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# Distinct Roles of c-Jun N-Terminal Kinase Isoforms in Neurite Initiation and Elongation during Axonal Regeneration

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c-Jun N-terminal kinases (JNKs) (comprising JNK1–3 isoforms) are members of the MAPK (mitogen-activated protein kinase) family, activated in response to various stimuli including growth factors and inflammatory cytokines. Their activation is facilitated by scaffold proteins, notably JNK-interacting protein-1 (JIP1). Originally considered to be mediators of neuronal degeneration in response to stress and injury, recent studies support a role of JNKs in early stages of neurite outgrowth, including adult axonal regeneration. However, the function of individual JNK isoforms, and their potential effector molecules, remained unknown. Here, we analyzed the role of JNK signaling during axonal regeneration from adult mouse dorsal root ganglion (DRG) neurons, combining pharmacological JNK inhibition and mice deficient for each JNK isoform and for JIP1. We demonstrate that neuritogenesis is delayed by lack of JNK2 and JNK3, but not JNK1. JNK signaling is further required for sustained neurite elongation, as pharmacological JNK inhibition resulted in massive neurite retraction. This function relies on JNK1 and JNK2. Neurite regeneration of *jip1*<sup>-/-</sup> DRG neurons is affected at both initiation and extension stages. Interestingly, activated JNKs (phospho-JNKs), as well as JIP1, are also present in the cytoplasm of sprouting or regenerating axons, suggesting a local action on cytoskeleton proteins. Indeed, we have shown that JNK1 and JNK2 regulate the phosphorylation state of microtubule-associated protein MAP1B, whose role in axonal regeneration was previously characterized. Moreover, lack of MAP1B prevents neurite retraction induced by JNK inhibition. Thus, signaling by individual JNKs is differentially implicated in the reorganization of the cytoskeleton, and neurite regeneration.

## Introduction

Regeneration of adult neurons requires their intrinsic capacity to survive a traumatic or degenerative lesion and activation of a cell-autonomous program that will end in plastic changes in neuronal circuitry. Integration of environmental signals leads to morphological and functional changes, depending both on regula-

tion of regeneration-related gene expression and reorganization of the cytoskeleton. The specific signaling pathways underlying neuronal regeneration are not yet fully elucidated.

c-Jun N-terminal kinases (JNKs) are members of a large group of serine/threonine-directed kinases known as mitogen-activated protein kinases (MAPKs) (Widmann et al., 1999; Davis, 2000). Mammalian JNKs are encoded by three distinct genes (*jnk1*, *jnk2*, and *jnk3*) giving rise to at least 10 different splice variants (Gupta et al., 1996). All variants share an epitope that needs to be dually phosphorylated for JNK activation (phospho-JNK). This is facilitated by scaffold proteins, notably the JNK-interacting proteins (JIP), that promote the sequential phosphorylation of cascade of MAP kinases (Morrison and Davis, 2003).

JNKs are best known as mediators of neuronal degeneration after stress and injury, but may be important also for neural development and survival (for review, see Waetzig et al., 2006). Recent studies support a role of JNK and JIP1 in the regulation of neurite outgrowth during development (Oliva et al., 2006; Dajas-Bailador et al., 2008). Moreover, JNKs are also implicated in axonal regeneration of adult DRG neurons, as their specific inhibition *in vitro* dramatically reduces neuritogenesis, without affecting neuronal survival (Kenney and Kocsis, 1998; Lindwall et al., 2004). In response to peripheral nerve injury, JNKs are rapidly

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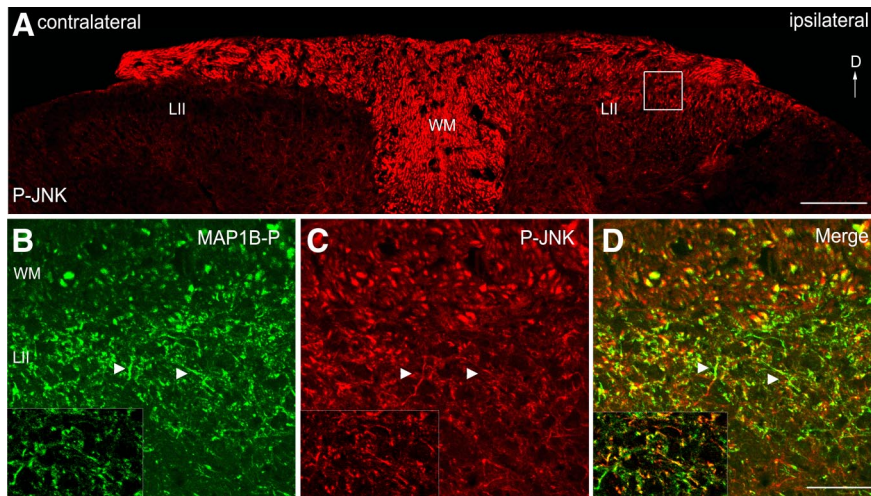
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**Figure 1.** JNK activation in central primary sensory fibers undergoing sprouting in response to a peripheral lesion. Shown are confocal microscope images. **A**, P-JNK immunostaining (red) in dorsal horn of adult mouse spinal cord, 1 week after unilateral sciatic nerve transection. P-JNK is detected in fibers in lamina II ipsilateral to the lesion (right), compared with the contralateral side (left), devoid of P-JNK staining. **B–D**, Higher magnification of the area marked by a white square in **A** reveals colocalization of P-JNK (red) with phosphorylated MAP1B (green) within the same fibers (arrowheads). The insets show single optical plane images. **D**, Dorsal; LII, lamina II; WM, white matter. Scale bars: **A**, 100  $\mu\text{m}$ ; **B–D**, insets, 40  $\mu\text{m}$ .

activated and retrogradely transported to the nuclei where they phosphorylate and, thereby, activate transcription factors such as c-Jun and activating transcription factor-3 (ATF3) (Lindwall et al., 2004; Cavalli et al., 2005). There is also a growing evidence for a cytosolic function of JNKs during neuronal development. In fact, JNKs may influence cytoskeletal reorganization via the phosphorylation of proteins regulating microtubule stability, including doublecortin (DCX) (Gdalyahu et al., 2004), stathmin family protein (SCG10) (Tataruk et al., 2006), and microtubule-associated proteins, MAP2 and MAP1B (Chang et al., 2003).

Whereas JNK1 and JNK2 are expressed in a large variety of tissues, JNK3 expression is restricted to the brain, heart, and testis (Gupta et al., 1996; Kuan et al., 1999), suggesting that the different JNK isoforms may be involved in distinct cellular processes. This prompted us to investigate the regulation and specific function of each JNK isoform during axonal regeneration and plasticity in the adult. Thus, using pharmacological and genetic approaches (use of JNK- and JIP1-deficient mice) on the paradigm of DRG neuron regeneration, we were able to attribute specific roles to individual JNK isoforms in either initiation or extension of regenerating neurites. We have further shown that JNK isoforms are important regulators of the phosphorylation state of MAP1B, a cytoskeletal protein for which we previously demonstrated a role in guidance, branching, and retraction of regenerating axons (Bouquet et al., 2004, 2007; Stroissnigg et al., 2007).

## Materials and Methods

**Animals.** Genetic inactivation of JNK1-, JNK2-, JNK3-, JIP1-, and MAP1B-deficient mice (male and female) has been described in detail previously (Yang et al., 1997, 1998; Dong et al., 1998; Meixner et al., 2000; Whitmarsh et al., 2001). Adult (>2 months) homozygous deficient mice and their control wild-type littermates were used after systematic genotyping. Genotyping of individual mouse was performed by isolating DNA from the tail, and PCR amplification was performed using the following primers: jnk1 forward, 5'-CGCCAGTCCAAAATCAAGATC-3'; 5'-GCCATTCTGGTAGAGGAAGTTTCTC-3'; jnk1 reverse, 5'-CCAGCTCATTCTCCACTCATG-3'; jnk2 forward, 5'-GGAGCCGATAGTATCGAGTTACC-3'; 5'-GTTAGACAATCCCAGAGGTTGTGTG-3'; jnk2 reverse, 5'-CCAGTTCATTCTCCACTCATG-3'; jnk3 forward, 5'-CCTGCTTCTCAGAAACACCCTTC-3'; 5'-CGTA-

ATCTTGTACAGAAATCCCATAC-3'; jnk3 reverse, 5'-CTCCAGACTGCCTTGGGAAAA-3'; jip1 forward, 5'-CGCGGTCTCAGGTGAGCAA-3'; 5'-CTGACTAGGCTGTAAGAC-3'; jip1 reverse, 5'-CTCCAGACTGCCTTGGGAAAA-3'; map1b forward, 5'-CTGGATTTCATC-GACTGTGGCCG-3'; 5'-ATAGGTGCTGTGCCTCGCTGA-3'; map1b reverse, 5'-TTAGGAAGGCAGAGGAGCAGCC-3'. PCR products were analyzed by agarose gel electrophoresis. All experimental procedures on animals were in accordance with the European Community directive (86/609/EEC; authorization number 91-78 to F. Nothias).

**Surgery.** Animals were anesthetized by intraperitoneal injection of a solution containing 7.4 mg/kg xylazine (Rompun, 2%; Bayer) and 146 mg/kg ketamine (Imalgène 500; Mérieux). For spinal cord analysis, right sciatic nerves were cut at midhigh levels and reapposed by a single 9-0 suture through the epineurium to allow nerve regeneration. The left (unoperated) spinal cords were used as normal controls, in addition to intact animals. After a survival time of 1 week after axotomy, animals were deeply anesthetized with pentobarbital and perfused transcardially with saline (0.9%

NaCl at 37°C) supplemented with heparin (50 U/ml; sanofi-aventis), followed by 4% paraformaldehyde (200 ml) in 0.1 M phosphate buffer (PB), pH 7.4. Spinal cords were dissected and cryoprotected in 30% sucrose in 0.1 M PB containing NaCl (0.9%; PB), and coronal sections (30  $\mu\text{m}$ ) were performed on a cryostat.

To analyze peripheral nerve regeneration *in vivo*, right sciatic nerves were crushed at midhigh levels with forceps for 10 s. After 3 d, animals were reanesthetized, and 3  $\mu\text{l}$  of anthra[1,9-*cd*]pyrazol-6(2*H*)-one (SP600125) (50  $\mu\text{M}$  in 5% DMSO/PBS;  $n = 2$ ) or 5% DMSO/PBS (for control animals;  $n = 2$ ) was applied onto the lesion site in a cuff. The injury site was closed, and animals were perfused as described above, 48 h after SP600125 or DMSO application. Nerves were dissected and cryoprotected in 15% sucrose in 0.1 M PB containing NaCl (0.9%; PB), and longitudinal sections (16  $\mu\text{m}$ ) were performed on a cryostat.

