate a suicide program upon withdrawal of NGF suggests that re-activation of mitogenic signals in a post-mitotic neuron can lead to cell death (Rubin *et al*, 1993). This idea is supported by the observations that cell cycle related genes such as cyclin D1, cyclin A, *c-jun* and *c-fos* are upregulated within hours following NGF withdrawal in sympathetic neurons (Estus *et al*, 1994; Freeman *et al*, 1994; Meikrantz *et al*, 1994; Kranenburg *et al*, 1996). In addition, the induction of dominant negative N17Ras (Ferrari and Greene, 1994) or microinjection of dominant negative FLAG $\Delta$ 169 c-Jun in neuronal PC12 cells or primary SCG neurons, respectively, just prior to NGF withdrawal can inhibit apoptosis (Ham *et al*, 1995). In many cell types, conflicting growth promoting and growth arresting signals can also induce apoptosis, as in the case where p53 transfection into E1A-transformed fibroblasts induces apoptosis (Lowe *et al*, 1993). Moreover, constitutive expression of *c-myc* in rat fibroblasts or Chinese hamster ovary (CHO) cells induces apoptosis in cells arrested by a variety of cytostatic manipulations, such as serum deprivation or addition of drugs which inhibit DNA replication (Bissonnette *et al*, 1992; Evan *et al*, 1992).

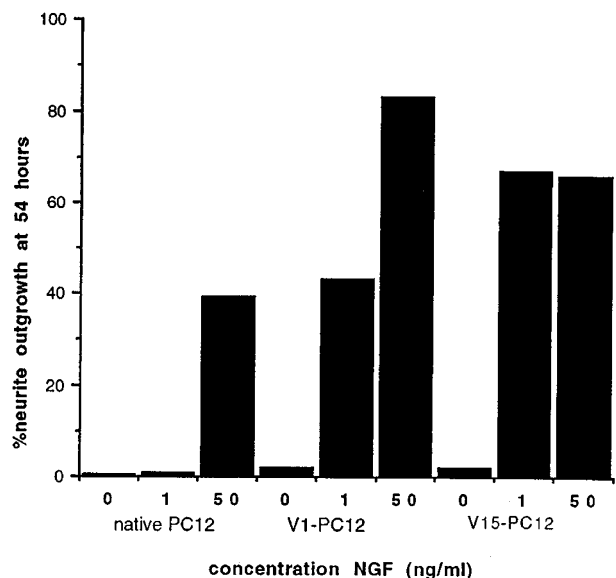
Adaptor proteins, which consist of Src Homology 2 and 3 (SH2 and SH3) domains, link receptor tyrosine kinases such as Trk A with downstream cytosolic effector proteins (Cantley *et al*, 1991) and are likely integral components of NGF signaling pathways (Park and Rhee, 1992; Rozakis-Adcock *et al*, 1992; Hempstead *et al*, 1994; Matsuda *et al*, 1994). SH2 domains are conserved motifs of approximately 100 amino acids which recognize and bind tyrosine phosphorylated proteins (Cantley *et al*, 1991); SH3 domains, comprised of approximately 60 amino acids, recognize proline rich motifs of the general consensus PXXP (Ren *et al*, 1993). The role of adaptor proteins in apoptotic signaling is largely uninvestigated, although microinjection of Grb3-3, an SH2 deleted isoform of Grb2, triggered apoptotic death in fibroblasts (Fath *et al*, 1994). We have recently shown that the v-Crk adapter protein potentiates TrkA signaling and accelerates NGF-induced neurite outgrowth in PC12 cells (Hempstead *et al*, 1994). The v-Crk protein contains avian retroviral Gag sequences fused in frame to cellular sequences containing one SH2 and one SH3 domain (Mayer *et al*, 1988) and is derived from an ubiquitously expressed cellular gene, called *c-crk*, which contains an SH2 domain followed by two SH3 domains (Reichman *et al*, 1992). To further investigate how SH2 and SH3 containing proteins influence neurotrophic pathways, we examined the short-term survival of PC12 cells expressing v-Crk following NGF deprivation. Here we report that v-Crk significantly delays apoptotic death in differentiated PC12 cells following NGF withdrawal. Using mutant v-Crk PC12 cells which are deficient in NGF signaling, we show that the ability of v-Crk to protect PC12 cells correlates with its ability to potentiate TrkA signaling. Moreover, re-stimulation of EGF Receptor in NGF-deprived v-Crk-expressing PC12 cells caused complete and long-term resistance to apoptosis. These results indicate that v-Crk can modulate tyrosine kinase signals in

PC12 cells and may function in a novel anti-apoptotic signaling pathway.

## Results

### Stable expression of v-Crk or TrkA delay cell death in NGF-deprived differentiated PC12 cells

Ectopic expression of v-Crk in rat PC12 cells augments the rate of neurite outgrowth following treatment with 50 ng/ml NGF (Hempstead *et al*, 1994) and potentiates phenotypic differentiation in response to sub-threshold (1 ng/ml) concentrations of NGF (Figure 1 and (Hempstead, 1994)). Cells maintained under conditions of long-term NGF treatment will be referred to as 'neuronal' PC12 cells throughout the paper. To assess whether v-Crk could also potentiate survival of neuronal PC12 cells as a result of short-term NGF withdrawal, a fluorescence live/death analysis of native PC12 cells or v-Crk-PC12 cells was performed following NGF deprivation (Figure 2c, d, g, h and summarized in Figure 3). In the NGF-deprived PC12 cultures, about 60% of the cells stained ethidium homodimer by 72 h, indicating the cells have lost membrane integrity in the late stages of cell death (Figure 3b). In contrast, more than 70% of the v-CrkPC12 cells were impermeable to ethidium homodimer and remained viable as manifested by intracellular esterase activity, assayed by fluorescent activation of calcein AM. Although viability was maintained in v-CrkPC12 cells after growth factor removal, neurite extension was not, with only 10–20% of cells still

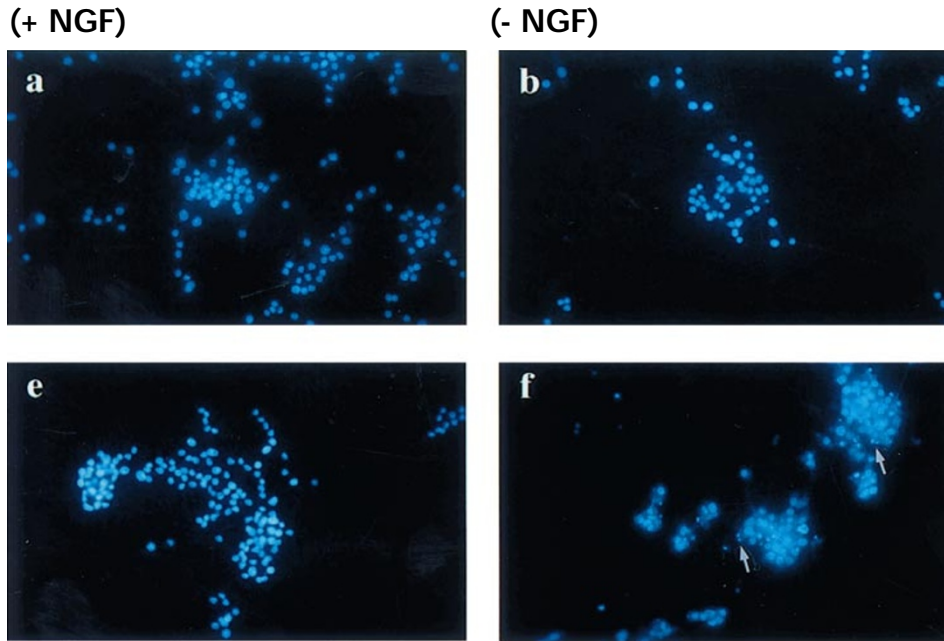


**Figure 1** v-Crk is an effector of the NGF signaling pathway at 1 ng/ml NGF. PC12 cells stably expressing v-Crk (clones V1 and V15) were maintained on rat-tail collagen coated plates in 3% serum (2% calf and 1% horse) and seeded into six well dishes after which either no ligand (0), 1 ng/ml NGF, or 50 ng/ml NGF was added for 48 h. Percent neurite outgrowth is expressed as the percentage of cells containing neurite processes greater than two cell bodies (approximately 20  $\mu$ m in diameter). Experiments are the average of triplicate wells.

