Ras activation occurs through stimulation of an upstream growth factor receptor such as epidermal growth factor receptor (EGFR). The ultimate effect of Ras is to induce nuclear transcription via a signaling pathway sequentially involving Raf, MAP kinase kinase (MEK), and mitogen-activated protein kinase (MAPK). To transform cells, Ras oncogenes must be posttranslationally modified with a farnesyl group in a reaction catalyzed by farnesyl protein transferase. Farnesyltransferase inhibitors, therefore, have been proposed as potent anticancer agents. This study demonstrates the growth-inhibitory effects of farnesyltransferase inhibitor SCH66336 on human glioblastoma cell lines U-251 MG, U-251/E4 MG (a stably transfected cell line with elevated EGFR expression), and U-87 MG. As determined by (3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium, inner salt) (MTS) and viability assays, the concentration required to achieve 50% inhibition (IC_{50}) ranged from 30 µM (single 24-h treatment) to 10 µM (5-day treatment). U-251/E4 MG with overexpression of EGFR were more sensitive than U-251 MG parental cells. These observations were also supported by soft agar analysis. Cells treated with SCH66336 underwent G2 arrest. Western blot analysis revealed a decrease in phospho-MAPK levels upon treatment with 10 µM SCH66336, whereas MAPK levels were unaffected by the drug. Interestingly, increased expression of EGFR was observed in U-251 MG and U-251/E4 MG but not in U-87 MG in the presence of the inhibitor. These results demonstrate that SCH66336 inhibits viability and anchorage-independent growth in a time- and dose-dependent manner in glioblastoma cell lines U-251 MG, U-251/E4 MG, and U-87 MG via a signal transduction pathway involving the down-regulation of phospho-MAPK. Overexpression of EGFR appears to alter cellular sensitivity to farnesyltransferase inhibitors. This may have a particularly important implication in glioblastoma, where over 50% of tumors have amplification and overexpression of EGFR.

**Neuro-Oncology 2, 151-158, 2000 (Posted to Neuro-Oncology [serial online], Doc. 00-020, June 1, 2000. URL <neuro-oncology.mc.duke.edu>**

Ras contributes to as many as 20% of all cancers and as many as 50% of the more common cancers, such as colon cancer (Bos, 1990). In many of these cancers, RAS is mutated. The role of Ras in cancer is not limited to its mutational status, however, as it has been found to be overactivated as well in various types of cancer, including astrocytomas and glioblastomas (Bredel et al., 1999; Gerosa et al., 1989; Gutmann et al., 1996; Orian et al., 1992; Pollack et al., 1998). This overactivation may be caused by the deregulation of upstream proteins including growth factors and growth factor receptors such as the EGFR (Prigent et al., 1996; Westermark et al., 1995). Several studies conducted in recent years have shown that EGFR is overexpressed in 50% of glioblastomas (Feldkamp et al., 1999; Libermann et al., 1983; Olson et al., 1995; Steck et al., 1988; Wong et al., 1987). Furthermore, it has been shown in human malignant astrocytomas that proliferative signals from EGFR lead to activation of the Ras mitogenic pathway and ultimately proliferation of these cells (Guha et al., 1997).
FTIs, which have become a major focus of therapeutics in recent years, were originally designed to inhibit the actions of oncogenic RAS. Binding of a growth factor such as EGF to its receptor (EGFR) triggers a series of events leading to the activation of Ras protein. When EGF binds to EGFR, a tyrosine kinase receptor, the receptor is activated through autophosphorylation, and growth factor receptor–bound protein-2 (Grb2) and son of sevenless (Sos) protein are recruited to the activated receptor to form a complex. This complex catalyzes the exchange of GDP for GTP on Ras, which activates Ras. Ras recruits Raf, a serine/threonine kinase, which is then activated and functions to phosphorylate MAP kinase kinase (MEK). MAP kinase kinase, in turn, phosphorylates MAPK on threonine and tyrosine residues. Once activated, MAPK translocates to the nucleus to phosphorylate various transcription factors and thus induces cellular proliferation (Lewin, 1997).

