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## Antiestrogen fulvestrant enhances the antiproliferative effects of epidermal growth factor receptor inhibitors in human non-small cell lung cancer

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### Abstract

**Introduction**—Estrogen receptor (ER) signaling and its interaction with epidermal growth factor receptor (EGFR) is a potential therapeutic target in non-small cell lung cancer (NSCLC). To explore cross-communication between ER and EGFR, we have correlated ER pathway gene and protein expression profiles and examined effects of antiestrogens with or without EGFR inhibitors in preclinical models of human NSCLC.

**Methods**—We evaluated 54 NSCLC cell lines for growth inhibition with EGFR inhibitors, antiestrogen treatment or the combination. Each line was evaluated for baseline ER pathway protein expression. The majority were also evaluated for baseline ER pathway gene expression. Human NSCLC xenografts were evaluated for effects of inhibition of each pathway either individually or in combination.

**Results**—The specific antiestrogen fulvestrant has modest single agent activity *in vitro*, but in many lines fulvestrant adds to effects of EGFR inhibitors, including synergy in the EGFR mutant, erlotinib-resistant H1975 line. ERa, ER $\beta$ , progesterone receptor (PR)-A, PR-B and aromatase proteins are expressed in all lines to varying degrees, with trends towards lower aromatase in more sensitive cell lines. Sensitivity to fulvestrant correlates with greater baseline ERa gene expression. Tumor stability is achieved in human tumor xenografts with either fulvestrant or EGFR inhibitors, but tumors regress significantly when both pathways are inhibited.

**Conclusions**—These data provide a rationale for further investigation of the antitumor activity of combined therapy with antiestrogen and anti-EGFR agents in the clinic. Future work should also evaluate dual ER and EGFR inhibition in the setting of secondary resistance to EGFR inhibition.

Conflicts of Interest: No conflicts of interest related to this work

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#### Keywords

epidermal growth factor receptor; estrogen; estrogen receptor; lung cancer; fulvestrant

### Introduction

Lung cancer is the leading cause of cancer death among both men and women in the United States.<sup>1</sup> A great deal of work has focused on the role of EGFR in NSCLC.<sup>2, 3</sup> Erlotinib is the only FDA-approved EGFR tyrosine kinase inhibitor (TKI), inducing objective tumor responses in approximately 10% of unselected NSCLC patients.<sup>4</sup> Patient selection is clearly important, with greatest benefit from erlotinib seen in patients whose tumors harbor EGFR mutations.<sup>2, 3, 5</sup> Even in this population, responses are often quite transient and resistance eventually emerges. Although there are several different mechanisms by which EGFR mutant NSCLC eventually become resistant to EGFR TKIs, resistance in approximately half of the cases is caused by a secondary mutation in exon 20 at position 790 (T790M).<sup>6</sup>

The ER pathway is among the most well studied pathways in breast cancer,<sup>7</sup> but its role in NSCLC has only been explored more recently as lung cancer rates among women have increased in recent decades.<sup>8</sup> New data from randomized, prospective clinical trials show a significant impact of hormone replacement therapy (HRT) in lung cancer. In the Women's Health Initiative, over 16,000 post-menopausal women with an intact uterus and no breast cancer history were randomly allocated to supplemental estrogen and progesterone or no HRT. After 5.6 years of study and 2.4 years of follow-up, the hazard ratio (HR) for lung cancer incidence in the HRT group was 1.28 (p = 0.12), with HR for death in NSCLC patients and death specifically from NSCLC of 1.61 (p = 0.02) and 1.87 (p = 0.004) respectively.<sup>9</sup> Further, a prospective cohort study confirmed an increased, dose-dependent lung cancer risk among women who received HRT.<sup>10</sup>

