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Induction of apoptosis in Neurofibromatosis Type 1 malignant peripheral nerve sheath tumor cell lines by a combination of novel farnesyl transferase inhibitors and lovastatin

Jonathan W. Wojtkowiak, Farid Fouad, Daniel T. LaLonde, Miriam D. Kleinman, Richard A. Gibbs, John J. Reiners Jr., Richard F. Borch, and Raymond R. Mattingly

Wayne State University School of Medicine, Department of Pharmacology (J.W.W., D.L., J.J.R., R.R.M), Detroit, Michigan; Barbara Ann Karmanos Cancer Institute, Programs in Molecular Biology and Human Genetics (R.R.M.) and in Proteases (J.J.R.), Detroit, Michigan; Wayne State University, Institute of Environmental Health Sciences (M.D.K., J.J.R.) and EHS Center in Molecular and Cellular Toxicology with Human Applications (J.J.R., R.R.M.), Detroit, Michigan; and Purdue University, Medicinal Chemistry and Molecular Pharmacology and Cancer Center (F.F., R.A.G., R.F.B.), West Lafayette, Indiana.

Abstract

Neurofibromatosis Type 1 (NF1) is a genetic disorder that is driven by the loss of neurofibromin (Nf) protein function. Nf contains a Ras GTPase activating domain (Ras-GAP), which directly regulates Ras signaling. Numerous clinical manifestations are associated with the loss of Nf and increased Ras activity. Ras proteins must be prenylated in order to traffic and functionally localize with target membranes. Hence, Ras is a potential therapeutic target for treating NF1. We have tested the efficacy of two novel farnesyl transferase inhibitors (FTI), **1** and **2**, alone or in combination with lovastatin, on two NF1 malignant peripheral nerve sheath tumor (MPNST) cell lines, NF90-8 and ST88-14. Single treatments of **1**, **2**, or lovastatin had no effect on MPNST cell proliferation. However, low micromolar combinations of **1** or **2** with lovastatin (FTI/lovastatin) reduced Ras prenylation in both MPNST cell lines. Further, this FTI/lovastatin combination treatment reduced cell proliferation and induced an apoptotic response as shown by morphological analysis, pro-caspase-3/-7 activation, loss of mitochondrial membrane potential, and accumulation of cells with sub G₁ DNA content. Little to no detectable toxicity was observed in normal rat Schwann cells following FTI/lovastatin combination treatment. These data support the hypothesis that combination FTI plus lovastatin therapy may be a potential treatment for NF1 MPNSTs.

Introduction

Neurofibromatosis Type 1 (NF1) is a genetically inherited syndrome that affects approximately 1:3000 individuals (Arun and Gutmann, 2004). NF1 presents with an array of clinical manifestations that can arise during early development through adulthood, including increased pigmentation of the skin (café au lait macules), Lisch nodules of the iris, learning disabilities, and abnormal development of the skeletal system (Lynch and Gutmann, 2002). NF1 is characterized by the development of benign peripheral nerve sheath tumors (BPNST) or neurofibromas. Approximately 10% of NF1 patients experience tumor transformation to the more aggressive malignant peripheral nerve sheath tumors (MPNST) (Ward and Gutmann, 2005). Progression toward MPNST is a leading cause of increased mortality for NF1 patients. Therapies are limited to excision of neurofibromas, radiation of plexiform

neurofibromas, and the use of cytotoxic compounds. Although excision of tumors is the primary form of treatment, the tumors tend to return (Packer et al., 2002). A molecularly targeted therapy designed against the molecular background of NF1 may reveal more effective approaches for treatment of NF1 (Dilworth et al., 2006).

The molecular pathogenesis of NF1 was better understood following the discovery of the *NF1* gene, which encodes the protein neurofibromin (Nf). Nf contains a Ras GTPase activating protein (Ras-GAP) domain (DeClue et al., 1991). This domain is responsible for controlling Ras signaling by increasing the intrinsic rate of Ras hydrolysis, thus converting the active Ras-GTP to the inactive Ras-GDP form (Eccleston et al., 1993). Germline mutations of the *NF1* gene result in reduced Nf expression and a loss of Ras-GAP activity. The consequence of losing Ras-GAP activity is aberrant Ras signaling that can potentially lead to the development of NF1 (Basu et al., 1992; Feldkamp et al., 1999). Our lab and others have previously targeted downstream signaling partners of Ras by treating MPNST cell lines with MEK inhibitors (Tang et al., 1998; Chadee and Kyriakis, 2004; Mattingly et al., 2006; Roth et al., 2007). We have shown that PD184352 (CI-1040) induced apoptosis in MPNST cell lines, confirming the dependence of the Ras-MAPK pathway in this disease (Mattingly et al., 2006).

Ras proteins are translated in the cytoplasm as inactive precursor molecules that must undergo a series of post-translational modifications before the protein can fully function (Gibbs et al., 2001). The first necessary step is the covalent addition of a prenyl group, either a 15C farnesyl or a 20C geranylgeranyl group, to the C-Terminal "CaaX" box (Basso et al., 2006).

Reducing the prenylation of proteins to treat NF1 has been recognized as a potential therapeutic approach. For example, the farnesyl transferase inhibitor (FTI) BMS-186511 reduces proliferation of MPNST cell line ST88-14 (Yan et al., 1995), and FTI L-739-749 reduces proliferation of Nf-deficient mouse Schwann cells (Kim et al., 1997). A phase I clinical trial utilizing FTI tipifarnib to treat plexiform neurofibromas was tolerated well in children, but no objective responses were achieved (Widemann et al., 2006). Although this study has advanced to an ongoing Phase II trial (NCT00029354), it is likely that further development of this treatment approach will be required.

Our lab is interested in utilizing FTIs and lovastatin, an inhibitor of the HMG-CoA reductase, to reduce prenylation of proteins as a potential therapy for numerous diseases. We have previously reported that lovastatin, in combination with FTI 3-allylfarnesol, induces relocation of RhoB from the membrane fraction to the cytosolic fraction following treatment in A10 vascular smooth muscle cells. The translocation of RhoB from the membrane to the cytosol is the result of inhibiting RhoB prenylation (Mattingly et al., 2002). A prodrug analog of 3-allylfarnesol phosphate was also shown to inhibit RhoB prenylation in STS-26T MPNST cells when used in combination with lovastatin, resulting in reduced cell proliferation (Clark et al., 2007).

Here we describe our efforts to characterize the effectiveness of two novel FTase inhibitors, **1** (Clark et al., 2007) and **2**, on human NF1 MPNST cell lines, NF90-8 and ST88-14. The prodrug structures are shown in Fig. 1. Prodrug **1** is highly lipophilic, so an analog **2** was prepared in which a carboxylate side chain, which would be ionized at physiologic pH, replaced the N-methyl group on the prodrug moiety. The entire prodrug moiety is released from the inhibitor following activation inside the cell (Clark et al., 2007), so this modification will have no effect on inhibitor affinity. Our objective was to determine the efficacy of these compounds on Ras prenylation and cell proliferation when used in combination with lovastatin. The data show reduction of Ras prenylation in both cell lines

with cell cycle G₁ arrest and increased caspase activity following FTI/lovastatin combination treatment, but lack of toxicity in normal rat Schwann cells.

Methods

Compounds and Reagents

The synthesis of prodrug **1** has been reported (Clark et al., 2007). Prodrug **2** was synthesized via an analogous route using aminoester **4** instead of methyl chlorobutylamine to generate prodrug ester **3**, followed by selective hydrolysis of the methyl ester (Fig. 1). Lovastatin (Sigma Aldrich, St Louis, MO) aliquots were prepared in dimethyl sulfoxide and stored at -80°C . JC-1 (5,5',6,6'-tetrachloro-1,1',3,3'-tetraethylbenzimidazolylcarbocyanine iodide) and Hoechst 33342 (Invitrogen Molecular Probes, Carlsbad, CA) were prepared in dimethyl sulfoxide and stored at 4°C . Mitotracker Orange CM-H2 TM Ros (Invitrogen Molecular Probes, Carlsbad, CA) aliquots were prepared in dimethyl sulfoxide and stored at -20°C .

Cell Culture

NF90-8 and ST88-14 MPNST cell lines were generously donated by T. Glover (University of Michigan, Ann Arbor, MI). Cells were maintained as adherent cultures in RPMI 1640 (Gibco, Carlsbad, CA) with 5% Fetal Bovine Serum (Hyclone Laboratories, Logan, UT), 100 units/ml penicillin, and 100 $\mu\text{g}/\text{ml}$ streptomycin (Invitrogen, Carlsbad, CA). Primary normal rat Schwann cells were isolated from the sciatic nerves of neonatal Sprague-Dawley Rats and grown on poly-D-lysine coated coverslips as described previously (Skoff et al., 1998). These cells were grown in Eagles medium with 10% calf serum prior to experimental manipulations. Normal, spontaneously immortal rat Schwann cell clones (iSC) isolated from sciatic nerves were a generous gift from E.M. Shooter (Stanford University, Stanford, CA) and described previously (Bolin et al., 1992). These cells were maintained in MEM supplemented with 10% horse serum, 100 units/ml penicillin, and 100 $\mu\text{g}/\text{ml}$ streptomycin. All cells were maintained in a humidified incubator under 5% CO₂. For all experiments, cells were plated 24 hours prior to drug treatment. Immediately before drug treatment, the medium was replaced with fresh growth medium for the duration of the experiment.

