Inhibition of Jun NH$_2$-Terminal Kinases Suppresses the Growth of Experimental Head and Neck Squamous Cell Carcinoma


Abstract

**Purpose:** This study was carried out to investigate whether c-Jun NH$_2$-terminal kinases (JNK) are potential targets for treating head and neck squamous cell carcinoma (HNSCC).

**Experimental Design:** JNK activity was first evaluated in 20 paired samples of human HNSCC. The antitumor activity of SP600125, a reversible nonselective ATP-competitive inhibitor of JNKs, was then investigated both in an HNSCC xenograft model and in vitro using immunohistochemistry, immunoblotting, enzyme immunoassay, flow cytometry, and a Matrigel assay of capillary tube formation. Complementary studies were carried out using small interfering RNA to JNK1/2.

**Results:** JNK activity was increased in human HNSCC compared with normal-appearing epithelium. Treatment of mice bearing HNSCC xenografts with SP600125 resulted in 760% inhibition of tumor growth relative to vehicle-treated animals. Inhibition of tumor growth was associated with significant reductions in both cell proliferation and microvessel density. SP600125 inhibited tumor cell proliferation by causing delays in both the S and G$_2$-M phases of the cell cycle. Inhibition of angiogenesis seemed to reflect effects on both tumor and endothelial cells. The JNK inhibitor suppressed the production of vascular endothelial growth factor and interleukin-8 by tumor cells and also inhibited endothelial cell proliferation and capillary tube formation. Reduced amounts and phosphorylation of epidermal growth factor receptor were found in tumor cells after treatment with SP600125. Small interfering RNA–mediated suppression of JNK1/2 led to reduced tumor cell proliferation and decreased levels of epidermal growth factor receptor, vascular endothelial growth factor, and interleukin-8.

**Conclusions:** JNK activity is commonly increased in HNSCC. Our preclinical results provide a rationale for evaluating JNK inhibition as an approach to treating HNSCC.

The c-Jun NH$_2$-terminal kinases (JNK) are members of the mitogen-activated protein kinase family that, when stimulated, regulate a variety of cellular activities, including proliferation, differentiation, and tumorigenesis (1, 2). There are three JNK genes. JNK1 and JNK2 gene expression is ubiquitous, whereas JNK3 expression is largely restricted to the heart, testis, and brain (3). The JNKs are activated in response to environmental stresses, inflammatory cytokines, and extracellular stimuli, including epidermal growth factor (4, 5). JNKs are serine/threonine protein kinases that phosphorylate a number of transcription factors, including the c-Jun component of the activator protein-1 (AP-1) transcription factor complex. AP-1 activation is important in cell transformation (6) and skin tumor formation in mice (7). In fact, JNK activation and c-Jun phosphorylation are required for transformation induced by Ras, an oncogene that is mutationally activated in ~30% of human malignancies (8) and frequently overexpressed in head and neck squamous cell carcinoma (HNSCC; ref. 9). Genetic and pharmacologic approaches have been used to evaluate the potential importance of JNKs in tumor formation and growth. Antisense oligonucleotides to JNKs suppressed the growth of tumor cells and inhibited the growth of PC3 prostate cancer xenografts (10). SP600125, a reversible, nonselective ATP-competitive inhibitor of JNK, suppressed the growth of both human prostate carcinoma xenografts and murine Lewis lung carcinoma (11). In another study, JNKs were found to be activated in a subset of human non–small cell lung cancers and to promote oncogenesis in the bronchial epithelium (12).

Although there is emerging preclinical evidence that inhibiting JNKs may be a useful approach to suppressing tumor growth, little is known about which tumor types to target. In the present study, we first determined that JNK activity was increased in human HNSCC compared with normal epithelium. Subsequently, we showed that SP600125 suppressed the growth of HNSCC xenografts by inhibiting tumor cell proliferation and angiogenesis. Complementary in vitro studies suggested that the JNK inhibitor acted on both tumor and endothelial cells to mediate tumor growth.
inhibition. Taken together, these data highlight the potential importance of targeting JNKs as a therapeutic strategy for treating HNSCC.