In order to transform cells, Ras oncoproteins must first be modified. Ras is initially formed as an inactive precursor protein in the cytoplasm of cells. A crucial step in the activation of Ras involves the transfer of a farnesyl group from farnesylpyrophosphate by the enzyme farnesyltransferase. The addition of the farnesyl group occurs on the carboxyl cysteine of a CAAX sequence, where C is a cysteine, A is an aliphatic amino acid, and X is any amino acid except leucine or isoleucine. This farnesyl group acts as a molecular hook that allows Ras to attach to the membrane. Without this farnesyl group, Ras is biologically inactive (Travis, 1993). FTIs were designed to target the enzyme farnesyltransferase and thus block the attachment of Ras to the membrane, rendering Ras biologically inactive.

FTIs have been shown to play an important role in the inhibition of oncogenic RAS. Studies have shown that cells transformed with oncogenic RAS morphologically resemble normal, untransformed cells in the presence of the FTI. Furthermore, Ha-ras-transformed Rat1 fibroblasts were able to revert to the normal phenotype and growth pattern when treated with inhibitors of farnesyltransferase (James et al., 1993). It has recently been noted that one mechanism by which FTIs exert their effect on cellular morphology is via the extension of microtubules in the cytoplasm of cells (Suzuki et al., 1998). It has been postulated that FTIs may exert their growth-inhibitory effect via apoptosis, as these inhibitors have been shown to induce apoptosis in v-Ha-ras-transformed Rat1 fibroblasts that have been denied substratum attachment (Lebowitz et al., 1997).

FTIs play an important role in human cancer cell lines. It has been demonstrated that FTIs are able to block anchorage-dependent and anchorage-independent growth in various human tumor cell lines, including breast, colon, prostate, pancreas, ovarian, and lung (Nagasu et al., 1995; Sepp-Lorenzino et al., 1995). Furthermore, studies have shown that FTIs are effective not only against tumors with mutant RAS but also against tumors with wild-type RAS (Sepp-Lorenzino et al., 1995). In vivo, FTIs have been found to inhibit the growth of various types of RAS-dependent murine and human tumors in animal models with no evidence of systemic toxicity (Kohl et al., 1994, 1995; Sun et al., 1995).

Although the exact mechanism by which FTIs work is not known, investigators have shown that FTIs have limited or no effect on normal cells in vitro and in vivo (Kohl et al., 1994, 1995). This is important because cells must maintain some level of Ras activity in order to function normally. Furthermore, many other key proteins must also be farnesylated to achieve biological functionality. With these findings, FTIs have become very promising agents for the treatment of many types of cancers, including glioblastoma.

In this study, we examined the effects of SCH66336, a specific inhibitor of the protein farnesyltransferase, on human glioblastoma cell lines U-251 MG, U251/E4 MG (a stably transfectect cell line with elevated EGFR expression), and U-87 MG. Here we report the anchorage-dependent and anchorage-independent growth inhibition by SCH66336 on glioblastoma cell lines U-251 MG, U251/E4 MG, and U-87 MG. The effects of SCH66336 on cell cycle phase distribution were also examined, demonstrating the inhibitor's ability to arrest cells in G2. We also elucidated the effect of SCH66336 on EGFR, phospho-MAPK, and MAPK activities.

**Materials and Methods**

**Cell culture**

Human glioblastoma cell lines U-251 MG and U-87 MG were obtained from the American Type Culture Collection (Manassas, Va.). The U251/E4 MG cell line was produced from parental U-251 MG as a stably transfected cell line with elevated EGFR expression. All cell lines were maintained in a 1:1 mixture of Dulbecco's modified Eagle's medium (DMEM) with Ham's F-12 supplemented with high glucose and 10% fetal calf serum (Life Technologies, Grand Island, N.Y.) in a humidified atmosphere containing 5% CO2 at 37°C.

**SCH66336**

The farnesyltransferase inhibitor SCH66336 was synthesized by Schering-Plough Research Institute (Kenilworth, N.J.). A stock solution of 10 mM was reconstituted in dimethylsulfoxide and stored at -20°C. Dilutions were then prepared in DMEM/F-12 (1:1)/high-glucose medium supplemented with 10% fetal calf serum (Life Technologies). Because dimethylsulfoxide at higher concentrations is known to have a toxic effect on cells, preliminary studies were conducted to rule out the effect of dimethylsulfoxide on the cell lines examined in our experiment (data not shown). Cell lines U-251 MG, U251/E4 MG, and U-87 MG were treated with various concentrations of dimethylsulfoxide ranging from 0% to 0.6%. No effect on cell growth was seen in this range of concentrations. Because our studies use SCH6636 that is reconstituted in 0.05% to 0.1% dimethylsulfoxide, it was safe to conclude that any effect observed on the cells was due strictly to SCH6636.

