

Phosphorylation of c-Jun Is Necessary for Apoptosis Induced by Survival Signal Withdrawal in Cerebellar Granule Neurons

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Cerebellar granule neurons die by apoptosis when deprived of survival signals. This death can be blocked by inhibitors of transcription or protein synthesis, suggesting that new gene expression is required. Here we show that *c-jun* mRNA and protein levels increase rapidly after survival signal withdrawal and that transfection of the neurons with an expression vector for a c-Jun dominant negative mutant protects them against apoptosis. Phosphorylation of serines 63 and 73 in the c-Jun transactivation domain is known to increase c-Jun activity. By using an antibody specific for c-Jun phosphorylated on serine 63, we show that this site is phosphorylated soon after survival signal withdrawal. To determine whether c-Jun phosphorylation is necessary for apoptosis, we have expressed c-Jun phosphorylation site mutants in granule neurons. c-Jun^{asp}, a constitutively active c-Jun mutant in which the known and potential

serine and threonine phosphoacceptor sites in the transactivation domain have been mutated to aspartic acid, induces apoptosis under all conditions tested. In contrast, c-Jun^{ala}, which cannot be phosphorylated because the same sites have been mutated to alanine, blocks apoptosis caused by survival signal withdrawal. Finally, we show that cerebellar granule neurons contain high levels of Jun kinase activity and low levels of p38 kinase activity, neither of which increases after survival signal withdrawal. Mitogen-activated protein kinase activity decreases under the same conditions. These results suggest that c-Jun levels and c-Jun phosphorylation may be regulated by novel mechanisms in cerebellar granule neurons.

Key words: AP-1; apoptosis; cerebellar granule neurons; c-Jun; Jun kinase; signal transduction; stress-activated protein kinases

During the development of the mammalian nervous system approximately half of the neurons that are formed subsequently die by apoptosis (Oppenheim, 1991). This death is usually the result of the limited availability of specific neurotrophic factors, which are required for the survival of developing neurons, and is thought to be a mechanism for ensuring that neuronal targets are innervated by the correct density of neurons (Barde, 1989). In the case of several different types of primary neuron cultured *in vitro*, inhibitors of transcription or protein synthesis block cell death caused by survival factor withdrawal (Martin et al., 1988; Scott and Davies, 1990; D'Mello et al., 1993; Milligan et al., 1994), suggesting that survival signal removal may activate genes whose products promote cell death (Johnson and Deckwerth, 1993). Initial studies of the transcriptional control of cell death in nerve growth factor (NGF)-dependent sympathetic neurons demonstrated that the transcription factor c-Jun plays a key role. Inhibition of the function of c-Jun, either by microinjection of antibodies against c-Jun or by expression of a c-Jun dominant negative mutant, protected sympathetic neurons from NGF withdrawal-induced death (Estus et al., 1994; Ham et al., 1995).

In the present study, we have investigated whether c-Jun is necessary for apoptosis in cerebellar granule neurons. When isolated from 8-d-old rats, cerebellar granule neurons can be maintained *in vitro* by adding 10% serum and 25 mM KCl to the culture medium (D'Mello et al., 1993). If, after 7 d *in vitro*, the serum is removed, and the KCl concentration is reduced from 25 to 5 mM, the granule neurons die by apoptosis, and this death is transcription-dependent (D'Mello et al., 1993). Miller and Johnson (1996) reported that the level of *c-jun* RNA increases in cerebellar granule neurons after KCl and serum deprivation but did not investigate whether the activity of c-Jun was required for cell death. Here we show that after survival signal withdrawal, *c-jun* RNA and protein levels increase before the transcriptional commitment point and that apoptosis can be inhibited by expressing a c-Jun dominant negative mutant. The transcriptional activity of c-Jun is increased by phosphorylation of serines 63 and 73 in the transactivation domain (Pulverer et al., 1991; Smeal et al., 1991). Using a phospho-c-Jun-specific antibody, we demonstrate that c-Jun is phosphorylated on serine 63 during apoptosis, and, by expressing c-Jun mutants in which specific phosphorylation sites have been altered, we show that phosphorylation of c-Jun is necessary for apoptosis to occur after survival signal withdrawal. Finally, we have measured the activity in granule neuron extracts of Jun amino terminal kinases (JNKs), also known as stress-activated protein kinases (SAPKs), which phosphorylate serines 63 and 73 in c-Jun (Dérjard et al., 1994; Kyriakis et al., 1994), and the activities of p38 kinase and mitogen-activated protein (MAP) kinase. The results of these

Received April 9, 1997; revised Oct. 1, 1997; accepted Nov. 5, 1997.

This work was financed by the Eisai Company of Japan. We thank Chantal Bazenet, Alan Hall, and Moshe Yaniv for useful discussions and critical reading of this manuscript and Joanne Taylor for advice on the preparation of cerebellar granule neurons. We are also grateful to Dr. Angel Nebreda for providing the mpk2/p38 antibody and to Dr. Dirk Bohmann for the expression vectors for c-Jun, c-Jun^{ala}, and c-Jun^{asp}.

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assays suggest that in cerebellar granule neurons c-Jun protein levels and c-Jun phosphorylation may be regulated by novel mechanisms.

MATERIALS AND METHODS

Cell culture. Cerebellar granule neurons were isolated from the cerebella of 8-d-old Sprague Dawley rats (supplied by the Biological Services Unit, University College London) as described by Taylor et al. (1997). The neurons were separated from non-neuronal cells by centrifugation at $1200 \times g$ for 20 min through 40.5% Percoll (Sigma, Poole, UK) and were plated in basal medium Eagle (BME; Life Technologies, Paisley, UK) supplemented with 10% fetal calf serum (Gibco, Esher, UK), 25 mM KCl, 35 mM glucose, and penicillin/streptomycin on polyornithine-coated dishes or glass coverslips. Cells were plated at a density of $5.6 \times 10^5/cm^2$. Approximately 24 hr after plating, cytosine arabinofuranoside (Sigma) was added to the culture medium to a final concentration of 10 μM to prevent the proliferation of any non-neuronal cells. Using this protocol 95–99% of the cultured cells were neurons (Hatten, 1985; Gao et al., 1991). Apoptosis was induced by reducing the extracellular potassium concentration from 25 to 5 mM as follows. Cells that had been cultured for 6–7 d *in vitro* were rinsed three times in serum-free BME containing 5 mM KCl supplemented with glucose and penicillin/streptomycin and then were maintained in the same medium. Control cultures were treated identically but were maintained in serum-free medium supplemented with KCl at 25 mM. Neuronal survival was assessed by MTT (Sigma) conversion to formazan by live cells (Mosmann, 1983) or, on the basis of nuclear morphology, visualized by staining paraformaldehyde-fixed cells with Hoechst dye (H33342, Calbiochem-Novabiochem UK Ltd.).

PC12 cells were cultured in a defined medium supplemented with 2% fetal calf serum and 10 $\mu g/ml$ insulin as described by Ham et al. (1995). HeLa and Rat1 cells were cultured in DMEM (Life Technologies) with 10% FCS. For treatment with UV radiation, HeLa cells were grown to confluence and then left in DMEM with 0.5% FCS overnight. The cells were exposed to short-wavelength UV radiation (254 nm) for 1 min using a hand-held UV lamp and were harvested 30 min later. Subconfluent Rat1 cells were treated in a similar manner.

Immunoblotting. Whole cell extracts were prepared from cultured cerebellar granule neurons by lysing cells in SDS buffer containing 1 mM PMSF, 1 $\mu g/ml$ pepstatin A, 5 $\mu g/ml$ leupeptin, and 2 $\mu g/ml$ aprotinin, as described by Ham et al. (1995). The resulting lysate was then cleared by centrifugation. Proteins were separated on 12.5% SDS polyacrylamide gels. Fifteen micrograms of extract were loaded per lane. After electrophoresis, the separated proteins were electroblotted onto Hibond ECL nitrocellulose (Amersham, Little Chalfont, UK). Jun and Fos family members were detected with affinity-purified rabbit polyclonal antibodies as described previously (Ham et al., 1995; Lallemand et al., 1997) using a horseradish peroxidase-conjugated anti-rabbit secondary antibody and the ECL system (Amersham). Relative levels of protein were determined by scanning autoradiographs on an imaging densitometer (Bio-Rad).

