

# A Combination of Two Receptor Tyrosine Kinase Inhibitors, Canertinib and PHA665752 Compromises Ovarian Cancer Cell Growth in 3D Cell Models

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Received: July 12, 2016 / Published online: September 27, 2016  
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## ABSTRACT

**Introduction:** Advanced ovarian cancer is often a fatal disease as chemotherapeutic drugs have limited effectiveness. Better targeted therapy is needed to improve the survival and quality of life for these women. Receptor tyrosine kinases including EGFR, Her-2 and c-Met are associated with a poor prognosis in ovarian cancer. Therefore, the co-activation of these receptors may be crucial for growth promoting activity. In this study, we explored the effect of combining two small molecule inhibitors that target the EGFR/Her-2 and c-Met receptor tyrosine kinases in two ovarian cancer cell lines. The aim of this study was to investigate the combined inhibition activity of a dual

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EGFR/Her-2 inhibitor (canertinib) and a c-Met inhibitor (PHA665752) in ovarian cancer cell lines in 3D cell aggregates.

**Methods:** OVCAR-5 and SKOV-3 ovarian cancer cell lines were cultured on a non-adherent surface to produce 3D cell clusters and aggregates. Cells were exposed to canertinib and PHA665752, both individually and in combination, for 48 h. The effect on growth, metabolism and the expression/phosphorylation of selective signaling proteins associated with EGFR, Her-2 and c-Met were investigated.

**Results:** The single drug treatments significantly decreased cell growth and altered the expression of signaling proteins in OVCAR-5 and SKOV-3 cell lines. The combination treatment showed greater reduction of cell numbers for both cell lines. Total expression and phosphorylation of signaling proteins were further reduced in the combination drug treatments, compared to the single inhibitor treatments.

**Conclusion:** Our findings suggest that the concurrent targeting of more than one receptor tyrosine kinase may be useful in developing more effective targeted drug

regimens for patients, who have EGFR, Her-2 and c-Met positive ovarian cancer cells.

**Keywords:** Canertinib; c-MET; Cell clusters; EGFR; Ovarian cancer; PHA665752; Tyrosine kinase inhibitors

## INTRODUCTION

Ovarian cancer is the leading cause of death from gynecological cancers in developed countries. The majority of ovarian cancer patients are diagnosed with an advanced stage of the disease, and as such they typically have a 5-year survival rate lower than 30% [1]. For the majority of women, cytotoxic chemotherapeutics have limited efficacy. Therefore, there is an urgent need for alternative anticancer agents that have the potential to control tumor growth, improve the quality of life, and increase the longevity of patients.

A number of targeted anticancer drugs have been studied; among these drugs are small molecule inhibitors of receptor tyrosine kinases. A subset of cancer cells is dependent on the over activation of epidermal growth factor receptor proteins including EGFR and Her-2. Therefore, an inhibitor that specifically blocks these receptors may have clinical activity. Specific inhibitors of EGFR and Her-2 including gefitinib, erlotinib, lapatinib, and a monoclonal antibody Herceptin have been approved by the FDA to treat different types of cancers that include breast cancer, colon cancer and non-small cell lung cancer, NSCLC [2, 3]. The EGFR and Her-2 inhibitors as a monotherapy show minimal clinical benefit in ovarian cancer trials [4, 5]. The lack of clinical efficacy of these inhibitors is perhaps due to activation of multiple receptor tyrosine kinases and the activation of several down stream

signaling proteins. In this study, we explore the possibility that targeting of multiple receptor tyrosine kinases will lead to greater effects on tumor growth potential.