**Growth Inhibition Assay—MTS Assay**

On the day before treatment began (referred to as day 0), cells were trypsinized and seeded in 3 triplicate sets of 96-well microtiter plates at 1 x 10^3 cells/well. On day 1, the medium was replaced with fresh medium containing SCH66336 at concentrations of 0, 5, 10, 25, or 50 µM.
Control cells received vehicle alone. In the first set of plates, the cells were treated with fresh medium on day 2 and allowed to grow untreated until day 11. In the second set of plates, the cells were treated daily on days 2 to 5 with the aforementioned concentrations of SCH66336. Fresh medium containing the drug was added with each daily drug application. After 5 days of treatment, fresh, inhibitor-free medium was added on day 6. Cells were allowed to grow untreated until day 11. Cells in the third set of plates were treated daily on days 2 to 10 in the same manner as the second set of plates. For all 3 sets, the number of viable cells in proliferation was determined on day 11 using a colorimetric cell proliferation assay (CellTiter 96 AQ_ueous Non-Radioactive Cell Proliferation Assay; Promega Corp., Madison, Wis.).

Cell Viability Assay

On day 0, cells were trypsinized and plated at $1 \times 10^4$ cells/well in triplicate in 6-well plates. On day 1, the medium was replaced with fresh medium containing 5 μM or 10 μM SCH66336. Control cells received vehicle alone. Daily treatment with the inhibitor continued on days 2 to 5. Cells were given fresh medium with each application of the drug. Viable cells were scored manually on day 6.

Anchorage-Independent Growth Assay

Cells were seeded at $1 \times 10^4$ cells/well in 6-well plates on day 0. On day 1, the medium was replaced with fresh medium containing 0, 5, or 10 μM SCH66336. Cells were treated with the inhibitor daily on days 2 to 5. Fresh medium was added with each application of the drug. On day 6, cells were washed with inhibitor-free medium and trypsinized, and $1 \times 10^3$ cells/well were seeded in 6-well plates in 0.65% agar over a 0.9% agar layer. After 2 weeks of growth, 1.5 ml medium was added to each well. Colonies were scored manually after 4 weeks of growth. All studies were conducted in triplicate.

Flow Cytometric Analysis

On day 0, cells were seeded in 100-mm plates at 1 to $2 \times 10^5$ cells/plate in 2 triplicate sets. Cells were treated daily with fresh medium containing 5 or 10 μM SCH66336 for 5 days beginning on day 1 and continuing through day 5. Control cells received vehicle alone. On day 6, 1 set of plates was harvested. The other set of plates was given fresh inhibitor-free medium and harvested on day 8. Upon collection, cells were washed twice with phosphate-buffered saline and fixed in 70% ethanol overnight. After fixation, the cells were washed twice with phosphate-buffered saline. The cells were then stained in phosphate-buffered saline containing 50 μg/ml propidium iodide and 20 μg/ml RNAse for 15 min at room temperature. Fluorescence-activated cell sorting was done using an EPICS profile flow cytometer (Coulter Corp., Hialeah, Fla.)

Western Blot Analysis

Cell lines U-251 MG, U251/E4 MG, and U-87 MG were seeded in 6-well plates on day 0 at $5 \times 10^4$ cells/well. All cell lines were treated daily with fresh medium containing 10 μM SCH66336 on days 1 to 5. Control cells received vehicle alone. On day 6, cells were washed 3 times with 2 ml DMEM/F-12 (1:1)/high glucose. After washing, 2 ml DMEM/F-12 (1:1)/high glucose was added to each well. Cells were incubated at 37°C for 6 h. After the 6-h incubation, the medium was removed and replaced with 1 ml DMEM/F-12 (1:1)/high glucose/2% fetal calf serum (Life Technologies). Cultures were incubated at 37°C for 24 h. After 24 h, cells were collected and washed in phosphate-buffered saline. Cells were lysed in RIPA buffer (150 mM NaCl; 1% NP-40; 0.5% DOC; 0.1% sodium dodecyl sulfate; 50 mM Tris, pH 8.0) containing protease inhibitors (protease inhibitor set; Boehringer Mannheim, Indianapolis, Ind.). Cell lysates were clarified by centrifugation and quantitated by Bradford assay. Cellular protein (30 μg) was heated at 98°C, immediately fractionated on a 10% Tris-HCl polyacrylamide gel (Bio-Rad, Richmond, Calif.), and then transferred to Hybrid ECL nitrocellulose membrane (Amersham Life Science, Arlington Heights, Ill.). The p44/42 anti-MAPK rabbit polyclonal immunoglobulin-G antibody and the phospho-specific p44/42 MAPK rabbit polyclonal immunoglobulin-G antibody (New England Biolabs, Beverly, Mass.) were used as recommended by the vendor, with a 1:2000 dilution of antirabbit immunoglobulin-G horseradish peroxidase–conjugated secondary antibody (Amersham Life Science). The anti-EGFR human (Ab-4, rabbit) antibody (Calbiochem, San Diego, Calif.) was used per manufacturer’s instructions in conjunction with a 1:1000 dilution of the aforementioned rabbit secondary antibody. Protein bands were visualized using a commercially available ECL kit (Amersham Life Science), and densitometry was performed using an Alpha Imager 2000 Documentation and Analysis System (Alpha Innotech Corp., San Leandro, Calif.).