In vitro-translated c-Jun phosphorylated on serine 63 was detected with an affinity-purified rabbit polyclonal phospho-c-Jun antibody that was raised against a peptide corresponding to mouse c-Jun amino acids 57–68 with a phosphorylated serine at position 63 (D. Lallemand, unpublished observations). To detect c-Jun phosphorylated on serine 63 in extracts prepared from cerebellar granule neurons, a sequential immunoprecipitation and immunoblotting experiment was performed. For these experiments, cells were lysed in high-salt lysis buffer (50 mM HEPES, pH 7.0, 500 mM NaCl, and 1% NP-40) supplemented with PMSF, leupeptin, aprotinin, 10 mM NaF, and 100 μM Na orthovanadate. After incubation on ice for 20 min, the cell extract was centrifuged at $13,000 \times g$ for 5 min, and the supernatant was transferred to a fresh tube and stored at $-80^\circ C$. For immunoprecipitations, protein A-agarose beads (Boehringer Mannheim) were washed three times in NET buffer (50 mM Tris, pH 7.5, 150 mM NaCl, 0.1% NP-40, and 1 mM EDTA). Three hundred micrograms of cell extract were then made up to a final volume of 700 μl and a final NaCl concentration of 150 mM using NET buffer with no NaCl. Fifty microliters of washed protein A beads were then added, and the tube was rotated for 1 hr at $4^\circ C$. The beads were then spun down, and the supernatant was transferred to a fresh tube. Phospho-c-Jun polyclonal antibody (1.5 μg) was then added together with 50 μl of washed protein A-agarose beads, and the tube was then rotated at $4^\circ C$ for 3 hr. The beads were then spun down and washed three times with NET buffer. After the final wash, the beads were resuspended in 30 μl of Laemmli

sample buffer, boiled for 5 min, and then spun for 15 min. The supernatant was loaded onto a 12.5% SDS-polyacrylamide gel. After electrophoresis, the gel was transferred to nitrocellulose, and immunoblotting was performed using a phospho-c-Jun monoclonal antibody (diluted 1:1000) that had been raised against the same phosphopeptide as that used for preparing the polyclonal antibody and that has a similar specificity (Lallemand, unpublished observations).

In vitro transcription and translation. c-Jun protein was produced *in vitro* using a TnT rabbit reticulocyte lysate or wheat germ extract system (Promega UK Ltd., Southampton, UK), according to the manufacturer's instructions. The plasmid pCdc-Jun (Ham et al., 1995) was used as template and contains the full-length mouse c-Jun open reading frame with a Kozak consensus initiation codon cloned downstream of the β -globin RNA leader sequence and bacteriophage T7 promoter. The β -globin leader has been shown to increase the efficiency of translation *in vitro* (Norman et al., 1988).

Immunofluorescence. Immunofluorescence experiments were performed with cells plated on glass coverslips. Cells were fixed in 3% paraformaldehyde for 30 min at room temperature, permeabilized with 0.5% Triton X-100, and then were blocked using 50% goat serum in 1% BSA in PBS. Primary and secondary antibodies were diluted in 10% goat serum in 1% BSA in PBS.

The polyclonal phospho-c-Jun antibody was used at a dilution of 1:250 for 16 hr at $4^\circ C$. FLAG Δ 169 was detected with the M2 monoclonal antibody (IBI Kodak, Cambridge, UK) diluted 1:200. Bcl-2 was detected with a monoclonal antibody (clone 124) diluted 1:50 (Dako, Glostrup, Denmark). β -Galactosidase was detected using a rabbit polyclonal antibody (5 Prime \rightarrow 3 Prime, Inc., Boulder, CO) diluted 1:500. c-Jun^{asp} and c-Jun^{ala} were detected with a hemagglutinin (HA) monoclonal antibody (clone 12CA5, Boehringer Mannheim) diluted 1:2000. Fluorescein- or rhodamine-conjugated goat anti-mouse and goat anti-rabbit secondary antibodies (Jackson ImmunoResearch, West Grove, PA) were used at a dilution of 1:100. After incubation with the primary and secondary antibodies, nuclei were stained with Hoechst dye (H33342) at 1 $\mu g/ml$. Coverslips were mounted in Citifluor (Citifluor, Canterbury, UK). Slides were viewed on a Nikon Microphot FXA fluorescence microscope. Color photographs were taken on Ektachrome 400x slide film (Kodak).

Northern analysis. Total RNA was prepared by the acid guanidinium thiocyanate phenol-chloroform method as described by Chomczynski and Saachi (1987). Twenty-microgram aliquots of total RNA were electrophoresed on 1.2% agarose gels containing formaldehyde, transferred to Hibond N⁺ membranes (Amersham), and hybridized with ³²P-labeled probes according to standard protocols (Sambrook et al., 1989). Membranes were stripped and reprobed with GAPDH to ensure equal loading between lanes. The probes used to detect the *c-jun* and GAPDH mRNAs were prepared from plasmids containing the mouse *c-jun* and rat GAPDH cDNAs, which were obtained from Moshe Yaniv (Institut Pasteur, Paris, France). The relative level of *c-jun* RNA was determined by scanning autoradiographs on an imaging densitometer (Bio-Rad).

Plasmids and transfection. Construction of pCDFLAG Δ 169 and pCDBcl-2 was described previously (Ham et al., 1995). Cytomegalovirus (CMV) expression vectors for HA epitope-tagged c-Jun, c-Jun^{asp}, and c-Jun^{ala} were provided by Dirk Bohmann (EMBL, Heidelberg, Germany), and CMVlacZ was provided by Art Alberts (ICRF, London, UK).

For transfection experiments, all plasmids were purified by centrifugation on two CsCl gradients. Transient transfection of granule neurons was performed by the calcium phosphate coprecipitation method (Graham and van der Eb, 1973) using a ProFection kit (Promega) according to the manufacturer's instructions. Briefly, neurons were cultured for 4 d *in vitro* on glass coverslips in 24-well dishes. pcDNA1, pCDFLAG Δ 169, pCDBcl-2, wild-type c-Jun, c-Jun^{asp}, or c-Jun^{ala} were added to the transfection mix at a final concentration of 5 $\mu g/ml$ together with 1 $\mu g/ml$ of an expression vector encoding β -galactosidase (CMVlacZ) to allow detection of the transfected cells. Calcium phosphate-DNA precipitates were prepared by standard procedures and were allowed to form at room temperature for 30 min. Fifty microliters of precipitate were added to the cells in 450 μl of the original conditioned medium. After 8 hr, the cells were subjected to a 15% DMSO shock for 90 sec at room temperature and then were washed three times with conditioned medium (from cells cultured for 4–7 d *in vitro*). The transfected cells were left for 36 hr in conditioned medium to allow expression of plasmid DNA and also to ensure that the cells were fully dependent on potassium for survival. The medium was changed after the 36 hr, and the cells were fixed an additional 24 hr later. The transfection efficiency was determined in immunofluorescence experiments by calculating the percentage of

Hoechst-stained cells that expressed β -galactosidase and was typically 3–7%. Furthermore, immunofluorescence analysis was performed to confirm that β -galactosidase was co-expressed with c-Jun, FLAG Δ 169, or Bcl-2, and that the proteins were localized correctly within neurons after transfection.

To analyze the effect of expression vectors on neuronal survival, the transfected cells were fixed and stained with an antibody against β -galactosidase and with Hoechst dye to visualize nuclear morphology. The viability of the transfected cells was calculated by scoring the percentage of β -galactosidase-expressing cells that had normal (nonpyknotic) nuclei. Coverslips were scored blind.

In vitro kinase assays. The cells were solubilized with lysis buffer (20 mM HEPES, pH 7.4, 2 mM EGTA, 1% Triton X-100, 10% glycerol, 1 mM DTT, 1 mM sodium orthovanadate, 50 mM β -glycerophosphate, 1 mM PMSF, 1 μ g/ml pepstatin A, 2 μ g/ml aprotinin, and 10 μ g/ml leupeptin) for 15 min on ice and then centrifuged at $13,000 \times g$ for 15 min at 4°C. For immunoprecipitations, 50 or 100 μ g of cell extract was precleared with 50 μ l of protein A-agarose (Boehringer Mannheim), diluted 1:1 in lysis buffer, for 1 hr at 4°C. Endogenous JNK, p38, or MAP kinase was immunoprecipitated with specific polyclonal antibodies and 50 μ l of protein A-agarose for 2–3 hr at 4°C. One to 2 μ l of anti-JNK (SAPK) antiserum (Kyriakis et al., 1994), 2 μ l of anti-p38 (*Xenopus* Mpk2) antibody (Rouse et al., 1994), and 1 μ g of anti-MAP kinase (MAPK) antibody (rat MAPK R2; Erk1-CT from Upstate Biotechnology, Lake Placid, NY) was used for 50–100 μ g of cell extract. The immunoprecipitates were washed three times with lysis buffer, two or three times with wash buffer I (100 mM Tris-HCl, pH 7.6, 500 mM LiCl₂, 0.1% Triton X-100, and 1 mM DTT), and two or three times with wash buffer II (25 mM HEPES, pH 7.5, 0.2% Triton X-100, and 1 mM EDTA) before they were suspended in 30 μ l of kinase buffer (in mM: 25 HEPES, pH 7.5, 20 MgCl₂, 20 β -glycerophosphate, 20 *p*-nitrophenylphosphate, 0.1 orthovanadate, and 2 DTT). MAPK and p38 immune complexes were washed once or twice with kinase buffer before the assay. The kinase reaction was started by the addition of 5 μ Ci of [γ -³²P]ATP (>5000 Ci/mmol), cold ATP to a final concentration of 20 μ M, and 2–3 μ g of glutathione *S*-transferase (GST)-cJun[1–169] (A. Eilers, unpublished observations), 2 μ g of GST-activating transcription factor 2 (ATF2) [1–96] (Santa Cruz Biotechnology, Santa Cruz, CA), or 10 μ g of myelin basic protein (Sigma) as substrate, respectively. After incubation for 15–30 min at 30°C, the reaction was terminated by the addition of Laemmli sample buffer. The phosphorylation of the substrate proteins was examined after SDS-PAGE by autoradiography.