**Culture of dissociated adult DRG neurons.** DRGs were dissected in F-12 medium and cut into small pieces before enzymatic dissociation by collagenase (90 min at 37°C; 4000 U/ml; Sigma-Aldrich), followed by treatments with trypsin/EDTA (Invitrogen) with DNase-1 (50  $\mu\text{g}/\text{ml}$ ; Sigma-Aldrich) for 10 min at 37°C, and by a final trituration using a narrowed Pasteur pipette. After multiple washes with F-12 medium, cells were resuspended in F-12 medium supplemented with N3 (Bottenstein and Sato, 1979) and penicillin/streptomycin (all reagents from Invitrogen), plated at a density of 100 cells/cm<sup>2</sup> on precoated glass coverslips, and placed in a 37°C, 5% CO<sub>2</sub> incubator. Coverslips were precoated with poly-L-lysine (10  $\mu\text{g}/\text{ml}$ ; Sigma-Aldrich) overnight at 37°C, followed by laminin (10  $\mu\text{g}/\text{ml}$ ; at least 3 h at 37°C; Sigma-Aldrich). To analyze the effect of JNK inhibition on neurite initiation, SP600125 (20  $\mu\text{M}$ ; BIOMOL International) was added to culture medium 4 h after plating (i.e., before any neurites emerged), and cell cultures were maintained for 24 and 48 h before fixation. The effect of SP600125 on neurite elongation was determined after adding the inhibitor to the culture medium 36 h after plating and further cultured for another 12 h.

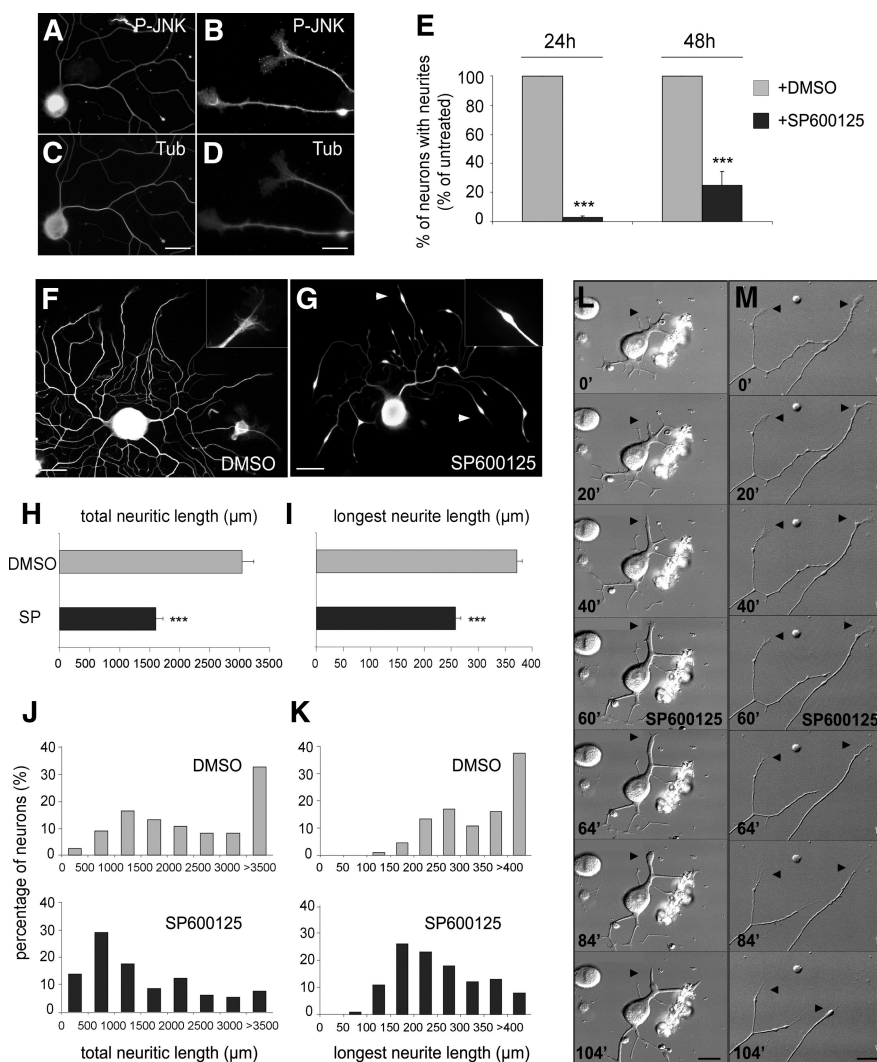
**Primary antibodies.** Mouse anti- $\beta$ 3 tubulin (Tuj-1) was obtained from Covance; mouse anti- $\beta$ -actin was obtained from Sigma-Aldrich; rat anti-L1 was obtained from Millipore Bioscience Research Reagents; rabbit anti-CGRP (calcitonin gene-related peptide) was from Peninsula; rabbit anti-detyrosinated tubulin (superlug) (Gundersen et al., 1984) was a gift from Dr. G. Gundersen (Columbia University, New York, NY); rabbit anti-total MAP1B (Fischer and Romano-Clarke, 1990) was provided by Dr. I. Fischer (Drexel University College of Medicine, Philadelphia, PA), and mouse anti-phosphorylated MAP1B (MAP1B-P) (Nothias et al., 1996) was homemade; rabbit polyclonal anti-total stath-

min (anti-peptide C) (Koppel et al., 1990) and anti-phospho-Ser25 stathmin (Gavet et al., 1998) were generously given by Dr. A. Sobel (Institut du Fer à Moulin, Paris, France); anti-phosphorylated JNK (P-JNK) was obtained from R&D Systems; mouse anti-JIP1 and mouse anti-glyceraldehyde-3-phosphate dehydrogenase (GAPDH) were obtained from Santa Cruz Biotechnology; mouse anti-JNK1 was obtained from BD Pharmingen; and anti-JNK2 and anti-JNK3 were obtained from Cell Signaling Technology.

**Immunostaining.** Cultured neurons were fixed with 4% paraformaldehyde/15% sucrose in PB, pH 7.4, for 20 min. For immunocytochemistry and histochemistry, specimens were permeabilized with 0.3% Triton X-100 in PB, and nonspecific binding sites were blocked with 10% normal goat serum (NGS) in PB, before incubation with primary antibodies diluted in 5% NGS in PB. After several rinses in 5% NGS in PB, coverslips were incubated with appropriate fluorochrome-labeled secondary antibodies (Jackson ImmunoResearch), rinsed again, and mounted in Mowiol.

**Morphological analysis.** Spinal cord coronal sections were immunostained for P-JNK and MAP1B-P and analyzed under confocal microscopy (SP5; Leica). Cells were plated on coverslips and visualized by tubulin staining. Images were taken on an Axiovert 200 inverted microscope equipped with an AxioCamera (Zeiss) and analyzed using NIH ImageJ (PC version: <http://rsbweb.nih.gov/ij/>) or Image ProPlus software (Media Cybernetics) for the following parameters: percentage of neurons with neurites, total neuritic length per neuron, and longest neurite per neuron. For analysis of the percentage of neurite-bearing neurons, cells were cultured for 24 and 48 h before fixation; all neurons on a coverslip were analyzed, and only neurites longer than one soma diameter were taken into account. For each condition, three independent experiments were performed, and per individual experiment at least 250 neurons were analyzed. For neuritic length analysis, only neurites longer than one soma diameter were manually traced and measured on size-calibrated images using Image ProPlus (three independent experiments with >50 neurons analyzed per condition). For neurite length distribution analysis, neurons were classified in different categories depending on their total neuritic length (ranging from 500 to >3500  $\mu\text{m}$ ) or length of their longest neurite (ranging from 50 to >400  $\mu\text{m}$ ), and the results were expressed as percentage of neurons in each category. Statistical significance was determined using independent *t* test and one-way ANOVA, and the levels of significance are indicated as follows: \*\*\**p* < 0.001, \*\**p* < 0.01, \**p* < 0.05. Compiled data are expressed as mean  $\pm$  SEM.

**Time-lapse video microscopy.** For time-lapse analysis, DRG neurons were plated on poly-L-lysine/laminin-coated coverslips. After 36 h, 10 mM HEPES, pH 7.4, was added to the medium, and cultures were placed in a chamber maintained at 37°C. Individual cells were imaged with a high-speed confocal microscope (resonant scanner SP5; 8000 Hz; Leica). Movies were composed of sequential 500 ms stills, taken at 2 or 4 min intervals over periods of 100 or 180 min. ImageJ software was used for quantitative evaluation of outgrowth and retraction velocities, by man-



**Figure 2.** JNK activation is required for both neuritogenesis and sustained extension of regenerating neurites. **A–D**, P-JNK and tubulin immunolabeling of DRG neurons cultured for 48 h. P-JNK is found in soma (including the nucleus), axon shafts, and growth cones. **E**, Quantification of neurite-bearing neurons cultured for 24 and 48 h, without (gray bars) or with chronic SP600125 treatment (black bars), reveals that neurite formation is blocked by JNK inhibition ( $n \geq 446$ ). **F, G**, Neurons treated with or without SP600125 and visualized by tubulin immunostaining. JNK inhibition induces loss of growth cones and leads to retraction bulb formation and trailing remnants along neurites (arrowheads); insets show higher magnifications of a typical growth cone under control conditions and a drug treatment-induced retraction bulb. **H, I**, Quantitative analysis reveals that JNK inhibition decreases total neuritic and longest neurite lengths of regenerating neurons ( $n \geq 55$ ). **J, K**, Distribution diagrams showing the percentage of neurons exhibiting a given total neuritic length and length of the longest neurite. Whereas the majority of neurons cultured under control conditions displays a total neurite length >3500  $\mu\text{m}$  and a longest neurite >400  $\mu\text{m}$ , JNK inhibition induces a shift of both values toward neurons with shorter neurites. **L, M**, Frames of time-lapse recordings illustrating the typical response of neurites to JNK inhibition: Before SP600125 addition ( $t = 60$  min), growth cones (arrowheads) are highly dynamic and neurites elongate steadily. Shortly after drug addition, growth cones collapse, and neurites stop extending and start to retract. Scale bars: **A, C, F, G, L, M**, 50  $\mu\text{m}$ ; **B, D**, 15  $\mu\text{m}$ . \*\*\**p* < 0.001. Error bars indicate SEM.

ually recording the distance traveled by the growth cones during 180 min, corresponding to the periods before and after application of SP600125. For each condition, three independent experiments were performed, and 10 neurites were analyzed per individual experiment.