Western Analysis

Lysates were prepared from monolayers of cells in 2 \times Laemmli sample buffer by boiling for 5 minutes and cleared by centrifugation (Mattingly et al., 2001). Samples were then separated on polyacrylamide-SDS gels and electrophoretically transferred to nitrocellulose. Ras was detected with a 1:250 dilution of anti-pan Ras monoclonal antibody (BD Transduction Laboratories, San Jose, CA). Caspase-3 was detected with a 1:1000 dilution of anti-Caspase-3 polyclonal antibody (Santa Cruz Biotechnology, Inc., Santa Cruz, CA) as described (Menard et al., 2005). Retinoblastoma protein (pRb) phosphorylated Serine 780 was detected with a 1:1000 dilution of anti-Phospho-Rb Ser 780 polyclonal antibody (Cell Signaling Technology, Danvers, MA). The nitrocellulose membrane was stripped and reprobed for total pRb with a 1:1000 dilution of a mouse monoclonal antibody (Cell Signaling Technology). α -Tubulin was immunoblotted with a 1:2000 dilution of the E7 monoclonal antibody (Developmental Studies Hybridoma Bank, Iowa City, IA).

Proliferation Assay

NF90-8 cells were plated at 20,000 cells/35-mm dish and ST88-14 cells were plated at 20,000 cells/60-mm dish 24 hours prior to drug treatment. Fresh medium was added before the drug treatment as described in the text. Attached cells were trypsinized and combined with media containing detached cells. The cells were collected by centrifugation for 5 minutes at 1000g and counted via a hemacytometer.

Flow Cytometry

NF90-8 and ST88-14 cells were treated and collected for DNA analysis as described previously (Reiners et al., 1999). DNA content was analyzed using a FACScalibur instrument (BD Biosciences, San Jose, CA). A minimum of 10^4 cells/sample were analyzed to determine the percentage of apoptotic cells and cells in G₁, S, and G₂/M phase (MODFIT; Variety Software, Topsham, ME).

DEVDase Activity Assay

Lysates of NF90-8 and ST88-14 cultures were prepared and used in DEVDase assays as previously described (Caruso et al., 2004). Changes in fluorescence over time were converted into picomoles of product by comparison with a standard curve made with 7-amino-4-methylcoumarin. DEVDase specific activities are reported as nanomoles of product per minute per milligram of protein. The bicinchoninic acid assay, using bovine serum albumin as a standard, was used to estimate protein concentrations.

Mitochondrial Membrane Potential and Nuclear Morphology

NF90-8 cells were plated at 100,000 cells/100-mm dish 24 hours prior to drug treatment. At the end of the drug treatment, mitochondrial membrane potential (ψ_m) was assayed by two independent methods. In one series of experiments, JC-1 was added directly to the medium at a final concentration of 5 μ g/ml, and incubated for 10 minutes at 37°C. Attached cells were trypsinized and combined with growth media containing detached cells. The cells were collected at 500 *g* for 5 minutes, washed twice in PBS, and suspended in PBS containing 1% FBS for analysis by excitation at 530 nm and emission at 590 nm. Cells with active mitochondria accumulate red emitting “JC-1 aggregates”. In cells having depolarized mitochondria, JC-1 is present as monomers, which exhibit green, not red, fluorescence emission.

In a second series of experiments, Mitotracker Orange CM-H2 TM Ros (excitation at 554 nM and emission at 576 nM) was added directly to the medium of treated cultures at a final concentration of 50 nM, and incubated for 10 minutes at 37°C. Hoëchst 33342 (excitation at 350 nM and emission at 461 nM) was added directly to the medium at a final concentration of 500 nM to reveal nuclear morphology, and co-incubated with the Mitotracker for 5 minutes at 37°C. Cultures were rinsed once with PBS and subsequently rinsed three times with fresh growth medium. Images were immediately acquired with an Axiovert 200M fluorescence microscope (Carl Zeiss). Cells with reduced ψ_m are unable to sequester and oxidize Mitotracker Orange, and will exhibit reduced fluorescence emission.

Statistical analysis

Two-tailed paired student t-test was employed to determine the statistical significance of a G₁ cell cycle arrest between FTI/lovastatin combination treatment cultures and the DMSO control cultures. The significance level for this analysis was set at $p < 0.05$.

Results

Our lab has previously reported that N-Ras is the predominant active Ras isoform expressed in the human derived MPNST cell lines NF90-8 and ST88-14 (Mattingly et al., 2006). N-Ras can be alternatively prenylated with a geranylgeranyl pyrophosphate in the presence of FTIs (Whyte et al., 1997). We therefore tested whether our novel FTIs (**1** and **2**) could reduce Ras prenylation in MPNST cell lines NF90-8 and ST88-14 (Fig. 2). Inhibition of prenylation is indicated by the slower mobility band or upshift by western analysis. Single 1 μ M treatments of **1**, **2**, or lovastatin had little to no detectable effect on Ras prenylation in

either cell line. However, a reduction of Ras prenylation was observed in the NF90-8 cells following combination treatment of 1 μM **1** or **2** with 1 μM lovastatin. Ras prenylation was also reduced in ST88-14 FTI/lovastatin combination treated cells compared to the single treatments of **1**, **2**, or lovastatin. The observed reduction of Ras prenylation in ST88-14 cells was moderate compared to that occurring in NF90-8 FTI/lovastatin treated cells.

Ras isoforms are known to regulate cell processes such as survival, growth, and proliferation (Rajalingam et al., 2007). Since our FTI/lovastatin combination treatment reduced Ras prenylation in NF90-8 and ST88-14 cells (Fig. 2), we next determined the effect of this treatment on cell proliferation. The results from proliferation experiments with NF90-8 cells are shown in Fig. 3A-B. Single treatments of 500 nM **1**, **2**, or lovastatin alone, had no effect on cell proliferation compared to cultures that were treated with DMSO. However, combination treatments of 500 nM **1** plus 500 nM lovastatin were cytostatic after 24 hours of treatment, and reduced total cell number below initial plating after 72 hours of treatment (Fig. 3A). A similar cytostatic response was observed following treatment with 500 nM **2** plus 500 nM lovastatin (Fig. 3B). Inhibition of proliferation also occurred in the presence of lower concentrations of lovastatin (33 and 100 nM) in combination with 500 nM **1** or **2**. Synergism of the FTI compounds with 33 nM lovastatin is notable because this dose of lovastatin is pharmacologically achievable in humans treated with anti-cholesterol doses of statins (Thibault et al., 1996). Proliferation data for ST88-14 cells treated with nanomolar combinations of FTI/lovastatin are shown in Fig. 3C-D. Single treatment with DMSO, **1**, **2**, or lovastatin had no effect on cell proliferation. However, when 500 nM **1** or **2** were used in combination with 500 nM lovastatin, we observed a reduction of ST88-14 proliferation. Unlike the NF90-8 cells, lower doses of lovastatin in combination with 500 nM **1** or **2** had less effect on cell proliferation.

We subsequently investigated the effects of FTI/lovastatin treatment on NF90-8 and ST88-14 cell cycle progression. Fig. 4 shows FACS analysis of NF90-8 cells treated with 500 nM **1**, or with 500 nM lovastatin, or a combination of the two drugs for 48 hours. Single treatments with these compounds yielded cell cycle profiles comparable to the DMSO control, which was also similar to untreated controls (Table 1 and Supplemental Fig. 1). The combination treatment results in an increased percentage of cells in G₁, and a significant percentage of cells undergoing apoptosis, compared to the control cultures. These data coincide with the proliferation data in Fig. 3A, where we observed a cytostatic affect of 500 nM **1** plus 500 nM lovastatin at 24 hours, and a cytotoxic effect at 48 hours. Similar results were observed with 500 nM **2** in combination with 500 nM lovastatin (Table 1 and Supplemental Fig. 1). As a control for these experiments, we also tested compound **4** (Fig. 1), an analog of compound **1** that is inactive as an FTI, and that has no effect on the proliferation of the spontaneous MPNST cell line, STS-26T [Compound **5d** in (Clark et al., 2007)]. The cell cycle distribution of NF90-8 and ST88-14 cells were not affected by treatment with 1 μM **4** alone or in combination with 1 μM lovastatin (data not shown). Cell cycle progression of ST88-14 cells was also analyzed at 24 and 48 hours following treatment with either 500 nM **1** or 500 nM **2** with or without 500 nM lovastatin (Table 2 and Supplemental Fig. 2). We observed a moderate G₁ cell cycle arrest at 24 hours that was maintained at 48 hours following FTI/lovastatin combination treatment. Although we observed an increased percentage of apoptotic cells at 48 hours, the effect was more modest than that observed in NF90-8 cultures.