Antiestrogens demonstrate impressive efficacy in breast cancer when tumors express estrogen receptors, with relatively modest toxicity.<sup>11</sup> Accumulating evidence now supports a potential stimulatory role for ER pathway in NSCLC pathogenesis and for ER inhibition in NSCLC suppression.<sup>12–17</sup> Significant ER expression is seen in lung cancer tissue and cell lines, particularly adenocarcinoma.<sup>14, 18</sup> Estrogens stimulate proliferation *in vitro* and progression *in vivo* in lung cancer models.<sup>19</sup> Data from mouse models shows that inhibition of this pathway can reduce the development of lung cancer.<sup>20</sup> In a retrospective analysis of a European database, a reduced risk of lung cancer death was seen in women with breast cancer if they received antiestrogen therapy, tamoxifen in most cases.<sup>21</sup> Molecular details of the interaction between ER and EGFR are emerging, and ER appears to be an important locus for signal convergence.<sup>22–24</sup> Clinical evaluation to date includes a phase I study showing safety and potential anti-tumor activity of gefitinib and fulvestrant in postmenopausal women,<sup>25</sup> and data demonstrating safety of erlotinib combined with fulvestrant.<sup>26–28</sup>

#### **Materials and Methods**

#### Cell lines, cell cultures and reagents

Erlotinib and fulvestrant were studied in 54 NSCLC cell lines *in vitro* (A427, A549, Calu-1, Calu-3, Calu-6, H23, H226, H358, H441, H460, H520, H522, H596, H661, H810, H838, H1155, H1299, H1385, H1435, H1437, H1563, H1568, H1581, H1623, H1651, H1666, H1703, H1734, H1755, H1793, H1836, H1838, H1869, H1944, H1975, H2023, H2030, H2073, H2106, H2110, H2122, H2126, H2135, H2172, H2228, H2286, H2291, H2342, H2405, H2444, HCC827, SHP-77, SK-LU-1). The breast cancer line MCF-7 was evaluated

in additional analyses. All lines were obtained from ATCC and identity was confirmed by genomic DNA, with comparison to the ATCC database, within 3 months of the described experiments.

A-549 was cultured in HAM's F12 (ATCC) supplemented with 5% heat-inactivated fetal bovine serum (FBS), 1% penicillin G-streptomycin-fungizone solution (PSF, Irvine Scientific, Santa Ana, CA), 5 mg/ml insulin (Sigma, Saint Louis, MO) and 1 mg/ml hydrocortisone (Sigma, Saint Louis, MO). Calu-3 and SK-LU-1 were grown in EMEM (ATCC) 10% FBS/PSF. H1155, H1435, H1581, H2286 and H2405 were grown in ACL-4 10% FBS/PSF. H1836, H2342, and H810 were grown in HITES 10% FBS/PSF. Remaining cell lines were cultured in RPMI 1640 (Cellgro, Manassas, VA) 10% FBS/PSF and 2 mmol/ L glutamine (Invitrogen, Carlsbad, CA). Each cell line was also grown in a second proliferation experiment using phenol red-free media with dextran-coated charcoal-stripped (DCC) FBS (same concentration as above). F-12K and EMEM were replaced by RPMI for these experiments and estrogen was not added to HITES, but the type of media was otherwise unchanged.

HCC827 harbors an EGFR mutation in exon 19, and H1975 harbors both an L858R mutation and a T790M mutation in EGFR. All other cell lines evaluated harbored wildtype EGFR genes. KRAS was mutant in the following lines: A427, A549, Calu-1, Calu-6, H23, H358, H441, H460, H1155, H1385, H1944, H2030, H2122, H2291, H2444, SHP77 and SK-LU-1 (sequenome data was unavailable for H1836). ATCC classifies the following cells as adenocarcinoma: Calu-3, H23, H522, H838, H1437, H1563, H1568, H1651, H1734, H1755, H1793, H1838, H1944, H1975, H2030, H2122, H2228, H2291, H2342, H2405, SK-LU-1; bronchoalveolar: H358, H441, H1666; large cell: H460, H661, H810, H1155, H1299, H1581; adenosquamous: H596, H1703; squamous: H226 and H520. Calu-6 is classified as anaplastic, SHP-77 is classified as neuroendocrine and the other cell lines do not have a specific histology allocated.