RESULTS

Transcriptional commitment point

The death of cerebellar granule neurons after KCl and serum deprivation can be delayed significantly when actinomycin D is included in the medium at concentrations that inhibit transcription (D'Mello et al., 1993; Galli et al., 1995; Schulz et al., 1996; Armstrong et al., 1997) (A. Watson, unpublished observations). As a first step toward understanding the transcriptional regulation of cell death in this system, we determined the transcriptional commitment point, defined as the time after removal of survival signals (serum and 25 mM KCl) at which only 50% of the neurons kept in 5 mM KCl can be rescued by the addition of actinomycin D. A previous report suggested that the transcriptional commitment point was relatively early in cerebellar granule neurons (Galli et al., 1995). To confirm that this was the case in our cell culture system and to determine the transcriptional commitment point more precisely, we performed a detailed analysis at early time points after the removal of survival signals (Fig. 1). Granule neurons were switched to serum-free medium containing 5 mM KCl, and actinomycin D or additional KCl was added to 1 μ g/ml and 25 mM, respectively, at different times after the medium had been changed. Survival was then assessed in an MTT assay 48 hr after the initial reduction in KCl concentration. As can be seen in Figure 1, 50% of the neurons could not be rescued by the inhibition of RNA synthesis after 3 hr in 5 mM KCl, suggesting that in these cells a gene or set of genes encoding proteins that

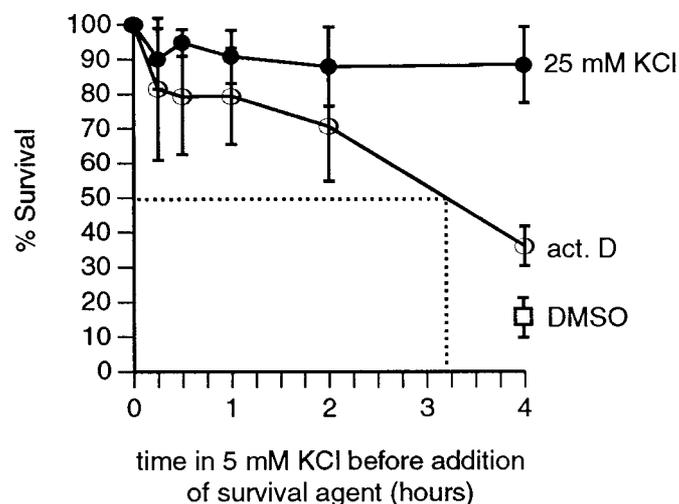


Figure 1. Determination of the transcriptional commitment point for cerebellar granule neurons undergoing apoptosis after withdrawal of survival signals. Cerebellar granule neurons plated in 96-well cell culture dishes were rinsed three times with serum-free BME containing 5 mM KCl and then were left in the same medium. At different times after reduction of the extracellular KCl concentration, actinomycin D (*act. D*; open circles) or additional KCl (filled circles) was added to final concentrations of 1 μ g/ml or 25 mM, respectively. As a control, the vehicle for actinomycin D, DMSO, was added to 0.1% (open square). Survival was assessed 48 hr later in an MTT assay. 100 corresponds to the MTT value after 48 hr for the 25 mM KCl 0 hr time point. The results shown represent the average of five independent experiments \pm SE.

trigger the cell death program had been transcribed by this time. In contrast, the addition of KCl to 25 mM could rescue 90% of the cells as late as 4 hr after the initial reduction in KCl concentration, because the KCl commitment point was later than 4 hr (~12 hr; data not shown).

c-Jun is selectively induced before the transcriptional commitment point

In experiments with NGF-dependent sympathetic neurons it has been established that *c-jun* RNA and protein levels increase after NGF withdrawal and that the activity of c-Jun is necessary for apoptosis (Estus et al., 1994; Ham et al., 1995). To determine whether this was also true in cerebellar granule neurons, we investigated which members of the Jun and Fos family were expressed and whether their pattern of expression changed on induction of apoptosis. To do this we performed immunoblotting experiments using affinity-purified polyclonal antibodies raised against the mouse Jun and Fos proteins (Lallemand et al., 1997). Cerebellar granule neurons were refed with serum-free medium containing 5 or 25 mM KCl. Whole-cell extracts were prepared at various times after the medium had been changed, and immunoblots were performed. Like sympathetic neurons, granule neurons expressed c-Jun, Jun B, and Jun D but not c-Fos or Fos B (Fig. 2A) (data not shown). When the cells were switched into medium containing 5 mM KCl, the level of c-Jun protein increased between 2 and 4 hr after the medium had been changed, but there was no change in cells maintained in 25 mM KCl, indicating that the increase was specifically attributable to the decrease in KCl concentration rather than the removal of serum. Increased levels of c-Jun protein were observed at 4 and 8 hr, but by 24 hr the amount of c-Jun had decreased to the same level as that in cells that had been kept in 25 mM KCl (Fig. 2B). In contrast, the levels of Jun B and Jun D did not increase when the cells were cultured

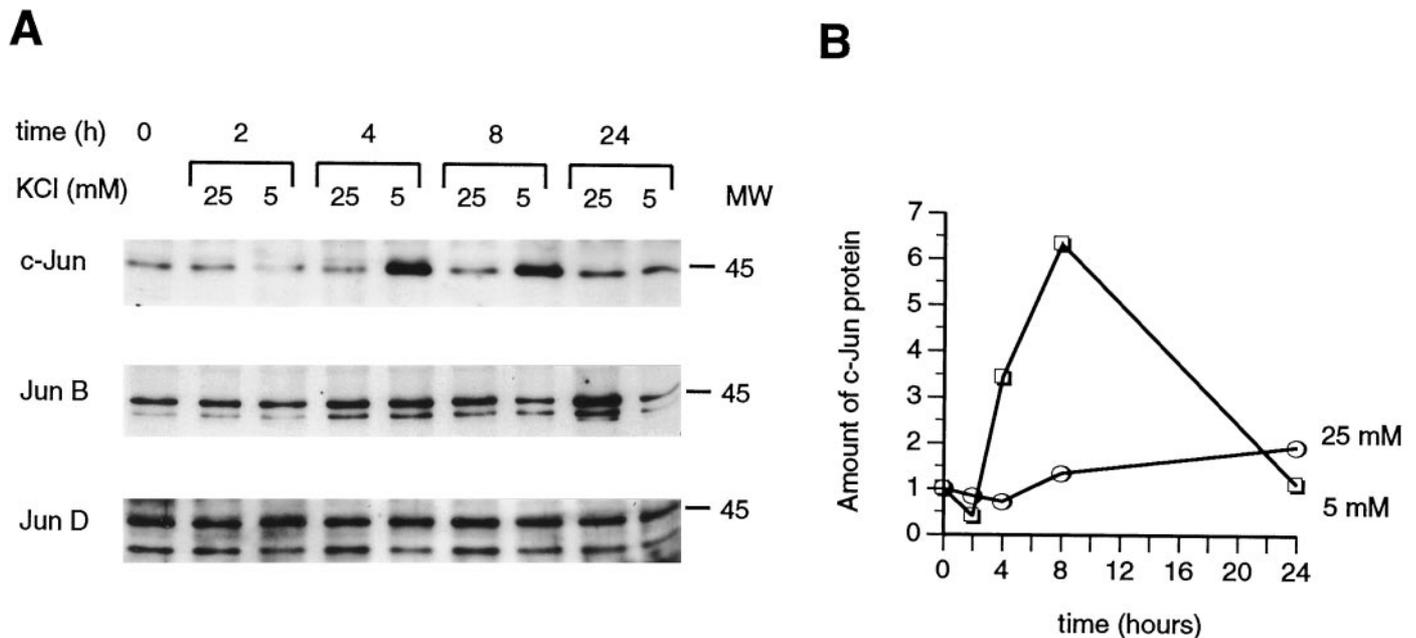


Figure 2. Pattern of expression of c-Jun, Jun B, and Jun D in cerebellar granule neurons deprived of survival signals. *A*, Cerebellar granule neurons were rinsed three times with serum-free BME supplemented with 5 mM KCl and were either maintained in the same medium or were refed with serum-free BME containing 25 mM KCl. At the times indicated, the cells were harvested, and whole-cell extracts were prepared. Proteins were separated on 12.5% SDS polyacrylamide gels, and immunoblotting was performed using affinity-purified polyclonal antibodies specific for c-Jun, Jun B, and Jun D (Lallemand et al., 1997). The positions and sizes in kilodaltons of molecular weight markers (*MW*) that were run in parallel are shown on the right. c-Fos and Fos B were not detected in granule neuron extracts, although the c-Fos and Fos B antibodies recognized *in vitro*-translated c-Fos and Fos B in immunoblotting experiments (data not shown). *B*, The c-Jun immunoblot shown in *A* was scanned on a densitometer, and the relative amount of c-Jun protein present in extracts from cells cultured in 25 mM KCl (open circles) or 5 mM KCl (open squares) was plotted against time.