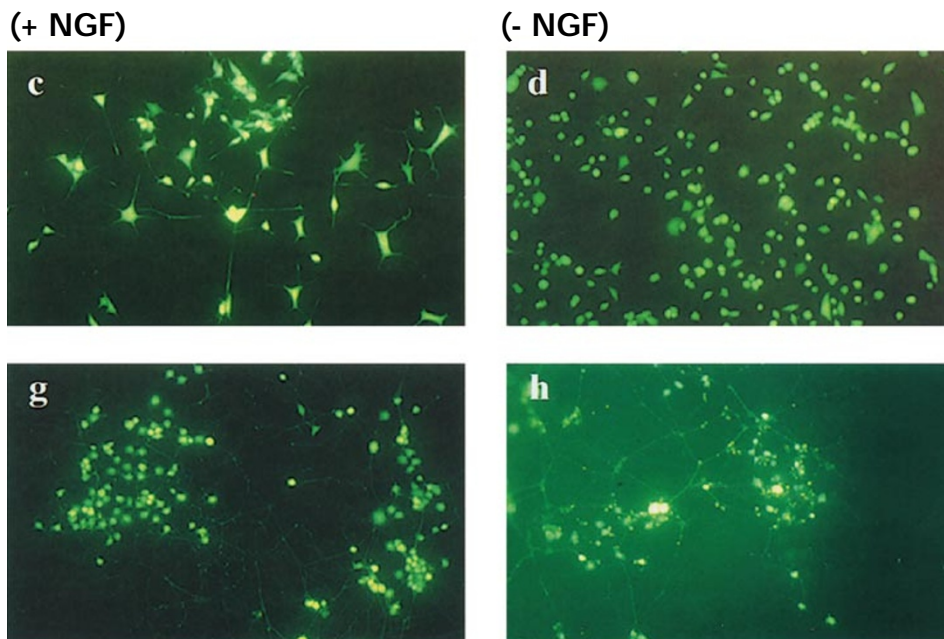
expressing neurites of two cell bodies after 72 h in NGF-free conditions. Cells were also stained with Hoechst 33258, a more conventional means to ascertain apoptotic chromatin condensation (Figure 2a, b, e, f). Herein, several PC12 cell

nuclei exhibited punctate chromatin staining following 36 h of growth factor deprivation, whereas parallel cultures of v-CrkPC12 nuclei had uniform, non-condensed chromatin staining (Figure 2 compare panels b versus f).

### Hoechst 33258



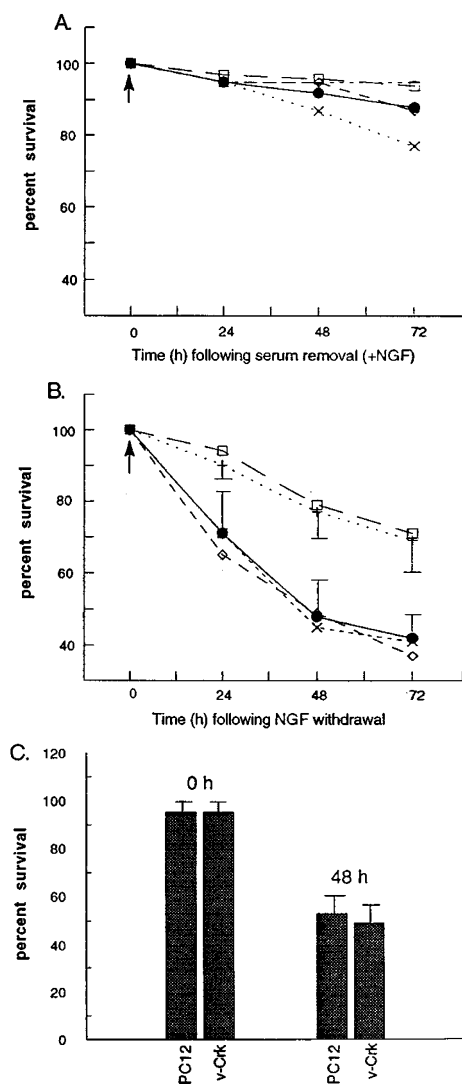
### Fluorescence live/death



**Figure 2** Morphological appearances of v-Crk-expressing PC12 cells and native PC12 cells during NGF withdrawal. Native or v-Crk-expressing PC12 cells were cultured on rat-tail collagen-coated plates and maintained in 3% serum plus 50 ng/ml NGF for 14 days (**a, c, e, g**) or transferred to serum-free media without NGF for 36 h (**b, d, f, h**). Live versus dead cells were either scored by a two-color fluorescence assay involving ethidium homodimer and calcein AM (**c, d, g, h**) or Hoechst 33258 staining (**a, b, e, f**). Similar results were obtained with v-Crk-expressing clone V15. Magnification is 20 × .

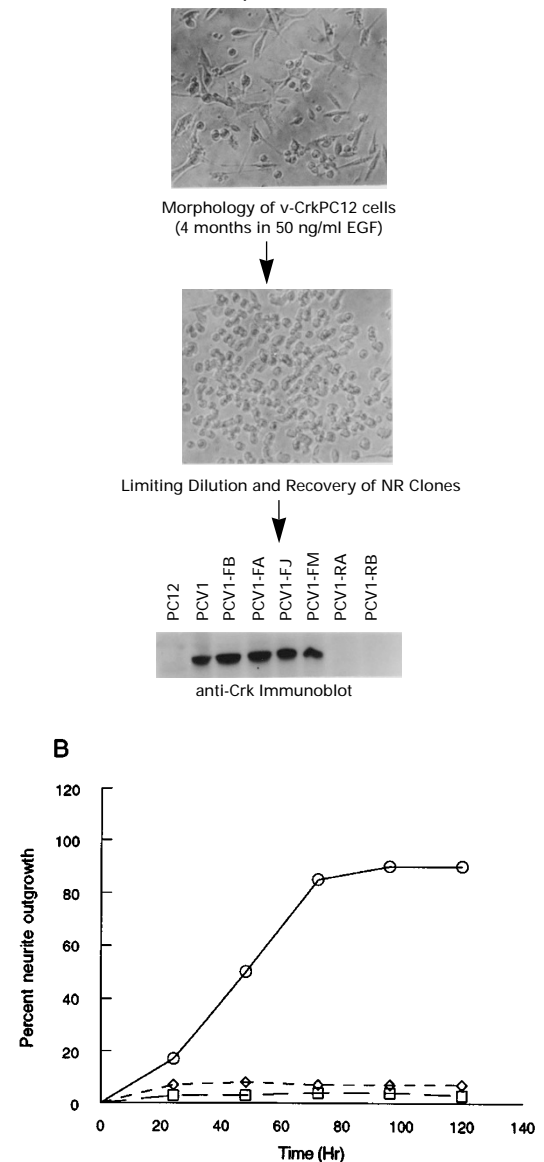
In contrast to the effects of v-Crk in NGF-deprived neuronal PC12 cells, v-Crk did not exert a survival benefit when naive undifferentiated PC12 cells were deprived of serum as a source of growth factors, as this condition led to comparable cell death to that of nontransfected PC12 cells (Figure 3C). A reciprocal test of the role of NGF-mediated differentiation in predicting v-Crk's survival advantage is to examine a cell line which cannot differentiate with exogenous NGF. For these experiments, we isolated a mutant v-CrkPC12 cell line, termed NR-v-Crk, or 'Non-Responder' v-Crk. NR-v-Crk PC12 cells were isolated by