One mechanism that could underlie a G₁ arrest would be if retinoblastoma protein (pRb) phosphorylation were reduced. We therefore investigated the phosphorylation pattern of pRb in the NF90-8 and ST88-14 cell lines (Fig. 5). Single treatments of 1 μM **1**, **2**, or lovastatin did not reduce hyperphosphorylation of pRb. However, combining 1 μM **1** or **2** with 1 μM lovastatin significantly reduced the phosphorylated pRb signal. Total pRb expression was

also reduced in **1** or **2** plus lovastatin treated cultures. As a further control for these experiments, we also tested the inactive control compound **4**. No effect on pRb phosphorylation was observed in either cell line following treatment with 1 μ M **4** alone or in combination with 1 μ M lovastatin (Fig. 5).

In addition to cell cycle arrest, FTI/lovastatin treatment induced an increase in the number of cells with sub-G₁ DNA content, which would be consistent with an induction of apoptosis. Fig. 6A-B is a morphological analysis of both cell lines treated with 500 nM **1** or **2** with 500 nM lovastatin. NF90-8 and ST88-14 cells treated singularly with **1**, **2**, or lovastatin at T=24 and T=48 have a flat, elongated morphology associated with healthy, proliferating cells. However, following 24 hours of 500 nM **1** or **2** with 500 nM lovastatin combination treatments, both cell lines became more refractile. By 48 hours of treatment, cells were shrunken, rounded, and blebbed.

The morphological characteristics of FTI/lovastatin treated MPNSTs, coupled with increased numbers of cells with sub-G₁ DNA contents, suggested the occurrence of apoptosis. To monitor for apoptosis, we used Ac-DEVD-AMC to assay for the activity of caspases-3 and -7 (Fig. 7) (Caruso et al., 2004). Treatment with 500 nM **1**, or **2**, or lovastatin resulted in no detectable activation of DEVDase. However, high amounts of active DEVDase were observed in NF90-8 cells (Fig. 7C-D), and moderate amounts in ST88-14 (Fig. 7A-B) cells, following treatment with 500 nM **1** or **2** and 500 nM lovastatin. In addition, 100 nM **1** with 500 nM lovastatin also activated DEVDase in NF90-8 cells. NF90-8 cells treated with 500 nM **1** plus 500 nM lovastatin also exhibited a time-dependent increase of cleaved pro-caspase-3 (Fig. 7E). The kinetics of procaspase-3 cleavage correlated with the time-dependent increases in DEVDase (Fig. 7C).

Changes in mitochondrial membrane potential (ψ_m) often precede/accompany the activation of pro-caspases, and are generally indicative of an activation of the intrinsic apoptotic pathway (REF). We employed two methods to monitor ψ_m . The first method involved live cell microscopy using Mitotracker Orange (Fig. 8A). NF90-8 cells treated with DMSO, 500 nM **1**, or 500 nM lovastatin exhibited tubular mitochondrial staining at 24 and 48 hours with little to no detectable chromatin condensation. However, following 24 hours of combination treatment with 500 nM **1** and 500 nM lovastatin, the cells exhibited less red fluorescence. Within 48 hours of treatment, both the cells and nuclei had shrunken, and there were dramatic reductions in red fluorescence (i.e., reduced ψ_m).

Flow cytometric analysis of ψ_m with JC-1 was employed to quantify mitochondrial membrane potential (Fig. 8B). NF90-8 cells were treated with drugs, and then incubated with JC-1, a compound that forms red fluorescent aggregates in cells having ψ_m . Following a loss of ψ_m , JC-1 does not aggregate and fluoresce red. As observed with Mitotracker Orange, treatment with DMSO, 500 nM lovastatin, or 500 nM **1** did not affect ψ_m in NF90-8 cells. However, following FTI/lovastatin combination treatment, a progressive loss of ψ_m occurred from 24 to 72 hours. Notably, a significant loss of ψ_m occurred at 24 hours, a time that precedes the activation of DEVDase (Fig. 7C) and pro-caspase-3 cleavage (Fig. 7E).

We have demonstrated that treatment of human MPNST cell lines NF90-8 and ST88-14 with 500 nM **1** or **2** in combination with 500 nM lovastatin greatly reduced cell proliferation and induced an apoptotic response. We tested the effects of this FTI/lovastatin treatment on normal primary Schwann cells isolated from the sciatic nerve of rat pups. Normal primary rat Schwann cells were treated for 72 hours, and DIC images were recorded (Fig. 9A). Once again, single treatments of DMSO, 500 nM **1** or **2**, or 500 nM lovastatin had no detectable toxicity. However, in a stark contrast to the effect on MPNST cells, combination treatments

of 500 nM **1** or **2** with 500 nM lovastatin had no observable toxicity on normal primary Schwann cells. To further examine whether the FTI/lovastatin treatment would be toxic, we tested our compounds on a spontaneously immortal rat Schwann cell clone (iSC) (Bolin et al., 1992). Single treatments of 500 nM **1**, or **2**, or lovastatin had no effect on proliferation (Fig. 9B). Combination treatments of 500 nM **1** or **2** with 500 nM lovastatin also did not significantly reduce proliferation of iSC cells.

Discussion

NF1 is a disease driven by the functional loss of Nf, a Ras-GAP protein (DeClue et al., 1991). The loss of Nf results in aberrant Ras signaling in cells derived from the neural crest. Ras must localize and attach to membranes where the protein functions and signals downstream. This membrane attachment requires numerous posttranslational modifications. The initial step is the covalent addition of a 15C farnesyl or a 20C geranylgeranyl group to the cysteine located on the C-Terminal “CaaX” box (Gibbs et al., 2001). This modification is critical for proper Ras function and has been investigated as a potential target for treating NF1 MPNST cells.

We used novel farnesyl transferase inhibitors, **1** and **2**, to reduce protein prenylation in MPNST cell lines NF90-8 and ST88-14. FTI **1** and **2** are FPP (Farnesyl diphosphate) based FTase inhibitors. To augment the inhibition of FTase, we co-treated the MPNST cell lines with lovastatin. Lovastatin is an inhibitor of hydroxymethylglutaryl coenzyme A reductase (HMG-CoA reductase), the rate limiting enzyme in cholesterol synthesis (Wong et al., 2002). In addition to blocking cholesterol synthesis, HMG-CoA reductase is responsible for the initial steps of FPP and GGPP (Geranylgeranyl diphosphate) synthesis (Morgan et al., 2003). Other labs have shown that micromolar doses of lovastatin reduce Ras prenylation and increase the recovery of the protein in cytosolic fractions (Mendola and Backer, 1990; Sebti et al., 1991; Rubins et al., 1998). In addition, lovastatin reverted the learning disabilities of NF +/- mice (Li et al., 2005). Recently, a clinical trial was opened to evaluate the safety of lovastatin in adults with NF1 (NCT00352599). In the current study, we used pharmacologically achievable (Thibault et al., 1996) nanomolar doses of lovastatin, which alone had no effect on NF90-8 and ST88-14 MPNST cell line proliferation, but showed synergy when used in combination with novel FTIs, **1** and **2**.

Our lab previously described N-Ras as the predominant active Ras isoform expressed in human-derived MPNST cell lines NF90-8 and ST88-14 (Mattingly et al., 2006). N-Ras is a member of a group of proteins that can undergo two types of prenylation, either farnesylation or geranylgeranylation. This characteristic could allow an escape mechanism in which N-Ras could be alternatively prenylated with a geranylgeranyl pyrophosphate to maintain proper N-Ras localization and function in the presence of FTIs (Whyte et al., 1997). Indeed, NF1 -/- hematopoietic cells confer a myeloproliferative disorder that is resistant to FTI L-744,832 treatment. This resistance occurs despite block of H-Ras prenylation, and is proposed to be due to lack of efficacy against N-Ras and for K-Ras (Mahgoub et al., 1999). Since N-Ras is commonly over expressed or mutated in cancer, designing a therapy that can reduce farnesylation and geranylgeranylation of N-Ras may be a logical approach.

Our data show that Ras prenylation was maintained following single treatments of **1**, **2**, or lovastatin in both MPNST cell lines. It is possible that these agents singularly inhibited FTase and reduced farnesylation. However, N-Ras may have undergone a compensatory alternative geranylgeranylation. Combination treatment with **1** or **2** with lovastatin induced a near complete inhibition of Ras prenylation in NF90-8 cells, and a moderate inhibition in ST88-14 cells. The combination treatment may have provided a more effective inhibition of

FTase, but it could also be that lovastatin may have reduced GGPP pools in the cell, and so impaired alternate prenylation of Ras (Morgan et al., 2005).