**Preparation of cell extracts and Western blotting.** Lysis of cultures of DRG neurons plated for 48 h was preceded by washing with ice-cold PB. For MAP1B analysis, cells were lysed in denaturing extraction buffer containing 100 mM Tris-HCl, pH 6.8, 4% SDS, 20% (v/v) glycerol, 12 mM EDTA, 0.2% bromophenol blue, 0.3% DTT, and protease inhibitors (tablets from Roche Diagnostics). Samples were sonicated (3 times 20 s) and heated at 95°C for 5 min before loading on a 4.5% SDS-polyacrylamide gel. For alkaline phosphatase (AP) treatment, N2A neuroblastoma cells were cultured on poly-L-lysine for 48 h in DMEM–10%

FCS (Invitrogen), at 37°C and 5% CO<sub>2</sub>. To some cultures, SP600125 (20 μM) was added for 24 h. Cells were then rinsed twice with PBS and lysed in extraction buffer (see above), and samples were sonicated, centrifuged, and divided into two aliquots. One aliquot was treated for 30 min at 37°C with 10 U of calf alkaline phosphatase (New England Biolabs) before loading, and the other was mock-treated. For JNK immunoblot analysis, cultured dissociated DRG neurons were solubilized in denaturing buffer containing 1% SDS, 1% protease inhibitor mixture, and 1 mM NaVO<sub>4</sub> (Sigma-Aldrich) and sonicated, and proteins were separated on 10% SDS-polyacrylamide gels. After transfer, nitrocellulose membranes were blocked with 5% nonfat dry milk and incubated with primary antibodies. After thorough washing, membranes were incubated with appropriate alkaline phosphatase-conjugated secondary antibodies followed by revelation of antigens by NBT (nitroblue tetrazolium)/BCIP (5-bromo-4-chloro-3-indolyl phosphate) staining, or with fluorochrome-labeled secondary antibodies followed by detection on an infrared imaging system (Odyssey; LI-COR).

## Results

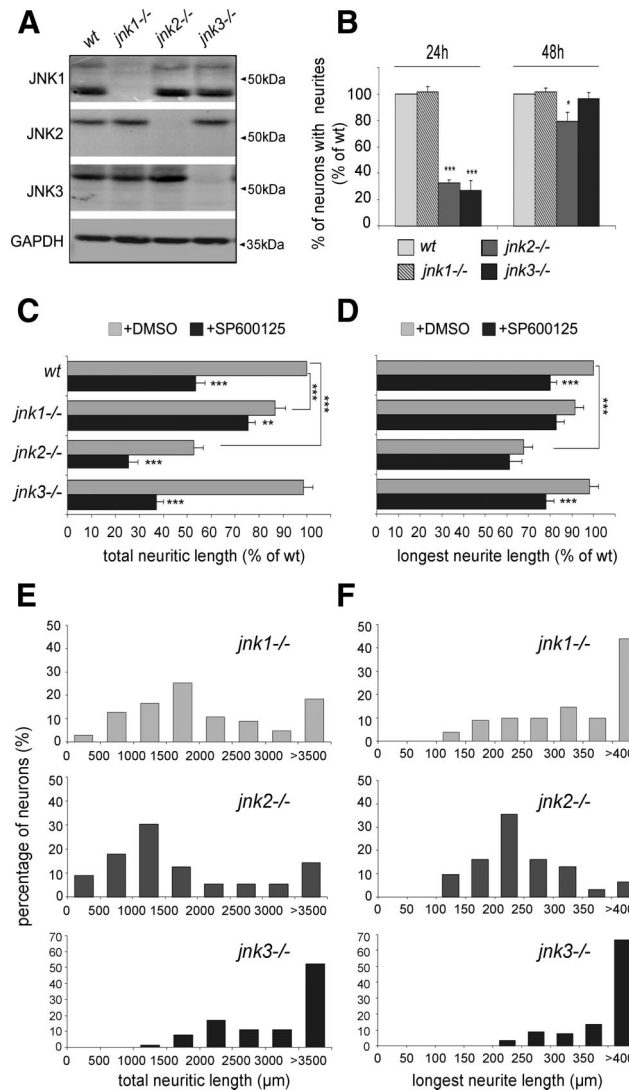
### Phospho-JNK is upregulated in axons undergoing plastic changes *in vivo*

We performed a lesion of sciatic nerve, known to induce a structural reorganization of axons within the central projection area of DRG neurons (i.e., sprouting in dorsal horn lamina II, most pronounced ~1 week after lesion in mice) (for details, see Soares et al., 2002). We previously showed that MAP1B-P increases within these fibers, and is well suited as marker for axonal sprouting and regeneration in the adult nervous system (Soares et al., 2002, 2007). In Figure 1A, fibers in lamina II ipsilateral to the sciatic nerve lesion display a strong increase in P-JNK immunostaining, whereas the contralateral dorsal horn is almost devoid of staining. Double immunostaining for MAP1B-P and P-JNK clearly shows that high levels of P-JNK are found in fibers that sprout in response to the peripheral lesion (Fig. 1B–D). These results demonstrate that, *in vivo*, DRG neurons, when entering the regenerative state in response to peripheral axotomy, exhibit a cytoplasmic increase of activated JNK.

### JNK activity is involved in both initiation and stabilization of regenerating neurons

Immunocytochemical analysis of primary cultures of adult DRG neurons, plated for 48 h, shows a distribution of P-JNK both in the nucleus and in the cytoplasm of soma and neurites of all DRG neuronal populations (Fig. 2A–D; supplemental Fig. S4A–D, available at www.jneurosci.org as supplemental material). Interestingly, P-JNK was present along distal neurite shafts and at growth cones (Fig. 2B), suggesting a local role in axonal regeneration. To test this hypothesis, we used an ATP-competitive inhibitor of all JNK isoforms, SP600125 (Bennett et al., 2001). Treatment of neurons with SP600125 4 h after plating, before neurite initiation, severely affected neuritogenesis since only 3.0 ± 0.7 and 24.9 ± 9.4% of SP600125-treated neurons had formed neurites after a culture period of 24 and 48 h, respectively, compared with untreated neurons set to 100% (Fig. 2E).

Given the presence of P-JNK in neurites during the whole culture period, we further investigated whether JNK activation is also required for sustained neurite extension. This was addressed by addition of JNK inhibitor for 12 h, at a time point when the majority of neurons had extended neurites (36 h after plating). JNK inhibition induced the formation of retraction bulbs and trailing remnants along neurites (Fig. 2F, G). A similar effect of SP600125 treatment was also observed *in vivo* on regenerating sciatic nerves after a lesion (supplemental Fig. S1, available at www.jneurosci.org as supplemental material). Whereas periph-



**Figure 3.** Differential involvement of JNK isoforms in neurite initiation and elongation. **A**, Western blots of protein extracts from wild-type and JNK-deficient mice DRG neurons, cultured 48 h, reacted with antibodies specific for individual JNK proteins. In regenerating wild-type neurons, all isoforms are expressed. **B**, Quantification of neurons bearing neurites after 24 and 48 h of culture: At 24 h, *jnk2*<sup>-/-</sup> and *jnk3*<sup>-/-</sup> neurons exhibit a strong decrease in the percentage of neurons with neurites compared with wild-type neurons. At 48 h, only *jnk2*<sup>-/-</sup> neurons still display a slight but significant difference. Neurite initiation is not affected in regenerating *jnk1*<sup>-/-</sup> neurons ( $n \geq 297$ ). **C, D**, Quantification of total neurite length and length of the longest neurite for wild-type versus JNK-deficient neurons without (gray bars) or with SP600125 treatment (black bars). Total neurite lengths for *jnk1*<sup>-/-</sup> and *jnk2*<sup>-/-</sup>, but not *jnk3*<sup>-/-</sup> neurons, are decreased compared with wild-type neurons; the average length of the longest neurite is significantly affected only by lack of JNK2. SP600125 treatment induces an additional reduction in total neurite length for neurons derived from all three *jnk*-ko mice, but induces a specific decrease in length of the longest neurite only in *jnk3*<sup>-/-</sup> neurons ( $n \geq 50$ ). **E, F**, The effect on neurite length of JNK1 and JNK2, but not JNK3 deficiency, is reflected in a leftward shift in the distribution diagrams for total neurite length; values for the length of the longest neurite per neuron display a leftward shift only for *jnk2*<sup>-/-</sup> neurons; those for *jnk1*<sup>-/-</sup> and *jnk3*<sup>-/-</sup> neurons are similar to wild type. \*\*\* $p < 0.001$ ; \*\* $p < 0.01$ ; \* $p < 0.05$ . Error bars indicate SEM.

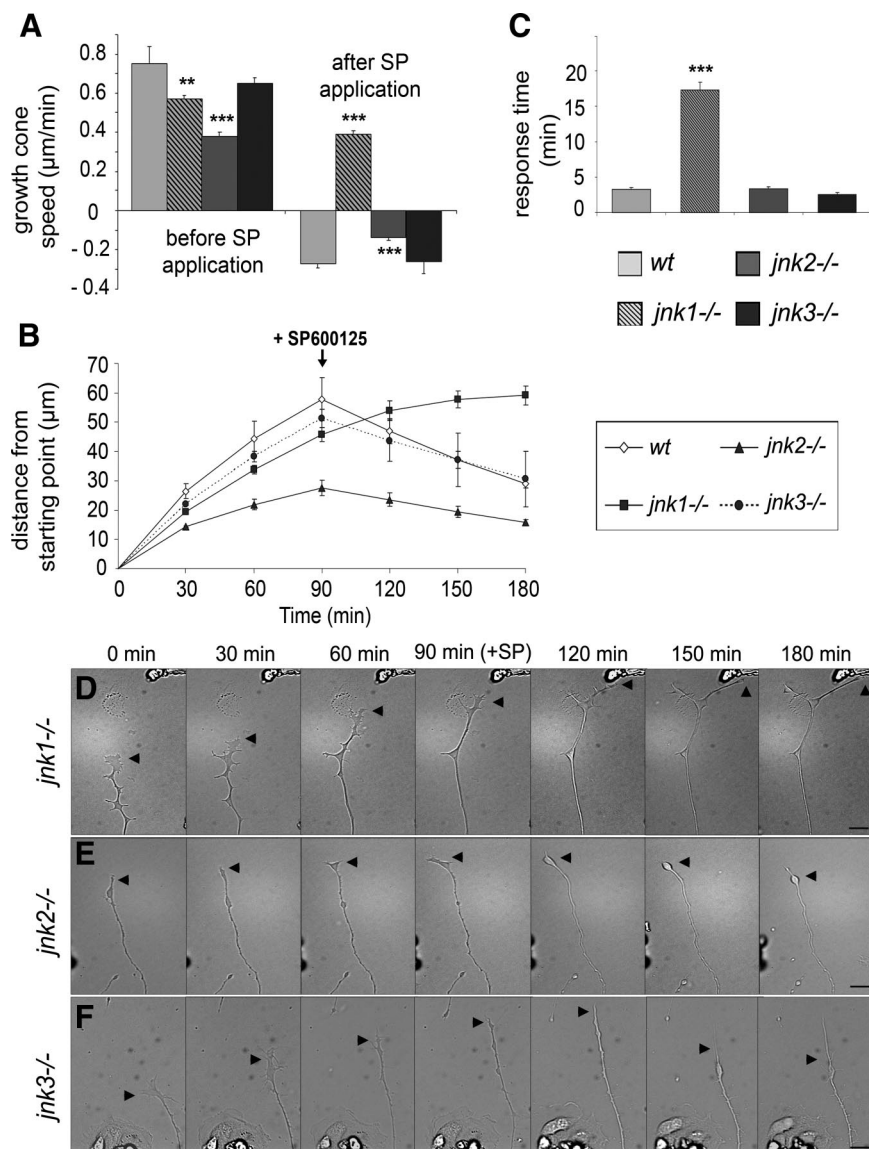
eral fibers grow linearly in condition control (supplemental Fig. S1A–D, I, available at www.jneurosci.org as supplemental material), the SP600125 addition severely disrupted and disorganized the regeneration profile (supplemental Fig. S1E–H, available at www.jneurosci.org as supplemental material) and caused the occurrence of retraction bulbs (supplemental Fig. S1J, available at