in 5 mM KCl medium. There was, however, some increase in the level of Jun B protein in cells cultured in 25 mM KCl, which might be a response to the removal of serum or to the addition of fresh medium. Finally, c-Fos and Fos B were still not detected after survival signal withdrawal (data not shown). Therefore, c-Jun was the only one of the AP-1 proteins examined that was affected significantly by reducing the KCl concentration, and the increase in c-Jun level started between 2 and 4 hr after the removal of survival signals.

To determine whether the increase in the level of c-Jun protein was preceded by an increase in the amount of *c-jun* mRNA, we performed northern blotting experiments with RNA isolated from granule neurons. Two *c-jun* mRNAs with sizes of 2.7 and 3.4 kb were detected (Fig. 3*A*). These have been detected previously in fibroblasts and PC12 cells and are thought to differ in their 3' ends (Ryseck et al., 1988; Bartel et al., 1989). Lowering the KCl concentration caused an increase in the level of *c-jun* mRNA within 1 hr, which reached a peak at 2 hr and then started to decline (Fig. 3*A,B*). The *c-jun* mRNA was therefore induced before the transcriptional commitment point (3 hr), and this induction was specific for cells in 5 mM KCl and was not accompanied by any change in GAPDH mRNA (Fig. 3*A*).

AP-1 activity is required for cell death in cerebellar granule neurons

We showed previously that microinjection of sympathetic neurons with an expression vector for a c-Jun dominant negative mutant (pCDFLAGΔ169) that lacks the transactivation domain and that can suppress AP-1 activity delayed cell death after NGF withdrawal (Ham et al., 1995). To determine whether interfering with c-Jun function would also inhibit apoptosis in granule neurons, we tested the effect of expressing FLAGΔ169 in these cells. As a

means of introducing expression vectors into cerebellar granule neurons, we developed a method for transiently transfecting them based on calcium phosphate co-precipitation. pCDFLAGΔ169 was transfected into granule neurons together with a β-galactosidase expression vector, CMVlacZ. The transfected cells could therefore be detected by staining with an anti-β-galactosidase antibody. As a positive control we used a Bcl-2 expression vector, because overexpression of Bcl-2 has been shown to slow the death of a number of neuronal types deprived of survival signals (Garcia et al., 1992; Allsopp et al., 1993). As a negative control we used the empty CMV expression vector pCDNA1. Figure 4*A* shows representative granule neurons transfected with CMVlacZ and pCDFLAGΔ169 or pCDBcl-2, which were fixed and stained with a β-galactosidase antibody and antibodies against the FLAG epitope or Bcl-2, respectively. Both proteins were correctly localized; FLAGΔ169 was found in the nucleus, and Bcl-2 was found in the cell body.

To determine whether expression of FLAGΔ169 or Bcl-2 could protect granule neurons against cell death induced by survival signal withdrawal, cells that had been cultured for 4 d *in vitro* were transfected with the appropriate expression vectors. Thirty-six hours after transfection, the cells were refed with serum-free medium containing 5 mM KCl. Twenty-four hours later, the cells were fixed, and neuronal viability was determined as follows; transfected cells were identified by staining with anti-β-galactosidase antibody, and nuclear morphology was visualized by Hoechst staining. Cells with condensed, pyknotic nuclei, shrunken cell bodies, and fragmented neurites were scored as apoptotic. Figure 4*B* shows cells that were transfected with CMVlacZ and pCDNA1 or pCDFLAGΔ169 and that were cultured in serum-free medium containing 5 mM KCl before being fixed and

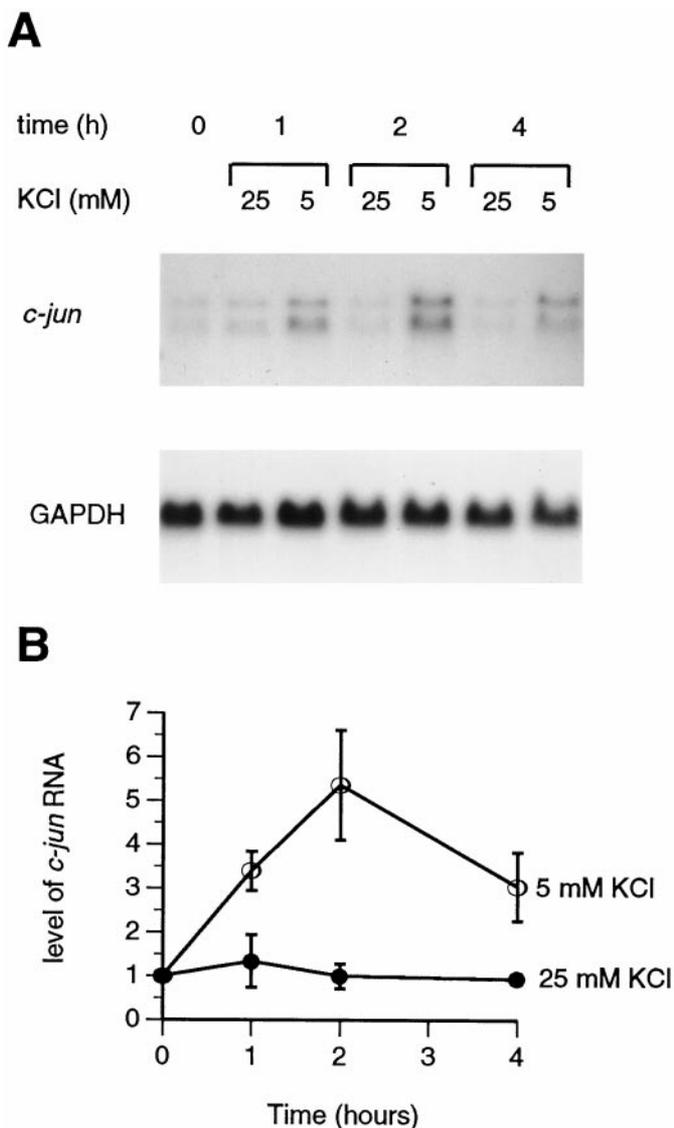


Figure 3. The *c-jun* RNA increases in level before the transcriptional commitment point. *A*, Cerebellar granule neurons were switched into serum-free BME containing either 25 or 5 mM KCl, and total RNA was isolated at the times indicated. The RNA samples were electrophoresed on a 1.2% agarose gel containing formaldehyde and then were transferred to a Hibond N⁺ membrane. The filter was hybridized with a ³²P-labeled *c-jun* probe. After autoradiography, the filter was stripped and rehybridized with a GAPDH probe. *B*, *c-jun* RNA levels were determined by scanning autoradiographs on a densitometer, and the relative level of *c-jun* RNA in either 25 mM KCl (filled circles) or 5 mM KCl (open circles) was plotted against time. The data shown represent the average of two independent Northern hybridization experiments ± SE.

stained. Examples of normal (Fig. 4*B*, *N*) and apoptotic (Fig. 4*B*, *A*) transfected cells are marked with arrows. After 24 hr in 5 mM KCl only 18% of the neurons transfected with pcDNA1 displayed the characteristics of normal cells, with round cell bodies and normal nuclei (Fig. 4*C*). In contrast, 93% of the cells transfected with pCDBcl-2 and 73% of the cells transfected with pCD-FLAGΔ169 were viable. Furthermore, expression of FLAGΔ169 or Bcl-2 had no adverse effect on neuronal viability in medium containing 25 mM KCl (Fig. 4*C*). These results demonstrate that AP-1 activity is necessary for cell death in granule neurons, because expression of a c-Jun dominant negative mutant, which

inhibits AP-1 activity (Ham et al., 1995), prevented cell death. Furthermore, because c-Jun was the only member of the AP-1 family that increased in level after survival signal withdrawal, and because this occurred before the transcriptional commitment point, it seems likely that c-Jun plays an important role in the death of these neurons.