#### Western blots

Cells growing in log-phase were washed in ice-cold PBS and lysed at 4°C in lysis buffer. Insoluble material was cleared by centrifugation at 10,000g for 10 min. Protein was quantitated using BCA (Pierce Biochemicals, Rockford, IL, USA), resolved by SDS-PAGE, and transferred to nitrocellulose membranes (Invitrogen, Carlsbad, CA, USA). Expression was detected by ERa (62A3) Ab: #2512 (Cell Signaling, Danvers, MA), ER $\beta$  Ab: #5513 (Cell Signaling), PR–A and PR–B (6A1) Ab: #3172 (Cell Signaling), Aromatase Ab: Abcam #ab18995 (Abcam, Cambridge, MA), and  $\alpha$ -Tubulin antibody #2144 (Cell Signaling).

#### Microarray analysis of cell lines

Agilent microarray analysis was performed to assess baseline gene expression as previously described.<sup>29</sup> Baseline microarrays were performed on Agilent Human 1A V2 chips with individual cell lines (labeled with cyanine-5) characterized by comparison to a NSCLC cell line mixed reference pool<sup>29</sup> (labeled with cyanine-3) on a single slide. Slides were read using an Agilent Scanner, and gene expression values were calculated using Agilent Feature Extraction software version 7.5. Extracted data was imported into Rosetta Resolver 5.1 to create expression profiles. Heat maps evaluated cell lines across probes corresponding to genes for ERa, ER $\beta$ , PR, and aromatase.

#### **Proliferation assays**

For experiments looking at erlotinib and fulvestrant individually, cells were seeded in duplicate at 5,000–10,000 cells per well. The day after plating, drug was added with a

starting concentration of 10  $\mu$ M with 2-fold dilutions over 12 concentrations. Cells were harvested by trypsinization and counted immediately using a Coulter Z2 particle counter (Beckman Coulter Inc.) Cells were counted on the day drug was added and five days later with percent growth inhibition defined as 100 × (1 – [generations in treated wells/ generations in untreated controls]). IC<sub>50</sub> was calculated using a four parameter logistic model. For synergy assays, the process was the same with both drugs added at a 1:1 ratio at a starting concentration of 10  $\mu$ M with 2-fold dilutions over 6 concentrations. Combination index was calculated based on method C as described elsewhere (Calcusyn, Biosoft).<sup>30</sup> For surveying the entire panel with both drugs, in duplicate experiments, fulvestrant was added with a starting concentration of 10  $\mu$ M with 2-fold dilutions over 6 concentrations in the presence of erlotinib 1  $\mu$ M.

#### **Apoptosis analysis**

Effects of erlotinib, fulvestrant and the combination on apoptosis were assessed in duplicate experiments using Nim-4', 6-diamidino-2-phenylindole (DAPI) staining. Cells were plated evenly in control and experimental wells and allowed to grow to log-phase, then treated with 1  $\mu$ M erlotinib, 200 nM fulvestrant or the combination for 72 hours. Cells were washed with PBS, and trypsin was applied, and then centrifuged at 3,000 rpm for 5 minutes. Supernatant was aspirated and cells were resuspended in 100  $\mu$ L of Nim-DAPI (NPE Systems) and gently vortexed. Cells were analyzed with UV using a Cell Lab Quanta SC flow cytometer (Beckman-Coulter).

#### In vivo analysis of EGFR TKIs plus fulvestrant

A549 cells ( $5 \times 10^7$  cells/mouse) were injected subcutaneously (SQ) in ovariectomized nude mice with estradiol-17 $\beta$  supplements in a biodegradable binder (1.7 mg/pellet; Innovative Research) as before.<sup>31</sup> When tumors reached 50–75 mm<sup>3</sup>, mice were randomized to treatment groups of 6–8 mice and administered vehicle control, gefitinib (160 mg/kg daily), fulvestrant 5 mg SQ weekly or a combination of gefitinib and fulvestrant at the doses above for 21 days. The experiment was continued to day 28. An independent experiment employed an identical design, but erlotinib (25 mg/kg by oral gavage daily for 21 days), replaced gefitinib, and the experiment was continued until tumors achieved a limiting volume of 500 mm<sup>3</sup> as before.<sup>32</sup> Data are presented as mean ± SEM for tumor volumes measured in mm<sup>3</sup>. Tumor volumes of mice in the single treatment and combination treatment arms were compared to controls and to each other. Data were analyzed using ANOVA and student's ttests.