Combination treatment of **1** or **2** with lovastatin impaired Ras prenylation in both NF1 MPNST cell lines, but to different degrees. The degree to which Ras prenylation was suppressed correlated with the anti-proliferative and pro-apoptotic effects of the combination treatment. Although inhibition of Ras prenylation correlated strongly with cellular response, Ras is not the exclusive target protein for FTIs (Lebowitz et al., 1997; Ashar et al., 2000; Basso et al., 2005; Clark et al., 2007). Approximately 0.5% of proteins in the cell must undergo prenylation in order to function properly (Epstein et al., 1991). Known farnesylated proteins include the Ras proteins, RhoB, Rheb, CENP-E, CENP-F, and the nuclear lamins (Tamanoi et al., 2001). The large list of potential FTI targets increases the difficulty in identifying the true therapeutic target, and which diseases would respond to an FTI-based therapy. Nevertheless, since Ras activation drives the NF1 phenotype, inhibition of Ras prenylation is likely to contribute to the efficacy of the combinatorial drug treatment.

Combination FTI/lovastatin induced G₁ arrest, coinciding with reduced pRb phosphorylation at 48 hours. Total pRb content was also reduced in combination treated cultures. Since pRb is a substrate for caspase-3 and -7 (Fattman et al., 1997; Fattman et al., 2001), and combination treatment resulted in the activation of procaspase-3 and -7, it is possible that pRb was cleaved by active caspases and degraded. This reduction of total pRb may also have contributed to reduced pRb phosphorylation and the observed G₁ arrest. Induction of G₁ arrest by prenylation inhibitors that include a lactone structure may be through a p21-dependent mechanism that includes inhibition of the proteasome (Efuet and Keyomarsi, 2006). The active compounds in the current study, however, do not contain a lactone moiety. They also induce a G₁ arrest in STS-26T cells (Clark et al., 2007), which do not express p21 protein (Mattingly et al., 2006).

Theoretically, toxicity to normal cells could be a concern following FTI treatment due to the large number of proteins that are prenylated. FTIs in combination with lovastatin may increase the number of proteins with impaired prenylation. However, we found a lack of detectable cytotoxicity in normal or immortalized rat Schwann cells following combined treatment. It is conceivable that the target protein of our FTIs responsible for the observed effects in the MPNST cell lines is not necessary for normal Schwann cell survival or proliferation.

NF1 MPNST cell lines have increased active Ras compared to non-NF1 cell lines. This makes Ras a rational therapeutic target for this disease. Ongoing clinical trials include one regarding lovastatin tolerance in adults with NF1 (NCT00352599), and the use of the peptide-competitive FTI tipifarnib on treating NF1 in children [(Widemann et al., 2006) and NCT00029354]. The novel FTIs described in this study are extremely effective against NF1 MPNST cell lines with a lack of toxicity against normal Schwann cells. We propose that a combination of FTI/statin treatment may be more efficacious in treatment of NF1 MPNSTs.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Abbreviations

NF1	Neurofibromatosis Type 1
Nf1	neurofibromin
BPNST	benign peripheral nerve sheath tumor
MPNST	malignant peripheral nerve sheath tumor
iSC	spontaneously immortal rat Schwann cell clone
FTase	farnesyl transferase
FTI	farnesyl transferase inhibitor
FACS	fluorescence activated cell sorting
GAP	GTPase activating protein
FPP	farnesyl diphosphate
GGPP	geranylgeranyl diphosphate
Rce1	Ras converting enzyme 1
ICMT	Isoprenylcysteine carboxyl methyltransferase
JC-1	5,5',6,6'-tetrachloro-1,1',3,3'-tetraethylbenzimidazolylcarbocyanine iodide
AMC	7-amino-4-methylcoumarin
m	mitochondrial membrane potential

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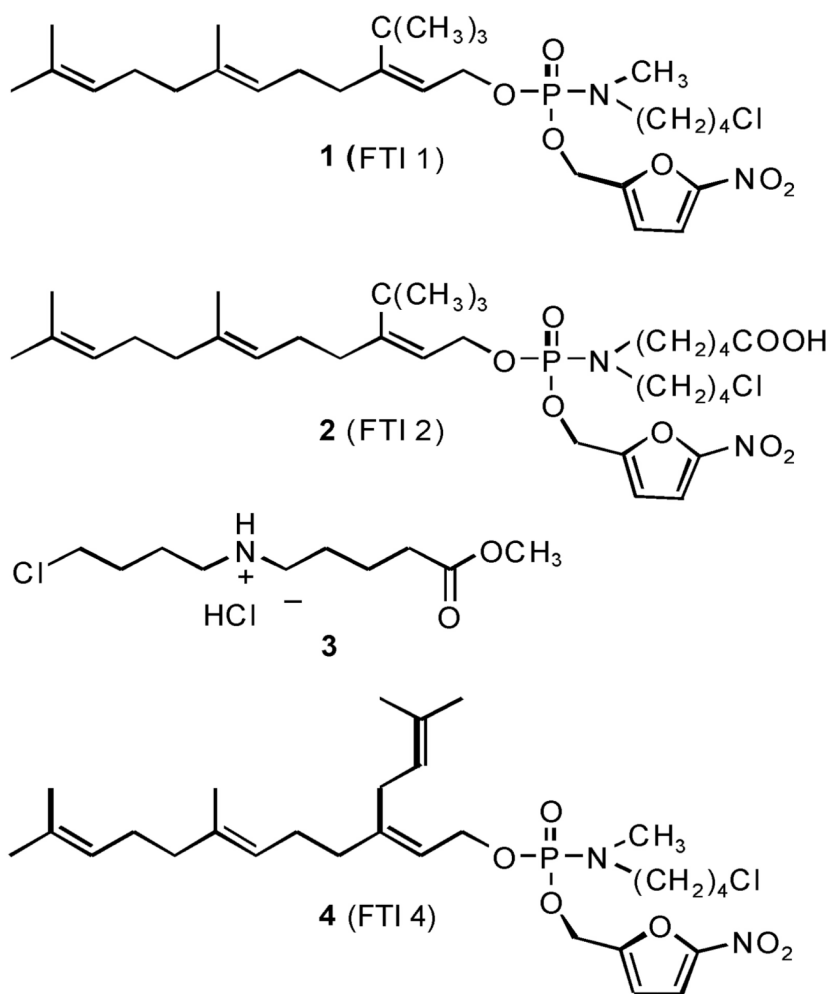


Figure 1.

The synthesis of prodrug **FTI 1**, which was previously reported as compound **5c**, has been described (Clark et al., 2007). Prodrug **FTI 2** was synthesized via an analogous route using aminoester **3** instead of methyl chlorobutylamine to generate the methyl ester of **FTI 1**, followed by selective hydrolysis to the free acid. Compounds **1** and **2** differ at the prodrug moiety, which is removed following entrance into the cell and prodrug activation. The prodrug moiety should not affect the inhibitory function of the farnesyl phosphate analog released. **FTI 4** is an analog of compound **1** that is inactive as an FTI (Clark et al., 2007).