Some advanced ovarian cancer cells have been shown to possess high levels of the receptors EGFR and Her-2 [6]. It is the activation of these receptors that may promote the growth of malignant cells, and the over expression of these receptors has been associated with a poor prognosis and invasive phenotypic changes [7, 8]. There is evidence to suggest that EGFR/Her-2 expressing malignant cells may not be sensitive to the EGFR and Her-2 inhibitors as ovarian cancer cells are able to use alternate a receptor tyrosine kinase for growth promotion [9]. The c-Met is also a receptor tyrosine kinase, which has elevated levels in ovarian cancer cells [10]. The c-Met receptors are activated when they are bound by the hepatocyte growth factor, HGF, which is elevated in the ascitic fluid of women with advanced stages of ovarian cancer [11]. The EGFR, Her-2 and c-Met receptors share common downstream signaling molecules that can promote cell survival, motility, and invasion and cross-communication between these three receptors may promote cancer progression and resistance to therapy [12]. Thus, c-Met could be an additional potential target for cancer treatment and raises the strategic possibility of combined targets of receptor tyrosine kinases concurrently.

Candidate molecules for such combined strategies are canertinib and PHA665752. Canertinib, also known as CI-1033, is an irreversible inhibitor of receptor tyrosine kinases, binding to the ATP-binding site. It has shown potent effects against several cancer cell lines targeting EGFR and Her-2. Clinical trials of canertinib have been conducted against a number of tumor types, including breast

cancer [13], NSCLC [3], and advanced ovarian cancer [5]. However, canertinib alone showed modest activities in clinical studies with ovarian cancer patients [5]. PHA665752 is a reversible inhibitor of c-Met, which at nano-molar concentrations has shown *in vivo* anti-tumor activity [14].

Advanced ovarian cancer patients often have ascitic fluid in the abdominal cavity. In the fluid, ovarian malignant cells form 3D structures; these microscopic clusters and aggregates are a major source of secondary growth potential [15, 16]. These clusters and aggregates then deposit on the peritoneal lining and the surface of internal organs to continue their growth. Therefore, it is appropriate to use cell clusters and aggregates as an *in vitro* cell model to replicate the behavior of ovarian cancer cells. We hypothesized that floating cell clusters and aggregates ovarian cancer cells may use the concurrent activation of EGFR, Her-2 and c-Met to sustain growth and cell survival. Combined inhibition of these receptors will produce a greater anti-growth effect in ovarian cancer cells. The aim of the present study was to test the effect of such combined inhibitors on 3D cell cultures of two ovarian cancer cell lines, OVCAR-5- and SKOV-3.

## METHODS

The human ovarian adenocarcinoma cell lines, OVCAR-5 and SKOV-3 were used. Both cell lines were obtained from Dr. Judith Mckenzie, Haematology Research group, University of Otago, Christchurch, New Zealand. OVCAR-5 and SKOV-3 cells were maintained in DMEM/F12 or MEM media (GIBCO<sup>®</sup>, Life Technologies, New Zealand), respectively, both supplemented with 5% fetal bovine serum (FBS) (GIBCO<sup>®</sup>, Life Technologies, New Zealand), PenStrep

(GIBCO<sup>®</sup>, Life Technologies, New Zealand) at a working concentration of 100 units/ml penicillin, 100 µg/ml streptomycin, 2 mM glutaMAX<sup>™</sup> (GIBCO<sup>®</sup>, Life Technologies, New Zealand), and 2 µg/ml Fungizone<sup>®</sup> (Life Technologies, New Zealand). The final concentration of glucose in the media was 5.5 mM. The respective supplemented media is henceforth referred to as working media. Cells were maintained at 37 °C in a humidified 5% CO<sub>2</sub> atmosphere. This article does not contain any studies with human or animal subjects performed by any of the authors.

### Generation of 3D Cell Aggregates

Twelve or twenty-four well culture plates were coated with 24 mg/ml poly (2-hydroxyethylmethacrylate) (poly-HEMA) (Sigma, New Zealand), a polymer that prevents cells from adhering to the surface of culture wells. The poly-HEMA was dissolved in 95% ethanol and heated to approximately 70 °C until completely dissolved. The cell culture plates were coated with 300 µL poly-HEMA solution and placed overnight on an orbital shaker at 37 °C and then left to cool down prior the cell culture. Plates were washed with PBS pH 7.4. Cell monolayers were incubated with diluted 1× Trypsin–EDTA (Thermo Fisher Scientific, New Zealand) for 20–30 min to detach cells from the flask. Cell numbers were determined with a hemocytometer to ensure that each well contained approximately 100,000 cells with a total volume of 1 ml working media. Cells were then incubated at 37 °C in a humidified 5% CO<sub>2</sub> atmosphere for 6 days to obtain cellular aggregates. Media were removed and replaced with 1 ml of fresh media every second day.