Results

SCH66336 Inhibits Cell Growth—MTS Assay

To examine the effect of SCH66336 on cell growth, U-251 MG, U251/E4 MG, and U-87 MG cells were treated with 0, 1, 5, 10, 25, or 50 μM SCH66336 for 1, 5, or 10 days (Fig. 1). Our initial studies using the MTS assay found that exposure to SCH66336 produced a dramatic dose- and time-dependent decrease in cellular proliferation. After treating U-87 MG cells for 24 h with this compound, growth of U-87 MG was inhibited by 36% in 10 μM SCH66336 and by 42% in 25 μM SCH66336 (Fig. 1C). Growth of U-87 MG cells was inhibited by 43% and 94% after 5 days in 10 or 25 μM SCH66336, respectively (Fig. 1C). Similar trends were observed in U-251 MG and U251/E4 MG (Fig. 1A and B). IC50 ranged from 30 μM (single 24-h treatment) to 10 μM (5-day treatment). Sensitivity varied among the cell lines, with U-87 MG exhibiting the most sensitivity to the drug, and U251/E4 MG cells exhibiting more sensitivity than the U-251 MG parental cells (Fig. 1).
**SCH66336 Inhibits Cell Viability**

To determine whether anchorage-dependent growth was inhibited by the FTI, cells were treated daily with fresh medium containing 5 or 10 μM SCH66336 for 5 days (Fig. 2). Cell viability was inhibited by 70% and 76% in U-251 MG when treated with 5 or 10 μM SCH66336, respectively. Similar trends were observed in U251/E4 MG cells, which were inhibited by 65% and 78% when treated with 5 μM or 10 μM SCH66336, respectively. The inhibitor’s effect was most pronounced in the U-87 MG cell line, which showed 93% growth inhibition when treated with 5 μM SCH66336 and 95% growth inhibition when treated with 10 μM SCH66336. Again, all cell lines demonstrated dose-dependent growth inhibition.

**SCH66336 Inhibits Anchorage-Independent Cell Growth**

We tested the effect SCH66336 had on the ability of human tumor cell lines to form colonies in soft agar (Fig. 3). Cells were treated with 5 or 10 μM SCH66336 in fresh medium daily for 5 days. Cells were harvested on days 6 and 8, stained with propidium iodide, and analyzed by fluorescence-activated cell sorting (Fig. 4). Addition of 10 μM SCH66336, a dose shown to have a pronounced effect on cell growth, had no significant effect on the percentage of cells in the sub G₀/G₁ phase of the cell cycle. However, cells treated with 5 and 10 μM of the drug showed a marked decrease in the number of cells in G₁. This decrease in G₁ cells was seen in cells that were harvested at both time points. A subsequent increase in the number of cells in the G₂ phase was seen in all 3 cell lines treated with both concentrations.