c-Jun is phosphorylated on serine 63 during apoptosis

The activity of the c-Jun protein is regulated by phosphorylation at specific sites (Karin, 1995). In particular, serines 63 and 73 in the transactivation domain play a key role. Phosphorylation of these residues increases the ability of c-Jun to activate the transcription of target genes (Pulverer et al., 1991; Smeal et al., 1991). To determine whether c-Jun was phosphorylated on serine 63 after survival signal withdrawal, we made use of a phospho-c-Jun-specific antibody that only recognizes c-Jun when it is phosphorylated at serine 63 and that does not recognize unphosphorylated c-Jun or phosphorylated Jun B or Jun D (D. Lallemand, unpublished observations). The specificity of this antibody is demonstrated in Figure 5*A*. When translated in a rabbit reticulocyte lysate, c-Jun becomes phosphorylated on serines 63 and 73 and threonines 91 and 93 and runs as a ladder of bands (Pulverer et al., 1991). In contrast, c-Jun protein synthesized in a wheat germ extract system is not phosphorylated and runs as a single band. In immunoblotting experiments with these *in vitro*-translated proteins, a normal c-Jun antibody raised against amino acids 1–58 of c-Jun (Lallemand et al., 1997) recognized both forms of c-Jun, whereas the phospho-c-Jun antibody only detected phosphorylated c-Jun (Fig. 5*A*, *c-Jun RRL*).

We then used the phospho-c-Jun antibody in immunofluorescence experiments with cerebellar granule neurons (Fig. 5*B*). Control cultures that had been maintained in 25 mM KCl in the absence of serum showed little, if any, staining with the phospho-c-Jun antibody (data not shown). This was also the case immediately after the cells had been switched into serum-free medium containing 5 mM KCl (Fig. 5*B*, *Oh*), whereas 1 hr after KCl and serum deprivation, phosphorylated c-Jun localized to the nucleus was readily detected (Fig. 5*B*). This staining pattern was not observed when the phospho-c-Jun antibody was incubated with the phosphopeptide that had been used to make the antibody but was unaffected by the addition of the equivalent nonphosphorylated peptide (data not shown). Thus, in granule neurons deprived of survival signals, specific nuclear phospho-c-Jun staining appeared before c-Jun protein levels had increased significantly (Fig. 2). The intensity of the staining had increased further at 4 hr, paralleling the increase in c-Jun protein detected by immunoblotting. As judged by Hoechst staining, the phosphorylation of c-Jun on serine 63 occurred before any of the nuclear changes characteristic of apoptotic cells.

The immunofluorescence results with the phospho-c-Jun antibody were confirmed by performing a sequential immunoprecipitation and immunoblotting experiment using extracts from granule neurons and polyclonal and monoclonal phospho-c-Jun antibodies (Fig. 5*C*). In pilot experiments we found that to obtain a satisfactory signal in immunoblots with cell extracts, it was necessary to enrich for phospho-c-Jun by first immunoprecipitating with a polyclonal phospho-c-Jun antibody before performing immunoblotting with a monoclonal phospho-c-Jun antibody. This may be because the phospho-c-Jun antibody recognizes its epitope more efficiently when the c-Jun protein is not denatured. As a control, we used extracts from unstimulated and UV-irradiated Rat1 cells. As expected, the phospho-c-Jun monoclo-

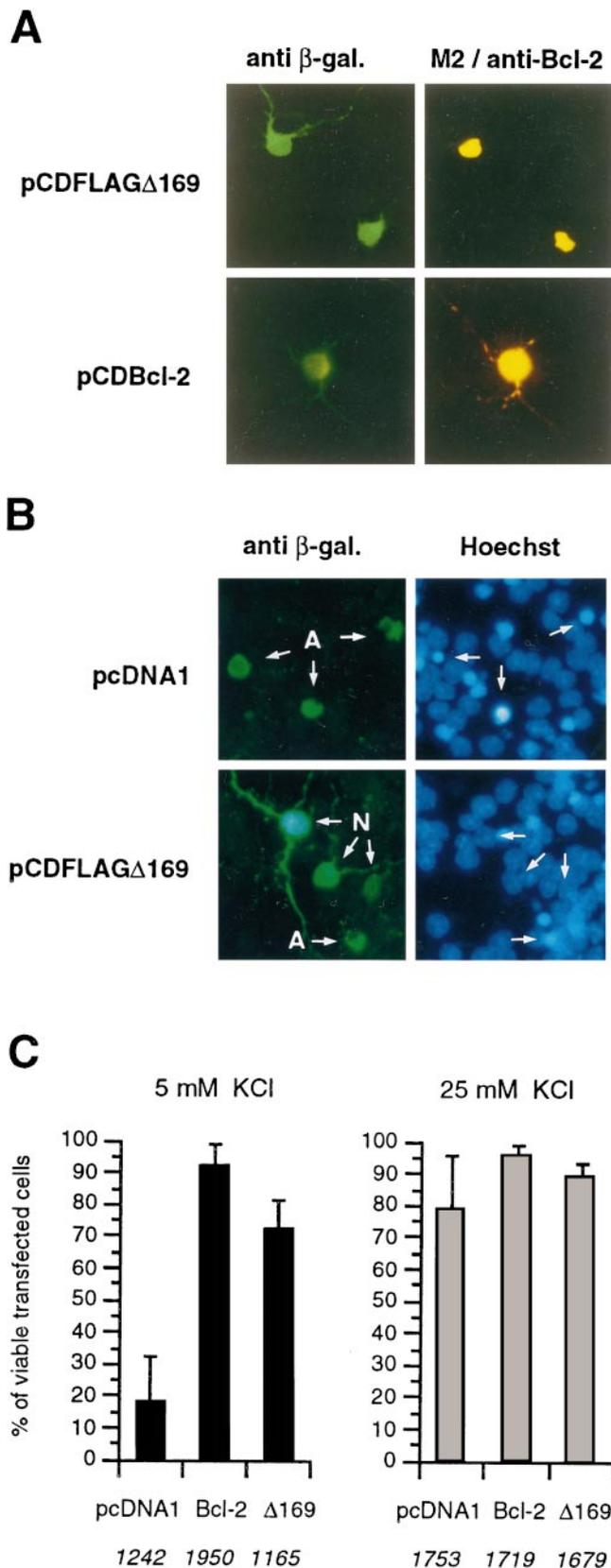


Figure 4. Overexpression of Bcl-2 or the c-Jun dominant negative mutant FLAG Δ 169 protects cerebellar granule neurons against cell death induced by survival signal withdrawal. *A*, Cerebellar granule neurons cultured on glass coverslips were transfected with pCDFLAG Δ 169 or pCDBcl-2 together with the β -galactosidase expression vector CMVlacZ, as described in Materials and Methods. The cells were subsequently fixed

and stained with an antibody against β -galactosidase to identify the transfected cells and with the FLAG-specific M2 antibody or an anti-Bcl-2 antibody. FLAG Δ 169 localized to the nucleus, as judged by comparison with Hoechst staining (results not shown). Bcl-2 localized to the cell body and neurites. *B*, Morphology of normal and apoptotic cerebellar granule neurons. Cells were transfected with CMVlacZ and pcDNA1 or pCDFLAG Δ 169. Thirty-six hours after transfection the cells were switched into 5 mM KCl medium. After fixation, the transfected cells were identified by staining with an anti- β -galactosidase antibody, and nuclear morphology was visualized by Hoechst staining. Apoptotic (*A*) and normal (*N*) cells are marked by white arrows. Apoptotic cells have a highly condensed, brightly staining nucleus. *C*, Cerebellar granule neurons were transfected with CMVlacZ plus pcDNA1 or pCDBcl-2 or pCDFLAG Δ 169. Thirty-six hours after transfection, the cells were refed with serum-free medium containing either 5 or 25 mM KCl. Twenty-four hours later the cells were fixed and stained with an anti- β -galactosidase antibody and Hoechst dye. The percentage of transfected cells that were viable, i.e., had a normal morphology, was then determined. The results shown are the average of nine independent experiments \pm SE. For each construct the total number of transfected cells that were scored is indicated. The coverslips were counted in a blinded manner.