www.jneurosci.org as supplemental material). This reproducible effect of SP600125 on the morphology of regenerating neurites, *in vitro*, was correlated with a reduction in total neurite length per neuron by 47.3% ( $1604.1 \pm 118.6 \mu\text{m}$  for SP600125-treated vs  $3041.2 \pm 190.4 \mu\text{m}$  for untreated neurons) (Fig. 2H) and with a 30.6% shortening of the length of the longest neurite ( $258.2 \pm 9.4$  vs  $372.1 \pm 12.6 \mu\text{m}$ ) (Fig. 2I). To determine whether JNK inhibition affected all growing neurites independently of their initial length, we established the distribution of neurons based on each parameter (Fig. 2J,K). Whereas a majority of untreated neurons (60%) exhibited a total neuritic length  $>2000 \mu\text{m}$ , this distribution was strongly shifted toward shorter neurites after SP600125 treatment: only 30% of neurons had a total neurite length  $>2000 \mu\text{m}$  (Fig. 2J). A similar shift was noted in the distribution of the length of the longest neurite: In control cultures, 64% of neurons show a longest neurite  $>300 \mu\text{m}$  versus 29% after drug treatment (Fig. 2K).

To assess the dynamics of these events, DRG neuron cultures were analyzed using time-lapse videomicroscopy (Fig. 2L,M). Image sequences, representing a neuron during the initial stages of neurite elongation (Fig. 2L) and a neuron having already extended longer neurites (Fig. 2M), are shown (see also supplemental Video 1, available at www.jneurosci.org as supplemental material). In both cases, neurites elongated steadily and possessed typical fan-shaped growth cones with filopodia and lamellipodia exploring the environment. Addition of SP600125 (at  $t = 60$  min) caused a rapid collapse of the growth cone (within 4–8 min), followed by neurite retraction (within 12–16 min) that could be reversed by removal of the inhibitor from the medium (data not shown). JNK inhibition-induced neurite retraction was observed on both short and long neurites, corroborating the analysis of neurite length distribution (Fig. 2J,K). Moreover, the rearward movement of neurites that followed the growth cone collapse was consistent with the neurite morphology observed on fixed preparations (Fig. 2G). Together, our analysis demonstrates that JNK activity is required for both neuritogenesis and sustained extension of regenerating neurites.

#### Differential involvement of individual JNK isoforms in neurite initiation and extension

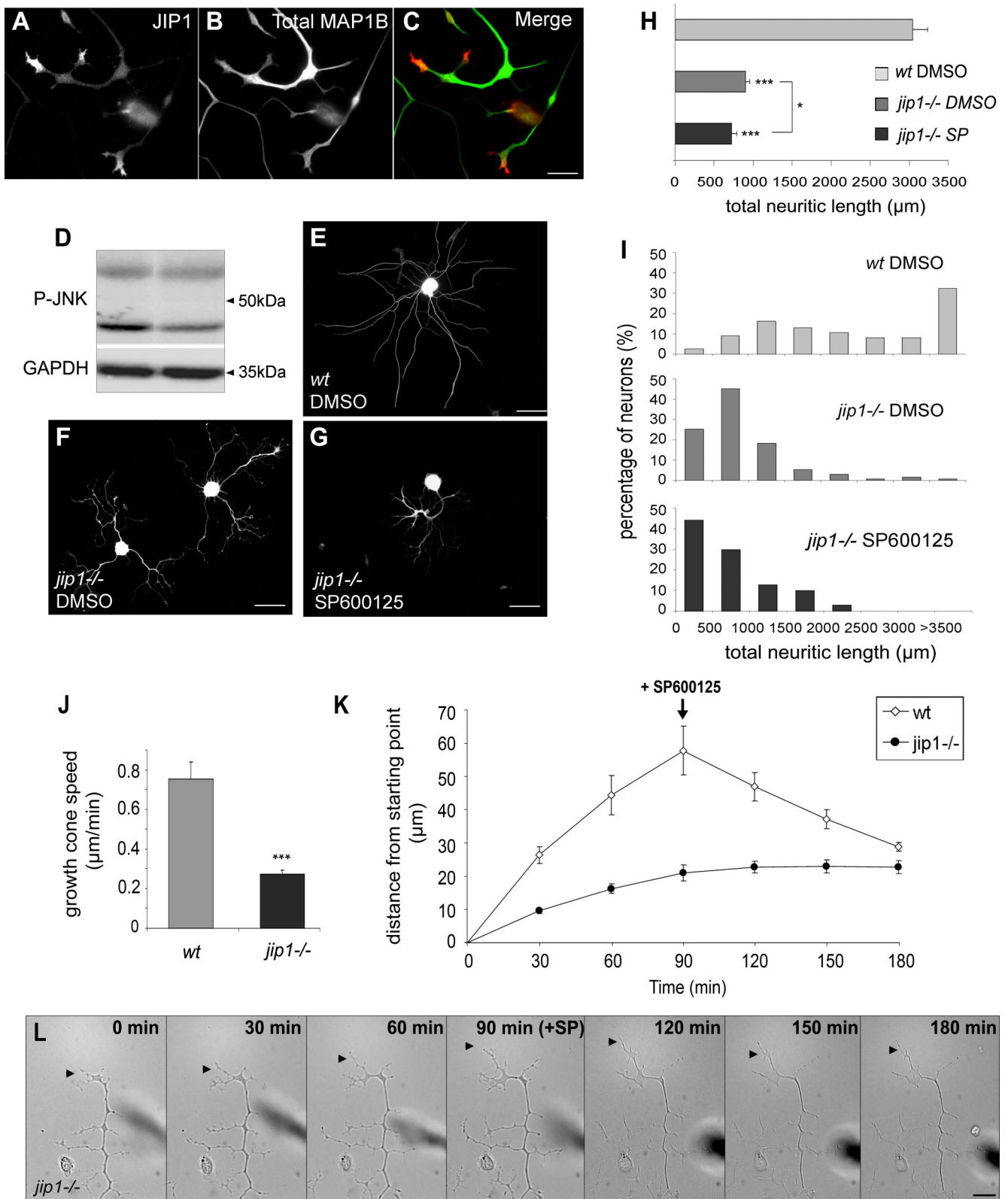
Since the pharmacological approach does not discriminate between individual JNK isoforms for which distinct, but also partly overlapping, functions had been described (Bogoyevitch, 2006), we examined the expression and potential contribution of each isoform to DRG neurite regeneration. Using antibodies specific for



**Figure 4.** Lack of JNK1 and JNK2 affects axonal elongation rate. **A**, Mean speed of growth cone advancement and retraction of wild-type and JNK-deficient DRG neurons. Note that *jnk1*<sup>-/-</sup> and *jnk2*<sup>-/-</sup>, but not *jnk3*<sup>-/-</sup>, neurite outgrowth rates are significantly decreased compared with wild type. After SP600125 application, the retraction rate of *jnk2*<sup>-/-</sup> neurites is decreased compared with *jnk3*<sup>-/-</sup> and wild-type neurites, and no retraction of *jnk1*<sup>-/-</sup> neurites occurs. **B**, Advancement or retraction of growth cones measured 90 min before and 90 min after SP600125 application. Within a few minutes after JNK inhibition, neurites from *jnk2*<sup>-/-</sup> and *jnk3*<sup>-/-</sup> neurons retract, whereas *jnk1*<sup>-/-</sup> neurites continue growing during the remaining recording period. Response time for growth cones to collapse after SP600125 application: *jnk2*<sup>-/-</sup> and *jnk3*<sup>-/-</sup> growth cones collapse rapidly, in contrast to *jnk1*<sup>-/-</sup> growth cones. **D–F**, Frames of time-lapse recordings illustrating the outgrowth and typical response of neurites to JNK inhibition from *jnk1*<sup>-/-</sup>, *jnk2*<sup>-/-</sup>, and *jnk3*<sup>-/-</sup> DRG neurons, 90 min before and after SP600125 addition. Arrowheads, Growth cones. Scale bars, 50  $\mu\text{m}$ .  $***p < 0.001$ ;  $**p < 0.01$ . Error bars indicate SEM.

each isoform, we found that JNK1, JNK2, and JNK3 were expressed in regenerating DRG neurons (Fig. 3A, Western blot). The specificity of each antibody was demonstrated by the absence of reactive protein bands when using DRG neuron extracts from mice deficient in the corresponding JNK isoform.

We then examined the implication of individual JNK isoforms in neuritogenesis by culturing neurons derived from *jnk1*<sup>-/-</sup>, *jnk2*<sup>-/-</sup>, and *jnk3*<sup>-/-</sup> mice for 24 and 48 h (Fig. 3B). At 24 h, neurite initiation of *jnk1*<sup>-/-</sup> neurons was similar to the one observed in wild-type neurons, whereas only  $32.8 \pm 2.1\%$  of *jnk2*<sup>-/-</sup> and  $27.1 \pm 7.4\%$  of *jnk3*<sup>-/-</sup> neurons had formed neurites compared with wild type. However, after 48 h in culture, no