Materials and Methods

**Materials.** DMEM/F12 and fetal bovine serum were from Invitrogen. Keratinocyte growth medium, EBM-2, and SingleQuots kits were from Clonetics Corp. Eagle’s MEM was from American Type Culture Collection, and RPMI 1640 was from Life Technologies. SP600125 was a gift from the Signal Research Division of Celgene Corp. Matrigel was from BD Biosciences and male immunodeficient nu/nu mice were from Charles River Laboratories. The stress-activated protein kinase/JNK assay kit was from Cell Signaling Technology. Antibodies to human phosphorylated c-Jun, JNK, total epidermal growth factor receptor (EGFR), and the phosphorylated c-Jun standard were from Santa Cruz Biotechnology. Anti–phosphorylated EGFR antiserum was from PharMingen. Anti–phosphorylated c-Jun, JNK activity was assessed in human and xenograft tissues using a stress-activated protein kinase/JNK assay kit. Lysates were prepared by probe sonication of tissue specimens in lysis buffer for 3 × 10 s on ice and centrifuging at 10,000 × g for 10 min to sediment particulate material. The protein concentration of the supernatant was measured by the method of Lowry et al. (19). Tissue lysate protein (100 µg) was diluted with lysis buffer to a volume of 250 µL and incubated overnight with 20 µL c-Jun fusion beads at 4 °C. Lysates were then re centrifuged and washed with lysis buffer (×2) and kinase buffer (×2) to eliminate nonspecific binding. Lysates were suspended in kinase buffer supplemented with 100 µmol/L ATP and incubated for 30 min at 30 °C. The reaction was terminated and c-Jun phosphorylation was assessed via Western blot analysis.

**Immunohistochemistry.** Proliferation and angiogenesis were evaluated by staining for Ki-67 and CD34 as described previously (20–22). Neutral buffered formalin-fixed tissue was embedded in paraffin. Tissue sections (5 µm) were prepared using a microtome and mounted on slides. Sections were deparaffinized in xylene, rehydrated in graded alcohols, and washed in distilled water. Endogenous peroxidase was quenched with 0.01% H2O2. Antigen retrieval was done by microwaving the sections in 10 mmol/L citric acid (pH 6.0) for 30 min. In addition, sections for Ki-67 analysis were treated with 0.05% trypsin, 0.05% CaCl2 in Tris-HCl (pH 7.6) for 5 min at 37 °C before microwave treatment. The slides were washed thrice in PBS and blocked for 30 min with 10% normal rabbit serum (CD34) or 10% normal horse serum (Ki-67). Tissue sections were then incubated with antisense to mouse CD34 at 25 µg/mL and antisense to Ki-67 at 1:5,000 dilution (2% bovine serum albumin in PBS) and incubated overnight at 4 °C. After being washed thrice with PBS, the sections were incubated with biotinylated anti-mouse immunoglobulins at 1:100 (CD34) or 1:500 (Ki-67) dilution for 30 min at room temperature. The slides were then washed thrice in PBS and labeled using 1:25 avidin-biotin peroxidase complexes (Vector Stain) for 30 min at room temperature. The reaction was visualized using 3,3-diaminobenzidine. Subsequently, the slides were rinsed in tap water and counterstained with hematoxylin. The slides were then dehydrated with ethanol, rinsed with xylene, and mounted.

Proliferation was assessed by counting the number of tumor cells with Ki-67–positive nuclei and the total number of tumor cells at ×400 magnification in three representative regions of the tumor. Results are expressed as the proliferation index: proportion of positively staining cells over the total number of cells. Microvessel density was assessed by counting the number of microvessels at ×400 magnification in three fields that had the highest vascularization. The results are expressed as an average number of microvessels per field.

**Western blotting.** Lysates were prepared by treating cells with lysis buffer [100 mmol/L Tris-HCl (pH 7.5), 50 mmol/L NaCl, 50 mmol/L NaF, 30 mmol/L sodium Ppi, 1 mmol/L EDTA, 1% Tween 20, 1 mmol/L NaVO3, 5 µg/mL aprotinin, complete mini protease inhibitor mixture, and 5 µmol/L 3,4-dichlorocoumarin]. Lysates were sonicated for 5 min on ice and centrifuged at 10,000 × g for 10 min to sediment the particulate material. The protein concentration of the supernatant was assessed by measurements of cell counts and lactate dehydrogenase release. There was no evidence of cytotoxicity in any of the experiments.