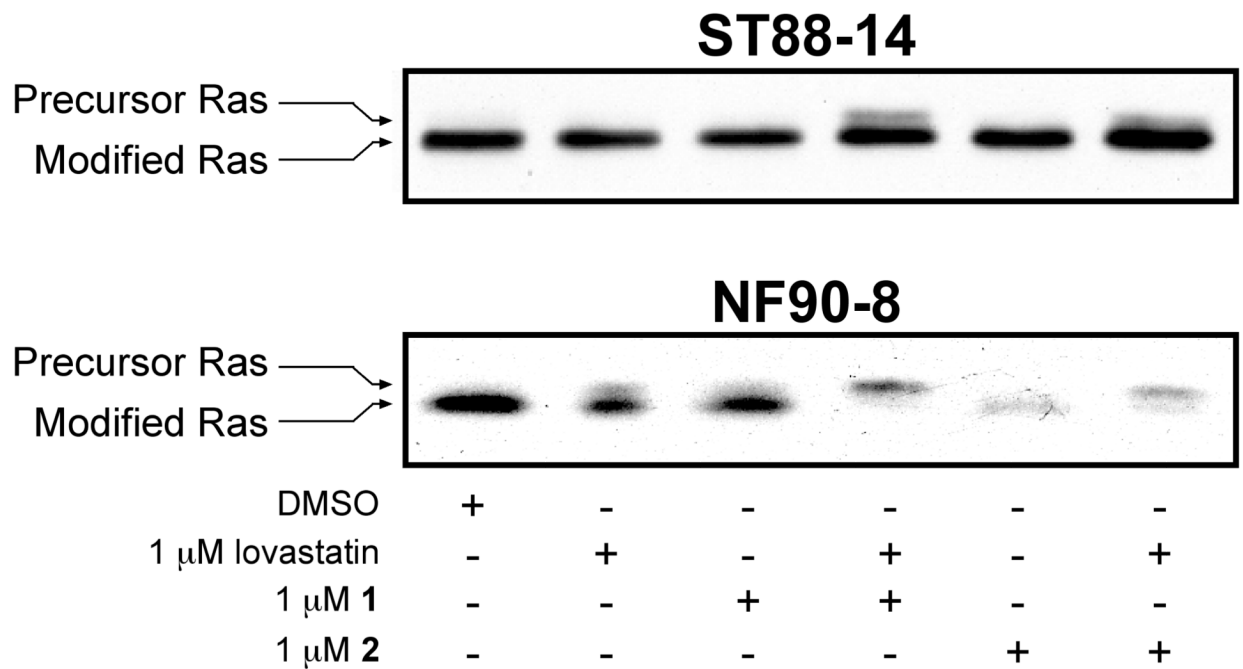
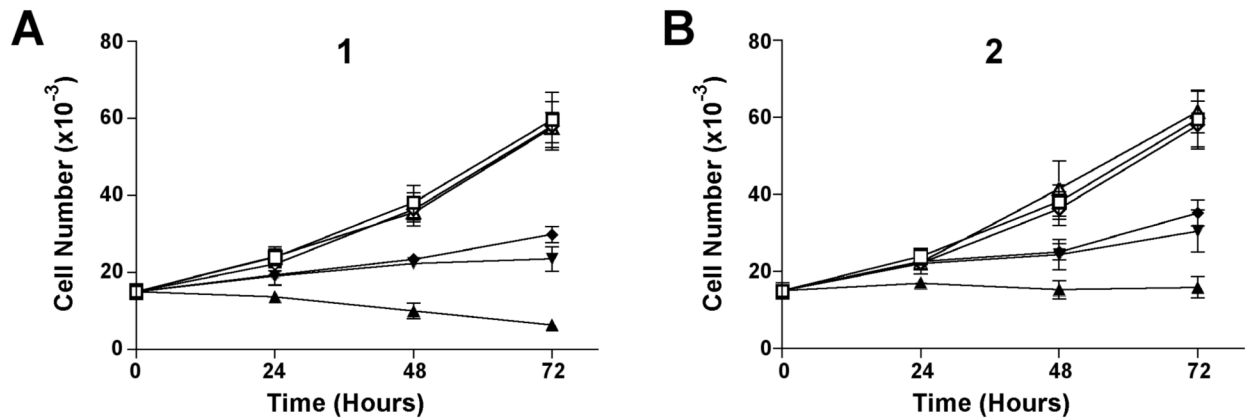


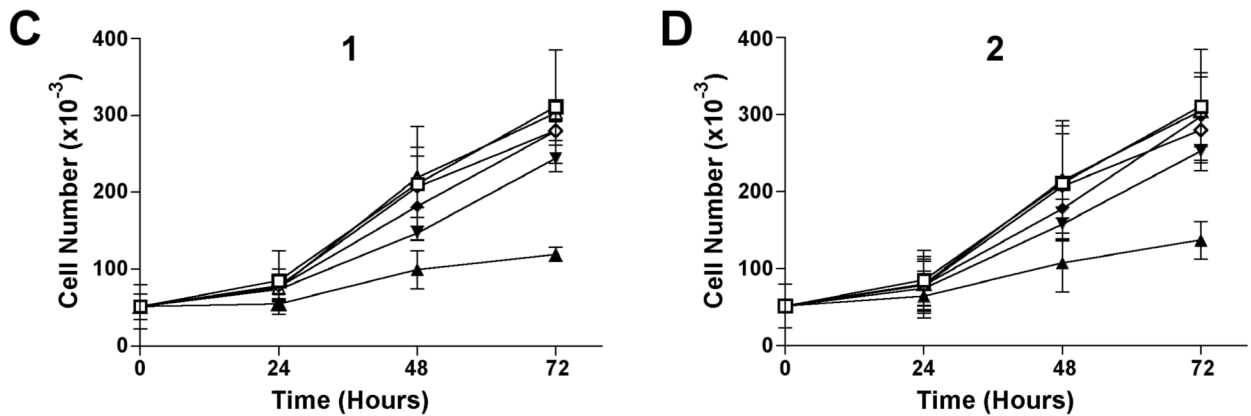
Figure 2. Inhibition of Ras prenylation in MPNST cells by FTI/lovastatin combination treatment

NF90-8 and ST88-14 cultures were treated as indicated for 24 hours. Whole cell lysates were probed for Ras prenylation status by monitoring changes in mobility via western analysis. Prenylated Ras forms migrate faster on SDS gels. FTI/lovastatin combination treatment reduced the expression of modified Ras in both cell lines. Results shown are representative of two independent experiments.

NF90-8



ST88-14



- DMSO
- ◇ 500 nM lovastatin
- △ 500 nM FTI
- ▲ 500 nM lovastatin + 500 nM FTI
- ▼ 100 nM lovastatin + 500 nM FTI
- ◆ 33 nM lovastatin + 500 nM FTI

Figure 3. Reduced MPNST proliferation following FTI/lovastatin treatment

NF90-8 (A-B) and ST88-14 (C-D) cell lines were treated as indicated in the figure. Samples were collected every 24 hours post treatment for analysis of cell number. Data represent means \pm S.D. of three independent experiments.

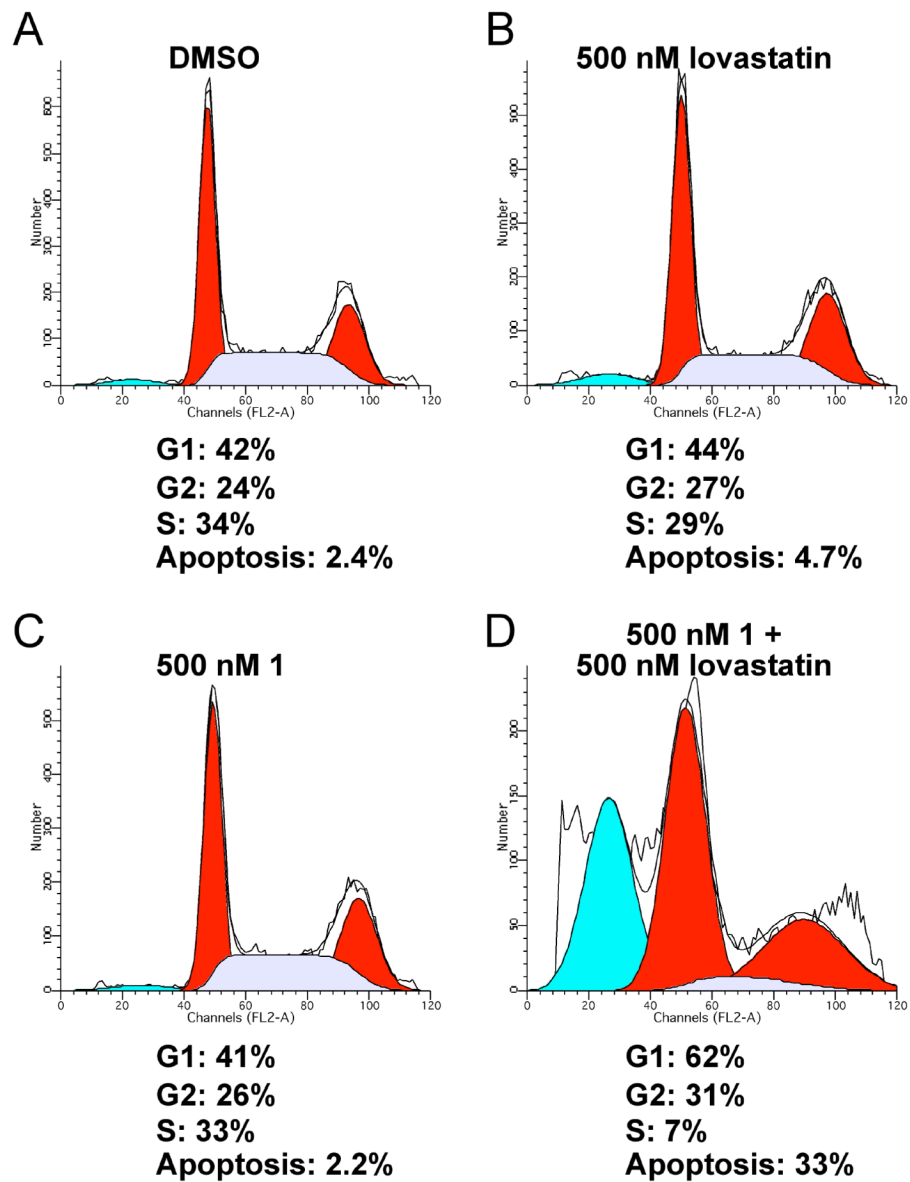


Figure 4. FTI/lovastatin combination treatment of NF90-8 cells increases G_1 and apoptotic cells
 NF90-8 cultures were treated 24 hours after plating according to the condition in figure. Cultures were harvested after 48 hours of treatment for DNA content stained with propidium iodide. Histograms represent 10^4 events and the cell cycle profile was determined using MODFIT. 500 nM **1** in combination with 500 nM lovastatin increased the number of cells in G_1 and greatly increased the presence of cells with sub- G_1 DNA content.