limiting dilution and clonal expression of V1 clones that no longer responded (via neurite outgrowth) to EGF, NGF, or bFGF, indicating a general defect in growth factor dependent neurite outgrowth (Figure 4). When examined for v-Crk levels by anti-Gag Western blotting, four out of the six clones analyzed continued to express v-Crk at levels equivalent to the parental v-Crk PC12 cell lines (Figure 4A)



**Figure 3** v-Crk promotes the survival of NGF-differentiated PC12 cells following NGF withdrawal. (a and b). Native PC12 (●), v-Crk-expressing PC12 clones V1 (+) or V15 (□), R273N-v-Crk (◇), and NR-v-Crk PC12 (x) cells were maintained in 3% serum plus 50 ng/ml NGF for 14 days and then placed in serum free media (indicated by the arrow) containing 50 ng/ml NGF (A) or serum-free media without added growth factor (B) for up to 72 h. Live versus dead cells were scored by the two-color fluorescence assay shown in Figure 2. Data show the mean  $\pm$  S.E. of at least six independent experiments. (c) native PC12 cells or v-CrkPC12 cells were grown in 3% serum without NGF for 1 week and then placed in serum free media for 48 h.

**A. Isolation of Non-responsive (NR) v-CrkPC12 Cells.**



**Figure 4** Isolation of Non-responder (NR) v-Crk PC12 cells. (A) v-Crk expressing PC12 clone V1, maintained for approximately 4 months in 3% serum plus 50 ng/ml EGF, was trypsinized and diluted to approximately ten cells per 10 cm dish. Cells were then expanded as clonal lines and individual clones not responding to EGF were isolated (middle panel). 40  $\mu$ g of detergent lysate was Western blotted with anti-Gag antibody (monoclonal antibody 3C2) to monitor v-Crk expression. The clone numbers are indicated on the top of the gel. (B) NR-v-Crk cells do not differentiate with EGF or NGF. Wild-type v-CrkPC12 cells (V1) treated with EGF (○) or clone NRv1-FA treated with 50 ng/ml EGF (□) or 50 ng/ml NGF (◇) for up to 120 h and were scored for neurite outgrowth as in Figure 1. Other NR-v-Crk clones gave the same results.

and therefore were chosen for further studies. Moreover, the v-Crk in these cells stably complexed with tyrosine phosphorylated EGF Receptor, p130<sup>cas</sup> and paxillin suggesting that v-Crk was wild-type in these cells (data not shown). Although NR-v-Crk cells do not differentiate in response to NGF, they do maintain viability under serum free conditions in the presence of 50 ng/ml NGF (Figure 3A). Thus, these cells represent a system where one can clearly dissociate survival and differentiative pathways. These latter characteristic distinguishes NR-v-Crk cells mechanistically from the PC12nnr5 cell mutants isolated by Greene and colleagues which are non-responsive to NGF due to lack of TrkA expression (Loeb and Greene, 1993). Importantly, NR-v-Crk cells did not retain the survival promoting action of v-Crk upon NGF deprivation (Figure 3B). Taken together, the results in Figure 3 and 4 demonstrate that PC12 cells expressing v-Crk require prior differentiation by NGF in order for v-Crk to exert a cytoprotective effect.

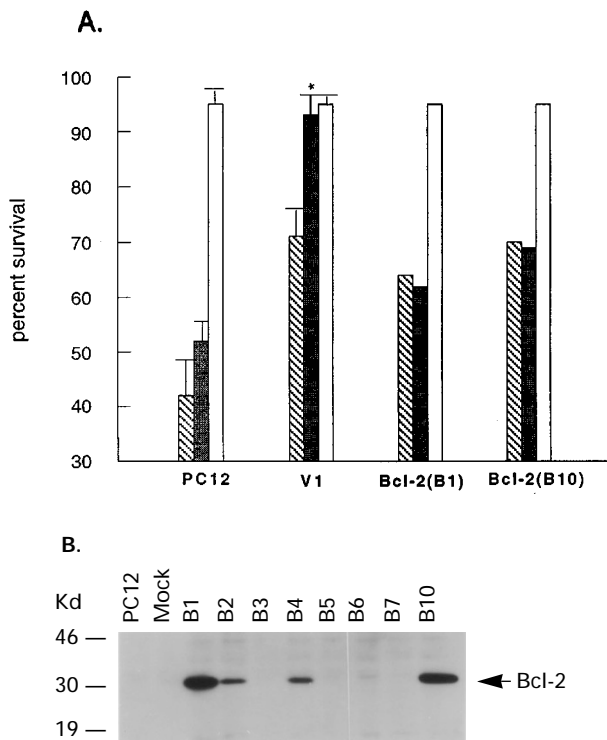
Given the fact that v-Crk potentiates TrkA signaling (Hempstead *et al*, 1994), and delays apoptosis only in cells with a neuronal phenotype, we hypothesized that over-

expression of TrkA itself may sustain viability in cell-free conditions. To address this question, we assessed survival in Trk 6-24 PC12 cells that exhibit 20-fold overexpression of the TrkA receptor (Hempstead *et al*, 1993). Figure 6B shows Hoechst staining of nuclei 48 h after NGF withdrawal in native PC12 cells and TrkA-expressing PC12 cells. We found no evidence of apoptotic condensed chromatin in any TrkA PC12 cells, by contrast to what we observed with native cells. Moreover, when cells were stained with calcein AM, over 95% of the Trk 6-24 cells survived following 72 h of NGF deprivation (Figure 6A). Hence, overexpression of Trk A in PC12 cells appears to confer growth factor independence in PC12 cells.