**SCH66336 Stimulates G₂ Arrest in Glioblastoma Cell Lines**

To test the effect of SCH66336 on cell cycling, cells were treated with 5 μM or 10 μM SCH66336 in fresh medium daily for 5 days. Cells were harvested on days 6 and 8, stained with propidium iodide, and analyzed by fluorescence-activated cell sorting (Fig. 4). Addition of 10 μM SCH66336, a dose shown to have a pronounced effect on cell growth, had no significant effect on the percentage of cells in the sub G₀/G₁ phase of the cell cycle. However, cells treated with 5 and 10 μM of the drug showed a marked decrease in the number of cells in G₁. This decrease in G₁ cells was seen in cells that were harvested at both time points. A subsequent increase in the number of cells in the G₂ phase was seen in all 3 cell lines treated with both concentrations.
centrations of the inhibitor. The increase in \(G_2\) was dose dependent, with U-87 MG exhibiting the most dramatic response to the FTI. In this cell line, the percentage of cells harvested on day 6 in \(G_2\) treated with 5 or 10 \(\mu M\) SCH66336, increased by 45% and 51%, respectively. U-87 MG cells harvested on day 8 and treated with 5 and 10 \(\mu M\) of the inhibitor showed an increase of 35% and 43%, respectively, in the number of cells in \(G_2\). A slight increase in the number of cells in S phase, when treated with either concentration of the FTI, was seen in U-251 MG and U251/E4 MG, but not in U-87 MG.

**SCH66336 Reduces Phospho-MAPK Activity but Not MAPK Activity**

To examine the effect of SCH66336 on MAPK and phospho-MAPK expression, cells were treated with 10 \(\mu M\) SCH66336 daily for 5 days. Fresh medium was added...
with each addition of the drug. Cells were washed, pre-conditioned with serum-free media, and conditioned with medium containing 2% fetal calf serum on day 6. Cells were collected on day 7 and analyzed via Western blot analysis (Fig. 5), which revealed no change in MAPK levels in the 3 cell lines in the presence of the drug. Furthermore, all of the cell lines exhibited similar levels of endogenous MAPK protein. However, a decrease in phospho-MAPK expression levels was observed in all of the cell lines upon exposure to 10 μM SCH66336. Up to a 2-fold decrease in the level of expression of phospho-MAPK levels was observed in cells treated with the drug. Basal phospho-MAPK levels in control cells varied among the cell lines, with U-87 MG exhibiting the lowest level of phospho-MAPK and U-251 MG exhibiting the highest. The effect of SCH66336 on EGFR protein levels was dependent on the cell line. The amount of EGFR protein in cell line U-87 MG remained unchanged in the presence of the drug. Cell lines U-251 MG and U251/E4 MG, however, exhibited an increase in EGFR levels upon treatment with the drug. Endogenous EGFR levels again varied among the cell lines, with the lowest expression level in U-87 MG and the highest in U251/E4 MG.

Discussion

The data presented here demonstrate that SCH66336 is capable of inhibiting anchorage-dependent growth in the 3 glioblastoma cell lines tested. Response of cells to the drug occurred in a time- and dose-dependent fashion. IC_{50}, according to the MTS assay, ranged from 30 μM (single 24-h treatment) to 10 μM (5-day treatment). It was determined that continuous drug exposure for several days was necessary to achieve optimum cytotoxic effects. Sensitivity among the cell lines differed, with cell line U-87 MG displaying the greatest sensitivity. Interestingly, cell line U251/E4 MG was more susceptible to the inhibitor than U-251 MG parental cells. Based on observations from the MTS assay, the optimal treatment condition appeared to be at a concentration of either 5 or 10 μM for a period of 5 days. Therefore, cells in the remaining experiments were treated in this manner prior to analysis. Viability assays confirmed the anchorage-dependent growth inhibition of SCH66336 on cell lines U-251 MG, U251/E4 MG, and U-87 MG.

Soft agar analysis showed that SCH66336 was also able to inhibit anchorage-independent growth in all 3 cell lines. Response of U251/E4 MG and U-87 MG to the drug was similar. Cell line U-251 MG was again the least sensitive to the drug. Clonogenic response of the cell lines to the drug occurred in a time- and dose-dependent fashion.

Results obtained from anchorage-dependent and -independent assays are consistent with previous experiments, which have demonstrated the ability of FTIs to block anchorage-dependent and -independent growth in various human tumor cell lines, including breast, prostate, colon, pancreas, ovarian, and lung, regardless of RAS status (Nagasu et al., 1995; Sepp-Lorenzino et al., 1995). Recent studies also support the observation of anchorage-dependent growth inhibition of glioblastoma cell lines by FTIs (Bredel et al., 1998).