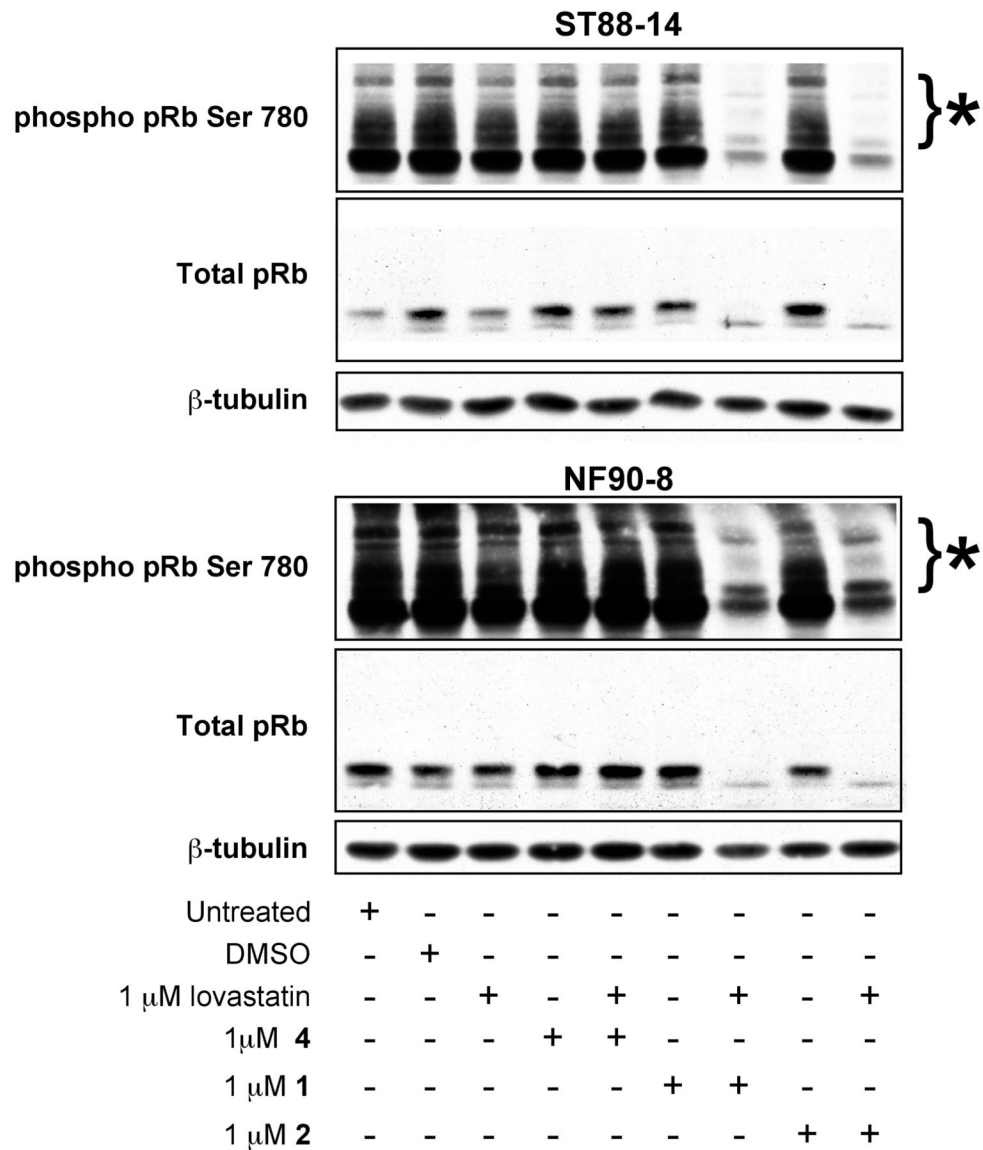


Figure 5. Reduction of pRb hyperphosphorylation in MPNST cells lines by FTI/lovastatin combination treatment

ST88-14 and NF90-8 MPNST cultures were treated as indicated for 48 hours. Whole cell lysates were probed for pRb phosphorylated at serine 780 by western analysis. Blots were stripped and reprobbed for total pRb and β -tubulin. Clear reduction of pRb phosphorylation was observed following treatment with FTI plus lovastatin. * Represents hyperphosphorylated pRb.

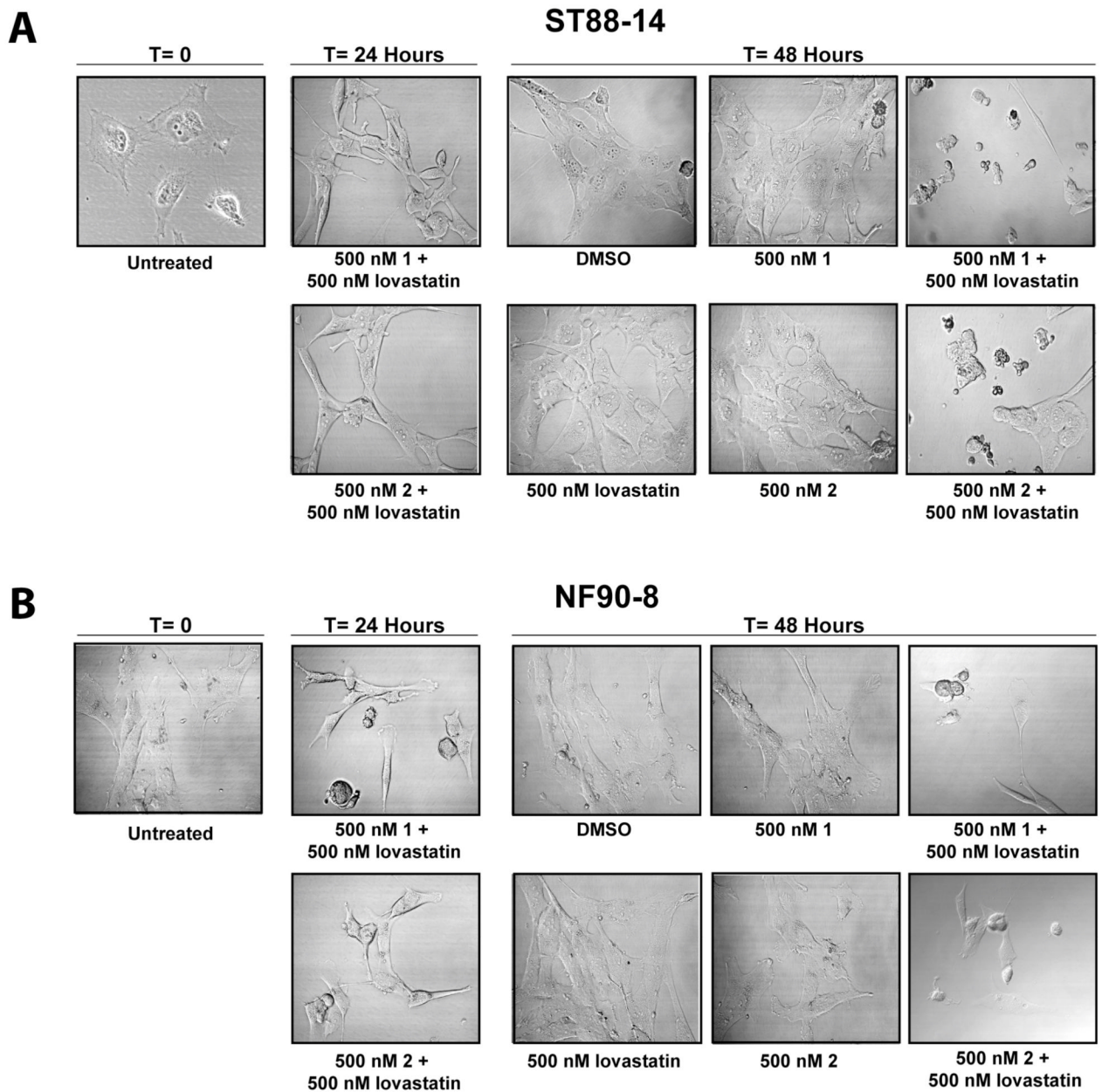


Figure 6. Morphological analysis of NF90-8 and ST88-14 cells following FTI/lovastatin combination treatment

(A) ST88-4 and (B) NF90-8 cells were treated as indicated at T = 0 and Differential Interference Contrast (DIC) images were captured at the 0, 24, and 48 hours time points.