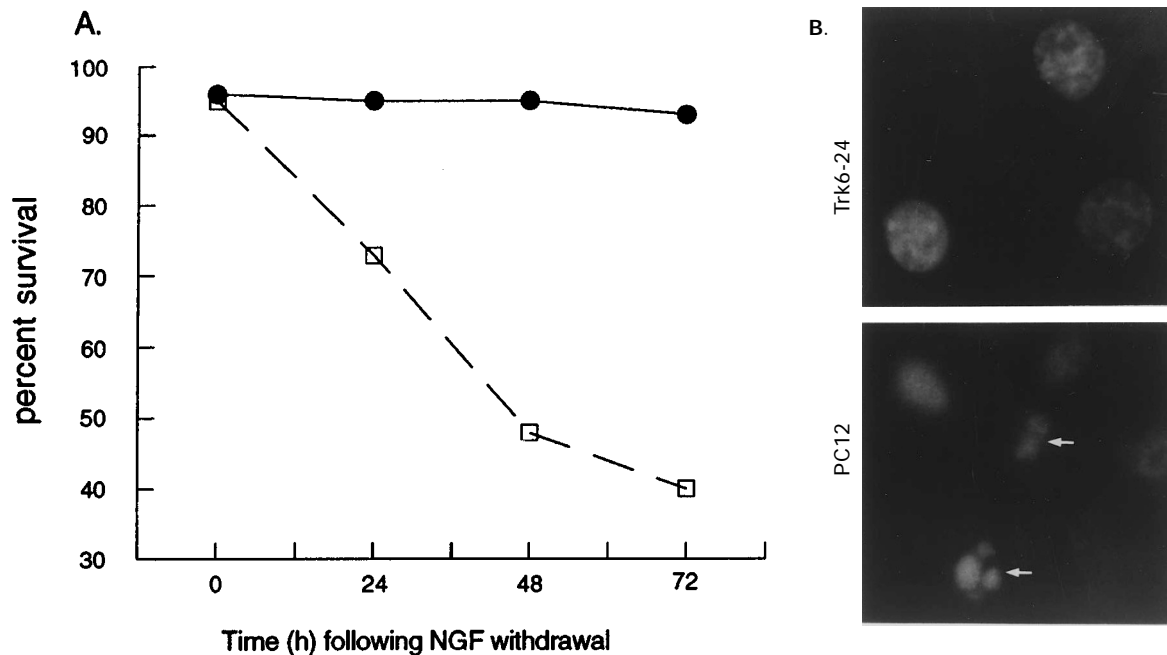
### EGF is a survival factor in v-Crk-expressing PC12 cells

Our previous findings demonstrated that EGF, a mitogenic growth factor in native PC12 cells, causes neurite outgrowth in v-Crk-expressing PC12 cells (Hempstead *et al*, 1994). To determine whether EGF could also rescue cells from apoptosis during NGF-deprivation, we replaced NGF with an identical concentration of EGF in long-term NGF-treated cells to assess its survival potential (Figure 5A). In two phenotypically differentiated v-Crk expressing cell lines, V1- and V15-PC12 cells, EGF was identical to NGF in acting as a neurotrophin; For example, 72 h after serum withdrawal, 93% of V1-PC12 cells were rescued in EGF alone, 94% in NGF alone, as compared to 71% in the absence of growth factor (Figure 5A). In fact, over 90% of the v-CrkPC12 cells were alive after 7 days in serum free media with EGF alone, while less than 10% survived in the absence of growth factor (not shown). EGF did not significantly augment survival in native PC12 cells, corroborating results by Greene *et al* (1991). To show that these effects were specific to v-Crk, we also stably expressed Bcl-2 in PC12 cells since Bcl-2 has been shown to confer resistance to apoptosis when overexpressed in PC12 cells (Mah *et al*, 1993) or SCG neurons (Garcia *et al*, 1992). Two independent PC12 clones, B1 and B10, showing highest levels of Bcl-2 expression (Figure 5B), were chosen for the viability assays. Consistent with previous findings (Mah *et al*, 1993), Bcl-2 partially protected differentiated PC12 cells from apoptosis in the absence of NGF. The degree of protection during NGF withdrawal in the v-Crk-expressing PC12 cells and Bcl-2-overexpressing PC12 cells were qualitatively similar, wherein approximately 70% of the cells in clone B10 were viable 72 h after NGF withdrawal (Figure 5A). Importantly, however, EGF replacement did not further enhance the survival benefits of Bcl-2 during NGF deprivation (Figure 5A) suggesting that v-Crk and Bcl-2 may operate through separate survival pathways.

Finally, to underscore that the effects of v-Crk were contingent on tyrosine kinase signaling, we utilized an SH2-inactive v-Crk mutant cell line R273N-v-Crk. R273N-v-Crk cannot bind phosphotyrosine-containing proteins and are slower than native PC12 cells in making neurites in response to 50 ng/ml of NGF (Teng *et al*, 1995). R273N-v-Crk PC12 cells are also susceptible to apoptosis in the absence of NGF and died at comparable rates to native PC12 cells following serum and NGF removal (Figure 3B).



**Figure 5** EGF replaces NGF as a neurotrophic factor in v-Crk-expressing PC12 cells but not Bcl-2-expressing PC12 cells. Native PC12 cells or PC12 cells expressing v-Crk (clone V1), human Bcl-2 (clones B1 and B10) or TrkA (clone 6-24) were maintained for 2 weeks in 3% serum plus 50 ng/ml NGF, after which serum and NGF were washed off, and replaced with 0% serum in the absence of growth factor (hashed symbol), 0% serum containing 50 ng/ml EGF (shaded symbol), or 50 ng/ml NGF (open symbol). The error bars in PC12 versus v-CrkPC12 cells denotes average  $\pm$  standard error of four independent experiments. **(B)** 50  $\mu$ g of total lysate from Human Bcl-2 transfected PC12 cell lines was Western blotted with an antibody specific for human Bcl-2 to indicate protein expression. The position of molecular weight standards (in kD) are indicated on the left and Bcl-2's migration is indicated on the right.



**Figure 6** Overexpression of TrkA suppress apoptosis. TrkA overexpressing cells (clone 6–24) (●) or native PC12 cells (□) were maintained as in Figure 2, and then transferred to NGF-depleted media for up to 72 h. (A) ethidium homodimer and calcein AM were utilized to determine the percentages of dead cells during NGF withdrawal. (B) the cells were grown on collagen-coated glass coverslips followed by NGF withdrawal for 48 h. Cells were DAPI stained as in Figure 2. The appearance of condensed chromatin in PC12 cells are indicated by the arrows.

Furthermore, EGF addition did not rescue R273N-v-Crk cells from apoptosis (data not shown). These data corroborate the above findings and suggest that PC12 survival advantage relies on intact signaling from receptor tyrosine kinases and may be related to the properties of v-Crk necessary for neurite outgrowth.

### v-Crk and Trk A delay apoptosis in the absence of cell cycle arrest

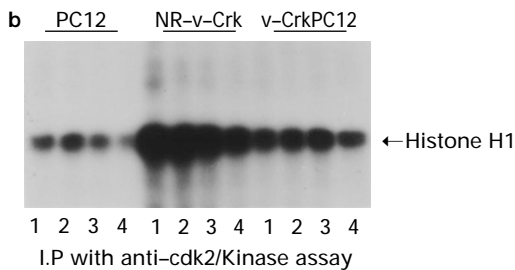
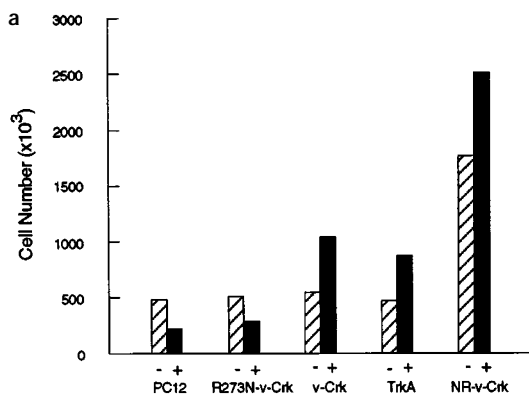
After several days of NGF treatment, PC12 cell proliferation typically ceases and differentiation occurs (Rudkin *et al*, 1989). Previous studies with G1-arrested PC12 cells or postmitotic sympathetic neurons support a hypothesis that postmitotic neurons and neuronal PC12 cells undergo apoptosis under NGF withdrawal by re-entering the cell cycle, an event referred to as a 'defective cell cycle' (Rubin *et al*, 1993). When PC12 cells were cultured for 11 days in 50 ng/ml NGF, NGF exerted an antiproliferative effect as evident from the lower number of total cells in 3% serum containing NGF as compared to 3% serum alone (Figure 7a). In contrast, v-Crk, but not mutant R273N-v-Crk, and NR-v-Crk cells increased in number upon NGF addition consistent with the mitogenic effects of wild type v-Crk. Trk 6-24 cells also proliferated in response to long-term NGF treatment. It is especially intriguing that greater than 90% of Trk 6-24 and v-Crk PC12 cells had processes of more than two cell bodies in diameter upon NGF treatment (data not shown). This suggests that significant crosstalk exists between differentiative and proliferative pathways in these cell lines.