**JNK activity.** JNK activity was assessed in human and xenograft tissues using a stress-activated protein kinase/JNK assay kit. Lysates were prepared by probe sonication of tissue specimens in lysis buffer for 3 × 10 s on ice and centrifuging at 10,000 × g for 10 min to sediment particulate material. The protein concentration of the supernatant was measured by the method of Lowry et al. (19). Tissue lysate protein (100 µg) was diluted with lysis buffer to a volume of 250 µL and incubated overnight with 20 µL c-Jun fusion beads at 4 °C. Lysates were then re centrifuged and washed with lysis buffer (×2) and kinase buffer (×2) to eliminate nonspecific binding. Lysates were suspended in kinase buffer supplemented with 100 µmol/L ATP and incubated for 30 min at 30 °C. The reaction was terminated and c-Jun phosphorylation was assessed via Western blot analysis.

**Fig. 1.** JNK activity is increased in HNSCC. JNK activity was measured by immunoprecipitation of JNK coupled with a kinase assay using c-Jun as substrate. JNK activity was consistently elevated in HNSCC (7) compared with adjacent normal-appearing epithelium (NT; subjects 1-5) or normal oral epithelium (NT) from healthy volunteers (subjects 6-8).

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measured by the method of Lowry et al. (19). SDS-PAGE was done under reducing conditions on 7.5% or 10% polyacrylamide gels as described by Laemmli (23). The resolved proteins were transferred onto nitrocellulose sheets as detailed by Towbin et al. (24). The nitrocellulose membrane was then incubated with primary antibodies. Secondary antibody to IgG conjugated to horseradish peroxidase was used. The blots were probed with enhanced chemiluminescence Western blot detection system according to the manufacturer’s instructions.

**Cell cycle analysis.** Cells were plated (2 × 10^5 per well) in 96-well plates and were allowed to adhere overnight before being treated with vehicle or SP600125. Fresh medium containing vehicle or the indicated concentration of SP600125 was added every 2 days. At each time point, the culture medium was removed and replaced with MIT (0.5 mg/mL). The reaction was stopped 3 h later by removing the medium and immediately solubilized by adding 100 μL DMSO. Absorbance was measured after 10 min at 560 nm in a 96-well plate reader.

**DNA synthesis.** Incorporation of [3H]thymidine was used to measure DNA synthesis. 1483 and D3 cells plated (2 × 10^5 per well) in 96-well plates were allowed to adhere overnight before being treated with vehicle or SP600125. After 18 h of treatment at 37°C, the medium was supplemented with [3H]thymidine (0.1 μCi/well) for 6 h. Cells were then washed thrice with PBS. Radioactivity was measured with a LS6800 liquid scintillation counter from Beckman.

**Cell cycle analysis.** 1483 cells were treated with vehicle (0.1% DMSO) or 25 μmol/L SP600125 for 24 h. The cells were released by treatment with trypsin-EDTA, washed, and suspended in ice-cold PBS (pH 7.4), counted, and fixed overnight in 50% ethanol at 4°C. The cells (10^6/mL) were then resuspended in 1.25% sodium citrate with RNase A (500 units/mL) for 30 min at 37°C. Propidium iodide (30 μg/mL) was added and the cells were maintained in the dark for 30 min at room temperature. The red fluorescence of single events was recorded using an argon ion laser at 488-nm excitation wavelength and 610-nm emission wavelength to measure DNA index on a Coulter Epics XL flow cytometer. The percentage of cells present in each phase of the cell cycle was determined using ModFitLT V2.0 software from Verity Software House.

**VEGF production.** Amount of VEGF produced was quantified with Quantikine human VEGF immunoassay kit according to the manufacturer’s instructions.

**Measurements of IL-8.** Total RNA was isolated using the RNeasy Mini Kit according to manufacturer’s instructions. Reverse transcription was done using 2 μg of RNA per 50 μL of reaction. The reaction mixture contained 1 × PCR Buffer II, 2.5 μmol/L MgCl2, 0.5 mmol/L deoxynucleotide triphosphates, 2.5 μmol/L oligo(dT)16 primer, 50 units RNase inhibitor, and 125 units murine leukemia virus reverse transcriptase (Roche Applied Science). Samples were amplified in a thermocycler for 10 min at 25°C, 15 min at 42°C, 5 min at 99°C, and 5 min at 5°C. The resulting cDNA was used for amplification. The volume of the PCR reaction was 25 μL and contained 5 μL of cDNA, 1 × PCR Buffer II, 2.5 μmol/L MgCl2, 0.4 μmol/L deoxynucleotide triphosphates, 400 nmol/L forward primer, 400 nmol/L reverse primer, and 2.5 units Taq polymerase (Applied Biosystems). Samples were denatured at 95°C for 2 min and then amplified for 35 cycles in a thermocycler under the following conditions: 95°C for 30 s, 62°C for 30 s, and 72°C for 45 s. Subsequently, the extension was carried out at 72°C for 10 min. Primer sequences were as follows: IL-8, sense 5'-AGGTTGCAAGATCCATAC-3', antisense 5'-AACACGAGGACGTGCAG-3', β-actin, sense 5'-GCTACCCACAGCCTCACC-3', antisense 5'-GGATGCCCCAGAATCCATG-3'. PCR products were subjected to electrophoresis on a 1% agarose gel with 0.5 μg/mL ethidium bromide. The identity of each PCR product was confirmed by DNA sequencing.