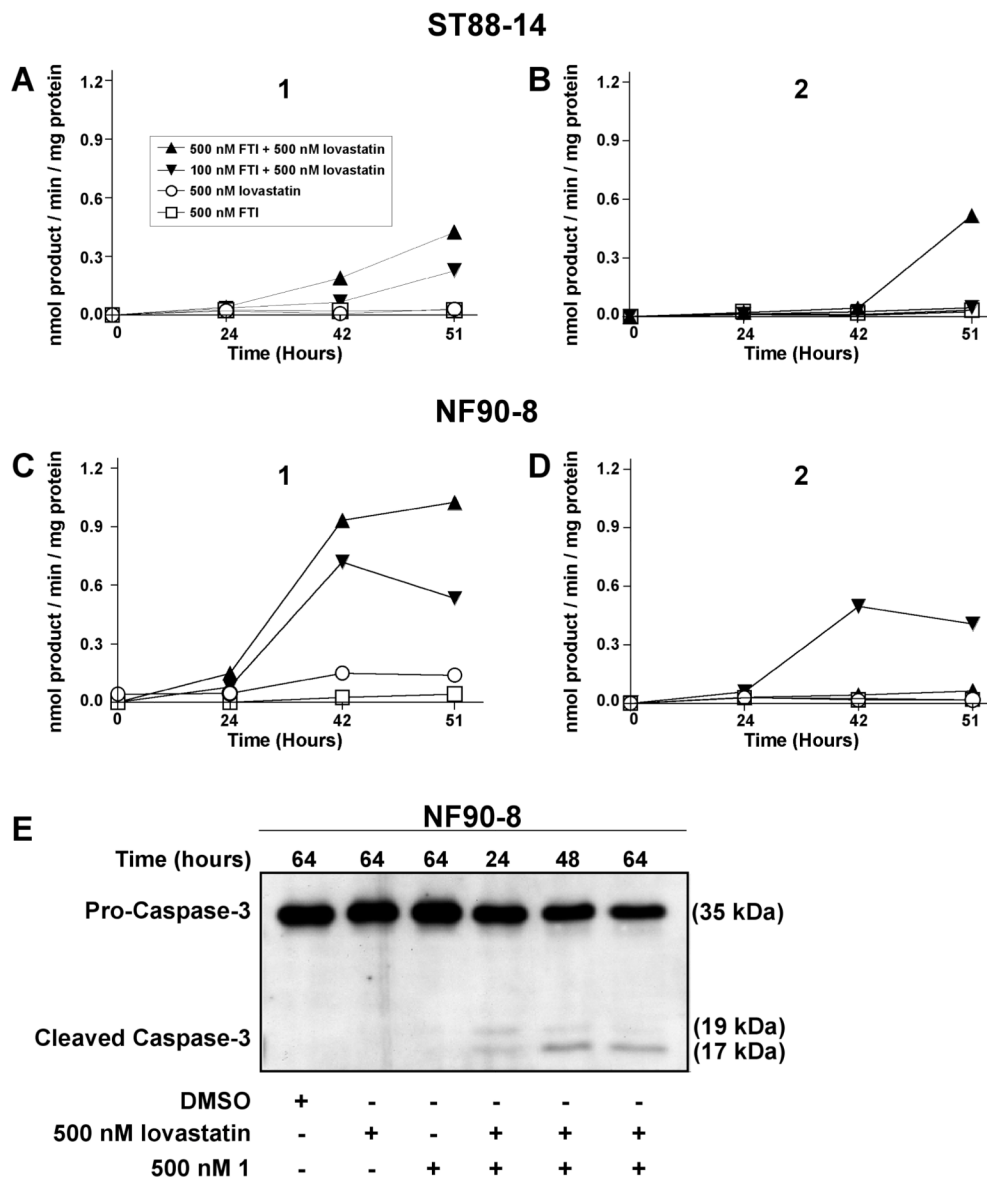


Figure 7. Increased DEVDase activity following FTI/lovastatin combination treatment (A-B) ST88-14 and (C-D) NF90-8 MPNST cells were treated as indicated in the figure. Increased DEVDase activity was observed in FTI/lovastatin combination treated samples. Data represent means of three samples, and are representative of two independent experiments. E, NF90-8 cells were treated as indicated in the figure. Attached and detached cells were collected together and whole cell lysates were separated for western analysis. Pro-caspase-3 cleavage occurred in a time-dependent manner in FTI/lovastatin combination treated cultures.

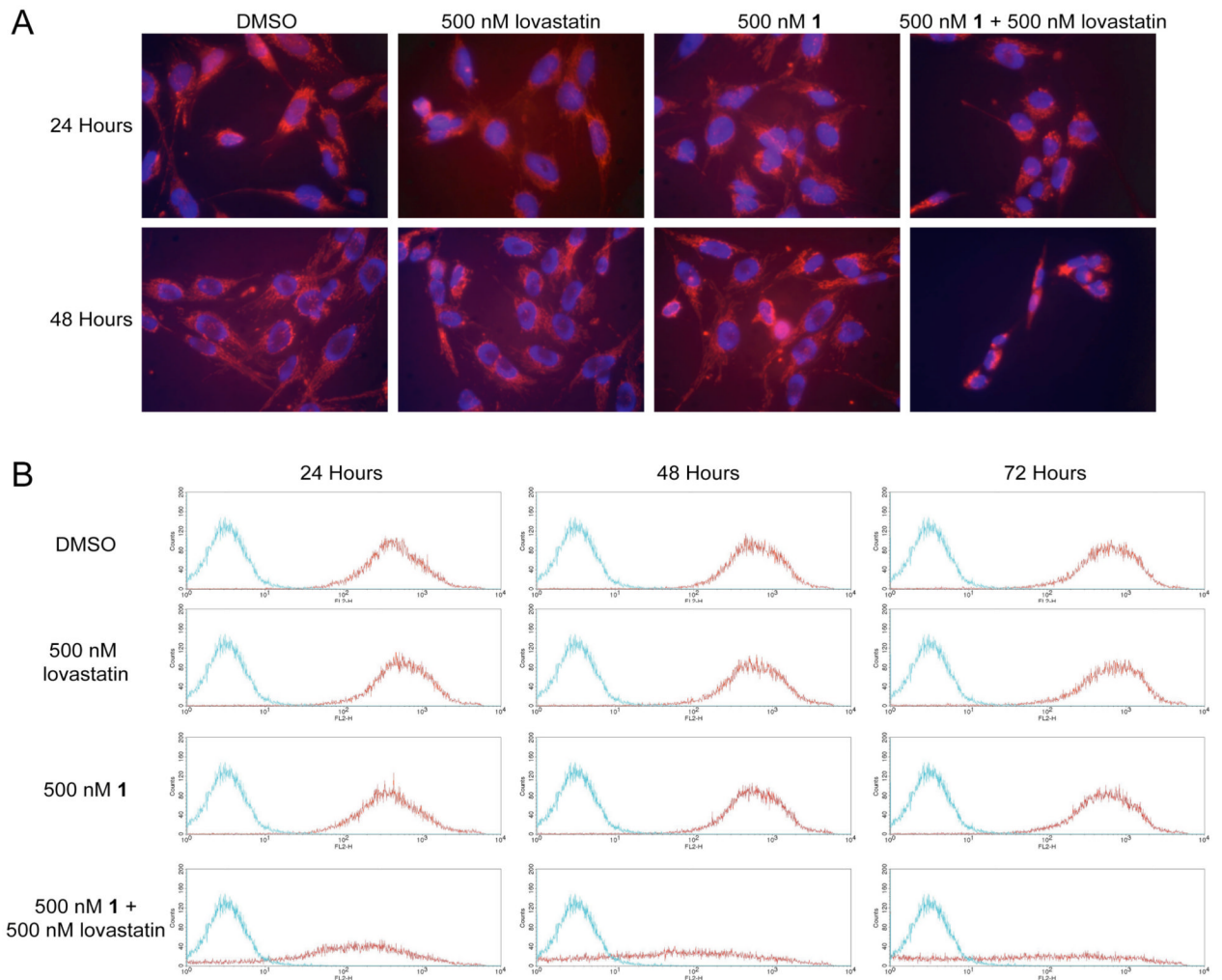


Figure 8.

A, NF90-8 cells were treated as indicated in the figure for 24 and 48 hours. Treated cultures were incubated with 50 nM Mitotracker Orange CM-H2 TM Ros and Hoëchst 33342 to monitor m and nuclear morphology, respectively. Cultures treated with 500 nM **1** in combination with 500 nM lovastatin have reduced m at 48 hours coinciding with chromatin condensation, as compared to the control cultures. B, NF90-8 cells were treated as indicated in the figure for 24, 48, and 72 hours and m was observed using JC-1 by flow cytometry. Histograms represent 2×10^4 events collected using the FL2-H channel. The red data points denote NF90-8 cells stained with JC-1, and the blue data points denote untreated NF90-8 cells not stained with JC-1. NF90-8 cultures treated with DMSO, 500 nM **1**, or 500 nM lovastatin maintained m as shown by the presence of red emission from 24 to 72 hours. However, 500 nM **1** in combination with 500 nM lovastatin resulted in a loss of m .