### Results

#### Sensitivity to erlotinib and fulvestrant differ among human NSCLC cell lines

Significant differences were found in the anti-proliferative effects of erlotinib across the cell lines in our panel, with  $IC_{50} < 1 \ \mu$ M in 6 lines, including the extremely sensitive EGFR mutant line HCC827, a line with a known deletion in exon 19 of the EGFR gene (Figure 1A). Seven lines had  $IC_{50} < 5 \ \mu$ M for fulvestrant in standard media (Figure 1B), and 8 had  $IC_{50} < 5 \ \mu$ M in phenol red-free media with DCC-FBS with a trend toward greater sensitivity across the cell panel in this estrogen-depleted media (Figure 1C). The breast cancer cell line MCF-7 was used as a positive control, and in phenol red-free media with DCC-FBS, the  $IC_{50}$  was below the lowest dose (325 nM) tested (data not shown). To assess combined effects of fulvestrant with erlotinib, sensitivity to fulvestrant in the setting of 1  $\mu$ M erlotinib was evaluated across the cell line panel. Assessing the panel at 2.5  $\mu$ M of fulvestrant plus 1  $\mu$ M erlotinib demonstrated several lines in which fulvestrant added significantly to the inhibition seen with erlotinib alone. This was shown particularly in lines with greater

sensitivity to fulvestrant as a single agent. (Figure 1D). The changes with lower doses of fulvestrant showed similar results with slightly less inhibition (Supplemental Figure)

# Western blot and microarray evaluation of NSCLC cell lines identify potential biomarkers for efficacy of ER inhibition in NSCLC

To assess baseline levels of relevant pathway genes and proteins, the cell line panel was evaluated by microarray analysis using a hormonal gene set and Western blots for ER $\alpha$ , ER $\beta$ , aromatase, PR–A and PR–B as compared to a tubulin control. No clear association was seen between sensitivity to fulvestrant and protein levels of ER $\alpha$ , ER $\beta$ , PR–A or PR–B; however, there was a trend toward lower aromatase levels in cell lines with greater sensitivity to fulvestrant (Figure 2A, B). Baseline gene expression data was available for 44 of the 54 lines. Microarray analysis demonstrated low variability across the panel in comparison to a similar analysis in our group's breast cancer panel.<sup>33</sup> There was a greater sensitivity in cell lines with higher ESR1 gene expression, the gene for ER $\alpha$ . One group of cell lines, including 20 of the 44 lines for which we had baseline expression data, were defined by elevated ESR1 expression. This group included all 4 of the lines with IC<sub>50</sub> < 5  $\mu$ M (p < 0.022) using standard media and 4 of 5 lines with IC<sub>50</sub> < 5  $\mu$ M (p < 0.1) in phenol red-free media with DCC-FBS (Figure 2C).

# *In vivo* evaluation of the combination of inhibition of EGFR and ER pathways demonstrates significant and sustained tumor inhibition

Mice harboring tumors from human NSCLC A549 line were randomly assigned to vehicle control, gefitinib 160 mg/kg daily; fulvestrant 5 mg SQ weekly; or gefitinib plus fulvestrant for a total of 21 days. Gefitinib alone delayed tumor growth, while fulvestrant alone led to tumor stabilization. Treatment with both gefitinib and fulvestrant elicited significantly reduced tumor volume as compared to all other treatments (p<0.001), with no visible tumors after several weeks of therapy (Figure 3A). When a similar study was conducted with erlotinib as the EGFR TKI as previously reported,<sup>31</sup> tumor volumes of mice in the combination arm (fulvestrant and erlotinib) were significantly different from all other treatments (p<0.001). We have now followed animals treated on the combination arm for 60 days, and tumor recurrence was not seen over that time (Figure 3B).