Amount of IL-8 protein produced was determined with a Quantikine human IL-8 immunoassay kit according to the manufacturer’s instructions.

**RNA interference.** JNK1/JNK2 targeting small interfering RNA (siRNA) was custom synthesized by Dharmacon. Sequences were as follows: sense 5'-TGAAGAAGTGCTCTGTCACATT-3' and antisense 5'-AAGGCACATCTTTCA-3' (25). Non-specific control siRNA was also obtained from Dharmacon. Cells were seeded in DMEM and 10% fetal bovine serum for 24 h before transfection. siRNA to JNK1/2 or non-specific siRNA (both 100 pmol/mL) was transfected using
DharmaFECT 4 transfection reagent according to the manufacturer's instructions.

**Angiogenesis.** Matrigel (150 μL/well) was added to 48-well plates and incubated at 37°C for at least 30 min before use. D3 cells grown to confluency were detached by treatment with trypsin-EDTA, counted, and resuspended in 1:4 diluted CT2X to a concentration of 1 × 10^4 cells/mL. The cells were then mixed with 0 to 20 μmol/L SP600125, plated onto the Matrigel surface of the 48-well plates, and incubated at 37°C for 18 h. Capillary tube formation was assessed. Representative images were captured digitally for each condition at ×40.

**Statistics.** Xenograft tumor growth was compared using repeated measures ANOVA to the cubed-root of volumes. Statistical significance of difference in growth rate between SP600125 and control was based on comparison of linear trends in the transformed scale. Results are expressed descriptively as the mean ± SE. All other comparisons between groups were made by the two-tailed Student's t test with results presented as the mean ± SD using Microsoft Excel 2000 (Microsoft Corp.). A difference between groups of P < 0.05 was considered significant.

**Results**

**JNK activity is increased in HNSCC.** JNK activity was evaluated by Western blot in 20-paired HNSCC specimens (tumor versus adjacent normal appearing mucosa). Activity was also determined in oral mucosal biopsies from 10 healthy volunteers. A representative Western blot is shown in Fig. 1.

JNK activity was elevated in HNSCC compared with normal mucosa. More specifically, JNK activity was easily detected in 95% of tumor samples but only 10% of adjacent normal-appearing mucosal samples. JNK activity was not detected in control specimens from healthy volunteers.

**Inhibiting JNK suppresses the growth of HNSCC xenografts.** To explore the role of JNKs as potential therapeutic targets, 1483 cell HNSCC xenografts were established and confirmed to have JNK activity similar to human tumors (Fig. 2A). Subsequently, nude mice xenografted with 1483 cells were randomized to treatment with either vehicle or SP600125, a JNK inhibitor. Daily treatment with SP600125 (50 mg/kg) for ~1 month resulted in >60% reduction in tumor volume (P < 0.001; Fig. 2B). Subsequently, measurements of cell proliferation and angiogenesis were carried out to explore the potential mechanisms underlying the antitumor activity of the JNK inhibitor. Nuclear staining of Ki-67 was used to assess cell proliferation. Treatment with SP600125 led to an ~36% decrease in cell proliferation (P = 0.001; Fig. 2Ca, Cb, and D).

Angiogenesis was evaluated by staining for CD34, an antigen present in endothelial cells. Microvessel density decreased by >50% in tumors from mice treated with SP600125 (P < 0.001; Fig. 2Cc, Cd, and D). Taken together, these findings suggest that JNK activity is an important determinant of both tumor cell proliferation and angiogenesis in HNSCC.