The results shown in Figure 7 suggest that v-CrkPC12 cells may have failed to arrest in G1 phase relative to their native PC12 counterparts in the events leading up to NGF withdrawal. To probe further into the relationship between cell cycle progression and apoptosis, we analyzed cell cycle parameters by (i) FACS using propidium iodide for DNA content (Table 1), and (ii) cdk2 activity (a cyclin-dependent kinase required at the G1/S transition) (Figure 7b) following NGF withdrawal. For the flow cytometry, cells were grown in media with 3% serum and 50 ng/ml of NGF for 14 days and then placed in one of four conditions: (i) 3% serum+NGF for an additional 24 h; (ii) 0% serum + 50 ng/ml NGF for 24 h; (iii) 0% serum and no NGF for 12 h or (iv) 0% serum and no NGF for 24 h. Several points are noteworthy from these data. First, congruent with the aforementioned effects of v-Crk on cell growth, v-Crk and NR-v-Crk PC12 cells consistently had a higher percentage of cells in S phase compared to the native PC12 cells (Table 1). For example, 24 h after NGF removal, v-Crk-PC12 cells were 22% in S phase, compared to 15% in native cells. As expected, NR-v-Crk cells, which are the most proliferative (Figure 7a), had the greatest percentage of cells in S-phase, and this steadily increases through 24 h of NGF withdrawal. It is interesting that NR-v-Crk cells appear to have a progressive G2/M block at 12 and 24 h in growth factor free conditions suggesting that they may undergo apoptosis during S-phase. Second, Trk6-24 cells had a high percentage of cells in G2/M. These cells do not appear to be growth arrested at G2/M since there was no diminution of cells in S phase (Table 1). Finally, with the

**Table 1** FACS analysis of DNA content

Cell	%G <sub>1</sub>	%S	%G <sub>2</sub> /M
<i>PC12</i>			
3%+NGF	73	13	14
0%+NGF	70	14	16
0%× 24 h	75	15	10
<i>V15-PC12</i>			
3%+NGF	64	21	15
0%+NGF	66	23	11
0%× 12 h	72	15	14
0%× 24 h	69	22	9
<i>NRV1-C</i>			
3%+NGF	66	29	5
0%+NGF	54	37	8
0%× 12 h	49	47	4
0%× 24 h	47	52	1
<i>Trk6-24</i>			
3%+NGF	60	16	24
0%+NGF	60	14	27
0%× 12 h	60	11	29
0%× 24 h	58	13	29

Cells were treated as in Figure 2 and then placed in one of four conditions; (i) cultured in 3% serum plus NGF for an additional 24 h (ii) transferred to 0% serum plus NGF for 24 h, or transferred to 0% serum without NGF for (iii) 12 h or (iv) 24 h. Cells were harvested and stained with propidium iodide, and the nuclei were isolated and analyzed by flow cytometry. Results are representative of several independent experiments. The data are summarized from the DNA frequency distribution histograms. G<sub>0</sub>/G<sub>1</sub>, S, and G<sub>2</sub>/M cell populations are indicated



**Figure 7** Analysis of cell proliferation in PC12 cells. (a) Cell doubling time. Native PC12, R273N-v-Crk, v-Crk, TrkA, or NR-v-Crk PC12 cells were trypsinized and 10,000 cells seeded subconfluently in either 3% serum (hashed symbol) or 3% serum containing 50 ng/ml NGF (shaded symbol) for 11 days. The wells were counted on a hemacytometer and data represent the average of triplicate wells. (b) Cdk2 activity. PC12, NR-v-Crk, or v-Crk PC12 cells maintained in 3% serum plus NGF for 14 days (lane 1) were deprived of serum and growth factor for 4, 8 or 12 h (lanes 2–4, respectively). For each condition, 400  $\mu$ g of protein lysate was immunoprecipitated with anti-cdk2 antibody and assayed for kinase activity with 5  $\mu$ g of histone H1 as a substrate. The migration of the phosphorylated substrate is indicated by the arrow.

exception of NR-v-Crk cells, there did not appear to be any major fluctuation in cell cycle distribution within each line as a result of NGF withdrawal.

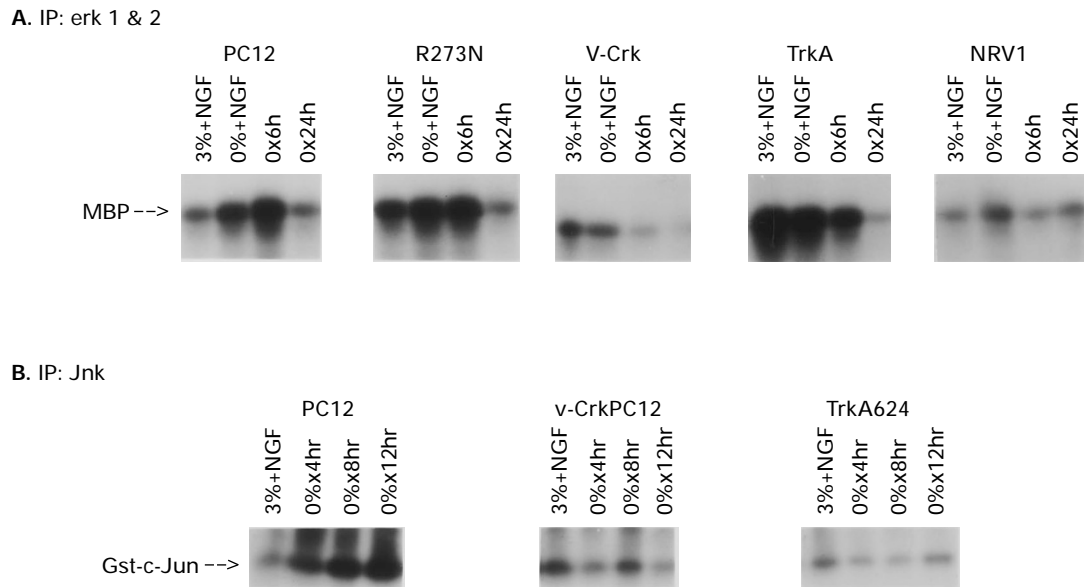
Conclusions from FACS are corroborated from the results of cdk2 kinase activity (Figure 7b). Native PC12 cells, which are known to down regulate kinase activity after exposure to NGF (Buchkovich and Ziff, 1994), show further diminution in activity after serum and NGF withdrawal. Most evident, however, is that the most striking differences in cdk2 activity occur between different cell lines. V-CrkPC12 cells had about twice the level of cdk2 activity per mg of cellular protein as native PC12 cells and, as expected, NR-v-Crk had dramatically higher cdk2 activity in all conditions. Similar to native PC12 cells, both NR-v-Crk cells and v-CrkPC12 cells exhibited modest diminution in cdk2 activity after serum or NGF withdrawal.