# Combination of erlotinib and fulvestrant is more effective than either agent alone in a model harboring a secondary EGFR mutation

To assess whether antiestrogen therapy can overcome secondary EGFR TKI resistance, dose response curves for the combination of erlotinib and fulvestrant were generated for the H1975 line. This line is resistant to EGFR TKIs and harbors two mutations in the EGFR gene, the sensitizing L858R mutation and the T790M mutation which is associated with EGFR TKIs resistance. In this model, the combination of erlotinib and fulvestrant demonstrated synergy (Figure 4A). When evaluating apoptosis in the H1975 cell line, low doses of fulvestrant (200 nM) led to increased apoptosis at 72 hours compared to control. A slight increase in apoptosis at 72 hours was seen in response to erlotinib 1  $\mu$ M, but the combination of erlotinib and fulvestrant led to a large increase in apoptosis (Figure 4B).

### Discussion

The role of ER signaling in NSCLC is an emerging area of interest based on laboratory discoveries and clinical data. We demonstrate that components of the hormone receptor signaling pathway (including ERa, ER $\beta$ , PR–A, PR–B and aromatase) are present ubiquitously, but to varying degrees, throughout a large panel of human NSCLC cell lines. To our knowledge, this is the most comprehensive evaluation to date of human NSCLC cell lines for gene and protein expression in this signaling pathway.

Fulvestrant was selected for use in these studies because it is generally regarded as a pure antiestrogen that selectively inhibits estrogen receptor signaling in target cells.<sup>34, 35</sup> Further, fulvestrant was shown to exhibit significant antitumor activity in NSCLC cells in vitro and in vivo, and the drug showed evidence of significant interactions with EGFR TKIs.14, 31 In the present studies, single agent antitumor activity with fulvestrant in vitro was modest, but was observed more frequently in NSCLC lines with increased ESR1 gene expression. The results achieved statistical significance for cells maintained in standard media and a trend in phenol red-free media with DCC-FBS. We note that the higher frequency of drug inhibitory activity was not directly related to the level of ESR1 RNA, but instead correlated with the "ESR1 high" group. This finding is considered hypothesis-generating. This data contrasts with some, but not all, data suggesting that ERB is more relevant in NSCLC, but the relative contribution of different components of this pathway remains an area of active investigation.<sup>16</sup> Patients with higher levels of estrogenic stimulus would be a group for whom there is a great clinical need, as a review of survival by estrogen levels among several Southwest Oncology Group trials showed lower survival in women with higher estrogen levels.<sup>36</sup> Similarly, Olivo-Marsten et al report a significant association of increased serum estrogen levels with poorer survival among male and female lung cancer patients.<sup>37</sup> In addition, our findings show a trend toward increased fulvestrant sensitivity in lines with lower levels of aromatase protein as reported by others,<sup>16</sup> although this trend is difficult to quantitate and cannot be considered definitive data on its own. Estrogen levels in lines with lower aromatase levels should be lower, thus allowing fulvestrant to compete with less endogenous estrogens for ER binding.

For proliferation assays, two sets of experiments were conducted. Cells were grown in either ATCC-recommended media or in phenol red-free media with DCC-FBS with a trend toward greater sensitivity in the latter analysis. As a potential screening tool, we also evaluated the entire panel for sensitivity to fulvestrant in the presence of 1 µM of erlotinib. Although doseresponse curves were generated across the panel, it was more representative to graph percent inhibition at a single point across the cell line panel. The combination data correlated well with sensitivity to erlotinib, but in cell lines in which the dual therapy significantly surpassed the response to erlotinib alone, the sensitivity to fulvestrant was generally high, as was the level of ESR1 gene expression. Of note, there are some lines for which the dual treatment appeared to have antagonistic effects. There is some clinical precedence for such an effect, in which the addition of an EGFR TKI elicits worse outcomes.<sup>38</sup> Further characterization of individual patients that may potentially experience harm will be important as clinical evaluation of this combination treatment regimen proceeds. It is possible that, in some subsets of NSCLC, erlotinib is not an appropriate partner for antiestrogen therapy. However, as reported previously in breast cancer, the synergy seen in some lung cancer cell lines with dual antiestrogen and EGFR/HER kinase inhibitors could well be attributable to biologic interactions between estrogen and EGFR signaling pathways.<sup>35, 39</sup> Indeed, significant molecular interactions of estrogen and EGFR signaling pathways in non-small cell lung cancer cells have now been reported in several independent investigations.14, 16, 21, 23, 24