### Treatments with Canertinib and PHA665752, HGF and EGF

Canertinib (CI-1033) was purchased from LC laboratories (Massachusetts, USA) and PHA665752 was obtained from Sigma (Auckland, New Zealand). Both compounds were dissolved in 100% DMSO. Final DMSO volumes were normalized for all experiments and were less than 0.5% (v/v). Cells were then synchronized in media without FBS for 24 h, before they were exposed to the inhibitors and growth factors. Cells were activated with HGF and EGF (GIBCO®, Life Technologies, New Zealand) during drug treatment. Fresh working media containing combined growth factors at concentrations of 0.2 and 20 ng/ml were used to activate cellular activity. In our dose-dependent stimulation of a single inhibitor, we used a range of (2–4  $\mu\text{M}$ ) canertinib and (0.5–2.0  $\mu\text{M}$ ) PHA665752.

For the combined treatments, cells were treated with both canertinib (3  $\mu\text{M}$ ) and PHA665752 (1  $\mu\text{M}$  for 24 h. After the first 24 h, PHA665752 was re-added to the combination-treated cells, and cells were incubated for a further 24 h before harvesting. At least three independent experiments were carried out and each individual experiment was carried out in triplicates.

### Analysis of Morphology and Cell Viability of Cells in Clusters and Aggregates

After culturing for 6 days, cell aggregates of both cell lines were imaged using an inverted light microscope equipped with a digital camera, using a 10 $\times$ /0.25 N.A. objective lens (ACCU-SCOPE® 3032). After treatment periods, cells in clusters and aggregates were harvested and digested with 1  $\times$  trypsin–EDTA to obtain

single cell suspensions, and cells were counted using a hemocytometer to assess cell viability.

### Immunofluorescent Detection of Receptors

For immunofluorescent imaging, OVCAR-5 and SKOV-3 cells were cultured for 6 days to generate cell clusters and aggregates. The media were refreshed every second day. Cell clusters and aggregates were harvested and fixed with ice-cold 50% (v/v) acetone/methanol solution. OVCAR-5 cell clusters were washed twice with ice-cold PBS, pH 7.4 and re-suspended in 200  $\mu\text{l}$  cold PBS. Clusters were then mounted on poly-lysine coated microscope slides and left to dry at 37 °C. Once cells were dry, cells were blocked with 4% BSA in PBS. Cells were stained overnight with a 1:200 dilution of primary antibody at 4 °C. Cells were then washed with ice-cold PBS pH 7.4. A 1:500 dilution of secondary antibody, conjugated with FITC was added to cells and further incubated for 60 min. The secondary antibody solution was removed and 500  $\mu\text{l}$  of 10  $\mu\text{g/ml}$  DAPI was added to the cells and left for 20 min in the dark. Cells were then washed with 0.1% Tween-20 in PBS pH 7.4 and were mounted in ice-cold anti-fading solution (2 mg/ml *p*-phenylenediamine in 80% glycerol, pH 7.8).

SKOV-3 aggregates were washed with PBS pH 7.4 and stained with aniline blue dye for 20 min. Cells were washed with milliQ water twice before they were embedded in a liquid OCT, an embedding medium used to aid sectioning of frozen tissue samples in a cryostat (Thermo Fisher Scientific, New Zealand). The liquid OCT blocks were frozen at –20 °C for at least 24 h. Sections (7  $\mu\text{m}$  thick) from the block of frozen OCT were cut using a CM186UV Cryostat (Lieca BIOSYSTEM,

Deutschland). Immunofluorescent staining was carried out on both cell lines with anti-EGFR (SC120), anti-Her-2 (SC-7301), and anti-c-Met (SC10) primary antibodies (Santa Cruz). Fluorescent images were captured using an epifluorescence microscope with a 40×/1.3 N.A. oil/DIC objective lens (AxioVision 4.5. Apotome software, Carl Zeiss, Oberkochen, Germany).