### Cell death in PC12 cells is associated with activation of JNK and MAP kinase

Recently, it has been reported that sustained c-Jun N-terminal kinase (JNK) activity and suppressed Erk/MAP kinase activity accompany PC12 apoptosis during NGF withdrawal (Xia *et al*, 1995). Taken together with our findings that v-Crk causes sustained activation of MAP kinase immediately after NGF stimulation (Teng *et al*, 1995), we examined the activation status of MAP kinase and JNK in Trk6-24, v-Crk, and native PC12 cells (Figure 8). Consistent with the results of Greenberg and colleagues, we observed a significant increase in JNK activity 6–8 h after NGF withdrawal in native PC12 cells. On the other hand, JNK activity was abated during NGF withdrawal in the Trk6-24 and v-Crk cells. Moreover, native PC12 cells and the R273N-v-Crk PC12 cells also had a significant increase in MAP kinase activity 6–8 h following NGF withdrawal while Trk6-24 and v-Crk PC12 cells did not show augmentation in MAP kinase activity. Thus, neuronal cell lines prone to apoptosis manifest an augmentation in MAP kinase and Jun Kinase activity within 8 h of growth factor removal, while cell lines resistant to apoptosis failed to demonstrate kinase activation (Table 2).

### Discussion

Both naive and NGF-differentiated (neuronal) PC12 cells undergo apoptotic cell death upon removal of serum and NGF, respectively (Lindenboim *et al*, 1995; Mesner *et al*, 1992). These properties of PC12 cells have therefore been proposed to mimic the dependence of mitotic and postmitotic neurons on growth factors in the developing nervous system. In our previous work, we established a model system to study NGF dependent neurogenesis by expressing the SH2/SH3-containing adapter protein v-Crk in PC12 cells (Hempstead *et al*, 1994; Teng *et al*, 1995). The present study documents the ability of v-Crk to delay apoptotic cell death in NGF-deprived neuronal PC12 cells. This capacity of v-Crk requires prior NGF priming since serum withdrawal on naive undifferentiated v-Crk-expressing PC12 cells did not yield a survival advantage over native PC12 cells. These data point to the importance of SH2/SH3-containing adapter proteins which can act on intracellular pathways of TrkA signaling and



**Figure 8** Activity of MAP Kinase and JNK during neuronal apoptosis. Native PC12 cells and the indicated transfectants were maintained in 3% serum plus 50 ng/ml of NGF for 2 weeks (lane 1 in each grouping) and then placed in either of three conditions: 0% serum+50 ng/ml NGF for an additional 24 h (lane 2); 0% serum without NGF for 6 h (lane 3); or 0% serum without NGF for 24 h (lane 4). 200  $\mu$ g and 400  $\mu$ g of total protein lysate for each condition was immunoprecipitated with specific antibodies for MAP Kinase (a) and JNK, respectively (b). The immunoprecipitate was then assayed for kinase activity using MBP (a) or a GST fusion protein containing the first 79 amino acids of c-Jun (b) as substrates. The positions of the phosphorylated substrates are indicated on the left.

**Table 2** Summary of cell lines utilized in this study. Relationships between kinetics of differentiation to NGF, S-phase response to NGF, and kinase activity and survival in absence of growth factor

Cell line	Kinetics of neurite outgrowth (+) NGF	Cell cycle response (+) NGF	JNK and MAPK activity (-) NGF	Survival (-) NGF 72 h
Native PC12	+	G <sub>1</sub>	increased	40%
R273N-v-crK	-	G <sub>1</sub>	increased	40%
NR-v-crK	0	S	no $\Delta$	35%
v-crK-PC12	++	S	decreased	70%
Trk 6-24	+++	S	decreased	95%

For native PC12 cells and the indicated transfectants, the following parameters were summarized: (i) kinetics of differentiation to NGF (column 1); (ii) cell cycle response to NGF (column 2); (iii) activities of JNK and MAP kinase after NGF withdrawal (column 3) and (iv) survival in the absence of NGF (column 4). In column 1, '+' signals a baseline velocity of neurite outgrowth, '++' and '+++ represent progressive acceleration, '-' indicates a dominant negative effect, and '0' indicates a total lack of NGF-induced neurite outgrowth. In column 2, 'G<sub>1</sub>' indicates more cell cycle arrest and 'S' indicates increased proliferation

transduce survival signals when cells are deprived of extracellular trophic support.

The ability of v-Crk to delay apoptosis specifically in growth factor-deprived neuronal PC12 cells, but not naive PC12 cells, suggests that v-Crk is an effector of the NGF signaling pathway and can act directly downstream to TrkA. This is not only supported by the fact that overexpression of TrkA completely renders cells NGF-independent for at least 5 days, as would be expected if TrkA were an upstream regulator of v-Crk, but by several other experimental observations including: (i) v-Crk induces neurite outgrowth in cells treated with a subthreshold concentration of NGF (Figure 1; Hempstead *et al*, 1994) (ii) v-Crk is tyrosine phosphorylated immediately following NGF stimulation (Hempstead *et al*, 1994) (iii) v-Crk augments NGF-induced GDP-GTP exchange on p21<sup>ras</sup>, a known downstream effector of TrkA (Teng *et al*, 1995) and (iv) expression of a dominant negative v-Crk (R273N-v-Crk), which carries a point mutation in the SH2 domain,

impairs NGF-induced neurite outgrowth (Teng *et al*, 1995), and as shown here, does not suppress apoptosis like wild-type v-Crk. Therefore, it is likely that v-Crk, by virtue of its ability to modulate TrkA tyrosine kinase activity/signaling after NGF removal, maintains a threshold of trophic support necessary for immediate PC12 survival. However, the augmentation of tyrosine kinase activity need not be restricted to TrkA, since restimulation of the EGF Receptor in NGF-deprived PC12 cells with EGF completely compensated for TrkA and yielded long-term survival. The fact that tyrosine kinase inhibitors staurosporine and K252A can induce apoptosis further argue that modulation of tyrosine kinase signaling can have striking effects on cell survival (Bertrand *et al*, 1994). Similarly, agonists which act on TrkA, such as GM1 ganglioside, have also been shown to significantly delay apoptosis (Ferrari *et al*, 1995). Recently, it has been shown that overexpression of wild-type PDGF-Receptor, but not a mutant PDGF-Receptor lacking the Phosphatidylinositol-3' Kinase (PI3'-Kinase)





binding site, confers PC12 survival, implicating a role for PI3-Kinase as a downstream effector of receptor kinases during protection from apoptosis (Yao and Cooper, 1995).

One hypothesis is that the pathways which augment differentiation via v-Crk may also play a role in enhancing v-Crk-mediated survival. This is supported by several lines of reasoning (Table 2). First, neuronal cell lines with accelerated kinetics of differentiation, such as Trk A or v-Crk-PC12 cells, also have enhanced survival curves in the absence of growth factor. In contrast, cells which are impaired in NGF-induced differentiation, such as R273N-v-Crk and NR-v-Crk PC12 cells, did not show any protection from neurotrophin-deprived apoptosis. Finally, EGF, which completely rescues v-CrkPC12 cells from apoptosis due to NGF deprivation, also promotes neurite outgrowth uniquely in v-CrkPC12 cells (Hempstead *et al*, 1994). Interestingly, long-term surviving Bcl-2 expressing cells maintained in the absence of NGF undergo neuronal differentiation suggesting a close mimicry between neuronal survival and neuronal differentiation (Sato *et al*, 1994). However, while NGF-differentiated PC12 cells expressing human Bcl-2 had delayed apoptosis comparable to v-Crk in the complete absence of growth factor, addition of EGF to these cultures did not further rescue them from apoptosis. Thus, we have provided evidence that one can predict the comparative viability after NGF withdrawal in PC12 transfectants based upon knowledge of their kinetics of differentiation to NGF.