Phosphorylation of c-Jun is necessary for cell death

To determine whether phosphorylation of the c-Jun transactivation domain was necessary for the induction of apoptosis in granule neurons, cells were transfected with expression vectors for either wild-type c-Jun or c-Jun^{asp}, a constitutively active mutant in which the known and potential serine and threonine phosphorylation sites in the activation domain have been mutated to aspartic acid, or c-Jun^{ala}, an inactive protein that cannot be activated by phosphorylation, because the same residues have been mutated to alanine (Fig. 6*A*) (Papavassiliou et al., 1995). HA epitope-tagged c-Jun, c-Jun^{asp}, and c-Jun^{ala} were expressed efficiently in transfected cells, as determined by immunofluorescence with an anti-HA antibody (data not shown). In the presence of 10% serum and 25 mM KCl, overexpression of wild-type c-Jun or c-Jun^{ala} had little effect on cell viability, whereas the majority (80%) of the cells transfected with c-Jun^{asp} had pyknotic nuclei. When serum was removed, expression of wild-type c-Jun induced apoptosis in 65% of the transfected cells, whereas the alanine mutant and pcDNA1 did not cause a significant increase in the percentage of apoptotic cells (Fig. 6*B*). Thus overexpression of wild-type c-Jun alone was sufficient in itself to induce apoptosis under certain conditions (25 mM KCl, no serum). In contrast, expression of the constitutively active c-Jun^{asp} induced apoptosis efficiently under all conditions, whereas c-Jun^{ala}, which cannot be phosphorylated, was unable to induce cell death under any of the conditions tested. Thus the ability of the various forms of c-Jun to induce apoptosis in granule neurons correlated with their potential phosphorylation status.

Previous work by Treier et al. (1995) and Peverali et al. (1996) showed that a *Drosophila* Jun^{ala} mutant could act as a dominant

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and stained with an antibody against β -galactosidase to identify the transfected cells and with the FLAG-specific M2 antibody or an anti-Bcl-2 antibody. FLAG Δ 169 localized to the nucleus, as judged by comparison with Hoechst staining (results not shown). Bcl-2 localized to the cell body and neurites. *B*, Morphology of normal and apoptotic cerebellar granule neurons. Cells were transfected with CMVlacZ and pcDNA1 or pCDFLAG Δ 169. Thirty-six hours after transfection the cells were switched into 5 mM KCl medium. After fixation, the transfected cells were identified by staining with an anti- β -galactosidase antibody, and nuclear morphology was visualized by Hoechst staining. Apoptotic (*A*) and normal (*N*) cells are marked by white arrows. Apoptotic cells have a highly condensed, brightly staining nucleus. *C*, Cerebellar granule neurons were transfected with CMVlacZ plus pcDNA1 or pCDBcl-2 or pCDFLAG Δ 169. Thirty-six hours after transfection, the cells were refed with serum-free medium containing either 5 or 25 mM KCl. Twenty-four hours later the cells were fixed and stained with an anti- β -galactosidase antibody and Hoechst dye. The percentage of transfected cells that were viable, i.e., had a normal morphology, was then determined. The results shown are the average of nine independent experiments \pm SE. For each construct the total number of transfected cells that were scored is indicated. The coverslips were counted in a blinded manner.

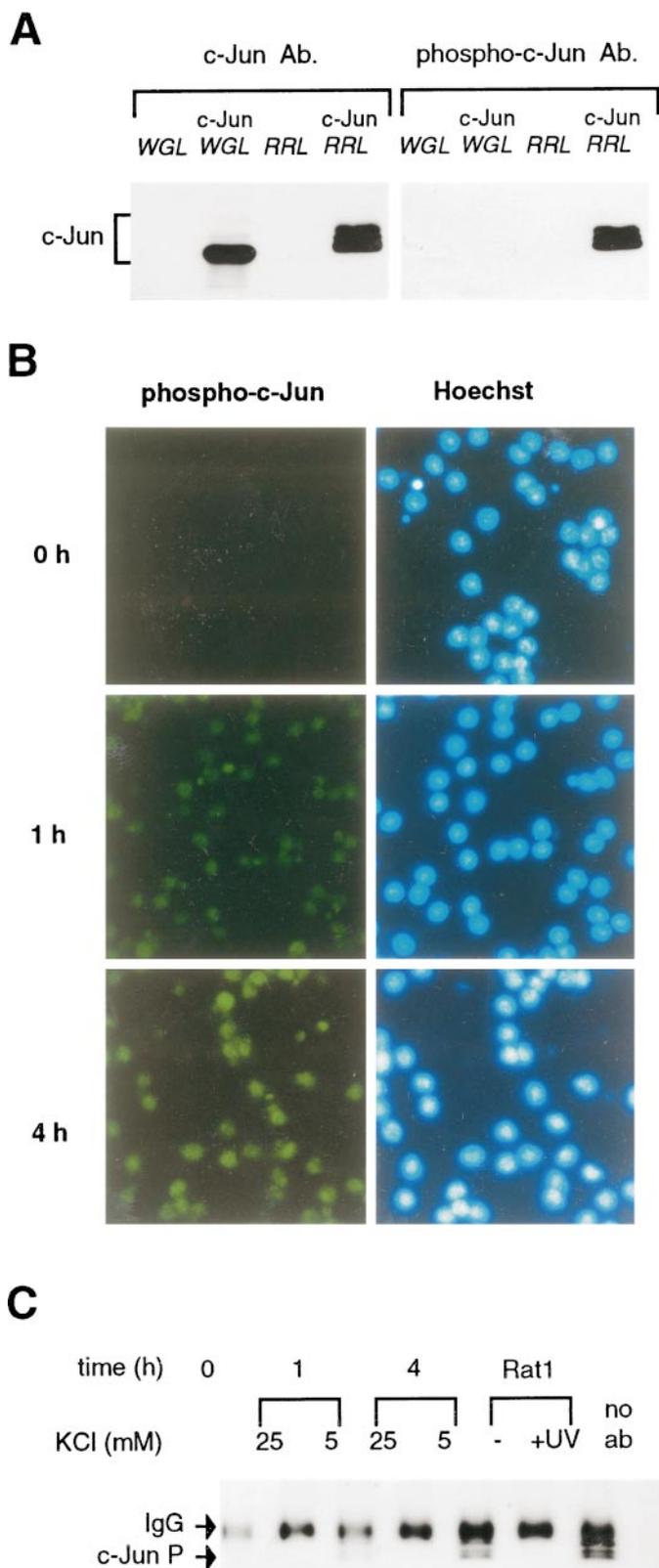


Figure 5. c-Jun becomes phosphorylated on serine 63 soon after cerebellar granule neurons are deprived of survival signals. *A*, c-Jun protein was synthesized in a wheat germ lysate transcription-translation system (*c-Jun WGL*) or a rabbit reticulocyte lysate (*c-Jun RRL*). c-Jun translated in the wheat germ lysate is not phosphorylated and runs as a single band, whereas c-Jun translated in the rabbit reticulocyte system is phosphorylated at serines 63 and 73 and threonines 91 and 93 and runs as a ladder of bands with a lower mobility in SDS polyacrylamide gels. The *in*