In an attempt to correlate sensitivity with protein levels of ER pathway genes, we evaluated the entire panel by Western Blot. For each protein, we selected a single antibody. The antibodies used have not been clinically validated, and in the context of the ongoing clinical trial, we will use clinically validated antibodies. We also evaluated most of the cell lines (all of the lines for which we have baseline gene expression data) for ER pathway genes, using the Agilent platform which compares a sample to a reference, namely pooled NSCLC cell lines. By definition, there will be cell lines with lower and higher expression than the pooled average, but the magnitude of differences was narrow when compared to a similar breast cancer panel in our laboratory. The evaluation of cell lines against a control of pooled cell

lines could be another factor responsible for differences between our observations and those of others with respect to the relative importance of ERa and ER $\beta$ . Our analysis doesn't assess differences between normal tissues and cancerous tissue, but instead assesses differences among different cancer models. This approach is designed to identify subgroups within a given histology in which disproportionate benefit would be expected.<sup>40, 41</sup>

In patients managed with EGFR inhibitors, drug resistance eventually occurs. Of patients with EGFR mutations (e.g. exon 19 deletions or the L858R mutation in exon 21), significant initial response to erlotinib is observed, but about half develop T790M mutations at the time of tumor progression. The H1975 cell line was chosen for further investigation based on molecular characteristics consistent with this clinical scenario. In the H1975 cell line, which has a T790M mutation, dual treatment with erlotinib and fulvestrant *in vitro* leads to significantly greater inhibition and induction of apoptosis than either agent given alone. The mechanism of this increased sensitivity to EGFR inhibition after the addition of an antiestrogen is not clear. Recent work by Xu et al demonstrates that estrogen can increase phoshpo-EGFR, and that this effect can be inhibited by gefitinb or fulvestrant, but it is inhibited to a greater degree when both agents are used.<sup>42</sup> We are planning further experiments to elucidate potential molecular mechanisms responsible for this effect. Such work will include evaluation of antiestrogens in the setting of cell lines that are initially sensitive to EGFR TKIs, but develop acquired resistance after exposure to erlotinib, rather than the *de novo* resistant H1975 cell lines.

In a NSCLC cell line for which neither erlotinib nor fulvestrant had appreciable single agent in vitro activity in these experiments, namely A549 cells, the combination of EGFR inhibition and ER inhibition led to significant tumor resolution in an *in vivo* setting. These results are striking and were not predicted by in vitro drug sensitivity or baseline expression levels of ER pathway protein or gene expression. The selection of the A549 cell line for in vivo work was made on the basis of data showing expression and activity of estrogen receptors among initially surveyed NSCLC cell lines.<sup>15</sup> It is notable that H23 lung tumor cells also demonstrate significant additive antitumor effects in vivo when fulvestrant is combined with a different EGFR kinase inhibitor, <sup>43</sup> despite an antagonistic effect *in vitro* The fact that synergy between antiestrogens and EGFR kinase inhibitors is significant in vivo but less significant in lung tumor cells used during *in vitro* assays of synergy poses interesting questions to be addressed more fully in future work. For example, in this study we explored alternative in vitro assay conditions (e.g. with or without steroid-depleted culture medium) to find if the *in vitro* assay may be less sensitive than the *in vivo* system due to competition for fulvestrant binding to tumor cell ER. As shown in the results, this may have been a small factor influencing antiestrogen activity. Another factor may be that the tumor microenvironment and neighboring cells (absent in the in vitro assay) may play a role in modulating either estrogen or EGFR signaling. In studies on the action of antiestrogens in preventing lung carcingenesis *in vivo*, Stabile et al. similarly report that the tumor niche may have an important effect on the *in vivo* activity of antiestrogens.<sup>17</sup> Finally, some actions of estrogens in lung in vivo may be due to interaction with subpopulations of tumor progenitor cells or to stimulation of tumor-associated angiogenesis,<sup>44</sup> actions that may be better assessed using in vivo models of tumor progression. One or more of these factors may explain differences observed between in vitro and in vivo assays of estrogen and EGFR signaling.