Flow cytometric studies showed that SCH66336 treatment induces G2 arrest in all 3 of the glioblastoma cell lines tested. The effects of FTIs on cell cycling have been previously documented. A recent study demonstrated that in lung carcinoma and fibrosarcoma cells, FTI-277 caused an accumulation of cells in G2/M (Vogt et al., 1997). Our experiments showed that G2 accumulation in cells treated with SCH66336 was dose-dependent. Furthermore, SCH66336 again had the most pronounced effect on cell line U-87 MG, where the percentage of cells in G2 on day 6 increased from 15% to 61% and 66% after treatment with 5 or 10 μM SCH66336, respectively. In our experiment, no apoptosis was observed in cells treated with 10 μM SCH66336, a dose that had previously been shown to have a notable effect on cell growth. In literature, the ability of FTIs to induce apoptosis varies according to conditions under which they are used. For example, recent studies have shown that FTIs do not induce apoptosis of Ras-transformed cells that are in contact with the substratum, but do cause apoptosis when these same cells are denied substratum attachment (Lebowitz et al., 1997).

To investigate the mechanisms accounting for the cellular effects of the FTI, we examined the effect of the inhibitor on expression levels of 3 proteins involved in the Ras signaling pathway. In all 3 glioblastoma cell lines, SCH66336 was found to cause a decrease in endogenous phospho-MAPK expression levels (the activated form of the enzyme). Our results are consistent with the report that treatment with FTI causes a reduction of MAPK activity (Nagase et al., 1996). SCH66336 was found to cause up to a 2-fold reduction in phospho-MAPK expression levels. These observations demonstrate that inhibition of farnesyltransferase blocks Ras processing, thus preventing the activation of downstream proteins that are
needed to phosphorylate and activate MAPK. Untreated cell line U-87 MG was found to express the lowest levels of endogenous phospho-MAPK among the cell lines. Levels of phospho-MAPK were essentially undetectable in U-87 MG parental cells, was found to possess lower endogenous levels of phospho-MAPK than the U-251 MG cells. The same explanation for differences in sensitivity would also apply here.

Next, we examined the levels of endogenous total MAPK protein in the glioblastoma cell lines. Upon treatment with the inhibitor, MAPK levels remained unchanged. Furthermore, intrinsic MAPK levels were similar for the 3 cell lines. The MAPK antibody is phosphorylation-state independent and thus recognizes both the phosphorylated (active) and unphosphorylated (inactive) forms of MAPK. Upon treatment with SCH66336, MAPK levels remained unchanged. This can be explained by the fact that although the total level of MAPK in the cells remained the same, most of the MAPK consisted of the unphosphorylated form. This explanation is supported by results obtained from the previously mentioned phospho-MAPK analysis.

Finally, EGFR protein levels in the 3 cell lines were assessed by Western blot analysis. EGFR expression levels were highest in cell line U251/E4 MG, as expected. U-87 MG expressed the lowest levels of EGFR of the 3 cell lines. Upon treatment with 10 μM SCH66336, EGFR levels increased in cell lines U-251 MG and U251/E4 MG. Previous studies have shown that EGFR expression is downregulated as part of a Ras-dependent negative feedback loop (Klarlund et al., 1995; Osterop et al., 1993). Thus, the increase in EGFR seen in cell lines U-251 MG and U251/E4 MG may function to compensate for the inhibition of Ras after treatment with 10 μM SCH66336. Interestingly, in the presence of 10 μM SCH66336, EGFR expression levels in U-87 MG remained unchanged. This phenomenon may be explained by the fact that the Ras pathway is not as active in this cell line (as previously shown here). Therefore, this negative feedback loop may not be as active and may not have the same degree of sensitivity to SCH66336 as cell lines U-251 MG and U251/E4 MG.

Throughout these experiments, we observed that U251/E4 MG cells with overexpression of EGFR are more sensitive to SCH66336 than are U-251 MG parental cells. Thus, overexpression of EGFR appears to alter cellular sensitivity to FTI. This observation may be of great clinical importance in the treatment of glioblastoma, as 40% to 50% of glioblastomas have been found to overexpress EGFR, which leads to elevated levels of Ras GTP (Feldkamp et al., 1999; Guha et al., 1997; Libermann et al., 1985; Olson et al., 1995; Steck et al., 1988; Wong et al., 1987). Therefore, FTIs offer a very feasible and promising approach in the treatment of glioblastoma. SCH66336 is presently being evaluated in phase I clinical trials.

References