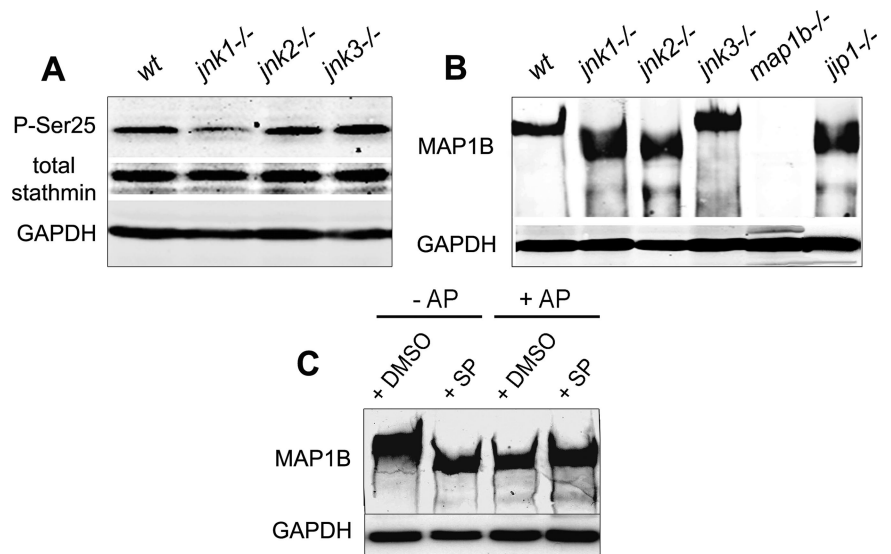


**Figure 5.** Scaffold protein JIP1 is required for neurite regeneration. *A, B*, and merge in *C*, JIP1 immunostaining is enriched in growth cones of regenerating DRG neurites, shafts of which are visualized by total MAP1B immunostaining. *D*, Western blot analysis of P-JNK in protein extracts from wild-type or *jip1*<sup>-/-</sup> DRG neurons cultured 48 h before lysis; the expression of GAPDH was monitored in the lower part of the same membrane as loading control. Lack of JIP1 results in a reduction of the 46 kDa form of P-JNK. *E–G*, Tubulin immunostaining of DRG neurons cultured for 48 h derived from wild-type (*E*), *jip1*<sup>-/-</sup> (*F*), and *jip1*<sup>-/-</sup> neurons with additional SP600125 treatment (*G*). Note the abnormal “curly” morphology of neurites from *jip1*<sup>-/-</sup> DRG exhibiting more terminal branching; additional SP600125 treatment has no effect on *jip1*<sup>-/-</sup> neurites. *H*, Quantification of total neuritic length of wild-type versus *jip1*<sup>-/-</sup> neurons reveals a difference of 70.4%; in contrast, additional SP600125 treatment has only a slight effect on neurite length of *jip1*<sup>-/-</sup> DRG neurons ( $n \geq 55$ ). *I*, Diagram showing the distribution of total neuritic lengths of wild-type versus *jip1*<sup>-/-</sup> DRG neurons, the latter cultured with or without additional SP600125 treatment. *J*, Mean speed of growth cone advancement from wild-type and JIP1-deficient DRG neurons showing that the outgrowth rate of *jip1*<sup>-/-</sup> neurite is drastically decreased. *K*, Growth cone advancement or retraction measured 90 min before and 90 min after SP600125 application, showing that no neurite retraction occurs from *jip1*<sup>-/-</sup> neurons during the remaining recording time, as illustrated in *L*, showing corresponding frames of time-lapse recordings. Arrowheads, Growth cones. Scale bars: *A, C*, 15 μm; *E–G*, 100 μm; *L*, 50 μm. \*\*\* $p < 0.001$ ; \* $p < 0.05$ . Error bars indicate SEM.

difference in the percentage of neurons bearing neurites were observed between *jnk3*<sup>-/-</sup> and wild-type neurons, whereas neuritogenesis of *jnk2*<sup>-/-</sup> neurons was partially restored but still significantly affected (79.4% of neurons with neurites). These results indicate that JNK2 and JNK3, but not JNK1, are involved in neurite initiation. Furthermore, the low percentage of *jnk2*<sup>-/-</sup> and *jnk3*<sup>-/-</sup> neurons bearing neurites at early stages was mainly attributable to a delay in neurite initiation.

To investigate the role of JNK isoforms in sustaining the extension of regenerating neurites over time, total neuritic length and length of the longest neurite per neuron were determined after culturing wild-type and JNK-deficient neurons for 48 h (Fig. 3C,D). Neurites of *jnk2*<sup>-/-</sup>, and to a lesser degree of *jnk1*<sup>-/-</sup>, neurons were significantly shorter than those of wild-type neurons ( $62.8 \pm 6.6$  and  $86.6 \pm 4.8\%$  of wild-type total neuritic length, respectively), whereas neurite outgrowth from *jnk3*<sup>-/-</sup> neurons was not affected (Fig. 3C). A difference in the average length of the longest neurite was, however, observed only for *jnk2*<sup>-/-</sup> neurons (Fig. 3D). Accordingly, the distribution of total neuritic length, and length of the longest neurite, exhibited a shift toward smaller values for both *jnk1*<sup>-/-</sup> and *jnk2*<sup>-/-</sup>, but not *jnk3*<sup>-/-</sup> neurons (Fig. 3E,F). To understand the basis for this decrease in neurite length, we monitored the neurite outgrowth speed of DRG neurons derived from the different JNK-deficient mice. Compared with wild-type neurites that advance at  $0.76 \pm 0.09 \mu\text{m}/\text{min}$ , the growth speed of *jnk1*<sup>-/-</sup>, and to an even higher degree of *jnk2*<sup>-/-</sup>, neurites was significantly lower ( $0.57 \pm 0.02$  and  $0.38 \pm 0.02 \mu\text{m}/\text{min}$ , respectively) (Fig. 4A,B). Slower growth rates are thus responsible for *jnk1*<sup>-/-</sup> and *jnk2*<sup>-/-</sup> neurites being shorter than their wild-type or *jnk3*<sup>-/-</sup> counterparts noted at the end of the culture period.

Finally, we assessed the additional effect of SP600125 treatment on the elongation of neurites from the individual JNK isoform-deficient neurons. Whereas complete JNK inhibition induces a decrease in total neurite length of *jnk2*<sup>-/-</sup> and *jnk3*<sup>-/-</sup> neurons by  $27.4 \pm 7.1$  and  $61.3 \pm 6.6\%$ , respectively, it has only a lower (although significant) effect on *jnk1*<sup>-/-</sup> neurites (Fig. 3C,D; supplemental Fig. S2, available at www.jneurosci.org as supplemental material). Dynamic analysis, by time-lapse videomicroscopy, again revealed a clear difference in the behavior of the different JNK-deficient neurons. Thus, similarly to wild-type neurons, *jnk2*<sup>-/-</sup> and *jnk3*<sup>-/-</sup> growth cones reacted to JNK inhibition by collapsing  $3.38 \pm 0.23$  and  $2.53 \pm 0.29$  min after drug addition, followed by neurite retraction after  $13.68 \pm 0.72$  and  $12.73 \pm 0.77$  min, respectively (Fig. 4C). Whereas *jnk3*<sup>-/-</sup> neurites retract with a rate comparable with wild type ( $0.26 \pm 0.06$  and  $0.27 \pm 0.02 \mu\text{m}/\text{min}$ , respectively), the rate of *jnk2*<sup>-/-</sup> neurite retraction is significantly reduced ( $0.14 \pm 0.01 \mu\text{m}/\text{min}$ ) (Fig. 4A,B; supplemental Videos 3, 4, available at www.jneurosci.org as supplemental material). The response of *jnk1*<sup>-/-</sup> neurites was delayed since, after SP600125 treatment, the growth cone collapse occurs within  $17.27 \pm 1.1$  min and no



**Figure 6.** Regulation of stathmin and MAP1B phosphorylation by JNKs: Western blot analysis with GAPDH used as internal loading control. **A**, Relative amounts of total stathmin and stathmin phosphorylated on serine 25 (P-Ser25) in lysates of wild-type and JNK-deficient DRG neurons cultured for 48 h; note the decrease in stathmin phosphorylation in regenerating *jnk1*<sup>-/-</sup> neurons. **B**, MAP1B staining of protein extracts from wild-type and JNK-deficient DRG neurons cultured for 48 h, showing a shift in the protein band representing MAP1B from higher to lower apparent molecular weight for *jnk1*<sup>-/-</sup> and *jnk2*<sup>-/-</sup>, but not *jnk3*<sup>-/-</sup> neurons. This shift reflects a reduced level of MAP1B phosphorylation as demonstrated in **C**, MAP1B in lysates of N2A neuroblastoma cells cultured for 48 h is phosphorylated under control conditions (lane 1); experimental dephosphorylation by AP treatment of lysates (lanes 3, 4) results in the same shift to a lower molecular weight as JNK inhibition by SP600125 (SP) addition to the N2A cultures (lane 2).

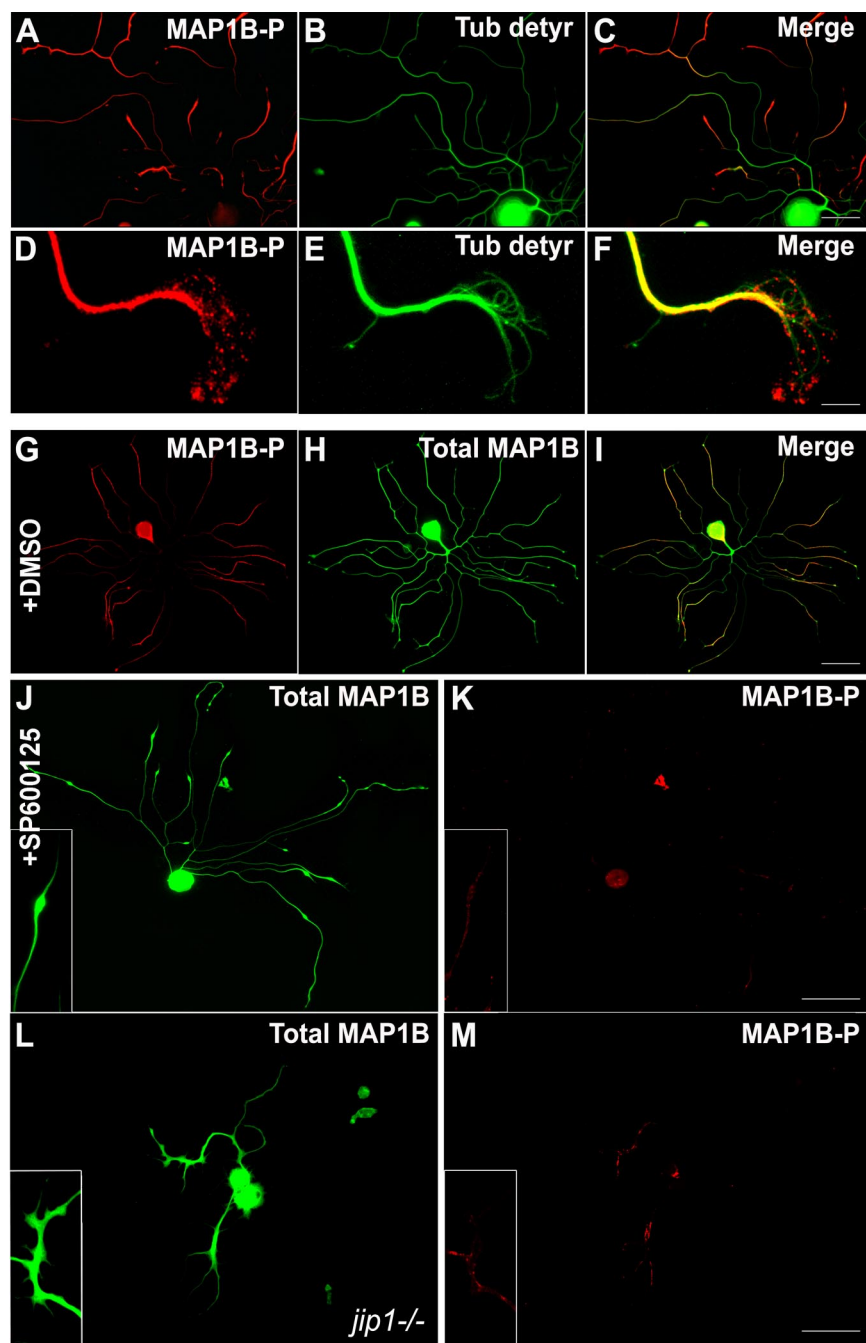
neurite retraction was yet observed during the 90 min that followed the drug addition (Fig. 4A–D; supplemental Video 2, available at www.jneurosci.org as supplemental material).