Presently, we do not know the molecular mechanisms by which v-Crk potentiates NGF's effects on cell survival. Immediately following NGF or EGF addition to v-Crk-PC12 cells, v-Crk binds to the tyrosine phosphorylated TrkA Receptor (Hempstead *et al*, 1994) or EGF Receptor, respectively (Teng *et al*, 1995) and causes a sustained activation of p21<sup>ras</sup>, which apparently plays a role in potentiating differentiation (Teng *et al*, 1995). While activation of the Ras/Raf/MAP Kinase pathway is critical for the differentiation of PC12 cells (Kaplan and Stephens, 1994), the role of Ras/MAP kinase during apoptosis is a matter of controversy. While expression of dominant negative RasN17 does not prevent the killing of naive PC12 cells during serum deprivation (Yao and Cooper, 1995), RasN17 expressed from an inducible dexamethasone-inducible promoter prevented neuronal PC12 cell apoptosis after NGF deprivation, presumably by blocking cell cycle progression (Ferrari and Greene, 1994). In contrast, induction of apoptosis in NGF-differentiated PC12 cells could be markedly suppressed by expression of a constitutively activated MKK1, a dual specificity kinase that specifically phosphorylates and activates Erk/MAP kinase (Xia *et al*, 1995). In the experiments described here, MAP Kinase had diminished activity during NGF withdrawal in TrkA and v-Crk-expressing PC12 cells, but increased activity in native PC12 cells that are more prone to apoptosis. These data suggest that MAP Kinase activity may be of lesser importance for v-Crk and TrkA induced survival than for their differentiation. It is certainly possible that both Ras-dependent and Ras-independent apoptosis occurs (Borasio *et al*, 1993) depending upon cellular factors such as the initial cell cycle distribution. The MEKK and JNK pathways also appears to be critical components of

the apoptotic machineries (Johnson *et al*, 1996; Xia *et al*, 1995). NGF withdrawal leads to sustained increases in JNK activity and dominant-interfering forms of the enzyme (Xia *et al*, 1995) or its substrate c-Jun (Ham *et al*, 1995) leads to abrogation of growth factor requirements in PC12 cells and sympathetic neurons, respectively. Consistent with these data, we have shown that lines relatively resistant to apoptosis manifest decreases in JNK activity after NGF removal. Thus, v-Crk or TrkA might impinge on downstream serine/threonine kinases such as MAP Kinase or JNK1, creating a cellular milieu impervious to growth factor withdrawal.

A second hypothesis is that TrkA and Crk affect cell cycle progression in such a way that NGF removal is not accompanied by programmed death. Native PC12 cells continue to proliferate for 4 to 5 days following NGF stimulation prior to their G<sub>1</sub> arrest and subsequent differentiation. An emerging model, mentioned previously, that may partially explain the mechanism of apoptosis in postmitotic neurons following withdrawal of trophic support postulates that cells attempt an abortive transversal of cell cycle progression (Rubin *et al*, 1993). Based upon these considerations, one might predict that v-Crk or TrkA, like N17Ras (Ferrari and Greene, 1994) or p16<sup>INK</sup> (an inhibitor of cyclin D-dependent kinases) (Kranenburg *et al*, 1996), would protect PC12 cells by promoting growth arrest during long-term NGF treatment. However, we observed the opposite, whereby neuronal TrkA or v-Crk PC12 cells, despite appearing phenotypically differentiated, appear to by-pass normal G<sub>1</sub> checkpoints and maintain more robust cell cycle activity after 14 days in continuous NGF as measured by cell doubling analysis, flow cytometry, and cdk2 activity.

Because v-Crk-PC12 cells appear to be more mitogenically active than native PC12 cells after long-term NGF treatment in low serum, the relationships between cessation or cell proliferation, differentiation, and apoptosis may be more accurately viewed in terms of the cell cycle hypothesis for proliferating neuroblasts. In this model, mitogenic neuroblasts would retain cell cycle proteins and hence do not require *de novo* RNA synthesis prior to apoptosis (Lindenboim *et al*, 1995). Therefore, v-Crk may activate signaling events which support PC12 cells through subsequent cell cycle progression upon NGF and serum withdrawal, thereby achieving a delay in apoptosis. We should point out that there was no evidence of dramatic perturbations in cell cycle distribution in v-CrkPC12 cells within 24 h following a switch to growth factor free conditions, suggesting that these parameters are not modulated as a means of resistance to an apoptotic stimulus. However, we must reconcile the fact that v-Crk does not have the capacity to spare undifferentiated fully mitotic PC12 cells nor NR-v-Crk PC12 cells from apoptosis during serum removal and, as noted above, molecules related to v-Crk's role in differentiation likely play a role in exerting v-Crk's cytoprotective effects. Relevantly, apoptosis in serum-deprived mitotic PC12 cells is not restricted to the G<sub>0</sub>-G<sub>1</sub> phase of the cell cycle, and death can occur at S and G<sub>2</sub>/M phase (Lindenboim *et al*, 1995). However, since v-CrkPC12 cells are not initially G<sub>1</sub> arrested upon

differentiation, v-Crk's effects on apoptosis need not be restricted to one phase of the cell cycle and thus modulation of the cell cycle parameters may be less important than v-Crk's effects on signaling molecules. Of course, v-Crk expression or TrkA overexpression might foster cellular viability through means other than manipulation of cell cycle dynamics, differentiative pathways, or kinase pathways. Among other possibilities, v-Crk might lead to an inhibition of interleukin-1 $\beta$ -converting enzyme (ICE) or CPP32 activity, or up regulation of anti-apoptotic genes, such as Bcl-2 or Bcl-x (Nicholson *et al*, 1995; Gonzalez-Garcia *et al*, 1995; Motoyama *et al*, 1995).

The present study was undertaken to describe the role of v-Crk in modulating neurotrophic activity in neuronal PC12 cells. We have found a novel function for v-Crk and propose that adapter proteins may provide an important mechanism to enhance neurotrophin-dependent tyrosine kinase activity to influence survival and differentiation in the developing nervous system. Moreover, the findings that EGF can replace NGF as a neurotrophic factor in differentiated v-CrkPC12 cells may have broader implications concerning the role of EGF and the EGF Receptor in nonproliferating neurons in the nervous system. EGF can act as a local survival and differentiation factor in certain cerebellar primary neurons (Tucker *et al*, 1994) and the EGF Receptor has been detected in adult neurons from the cerebellum, cortex, and hippocampus (Gomez-Pinilla *et al*, 1988). Interestingly, our results here suggest that in the context of v-Crk, EGF Receptors and TrkA Receptors seem to perform identical functions with respect to neurotrophism. Therefore, an attractive hypothesis suggests that adaptor proteins, like Crk, which act stoichiometrically to strengthen receptor tyrosine kinases, may control the signaling events leading to mitosis, differentiation and survival. Future studies examining the role of Crk and other adapters as regulators of apoptosis *in vivo* should further our understanding of signal transduction.