In addition to independent evidence on significant interactions of estrogen and EGFR signaling in NSCLC<sup>20, 45</sup> results described in this manuscript helped to develop and launch an ongoing, randomized phase II clinical trial of dual erlotinib and fulvestrant therapy as compared to erlotinib alone in previously-treated NSCLC patients. All patients participating in the clinical trial were required to provide a tissue sample, and we plan to analyze these

specimens for EGFR mutation status and expression of ER and aromatase, with a focus on identification of biomarkers that may identify an optimal patient population for further development of antiestrogen therapy in NSCLC.

#### Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Figure 1. *In vitro* sensitivity to erlotinib, fulvestrant and the combination of both agents 54 NSCLC cell lines with  $IC_{50}$  for (A) erlotinib and (B) fulvestrant using ATCC-recommended media are presented in  $\mu$ M. (C)  $IC_{50}$  values for fulvestrant in lines grown in phenol red-free media with DCC-FBS are shown. (D) In phenol red-free media with DCC-FBS, sensitivity to combined erlotinib 1  $\mu$ M and fulvestrant 2.5  $\mu$ M is shown, with fulvestrant (white bars), erlotinib (gray bars) and the combination (black bars). Lines are arranged in descending order of the difference in  $IC_{50}$  between combination treatment and erlotinib, with boxes around cell lines with  $IC_{50}$  for fulvestrant < 5  $\mu$ M. Error bars indicate standard error based on duplicate experiments.





Figure 2. Baseline expression of hormone receptor genes and proteins

For each cell line, protein levels of ER $\alpha$ , ER $\beta$ , PR–A, PR–B, aromatase and tubulin were evaluated by established Western Blot methods (A and B). Green bars by the cell line name indicate an IC<sub>50</sub> < 5  $\mu$ M in the ATCC recommended media, while red bars by the cell line name indicate an IC<sub>50</sub> < 5  $\mu$ M in phenol red-free media with DCC-FBS. In a subset of evaluated cell lines, baseline gene expression data was available, and expression of a hormonal gene set is shown with an X marking lines with an IC50 < 5  $\mu$ M (C).





(A) Ovariectomized nude mice with estradiol supplements and A-549 tumors received 21 days of vehicle control (squares); gefitinib (circles) 160 mg daily; fulvestrant (triangles) 5 mg SQ weekly; or the combination of gefitinib and fulvestrant (diamonds). (B) The same model was assessed with erlotinib (circles) as the EGFR TKI. Mice were treated with assigned therapy for 21 days, and then followed for 60 days to assess outcomes.

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Figure 4. *In vitro* effects of erlotinib, fulvestrant or the combination of both agents on EGFR-TKI resistant, EGFR-mutant lung cancer cell proliferation and viability

Growth curves of the H1975 line after 6 days of treatment with erlotinib (diamonds), fulvestrant (squares), or the combination of the two drugs (triangles) are shown at the concentrations listed. Combination index (CI) for the combination at constant ratio is shown (A). Apoptosis at 72 hours (B) was assessed in H1975 cells with no therapy (white), erlotinib 1  $\mu$ M (stripes), fulvestrant 200 nM (black) or the combination (checked). Error bars indicate standard error based on duplicate experiments.