To summarize, our results indicate that the combination of JNK1 and JNK2 activities is crucial for neurite extension.

### JIP1 is required for axonal regeneration through signaling that involves JNK

The JIP1 scaffolding protein interacts with all three JNK isoforms. It contributes to their activation in response to stress stimuli (Whitmarsh and Davis, 1998) by facilitating the phosphorylation cascade based on MAP kinases that ultimately leads to JNK activation. JIP1 appears to be involved in axonal outgrowth during development (Dajas-Bailador et al., 2008). Here, we found JIP1 to be enriched in growth cones of adult regenerating DRG neurons (Fig. 5A–C). Thus, we examined the neuritogenesis of DRG neurons from *jip1*<sup>-/-</sup> mice. Twenty-four hours after plating, only  $32.9 \pm 13.9\%$  *jip1*<sup>-/-</sup> neurons were bearing neurites, compared with wild-type neurons set to 100% (supplemental Fig. S3A, available at www.jneurosci.org as supplemental material). However, after 48 h, the percentage of *jip1*<sup>-/-</sup> neurons bearing neurites was similar to wild type, suggesting that JIP1 contributes to neurite initiation. Although neurite initiation was restored at 48 h, *jip1*<sup>-/-</sup> neurons exhibit shorter neurites (by 70.4%) compared with wild-type neurons ( $900.4 \pm 56$  vs  $3041.2 \pm 190.4 \mu\text{m}$ ) (Fig. 5E,F,H) [see also the drastic leftward shift in the length distribution graph (Fig. 5I)]. Similar reductions were observed for the length of the longest neurite (supplemental Fig. S3C, available at www.jneurosci.org as supplemental material), associated to a drastic decrease in neurite growth rate ( $0.27 \pm 0.02$  vs  $0.76 \pm 0.09 \mu\text{m}/\text{min}$  for wild-type neurites) (Fig. 5J–L). Moreover, the morphology of *jip1*<sup>-/-</sup> neurons was abnormal, their neurites being curled, and exhibiting a higher degree of terminal branching (Fig. 5F; quantitative supplemental Fig. S3B, available at





**Figure 7.** JNK inhibition affects MAP1B phosphorylation in regenerating neurons. **A–C**, In control DRG neurons, distal parts of neurites are enriched in MAP1B-P and characterized by low levels of detyrosinated tubulin (representing stable microtubules). **D–F**, Higher magnification of a growth cone showing the presence of MAP1B-P at the leading edge, where no colocalization with detyrosinated tubulin can be found. **G–I**, Double immunostaining reveals that total MAP1B is distributed in all parts of the neuron (green), whereas MAP1B-P (red) is mainly localized in neurites and exhibits a proximo-distal gradient. **J, K**, Application of SP600125, although not altering total MAP1B levels, dramatically decreases the neurite content in MAP1B-P, concomitantly with growth cone retraction (see also insets). **L, M**, Immunostaining for MAP1B-P, compared with total MAP1B, on regenerating *jip1*<sup>-/-</sup> neurons reveals that lack of JIP strongly affects the level of MAP1B phosphorylation. Scale bars: **A–C**, 50  $\mu$ m; **D–F**, 10  $\mu$ m; **G–M**, 100  $\mu$ m.

www.jneurosci.org as supplemental material). We further examined DRG neurons from *jip1*<sup>-/-</sup> mice with respect to alterations in JNK phosphorylation. Western blot of protein extract from cultured *jip1*<sup>-/-</sup> DRG neurons shows an obvious decrease of the 46 kDa band corresponding mainly to phospho-JNK1 (Fig. 5D), probably related to the fact that JNK1 is the most active isoform during later phases of neurite extension (here, at 48 h of culture).

Since JNK phosphorylation is not completely abolished in *jip1*<sup>-/-</sup> neurons, we examined a potential additional effect of SP600125 treatment on sustained neurite extension in *jip1*<sup>-/-</sup> neurons. Whereas the morphology of *jip1*<sup>-/-</sup> neurons cultured for 48 h was apparently only little affected by JNK inhibition (Fig. 5G), only a small decrease (19.5%) was noted for their total neurite length and length of the longest neurite (Fig. 5H; supplemental Fig. S3C, available at www.jneurosci.org as supplemental material) (compare with wild-type data in Fig. 2H,I). The effect of SP treatment on *jip1*<sup>-/-</sup> neurons was also reflected by a slight leftward shift in the distribution diagram of total neurite lengths (Fig. 5I; length of longest neurites, supplemental Fig. S3D, available at www.jneurosci.org as supplemental material). Dynamic analysis by time-lapse videomicroscopy again revealed that no neurite retraction was observed during the 90 min that follow the drug addition (Fig. 5K,L; supplemental Video 5, available at www.jneurosci.org as supplemental material).

Our results demonstrate that JIP1 deficiency in adult DRG neurons results in reduced JNK phosphorylation and a drastically reduced neurite regrowth capacity.

#### JNK signaling in regenerating axons acts on the cytoskeletal protein MAP1B

It is now widely accepted that the regulation of neurite outgrowth in response to extracellular factors passes through signaling cascades that converge onto a reorganization of the neuronal cytoskeleton. Thus, extracellular signaling will ultimately trigger posttranslational modifications (such as phosphorylation) of microtubule-interacting proteins, including stabilizing and destabilizing factors, and thereby regulate microtubule dynamics. During development, MAP1B and the stathmin family members are examples for microtubule-associated proteins that might be influenced by JNK signaling. Stathmin family proteins, when activated, increase the rate of microtubule catastrophes, either through tubulin sequestration or by acting on microtubule plus ends. As shown by Tataruk et al. (2006), JNKs may negatively regulate the activity of these destabilizing proteins (e.g., SCG10) during development, resulting in maintaining microtubule stability and efficient neurite elongation.

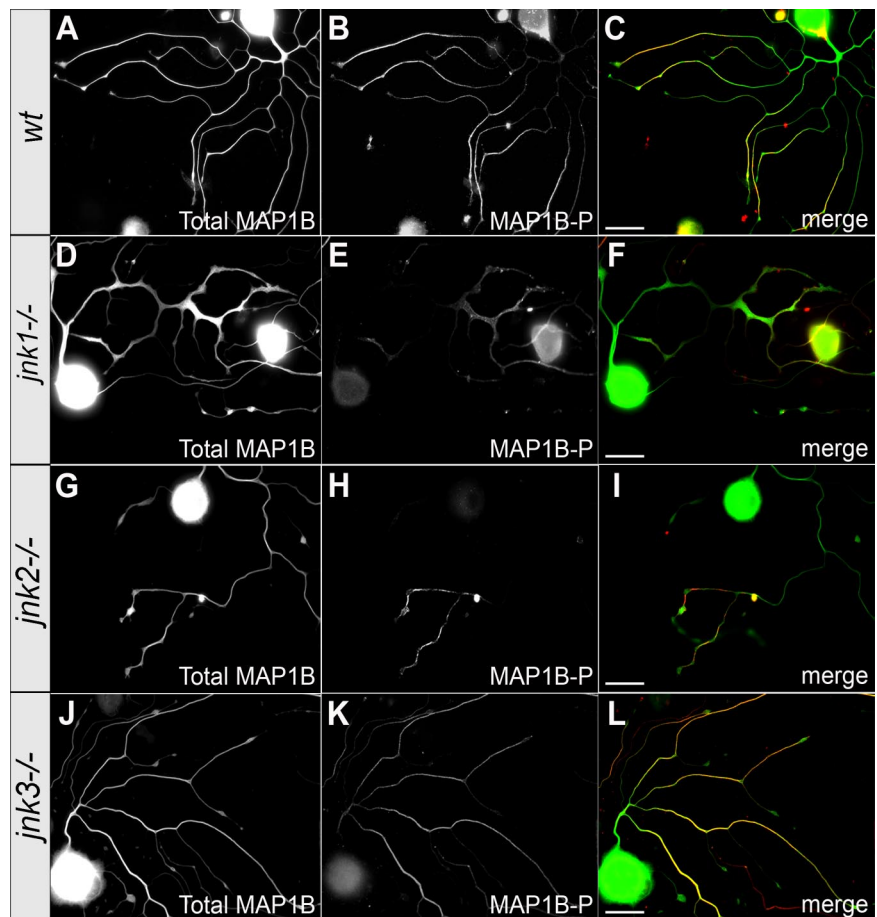
Here, we analyzed by Western blotting the phosphorylation state of both stathmin and MAP1B in extracts from cultured adult regenerating DRG neurons deficient for each JNK isoform. Phosphorylation of stathmin was monitored using an antibody specifically recognizing stathmin protein phosphorylated on serine 25, a site known to be predominantly phosphorylated by

MAPK (Leighton et al., 1993; Hayashi et al., 2006). Hypophosphorylation of stathmin was found in *jnk1*<sup>-/-</sup> DRG neurons, whereas no changes were noted for regenerating *jnk2*<sup>-/-</sup> or *jnk3*<sup>-/-</sup> neurons (Fig. 6A). Using the same protein extracts, we then showed a hypophosphorylation of MAP1B in *jnk1*<sup>-/-</sup> and *jnk2*<sup>-/-</sup> neurons, as indicated by a lower apparent molecular weight of the MAP1B protein band, compared with highly phosphorylated MAP1B from wild-type and *jnk3*<sup>-/-</sup> neurons (Fig. 6B). The downward shift in the protein band representing MAP1B is indeed characteristic of hypophosphorylation, described previously (Nothias et al., 1996; Alfei et al., 2004; Soares et al., 2007) and demonstrated here in Figure 6C: SP600125 treatment (lane 2) and dephosphorylation by AP of protein extract from cells that had not been treated with SP (lane 3) result in the same shift of the band representing MAP1B to a lower apparent molecular weight.

These results suggest that JNK1 regulates the phosphorylation of stathmin, whereas both JNK1 and JNK2 regulate MAP1B phosphorylation.