**Inhibition of JNKs suppresses tumor cell proliferation and angiogenesis.** Experiments were next carried out to further...
Fig. 4. JNK plays a role in tumor cell production of VEGF and IL-8 and endothelial capillary tube formation. A, 1483 cells were treated with 0 to 50 μmol/L SP600125 for 18 h before measuring VEGF protein production. B, VEGF production by 1483 cells is reduced after treatment with siRNA to JNK1 and JNK2. Following transfection with either nonspecific siRNA or siRNA to JNK1 and JNK2, VEGF production was measured. C, 1483 cells were treated with 0 to 25 μmol/L SP600125 for 24 h (top) or transfected with nonspecific siRNA or siRNA to JNK1 and JNK2 for 36 h (bottom). Subsequently, amounts of IL-8 and β-actin mRNA were determined by reverse transcription-PCR. D, 1483 cells were treated with 0 to 25 μmol/L SP600125 for 24 h before measuring the amount of IL-8 protein released into the medium. E, D3 endothelial cells were treated with 0 to 20 μmol/L SP600125 and plated onto Matrigel. Capillary tube formation was assessed qualitatively after 18 h of treatment. Representative images are presented at × 40 magnification. F, effect of SP600125 on endothelial cell proliferation. Cells were treated with the indicated concentration of SP600125 for 5 d. Cell number was assessed by MTT assay and expressed relative to vehicle-treated cell number. G, effect of SP600125 on DNA synthesis in endothelial cells. Cells were treated with 0 to 20 μmol/L SP600125. DNA synthesis was quantified after 18 h of treatment. [3H]thymidine incorporation was expressed relative to vehicle-treated cells. Columns (A, B, D, and G), mean; bars, SD; n = 6; *P < 0.05, **P < 0.01, ***P < 0.005, and ****P < 0.001.
In the current study, we focused on the potential significance of JNKs as therapeutic targets for the treatment of HNSCC. The choice to investigate JNKs was based on prior evidence that JNKs and the AP-1 transcription factor complex are important for cell transformation, growth, differentiation, and angiogenesis. Interestingly, tobacco smoke can activate JNKs, at least in part, by oxidative stress (39). We detected a marked increase in JNK activity in HNSCC (37). Several potential mechanisms can explain the increased JNK activity in HNSCC. JNKs are activated by receptor tyrosine kinases including the EGFR (4). For example, activation of EGFR signaling stimulates the Ras-Raf-mitogen-activated protein kinase subfamily, activated extracellular signal-regulated kinases (ERK1/2), an increase in JNK activity in HNSCC. JNKs are activated by receptor tyrosine kinases, cellular stress and cytokines activate JNKs (4). The fact that the survival benefits were modest underscores the need for additional therapies.

Targeted molecular therapies hold great promise for cancer treatment. In the case of aerodigestive malignancies, therapies that target the EGFR were recently found to improve the survival of patients with advanced-stage HNSCC (32) and non–small cell lung cancer (33). The survival benefits were modest underscores the need for additional therapies.

In vitro studies were carried out to evaluate the role of JNKs in angiogenesis. In the current study, we focused on the potential significance of JNKs as therapeutic targets for the treatment of HNSCC. The choice to investigate JNKs was based on prior evidence that JNKs and the AP-1 transcription factor complex are important for cell transformation, growth, differentiation, and angiogenesis (1, 34–36). We detected a marked increase in JNK activity in HNSCC (37). Several potential mechanisms can explain the increased JNK activity in HNSCC. JNKs are activated by receptor tyrosine kinases including the EGFR (4). For example, activation of EGFR signaling stimulates the Ras-Raf-mitogen-activated protein kinase pathway. Because EGFR signaling is commonly activated in HNSCC (38), this is likely to contribute to the observed increase in JNK activity. In addition to receptor tyrosine kinases, cellular stress and cytokines activate JNKs (4). Interestingly, tobacco smoke can activate JNKs, at least in part, by oxidative stress (39). Each of these factors may contribute to the increase in JNK activity found in HNSCC.