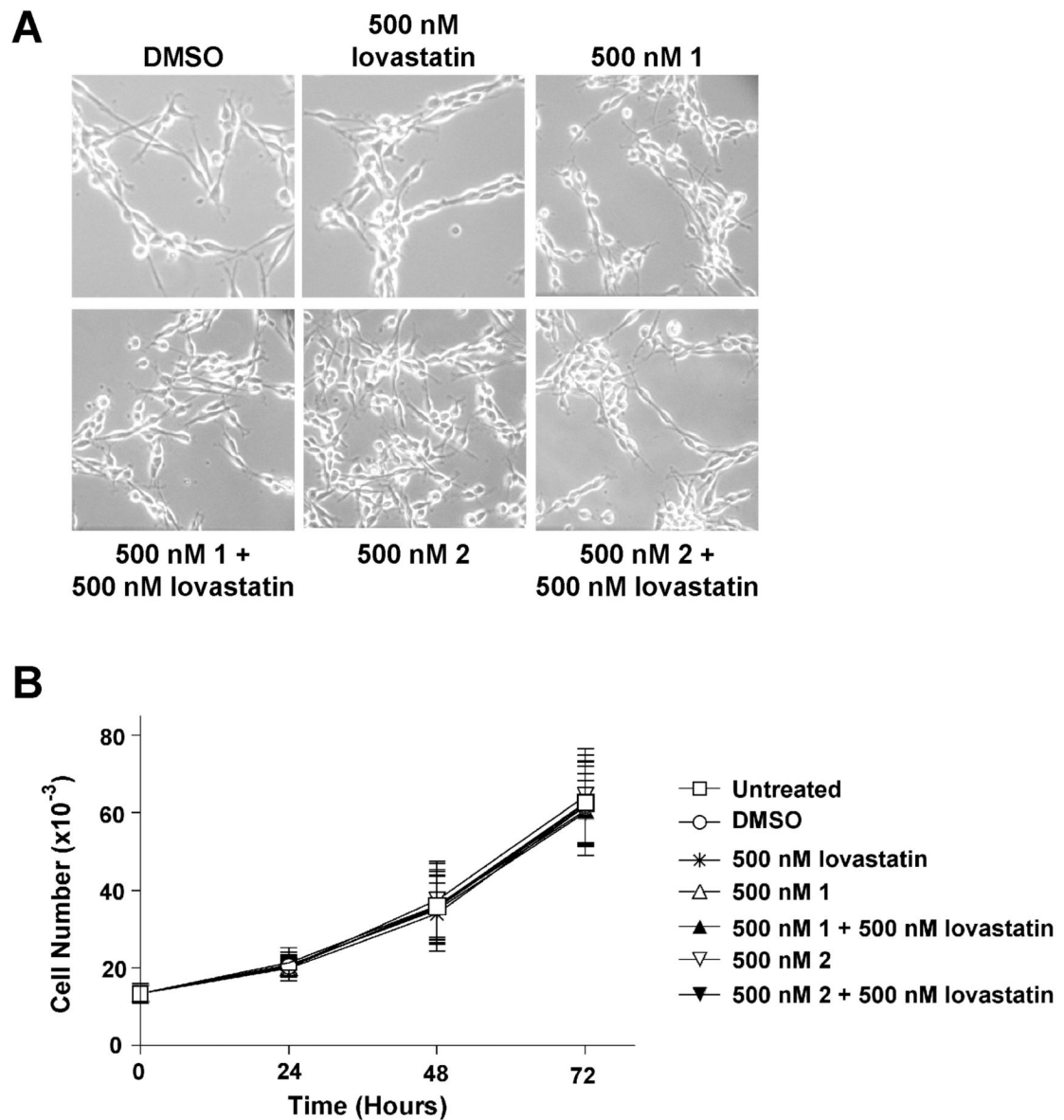


Figure 9. Lack of FTI/lovastatin cytotoxicity in normal rat Schwann cells

A, Primary normal rat Schwann cells were treated as indicated and observed every 24 hours post treatment. These data represent the 72-hour time point and show a lack of observable cytotoxicity. B, iSC cells were treated as indicated. Samples were collected as shown and total cell number was analyzed. All treatments lacked the ability to reduce the proliferation of immortalized normal Rat Schwann cells. Data represent the mean \pm SD of three independent experiments.

Table 1**NF90-8 cell cycle analysis**

NF90-8 cells were treated as indicated above for 24 or 48 hours. Data represent the mean \pm SD of three independent experiments.

DNA analysis of NF90-8 MPNST cells				
	G ₁	G ₂	S	Apoptosis
0 Hours				
Untreated	32.0% \pm 3.7	37.4% \pm 0.8	30.6% \pm 4.3	0.2% \pm 0.4
24 Hours				
DMSO	40.0% \pm 2.1	29.0% \pm 2.7	30.8% \pm 5.2	1.3% \pm 0.3
500 nM lovastatin	40.0% \pm 2.3	24.5% \pm 2.8	35.3% \pm 1.2	2.0% \pm 0.4
500 nM 1	41.1% \pm 1.8	24.7% \pm 2.9	34.2% \pm 1.3	1.7% \pm 0.3
500 nM 1 + 500 nM lovastatin	50.2% \pm 2.0 *	25.9% \pm 2.0	23.9% \pm 3.0	12.6% \pm 1.8 *
500 nM 2	41.5% \pm 1.6	26.5% \pm 2.2	32.0% \pm 3.6	1.5% \pm 0.5
500 nM 2 + 500 nM lovastatin	49.6% \pm 3.4 *	26.2% \pm 1.3	24.2% \pm 2.7	5.9% \pm 0.9
48 Hours				
DMSO	44.4% \pm 2.4	23.3% \pm 2.3	32.3% \pm 1.9	2.7% \pm 0.4
500 nM lovastatin	46.7% \pm 2.6	26.4% \pm 1.4	26.9% \pm 1.9	3.6% \pm 1.0
500 nM 1	43.0% \pm 2.5	24.4% \pm 1.6	32.6% \pm 2.4	3.0% \pm 0.7
500 nM 1 + 500 nM lovastatin	62.7% \pm 2.9 *	31.0% \pm 2.0	6.3% \pm 1.1 *	31.4% \pm 6.3 *
500 nM 2	40.2% \pm 1.3	26.0% \pm 1.6	33.9% \pm 0.7	1.8% \pm 0.7
500 nM 2 + 500 nM lovastatin	55.0% \pm 1.0 *	27.0% \pm 5.2	18.0% \pm 14.8	30.3% \pm 3.4 *

* Statistical significance was determined between the FTI/lovastatin treated cultures and the DMSO controls. Significance was set at $p < 0.05$.

Table 2**ST88-14 cell cycle analysis**

ST88-14 cells were treated as indicated above for 24 or 48 hours. Data represent the mean \pm SD of three independent experiments.

DNA analysis of ST88-14 MPNST cells				
	G ₁	G ₂	S	Apoptosis
0 Hours				
Untreated	37.7% \pm 10.3	29.8% \pm 9.1	32.6% \pm 2.2	0.1% \pm 0.6
24 Hours				
DMSO	51.7% \pm 1.3	19.8% \pm 1.2	28.6% \pm 1.9	0.1% \pm 0.0
500 nM lovastatin	50.9% \pm 2.0	20.5% \pm 1.8	30.5% \pm 1.8	0.2% \pm 0.1
500 nM 1	51.7% \pm 3.3	20.0% \pm 1.2	28.4% \pm 3.0	0.2% \pm 0.1
500 nM 1 + 500 nM lovastatin	58.7% \pm 1.6 *	19.0% \pm 1.5	22.4% \pm 3.1 *	0.2% \pm 0.1
500 nM 2	49.5% \pm 1.8	20.0% \pm 0.7	30.6% \pm 1.3	0.1% \pm 0.1
500 nM 2 + 500 nM lovastatin	60.0% \pm 0.9 *	19.2% \pm 2.4	22.8% \pm 2.6 *	0.2% \pm 0.0
48 Hours				
DMSO	60.9% \pm 5.8	13.1% \pm 1.8	26.0% \pm 4.4	0.4% \pm 0.2
500 nM lovastatin	56.7% \pm 5.6	13.8% \pm 1.6	29.5% \pm 4.3	0.6% \pm 0.4
500 nM 1	59.2% \pm 3.2	15.2% \pm 2.1	25.6% \pm 3.0	0.5% \pm 0.5
500 nM 1 + 500 nM lovastatin	77.2% \pm 3.0	13.1% \pm 2.1	9.7% \pm 3.0 *	8.9% \pm 5.2 *
500 nM 2	57.0% \pm 2.7	14.3% \pm 1.2	28.8% \pm 2.0	0.7% \pm 0.5
500 nM 2 + 500 nM lovastatin	74.7% \pm 1.6	11.5% \pm 0.5	13.9% \pm 1.3 *	5.7% \pm 1.7 *

* Statistical significance was determined between the FTI/lovastatin treated cultures and the DMSO controls. Significance was set at $p < 0.05$.