Since the deletion of both JNK1 and JNK2 appeared to affect axonal regeneration in DRG neurons, we focused on the relationship between JNK signaling and the activity of MAP1B for which we had already shown a role in axonal regeneration. In addition, the codistribution of MAP1B-P and P-JNK in sprouting or regenerating fibers, reported here (Fig. 1A–D), suggested that MAP1B might be phosphorylated through JNK signaling during remodeling of the neuronal cytoskeleton. As shown in Figure 7A–C, MAP1B-P staining exhibits a proximo-distal gradient in regenerating neurites, from all DRG neuronal populations (small to large size) (supplemental Fig. S4E–L, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material). The highest MAP1B-P levels are observed in the tips of growing neurites where stable microtubules are rare, as indicated here by the low proportion of detyrosinated tubulin. MAP1B-P, but no detyrosinated tubulin, is found at sites where dynamic “pioneer” microtubules (tyrosinated microtubules) (Bouquet et al., 2004, 2007) invade the peripheral growth cone domain (Fig. 7D–F). In contrast to total MAP1B, displaying a homogeneous distribution in all neuronal compartments (Fig. 7G) that is unchanged after JNK inhibition (Fig. 7J), the neurite localization of MAP1B-P was dramatically altered in response to JNK inhibition by SP600125 (Fig. 7K). The absence of MAP1B-P in neurites after JNK inhibition can be correlated with the strong reduction in levels of phosphorylated MAP1B after SP600125 treatment revealed by Western blotting (Fig. 6C). A role for JNK signaling in MAP1B phosphorylation was further confirmed by analysis of neurons derived from *jip1*<sup>-/-</sup> mice, in which we noted strong reduction in MAP1B-P levels, as shown by Western blotting (Fig. 6B) and immunocytochemistry (Fig. 7L, M).

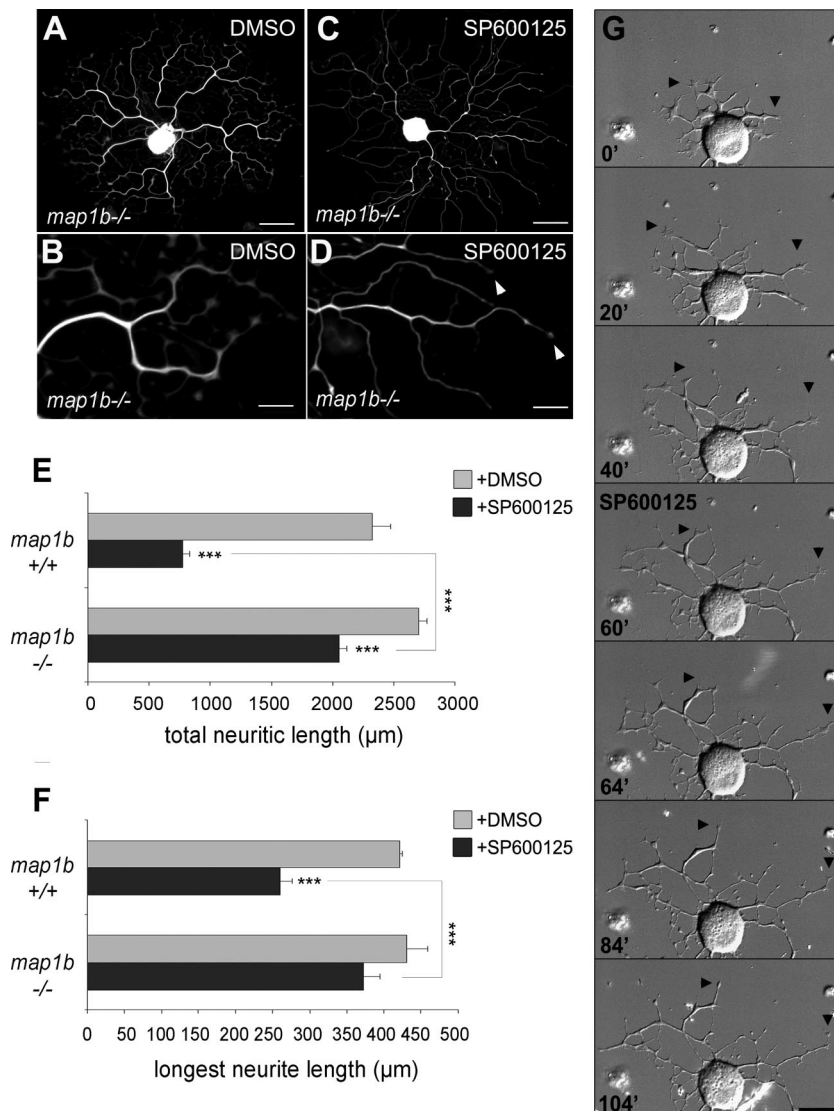
Next, we examined whether the overall levels of MAP1B-P, or its proximo-distal distribution along the neurite, depended on a



**Figure 8.** JNK1 and JNK2 regulate MAP1B phosphorylation. Immunostaining for total MAP1B (left row) and MAP1B-P (middle row; the right row represents merge images) of regenerating neurons from wild-type (A–C) and JNK-deficient mice. MAP1B-P levels are strongly reduced in neurites elongating from *jnk1*<sup>-/-</sup> (D–F) and *jnk2*<sup>-/-</sup> (G–I) neurons, whereas both the level and the proximo-distal distribution of MAP1B-P in *jnk3*<sup>-/-</sup> neurites (J–L) are similar to those of wild type. Scale bars, 50  $\mu$ m.

specific JNK isoform. Compared with wild-type neurons (Fig. 8A–C), immunocytochemical staining for MAP1B-P was markedly reduced in regenerating *jnk1*<sup>-/-</sup> and *jnk2*<sup>-/-</sup> neurons (Fig. 8D–I) but unchanged in *jnk3*<sup>-/-</sup> neurites (Fig. 8J–L). These results demonstrate that MAP1B phosphorylation is regulated through signaling by JNK1 and JNK2.

Finally, we used DRG neuron cultures derived from *map1b*<sup>-/-</sup> mice to investigate the functional relationship between MAP1B and JNK signaling during axonal regeneration. Whereas JNK inhibition by SP600125 treatment in wild-type neurons led to a rapid retraction of neurites (compare Fig. 2), no retraction was observed under the same conditions for *map1b*<sup>-/-</sup> neurons, as illustrated in Figure 9A–D. Quantitative analysis shows that JNK inhibition in *map1b*<sup>-/-</sup> neurons reduced their total neuritic length by only 24% (Fig. 9E) and their longest neurite by 3.6% (Fig. 9F), in contrast to wild-type neurons showing a reduction by 66.7 and 38.2%, respectively. The rather small effect of SP600125 treatment on *map1b*<sup>-/-</sup> neurons is also reflected in the distribution diagrams of total neurite length and longest neurite per neuron (supplemental Fig. S5, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material). The response of *map1b*<sup>-/-</sup> neurites to SP600125 treatment was further examined by time-lapse videomicroscopy (Fig. 9G; supplemental Video 6, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material). Before the addition of JNK inhibitor, *map1b*<sup>-/-</sup> neurites exhibit dynamic growth cones and elongate continuously.



**Figure 9.** MAP1B is required for neurite retraction induced by JNK inhibition. **A–D**, *map1b*<sup>-/-</sup> neurons after 48 h in culture, with or without SP600125 treatment, visualized by tubulin immunostaining. Note the increased number of branching points on the neurite tree of *map1b*<sup>-/-</sup> neurons. JNK inhibition by SP600125 addition to the culture medium induces growth arrest and loss of growth cones (arrowheads), but no retraction bulbs are observed along the neurites, in contrast to wild-type neurons (compare Fig. 2*G*). **E, F**, Quantitative analysis reveals no significant differences in total neurite length and length of longest neurites between *map1b*<sup>+/+</sup> and *map1b*<sup>-/-</sup> neurons under control conditions (gray bars); however, whereas SP600125 application (black bars) dramatically reduces total neurite length and length of longest neurites on *map1b*<sup>+/+</sup> neurons, JNK inhibition has only very limited effect on *map1b*<sup>-/-</sup> neurites ( $n \geq 52$ ). **G**, Time-lapse recording illustrating the typical response of a *map1b*<sup>-/-</sup> neurite: After SP600125 addition (at  $t = 60$  min), growth cones collapse, neurites stop extending, but no retraction is observed (arrowheads) (see movie in supplemental data S6, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material). Scale bars: **A, C**, 50 μm; **B, D**, 15 μm; **G**, 25 μm. \*\*\* $p < 0.001$ . Error bars indicate SEM.

Extension and retraction of filopodia is very similar to wild-type neurons, although the main neurites of *map1b*<sup>-/-</sup> neurons are highly branched (Bouquet et al., 2004). On SP600125 addition, growth cone filopodia collapse (that could explain the small reduction in total neuritic length), but in contrast to wild-type neurons, no neurite retraction occurs. Thus, the neurite retraction resulting from JNK inhibition depends on MAP1B.

## Discussion

Using both pharmacological JNK inhibition, and mice deficient for each JNK isoform and JIP1, we provide evidence that JNK signaling is involved in two processes crucial for axonal regeneration: initiation and elongation.

All three JNK isoforms are expressed in adult DRG neurons, but their specific roles in the regeneration process obviously differ. Lack of JNK2 or JNK3, but not JNK1, delays neurite initiation. Whereas JNK1 is present mainly in the cytoplasm of neuronal processes (Coffey et al., 2002), JNK2 and JNK3 can be translocated to the nucleus (Fig. 10*a*) (for review, see Raivich and Makwana, 2007) and affect transcription of regeneration-associated genes such as *c-jun*, whose role in promoting neuronal differentiation and neurite outgrowth is well characterized. JNK2 and JNK3 may act synergistically in phosphorylating numerous transcription factors including *c-Jun*, *junD*, *ATF3*, and *STAT3* (Kenney and Kocsis, 1998; Sheu et al., 2000; Lindwall et al., 2004). Their different functions in neurite initiation are likely attributable to different affinities for individual substrates (Kallunki et al., 1994; Gupta et al., 1996). During later phases of neurite extension, JNK2 and JNK1 (but not JNK3) are important. JNK1 deficiency only affects extension of preformed neurites, which indicates that JNK1 may play a major role in elongation by phosphorylating proteins that regulate microtubule dynamics, such as MAP1B, MAP2, and stathmin family proteins (Chang et al., 2003; Björklom et al., 2005; Tararuk et al., 2006; present data) (see below). Moreover, during later phases of neurite regeneration, JNK1 activity appears to be particularly affected by JIP1 deletion and *jnk1*<sup>-/-</sup> neurites are less sensitive to additional SP600125 treatment. Only JNK2 appears to act on both neurite initiation and elongation, the latter activity being at least in part related to phosphorylation of MAP1B. Not surprisingly, neurite regeneration is most severely affected in *jnk2*<sup>-/-</sup> mice.