To evaluate the potential significance of increased JNK activity in HNSCC, we used an HNSCC xenograft model.
Treatment with SP600125 caused a significant reduction in tumor growth. To investigate the mechanism underlying the growth-inhibitory effects of SP600125, measurements of cell proliferation and microvesSEL density were done. Treatment with SP600125 caused a significant reduction in both tumor cell proliferation and microvesSEL density. Subsequently, in vitro studies were carried out to further evaluate the importance of JNK activity in regulating tumor cell growth and angiogenesis. Tumor cell proliferation and [3H]thymidine incorporation were suppressed by SP600125. In fact, the JNK inhibitor suppressed the growth of several cell lines, including those derived from leukoplakia, HNSCC, and a lymph node metastasis. Because small-molecule inhibitors can have off-target effects, siRNA to JNK1/2 was also used. Consistent with the findings for SP600125, down-regulation of JNK1 and JNK2 led to a significant reduction in DNA synthesis. The JNK inhibitor also caused an increase in the percentage of cells in the S and G2-M phases of the cell cycle. JNK activity, which is increased in late S and G2, has been shown to play a key role in cell cycle progression (34). Notably, treatment with SP600125 results in a reproducible slowing of cell cycle progression through the S and G2 phases and induction of G2-M arrest in multiple other models (11, 25). The observed decrease in microvesSEL density in xenografts from mice treated with SP600125 could reflect effects on either tumor or endothelial cells. Notably, the AP-1 transcription factor complex can stimulate VEGF gene expression, a proangiogenic factor that is a recognized therapeutic target (40). We showed that treatment with either SP600125 or siRNA to JNK1/2 led to a significant reduction in VEGF production by 1483 cells. This finding is consistent with previous evidence that JNKs can regulate VEGF production in human gingival fibroblasts (41). Because VEGF can also stimulate JNK activity, these results are consistent with the idea that JNKs can regulate VEGF expression by mitogen-activated protein kinase (MAPK) signaling. The regulation of AP-1 activity by mitogen-activated protein kinase (MAPK) signaling has been shown to be mediated by the JNK family of enzymes, which play a key role in the regulation of cell proliferation and apoptosis (35). It is possible that a JNK inhibitor could disrupt the positive feedback loop that is important for tumor growth in addition to suppressing the production of VEGF, inhibition of JNKs led to reduced production of IL-8, a chemokine implicated in both HNSCC and angiogenesis (26–28). This finding is consistent with previous evidence that JNK activity can regulate IL-8 expression (42). Additionally, the JNK inhibitor had direct effects on endothelial cells. More specifically, endothelial cell proliferation and capillary tube formation were inhibited by treatment with the JNK inhibitor. Activation of EGFR signaling can activate JNKs and stimulate tumor cell proliferation and angiogenesis (4). Conversely, increased JNK signaling can induce the expression of EGF, suggesting the existence of a positive feedback loop (29, 30). We showed that inhibition of JNKs led to reduced expression and phosphorylation of EGF. The strong interdependence of these two molecules lends further support to the notion of targeting JNK activity as a strategy to inhibit tumor growth.

JNK activity is important for cell proliferation and cell cycle progression (34). As mentioned above, we found that JNK inhibition resulted in an increased percentage of cells in the S and G2-M phases of the cell cycle. Tumor cells in G2-M are highly sensitive to radiation (43). In fact, pharmacologic inhibition of JNKs has been reported to inhibit the repair of radiation-induced DNA damage and to increase radiation sensitivity (44). Both cisplatin and radiation, which are commonly used in combination for treatment of HNSCC, may be efficacious in cell killing in the S phase (45, 46). JNKs can be activated by radiation (47) and DNA-damaging agents such as cisplatin (48). Furthermore, activation of JNKs has been linked to DNA repair, raising the possibility that inhibition of JNKs could augment the activity of therapies that induce DNA damage (49). Based on the current findings, it would be logical to determine whether inhibition of JNKs before concurrent chemoradiation increased cell kill in HNSCC.

Collectively, these data provide a strong rationale for pursuing additional studies of JNKs as potential therapeutic targets in HNSCC and possibly other malignancies. Other studies have suggested that JNKs can act as a tumor suppressor in fibroblast transformation (50). Hence, it will be important to gain a deeper understanding of the mechanism for the different roles of JNKs in tumors if the full potential of JNK inhibitors as anticancer therapies is to be understood. As for many other treatments, inhibitors of JNKs may prove useful in the treatment of certain tumor types but not others. The development of compounds with greater selectivity for individual JNK isoforms than SP600125 will provide new insights into the role of JNK1, JNK2, and JNK3 in carcinogenesis. Because JNKs are relevant to cell biology in general, developing agents with an acceptable therapeutic index may also prove challenging until more selective agents are developed.

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References


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