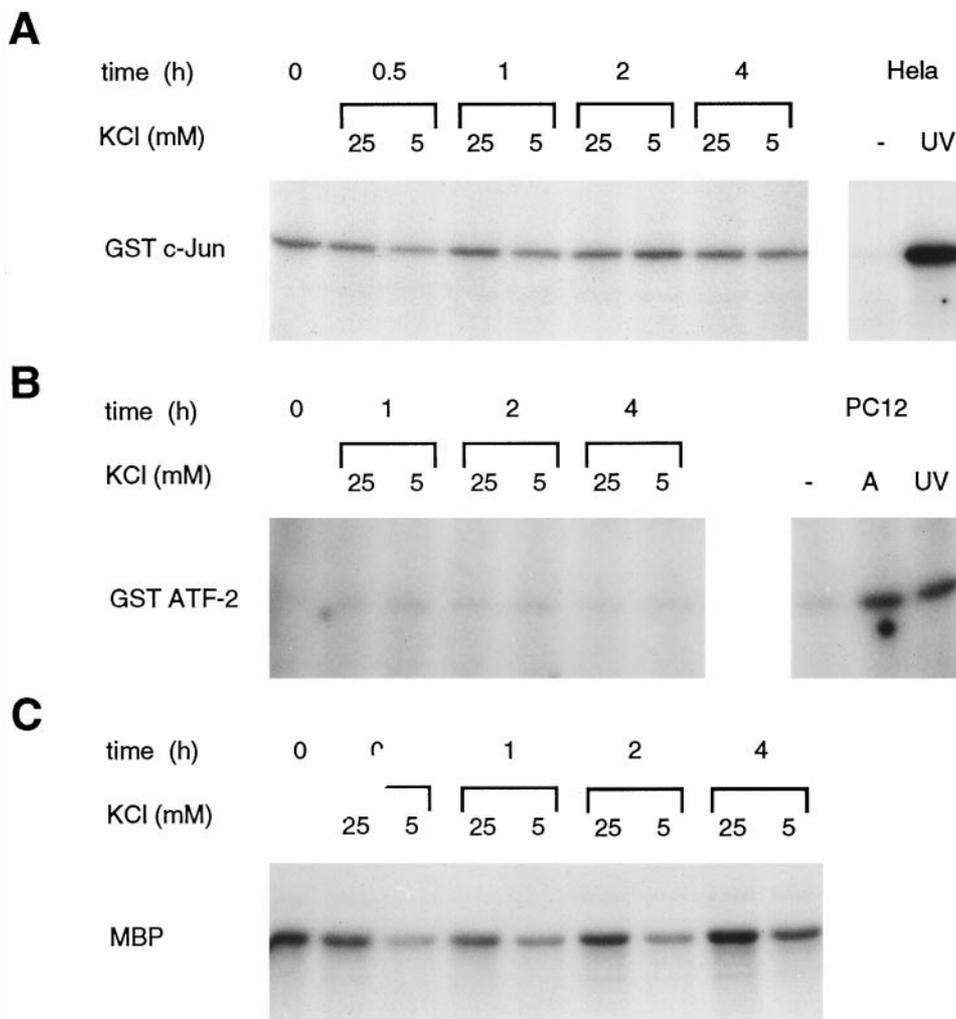
inhibitor of R7 photoreceptor differentiation. To determine whether c-Jun^{ala} could block apoptosis in granule neurons, cells were transfected with c-Jun^{ala} and then were induced to die by removing serum and lowering the extracellular KCl concentration. After 24 hr in 5 mM KCl, only a minority of the cells transfected with pcDNA1, wild-type c-Jun or c-Jun^{asp} were viable (10% in this series of experiments). In contrast, 50% of the cells transfected with c-Jun^{ala} were viable (Fig. 6*B*). c-Jun^{ala} appeared to block cell death less efficiently than the c-Jun deletion mutant FLAGΔ169 (compare Figs. 4*C*, 6*B*). This may be because c-Jun^{ala} can still activate transcription weakly (Metivier et al., 1993), whereas FLAGΔ169 cannot. In conclusion, these results indicate that expression of c-Jun, in particular the activated form c-Jun^{asp}, is sufficient in itself to induce apoptosis in cerebellar granule neurons cultured in the presence of survival signals. Furthermore, it appears that phosphorylation of c-Jun is necessary for apoptosis induced by survival signal withdrawal, because overexpression of c-Jun^{ala}, which cannot be phosphorylated, delays cell death.

Cerebellar granule neurons contain a high level of Jun kinase activity, which does not increase after survival signal withdrawal

Members of the MAPK superfamily are known to regulate the activity of the *c-jun* promoter and c-Jun protein (Karin, 1995). Jun kinases (JNK/SAPKs) bind directly to c-Jun with high affinity and phosphorylate serines 63 and 73, thereby increasing the transcriptional activity of the c-Jun protein (Pulverer et al., 1991; Smeal et al., 1991; Dérijard et al., 1994; Kyriakis et al., 1994). Jun kinases can also increase the rate of transcription of the *c-jun* gene, because c-Jun binds as a heterodimer with ATF-2 to TRE-like sites in the *c-jun* promoter, and the transactivation domains of both proteins are phosphorylated by Jun kinase (Van Dam et al., 1993, 1995; Herr et al., 1994). Three JNK genes have been identified, which through alternative splicing give rise to multiple isoforms (Kyriakis et al., 1994; Gupta et al., 1996). p38 kinase, which can also be activated by stress, does not phosphorylate the

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in vitro-translated proteins and corresponding unprogrammed lysates (*WGL* and *RRL*) were electrophoresed on a 10% SDS polyacrylamide gel. After transfer to nitrocellulose, immunoblots were performed with a c-Jun antibody that was raised against amino acids 1–58 of c-Jun (*c-Jun Ab.*) diluted 1:400 or with the polyclonal phospho-c-Jun-specific antibody diluted 1:2000. *B*, Cerebellar granule neurons cultured on glass coverslips were rinsed with serum-free BME containing 5 mM KCl and then were maintained in the same medium. At the time points indicated, the cells were fixed and stained with a phospho-c-Jun-specific antibody and Hoechst dye. Nuclear phospho-c-Jun staining was clearly visible 1 hr after survival signal withdrawal and had increased in intensity at 4 hr. The specificity of the staining pattern was confirmed by peptide competition (data not shown). Furthermore, granule neurons cultured in BME plus 10% FCS plus 25 mM KCl did not stain with the phospho-c-Jun antibody (data not shown). *C*, Cerebellar granule neurons were rinsed with serum-free BME containing 5 mM KCl and then were maintained in the same medium or in medium supplemented with 25 mM KCl. At the time points indicated, extracts were prepared, and phospho-c-Jun was immunoprecipitated using a polyclonal phospho-c-Jun antibody, as described in Materials and Methods. The resulting immunoprecipitates were electrophoresed on a 12.5% SDS-polyacrylamide gel and transferred to nitrocellulose, and immunoblots were performed using a monoclonal phospho-c-Jun antibody. The position of phosphorylated c-Jun (*c-Jun P*) and of a nonspecific band that probably corresponds to IgG is indicated. Extracts from untreated and UV-irradiated Rat1 cells were used as controls. Antibody was also omitted from a duplicate Rat1 (+UV) immunoprecipitation as a further control (*no ab*).

Figure 7. Measurement of Jun kinase, p38 kinase, and MAP kinase activity in extracts from cerebellar granule neurons deprived of survival signals. Cerebellar granule neurons were rinsed three times and then were maintained in serum-free BME containing either 25 or 5 mM KCl. At the times indicated, extracts for kinase assays were prepared as described in Materials and Methods. **A**, Jun kinases were immunoprecipitated from cell extracts using a pan-SAPK antibody that recognizes all of the different rat Jun kinase isoforms (Kyriakis et al., 1994). After washing of the immunoprecipitates, the activity of the kinases was assayed using GST-c-Jun[1–169] as substrate. As a control, extracts from quiescent (–) or UV-treated (UV) HeLa cells were assayed in parallel. **B**, p38 kinase was immunoprecipitated from cell extracts using a *Xenopus* mpk2/p38 antibody, which also recognizes mammalian p38 (Rouse et al., 1994). After the immunoprecipitates had been washed, p38 kinase activity was assayed using GST-ATF-2[1–96] as substrate. Control extracts were prepared from undifferentiated PC12 cells, which had been pretreated with anisomycin at 10 μ g/ml for 30 min (A) or irradiated with short-wavelength (254 nm) UV light (UV). **C**, MAP kinases were immunoprecipitated from cell extracts using an antibody that recognizes multiple ERK isoforms (rat MAPK R2; Erk1-CT from Upstate Biotechnology). MAP kinase activity was assayed using myelin basic protein (MBP) as substrate.



against SAPK β (JNK3) that also recognizes the other members of the rat JNK/SAPK family (Kyriakis et al., 1994). The activity of the immunoprecipitated kinase was assayed using purified GST-c-Jun[1–169] as substrate (Fig. 7A). As controls we used extracts from quiescent and UV-irradiated HeLa cells, because Jun kinase is known to be strongly activated by UV treatment in these cells (Hibi et al., 1993). As predicted, we found that quiescent HeLa cells contained very little Jun kinase activity, whereas UV treatment caused a strong induction of JNK activity (Fig. 7A). When we assayed extracts from granule neurons, we found, surprisingly, that these cells had high basal levels of JNK activity and that this did not increase after KCl deprivation but, instead, remained more or less constant. The autoradiographs obtained in three independent experiments were scanned on an imaging densitometer, and the average values were calculated (A. Watson, unpublished observations). On average there was no significant increase in the high basal level of JNK activity after survival signal withdrawal. Similar results were obtained when we assayed nuclear, rather than whole-cell, extracts prepared from granule neurons (data not shown).

To immunoprecipitate p38 kinase we used an antibody prepared against the *Xenopus* mpk2/p38, which also recognizes mammalian p38 (Rouse et al., 1994). To measure the activity of the immunoprecipitated kinase we used GST-ATF-2[1–96] as substrate. p38 kinase activity can be induced strongly by treating cells

with anisomycin or UV radiation, and we found that this was the case when we tested extracts from PC12 cells that had been treated in this way (Fig. 7B). Like unstimulated PC12 cells, cerebellar granule neurons only had low levels of p38 kinase activity. However, this did not increase when the cells were refed with serum-free medium containing 5 mM KCl. Finally, we performed MAP kinase assays using a pan-MAPK antibody, which recognizes several members of the MAPK family, and myelin basic protein as the substrate. We found that granule neurons contained readily detectable levels of MAPK activity, which decreased after survival signal withdrawal (Fig. 7C). In conclusion, none of the kinases examined was activated when granule neurons were refed with low KCl medium. Rather, the cells appeared to contain very high constitutive levels of JNK activity and low levels of p38 activity, neither of which increased when the extracellular KCl concentration was reduced. In contrast, MAP kinase activity decreased after survival signal withdrawal.