## Materials and Methods

### Cell culture and transfection

Rat pheochromocytoma cells (PC12) were a kind gift from Dr. David Kaplan (NCI-Frederick) and maintained in Dulbecco's modified Eagle's medium (DMEM) supplemented with 7% heat inactivated calf serum and 3.5% heat inactivated horse serum at 37°C in a humidified atmosphere of 5% CO<sub>2</sub>. Stable PC12 transfectants expressing v-Crk (Hempstead, 1994), R273N-v-Crk (Teng *et al*, 1995), and TrkA (Hempstead *et al*, 1992) were generated as described. Parental cells plated on a rat tail collagen coated 100 mm plate were also transfected with 20  $\mu$ g of pSFFV *bcl-2nl* (obtained from Dr. Stanley Korsmeyer, Washington University) along with 20  $\mu$ g of salmon sperm DNA, and 50  $\mu$ l of lipofectamine (diluted 1:7 from the original stock) (Gibco). The plasmid pSFFV *bcl-2nl* contains a spleen focus-forming virus long terminal repeat driving the constitutive expression of a human *bcl-2* complementary DNA. Cells were selected in G418 (500  $\mu$ g/ml), subcloned and expanded. Nonresponding v-Crk PC12 cells (NR-v-Crk PC12) were isolated through limiting dilution from v-Crk PC12 cells maintained for approximately 4 months in DMEM supplemented with 2% calf

serum, 1% horse serum, and 50 ng/ml of EGF. Cells were diluted into a 96-well rat tail collagen coated plate at a density of approximately 1–10 cells/well, and cultured in 10% serum (6.5% calf, 3.5% horse) DMEM until individual colonies could be recovered and expanded. Those lines which did not differentiate in the presence of EGF were subcloned and expanded. In all cell cultures, the media was replaced every 48 h. As indicated in the text, media was supplemented with 50 ng/ml of either receptor grade mouse 2.5S NGF (Bioproducts for Science) or receptor grade mouse EGF (Bioproducts for Science). Prior to placement in growth factor-free media, cells were gently washed twice in large volumes of unsupplemented DMEM while attached to the collagen-coated plate.

### Viability/death assays

Percentages of live and dead cells were obtained through a two-color fluorescence assay (Molecular Probes, Eugene, OR) that measures (i) intracellular esterase activity (a sign of viability) via the enzymatic conversion of non fluorescent cell-permeant calcein AM to the intensely green calcein; and (ii) plasma membrane integrity by means of ethidium homodimer which enters damaged membranes and undergoes a 40  $\times$  enhancement of red fluorescence upon binding to nucleic acids. The assay was done on a 24 well rat tail collagen coated plate (Bioproducts for Science). For each condition, triplicate wells of adherent cells were gently washed in Phosphate Buffered Saline (PBS) supplemented with 9 mM CaCl<sub>2</sub> and 5 mM MgCl<sub>2</sub> (PBS<sup>+</sup>). Cells were then incubated at 37°C for 20 min in 250  $\mu$ l of 2 mM calcein AM and 4 mM ethidium homodimer in PBS<sup>+</sup>. After aspiration of the solutions, percentages of live and dead cells were obtained by counting approximately 200 cells/well from several random fields using a conventional 485 nm fluorescein excitation filter. For Hoechst staining of chromatin, cells were differentiated on Poly-D-Lysine 12 mm coverslips (Collaborative Biomedical #40425), after which cells were fixed with 3% paraformaldehyde, washed twice with PBS, and stained for 5 min at room temperature with an 8  $\mu$ g/ml solution of Hoechst 33258 (Sigma) in PBS. After a final wash, DAPI auto fluorescence was measured using a 365 nm Excitation light source with a BA420 Barrier filter and photographed on a Nikon microscope.

### Immunoprecipitation, Western blotting and kinase assays

For Western immunoblot analysis, cells were lysed in RIPA buffer (10 mM Tris HCl, 5 mM EDTA, 150 mM NaCl, 10% glycerol, 1% Triton X-100, 1% Deoxycholic acid and 0.1% SDS) containing 1 mM sodium vanadate, 1 mM phenylmethylsulfonyl fluoride (PMSF), 1% aprotinin, and 100  $\mu$ M molybdate. Equivalent concentrations of protein, as measured by Bio-Rad Protein Assay using bovine serum albumin (BSA) as a standard, were resolved by sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and transferred to Immobilon-P (Millipore). The membrane was blocked in Tris-Buffered Saline (TBS) containing 1% BSA and then incubated with the indicated primary antibody for approximately 3 h at room temperature. Blots were washed with TBS containing 0.5% Tween 20 (TBS-T), incubated with peroxidase-conjugated secondary antibody, and washed again with TBS-T. All blots were developed by using the Renaissance Enhanced Chemiluminescence system (Dupont). Antibodies against Crk or Gag were used as previously described (Teng *et al*, 1995) while mouse monoclonal anti-human Bcl-2 antisera was obtained from DAKO.

For immunoprecipitations followed by kinase assays, cells were lysed in nonionic lysis buffer (20 mM HEPES, 150 mM NaCl, 10%

glycerol) containing 1.0% Triton X-100 (HNTG buffer). 200  $\mu$ g of protein from each lysate was immunoprecipitated with one of the following antibodies conjugated to Protein A sepharose beads: 15  $\mu$ l of monoclonal antibody of cdk2 (Santa Cruz), 3  $\mu$ l of a 1:1 mix of rabbit polyclonal antibodies to Erk 1 and Erk2 (sc-163, Santa Cruz), or 3  $\mu$ l of a monoclonal antibody to JNK1 (Pharmingen). After being washed five times with HNTG buffer adjusted to 0.1% Triton X-100, the beads were resuspended in 50  $\mu$ l of wash buffer containing 5  $\mu$ Ci of  $^{32}$ P-ATP, 5  $\mu$ M cold ATP, 10 mM MnCl<sub>2</sub>, and 5  $\mu$ g of the substrate of interest (Histone H1 for cdk2, Myelin Basic Protein for Erk 1 and 2, and Gst-c-Jun residues 1–79 for JNK1). The mixture was rocked for 45 min at room temperature, followed by centrifugation in a microfuge. The supernatant was run on 12.5% SDS-polyacrylamide gel and radioactivity was detected by autoradiography of the fixed gels.

### Cell cycle analysis

For analysis of cell doubling times, trypsinized cells were plated on a 6-well rat-tail collagen-coated plates at 10,000 cells per well and were cultured as indicated in the presence or absence of exogenous growth factors. At the indicated times, cells were removed from the plate with phosphate-buffered saline (PBS) containing trypsin-EDTA, and counted in a hemacytometer. To measure the cell cycle parameters, fluorescence activated cell sorting (FACS) was performed. To circumvent the problem of cell clumping during PC12 differentiation, nuclei were isolated from the cells prior to staining. Cells were grown in 150 cm plates, trypsinized, and centrifuged at 1000 rpm for 10 min. They were resuspended to a concentration of  $2 \times 10^6$  cells/ml in a 4 mM citrate buffer, pH 7.8, containing 50  $\mu$ g/ml of propidium iodide (PI, Sigma), 180 units/ml of RNase (Worthington), 0.1% Triton-X 100, and 30  $\mu$ g/ml polyethylene glycol (PEG). After a 20 min incubation at 37°C, an equal volume of a 0.04 M NaCl buffer containing 50  $\mu$ g/ml PI, 0.1% Triton-X, and 30 mg/ml PEG was added. The mixture was kept in the dark for at least 6 h at 4°C. Cellular DNA was measured on a Coulter Eliter Analyzer using 488 nm excitation and a 630 long pass filter for the collection of PI fluorescence. The fluorescence was gated on single cells to minimize the effect of clumping on fluorescence intensity. Cell cycle analysis was performed with Multicycle software (Phoenix Flow Systems, San Diego, CA); the curve fitting program permitted the measurement of each phase of the cell cycle after the histograms were corrected for a background consisting of cellular debris.

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