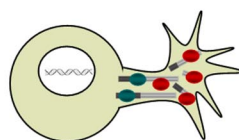
Although one may argue about the specificity of SP600125, our data clearly show that it does inhibit JNKs efficiently and has only very little effect on *jip1*<sup>-/-</sup> neurons. A previous study reported that this drug exhibited a higher affinity for JNK1 and JNK2 than for JNK3 (Bennett et al., 2001), corroborating our observations. The relatively strong effect of SP600125 seen in *jnk2*<sup>-/-</sup> and *jnk3*<sup>-/-</sup> neurites may be related, even partially, to changes in expression and/or activity of the remaining JNK isoforms. This phenomenon has been previously described for *jnk2*<sup>-/-</sup> mouse embryo fibroblasts in which increased JNK1 activity was noted (Jaeschke et al., 2006).

Interestingly, we did not observe any differences in neurite extension when comparing wild-type and JNK-deficient DRG neurons from embryonic day 13.5 embryos, despite a rather pronounced effect of pharmacological JNK inhibition on these neurons (data not shown). During development, lack of individual JNK proteins may be completely compensated by other isoforms,

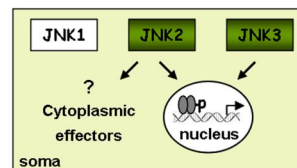
or even other signaling pathways/kinases. Accordingly, no specific function in neurite outgrowth from developing hippocampal or cortical neurons could be attributed to individual JNK isoforms (Eminel et al., 2008). The differential functions for individual JNKs, reported here, may thus be specific to adult regenerating neurites and particularly axons. Indeed, the DRG neurites examined in our study are most likely axons: they are all immunopositive for axon-specific cytoskeleton proteins such as tau and phosphorylated MAP1B (Soares et al., 2002; present data), but not stained for the dendrite marker protein MAP2. Furthermore, all neurites are immunostained by antibodies to JIP1, shown to be specifically enriched in nascent and mature axons (Dajas-Bailador et al., 2008). Lack of JIP1, in adult neurons, results in reduction in the phosphorylation and hence activation of JNK, inducing dramatic effects on neuritegenesis. However, abnormal neuritegenesis of *jip1*<sup>-/-</sup> neurons may not simply be a consequence of altered JNK signaling. JIP1 was shown to interact with kinesin-1 and doublecortin, thereby regulating polarization, transport, and axon outgrowth of cortical neurons (Gdalyahu et al., 2004; Horiuchi et al., 2005; Dajas-Bailador et al., 2008).

Phosphorylated JNK, and even more JIP1, are enriched in the distal part of growing neurites both during development (Whitmarsh et al., 2001; Gdalyahu et al., 2004; Tararuk et al., 2006) and axonal regeneration (present data). This is suggestive of a direct action of phospho-JNK on local molecular effectors, for which the following microtubule stability-regulating proteins are good candidates: DCX (Gdalyahu et al., 2004), tau (Yoshida et al., 2004), MAP2 and MAP1B (Chang et al., 2003), and microtubule-destabilizing protein SCG10 (Tararuk et al., 2006). Evidence for a direct action of JNKs on cytoskeleton dynamics is provided by our observation that regenerating growth cones and axon shafts respond to local application of JNK inhibitor within a few minutes, excluding transcriptional regulation. This response is associated with a rapid retraction, rather than a depolymerization of the microtubule network (Ahmad et al., 2000; Baas and Ahmad, 2001; Bouquet et al., 2007; Ertürk et al., 2007). A regulation of microtubule dynamics by JNKs in regenerating neurons is further supported by our *in vivo* data showing activation of JNKs in fibers undergoing lesion-induced sprouting. Increased levels of cytoplasmic phospho-JNK are maintained during the whole period of axonal remodeling, suggesting involvement of other JNK effectors, in addition to nuclear *c-jun* activation. Moreover, the clear association between JNK activation and MAP1B phosphorylation within sprouting fibers and, conversely, the decrease in MAP1B-P levels after JNK inhibition (and in JNK1-, JNK2-, and JIP1-deficient mice) argue for MAP1B being an effector of JNK

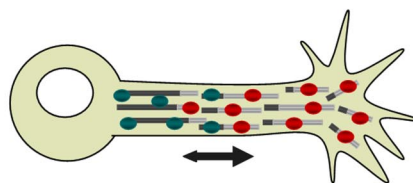
### a axonal initiation: involvement of JNK2 and JNK3



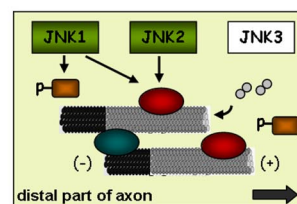
Nuclear effectors :  
c-jun, ATF2, STAT3



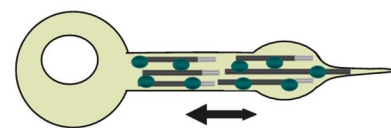
### b axonal outgrowth: involvement of JNK1 and JNK2



Cytoplasmic effectors: MAP1B, SCG10, DCX, kinesin, tau



### c axonal retraction induced by JNK inhibition: involvement of MAP1B

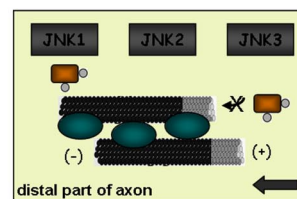


#### Microtubule

← stable domain  
← dynamic domain  
● Tubulin

● Unphosphorylated MAP1B  
● Phosphorylated MAP1B  
● Unphosphorylated stathmin  
● Phosphorylated stathmin

● Transcription factors  
➡ Extension forces (driven by dynein)  
➡ Retraction forces (actin-myosin contraction)



**Figure 10.** Model of JNK functions in axonal regeneration. This model is based on previous studies (see Discussion) and incorporates our present findings and hypotheses. **a**, Axotomy provokes an activation of JNK signaling leading to phosphorylation of transcription factors (including *c-jun*, ATF3, and STAT3) that, in turn, will regulate the expression of regeneration-associated genes. During early stages of DRG neuron regeneration, specific activation of JNK2 and JNK3 may mediate transcriptional events required for neurite growth initiation in the “correct” time window. **b**, JNK activation is maintained during neurite extension. Signaling by P-JNK1 and -2, present in axon shaft and growth cone, regulates cytoplasmic effectors that finally influence microtubule (MT) dynamics. Among cytoplasmic effectors of JNKs are MT-destabilizing (stathmin) and -stabilizing proteins (MAP1B). Phosphorylation of stathmin (by JNK1) family members overcomes their destabilizing function on MT plus ends, whereas phosphorylation of MAP1B (by JNK1 and JNK2) affects its binding to MTs or bind more labile MTs. **c**, JNK inhibition induces retraction of previously formed axons. In parallel, a decrease in MAP1B phosphorylation will increase the binding of MAP1B to MTs, enhancing their stabilization and protecting them from depolymerization. Process retraction, crucial for correct axonal pathfinding, is based on two independent molecular mechanisms: Extension forces generated by the action of dynein on MTs are counterbalanced by actin-myosin contractility resulting in axon retraction. The dynamic interaction between the two opposed forces likely involves competition between MAP1B and dynein.

signaling. The function of MAP1B in stabilizing axonal microtubules is regulated by phosphorylation through two mechanisms: mode II phosphorylation via casein kinase II is observed until adulthood. Mode I is developmentally regulated and implies PD-PKs (proline-directed protein kinases), such as GSK-3 $\beta$ , Cdk-5, and MAPK (JNKs) (Ulloa et al., 1993; Kawachi et al., 2003, 2005; Trivedi et al., 2005) (for references and review, see Riederer, 2007). Nonphosphorylated and mode II phosphorylated MAP1B is found in all compartments of neuronal cytoplasm. In contrast, mode I phosphorylated MAP1B (termed MAP1B-P) is absent from dendrites, preferentially localized to distal regions of the axon, and its distribution in growth cones resembles that of newly formed (mainly tyrosinated), highly dynamic microtubules (Baas and Black, 1990; Ahmad et al., 1993; Bouquet et al., 2004, 2007). These observations, as well as the data provided by Mack et al. (2000) showing that phosphorylated MAP1B is required for microtubule stabilization in the growth cone, would suggest that MAP1B-P is implicated in regulating microtubule dynamics in

highly motile regions of neurites (Scales et al., 2009). In regenerating DRG neurites, this MAP1B phosphorylation should be mainly regulated by JNK1 and JNK2 since its level is not affected by lack of JNK3 (Fig. 10b).

Moreover, the requirement of MAP1B for neurite retraction after JNK inhibition is consistent with MAP1B being a crucial downstream effector of JNK signaling during reorganization of neuronal cytoskeleton in the adult nervous system. We previously demonstrated that MAP1B regulates growth cone turning and branching of regenerating neurites (Bouquet et al., 2004) and provided physiological evidence for its requirement in coupling microtubule and actin filament dynamics during the retraction of cell processes in response to extracellular stimuli involving the RhoA/ROCK GTPases (Bouquet et al., 2007; Stroissnigg et al., 2007). This MAP1B-dependent neurite retraction does not result in microtubule depolymerization, which indicates that axon extension and retraction are based on two interconnected molecular mechanisms: The forces driving axon extension are generated by the action of dynein on microtubules, counterbalanced by contractile forces based on actin–myosin activity leading to axon retraction (Ahmad et al., 2000). This dynamic interaction between the two opposed forces might involve a competition between MAP1B and dynein, since the microtubule-binding sequence of these two proteins is similar (Koonce and Tikhonenko, 2000) [see also discussion in the study by Stroissnigg et al. (2007)]. The function of MAP1B in axon retraction appears to be regulated by posttranslational modifications, notably nitrosylation induced by nNOS (neuronal nitric oxide synthase) activation (Stroissnigg et al., 2007). Here, we extend this notion by demonstrating that decreasing the phosphorylation level of MAP1B by JNK inhibition further confirms its involvement in neurite retraction, whereas neurite extension is obviously facilitated by JNK signaling (probably activated by Rac1) (Weston and Davis, 2002) that converges on MAP1B. Moreover, the two opposed processes (i.e., MAP1B nitrosylation leading to neurite retraction and JNK-induced phosphorylation leading to extension) are likely to be interdependent since exogenous nitric oxide negatively regulates the activity of JNKs (Pei et al., 2008). This fine-tuning of the function of MAP1B seems a good example for the convergence of a wide range of signaling pathways, activated in response to extracellular guidance cues, on the regulation of cytoskeletal dynamics. At the same time, a single signaling pathway may act on cytoskeleton-associated proteins exerting opposite functions: As we show here (Fig. 10), JNK signaling affects both the microtubule-stabilizing protein MAP1B and the destabilizing stathmin family proteins.

Together, our data clearly demonstrate the differential involvement of JNK isoforms in controlling cytoskeleton reorganization, notably by regulating the function of MAP1B. Our study therefore provides new insight into how extracellular cues will eventually regulate guidance, branching, and retraction of neurites through orchestrated rearrangement of the cytoskeleton, triggered by specific intracellular signaling cascades.

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