DISCUSSION

In the case of several different types of primary neuron, cell death induced by survival signal withdrawal can be delayed dramatically by the addition of actinomycin D at concentrations that will inhibit transcription. This has suggested the hypothesis that the withdrawal of survival signals leads to the transcriptional induction of genes that encode proteins that promote cell death. Are

similar genes activated in different kinds of neurons after survival signal withdrawal? We and others previously have studied the pattern of expression of members of the AP-1 family in sympathetic neurons undergoing apoptosis and found that *c-jun* RNA and protein levels increased selectively after NGF withdrawal (Estus et al., 1994; Ham et al., 1995). *c-jun* RNA levels also increased in differentiated PC12 cells deprived of NGF (Mesner et al., 1995) and in cerebellar granule neurons after KCl and serum deprivation (Miller and Johnson, 1996). Here we have demonstrated that in cerebellar granule neurons the *c-jun* RNA is induced rapidly after survival signal withdrawal, peaking at 2 hr, before the transcriptional commitment point (3 hr; Fig. 1), and that the level of c-Jun protein increases. Furthermore, we showed that AP-1 activity was necessary for cell death, because expression of FLAG Δ 169, a c-Jun dominant negative mutant that inhibits AP-1 activity, increased the percentage of viable cells after survival signal withdrawal. Expression of FLAG Δ 169 also protected sympathetic neurons or PC12 cells against NGF withdrawal-induced death (Ham et al., 1995; Xia et al., 1995). In *c-jun* $-/-$ knockout mice, programmed cell death occurs normally in dorsal root ganglion (DRG) neurons at least up to embryonic day 11.5 (Roffler-Tarlov et al., 1996). However, because these mice die during midgestation, it has not been possible to use them to study apoptosis in postnatal sympathetic neurons or cerebellar granule neurons. It is possible that the death of embryonic DRG neurons may not be affected in these animals, because other members of the AP-1 family or other transcription factors, such as ATF-2, can substitute for c-Jun. Alternatively, the signal transduction pathways that activate programmed cell death in embryonic DRG neurons may be different from those functioning in postnatal sympathetic and cerebellar granule neurons.

The transcriptional activity of the c-Jun protein is increased by phosphorylation of serines 63 and 73 in the transactivation domain. Using a phospho-c-Jun-specific antibody we showed that in granule neurons c-Jun was phosphorylated on serine 63 soon after the extracellular KCl concentration had been lowered, suggesting that c-Jun activity increases during apoptosis. To determine whether phosphorylation of c-Jun was important for cell death in granule neurons, we investigated the effect of overexpressing different c-Jun mutants in which the phosphorylation sites had been altered. A constitutively active mutant in which the serine and threonine phosphoacceptor sites in the transactivation domain had been mutated to aspartic acid was able to induce apoptosis when expressed in granule neurons cultured in medium containing 10% serum and 25 mM KCl, whereas the wild-type c-Jun protein could not. On the other hand, when wild-type c-Jun was overexpressed in cells cultured in serum-free medium containing 25 mM KCl, it was able to induce cell death. This result suggests that serum may contain a factor that prevents wild-type c-Jun from killing granule neurons. c-Jun^{ala}, an inactive c-Jun mutant, which cannot be phosphorylated, did not induce apoptosis under any conditions. However, expression of c-Jun^{ala} protected granule neurons against apoptosis after survival signal withdrawal. Because the only difference between wild-type c-Jun and c-Jun^{ala} is that the latter lacks the phosphoacceptor sites, this result suggests that phosphorylation of c-Jun is necessary for cell death in granule neurons. c-Jun^{ala} has a reduced ability to activate transcription (Smeal et al., 1991), and c-Jun^{ala} isolated from mammalian cells is unable to bind to a probe containing a TRE site in electrophoretic mobility shift assays (Papavassiliou et al., 1995). Overexpressed c-Jun^{ala} may act as a dominant negative mutant by sequestering kinases, such as the JNK/SAPKs, that

normally would directly bind to and activate the endogenous c-Jun protein. Alternatively, c-Jun^{ala} might interact with co-activator proteins that potentiate transactivation by c-Jun, such as CBP or JAB1 (Arias et al., 1994; Claret et al., 1996). c-Jun^{ala} appeared to inhibit apoptosis less efficiently than the c-Jun deletion mutant FLAG Δ 169 (compare Figs. 4, 6), which may be because it is expressed less efficiently or because it is still partially active (Metivier et al., 1993). Jun mutants similar to those described here have previously been used to show that *Drosophila* Jun (D-Jun) plays an important role in photoreceptor differentiation in the developing compound eye (Bohmann et al., 1994; Treier et al., 1995; Peverali et al., 1996). A D-Jun^{asp} mutant could induce R7 photoreceptor differentiation, whereas D-Jun^{ala} acted as a dominant negative mutant (Treier et al., 1995; Peverali et al., 1996).

Because we had obtained evidence that c-Jun phosphorylation is important for cell death in granule neurons, we investigated whether there were changes in the activity of any of the members of the MAP kinase superfamily that are known to regulate *c-jun* gene expression and c-Jun phosphorylation. We found that extracts from granule neurons contained high levels of Jun kinase activity, low levels of p38 kinase activity, and high levels of MAP kinase activity. After survival signal withdrawal, there was no increase in the level of Jun kinase or p38 kinase activity, whereas MAP kinase activity decreased. These results are different to those obtained for differentiated PC12 cells, in which both Jun kinase and p38 kinase were activated after NGF withdrawal (Xia et al., 1995) (Eilers, unpublished observations), and suggest that in cerebellar granule neurons novel mechanisms may exist for regulating c-Jun levels and c-Jun phosphorylation. One possibility is that granule neurons may contain a Jun kinase isoform that is not recognized by the pan-SAPK antibody that we used for immune complex kinase assays. Alternatively, given that granule neurons contain very high basal levels of Jun kinase activity, it is possible that in the presence of survival signals there are phosphatase activities that prevent the c-Jun transactivation domain from remaining phosphorylated. After survival signal withdrawal, the level of phosphatase activity might decrease, thereby allowing c-Jun phosphorylated by JNK/SAPKs to accumulate. Because MAP kinase activity declines in granule neurons after survival signal withdrawal, another possibility is that it is the balance between the level of MAP kinase and Jun kinase activity that influences cell survival, as has been reported for differentiated PC12 cells deprived of NGF (Xia et al., 1995).

An alternative explanation for our results is that perhaps, in addition to Jun kinase, cerebellar granule neurons may contain a novel kinase that is activated by lowering the extracellular KCl concentration and that stimulates the *c-jun* promoter or phosphorylates c-Jun. In the developing *Drosophila* eye, in which Jun phosphorylation is important for photoreceptor differentiation, D-Jun was a substrate for the ERK-related MAP kinase Rolled, which is part of the signal transduction pathway that triggers R7 photoreceptor differentiation (Peverali et al., 1996). Furthermore, although *Drosophila* Jun kinase (D-JNK) can phosphorylate D-Jun *in vitro*, fly ommatidia that lack D-JNK can develop normally (Riesgo-Escovar et al., 1996).

The transcriptional activation function of c-Jun and its partners seems to be important for apoptosis in cerebellar granule neurons, because the two c-Jun mutants that blocked cell death in these cells are both unable to activate transcription; FLAG Δ 169 lacks the transactivation domain, and c-Jun^{ala} only activates transcription weakly. The c-Jun target genes that are important for

neuronal cell death have yet to be identified. One may be the *c-jun* gene itself, because c-Jun is known to be able to activate transcription of the *c-jun* promoter (Angel et al., 1988). Cerebellar granule neurons appear to contain adequate levels of the effectors of apoptosis, because they can be killed by high concentrations of the broad-spectrum kinase inhibitor staurosporine in the presence of inhibitors of protein synthesis (Taylor et al., 1997). Therefore, after survival signal withdrawal, c-Jun may not activate the expression of genes encoding apoptosis effector proteins but, rather, may induce genes that code for molecules that activate the effectors of cell death. The identification of c-Jun target genes that are important for cell death will help resolve the apparent paradox that, as well as playing a role in neuronal cell death, c-Jun is also implicated in cell proliferation and neuronal regeneration. Furthermore, if c-Jun proves to be important for cell death *in vivo*, an understanding of c-Jun regulation and the mechanism by which c-Jun induces cell death may lead to the development of novel strategies for treating diseases of the nervous system in which neuronal apoptosis occurs.

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