### Immunoblotting Analysis

Cell clusters and aggregates were harvested by centrifugation at 400g for 5 min, and the cell pellets were lysed in cold RIPA buffer (50 mM Tris-HCl pH 7.4, 100 mM NaCl, 5 mM EDTA, 1% NP-40, 0.1% SDS, 0.5% Na deoxycholate, 1 mM Na<sub>2</sub>VO<sub>3</sub> containing protease inhibitor cocktail tablets (Complete Mini, Roche, New Zealand). The cell lysates were left on ice for a further 30 min. Sample buffer [0.2% (v/v) bromophenol blue, 25% (v/v) glycerol, 10% SDS in Tris-HCl, and pH 6.8] was added and protein lysates were boiled for 10 min. Prior to loading, the cell lysates were mixed and centrifuged at 9700g for 5 min. A total of 10 µg protein lysate was loaded and separated by SDS-PAGE using a 7% stacking gel and a 10% separating gel. The SDS-PAGE was run at 120 V using Tris-glycine running buffer. The SDS-PAGE markers used were MagicMark™ XP Western Standard (Thermo Fisher Scientific, New Zealand) and Precision Plus Protein standard (Bio-Rad, Hercules, USA). Separated proteins were electro-blotted onto a poly-vinyl membrane (PVDF) (GE Healthcare Life Sciences, New Zealand). The electro blot was run at 100 V for 60 min in cold Tris-glycine running buffer containing 10% v/v methanol. The membranes were blocked for 60 min with either 5% (w/v) non-fat skim milk (Pams brand, New World, New Zealand) or 1% (w/v)

bovine serum albumin (Thermo Fisher Scientific, New Zealand) made up in TBS-T buffer or with Pierce Protein-Free Blocking Buffer (Thermo Scientific, New Zealand). Antibodies (detailed below) were diluted from 1:500 to 1:1000 with the appropriate blocking solution. Membranes were incubated with the primary antibodies overnight at 4 °C. The membranes were washed with TBS-T buffer on an orbital shaker for 4 × 10 min and then incubated with secondary antibody on an orbital shaker for 90 min at room temperature. Membranes were further washed four times with TBS-T. Antibody localization was determined using a chemiluminescent detection kit (Amersham ECL Prime Western Blotting Detection Reagent Kit, GE Healthcare). The protein bands were visualized and a densitometry analysis was performed using Alliance 4.7, Unitec (Cambridge, UK). Cell lysates were collected from at least three separated cell culture experiments. The primary antibodies used in this study were anti-PCNA (sc-25280), anti-GAPDH (sc-25778), anti-EGFR (sc-03), anti-p-EGFR (sc-101668), anti-pHER2 (sc-12352-R), anti-ERK (sc-94), anti-pERK1/2 (sc-7383), Anti-c-Met (sc-10), anti-p-Met (sc-101736), anti-Akt (sc-8312), and anti-p-Akt (sc-101629) (all from Santa Cruz Biotechnology Inc., Santa Cruz, CA, USA, apart from the anti-Her2 which was purchased from BD Biosciences (Auckland, New Zealand)). The two secondary antibodies used in this study were bovine anti-rabbit IgG-HRP (sc-2385) and bovine anti-mouse IgG-HRP (sc-2380).

### Statistical Analysis

Statistical analysis of data was carried out using Prism graph pad. One-way ANOVAs were carried out where  $p < 0.05$  (\*),  $p < 0.01$  (\*\*),



$p < 0.001$  (\*\*\*) and  $p < 0.0001$  (\*\*\*\*) were considered to indicate levels of statistical significance. Prior to analysis percentile and ratio-metric values were logarithmically transformed. All data are presented as mean  $\pm$  SEM. Each experiment was repeated at least three times.

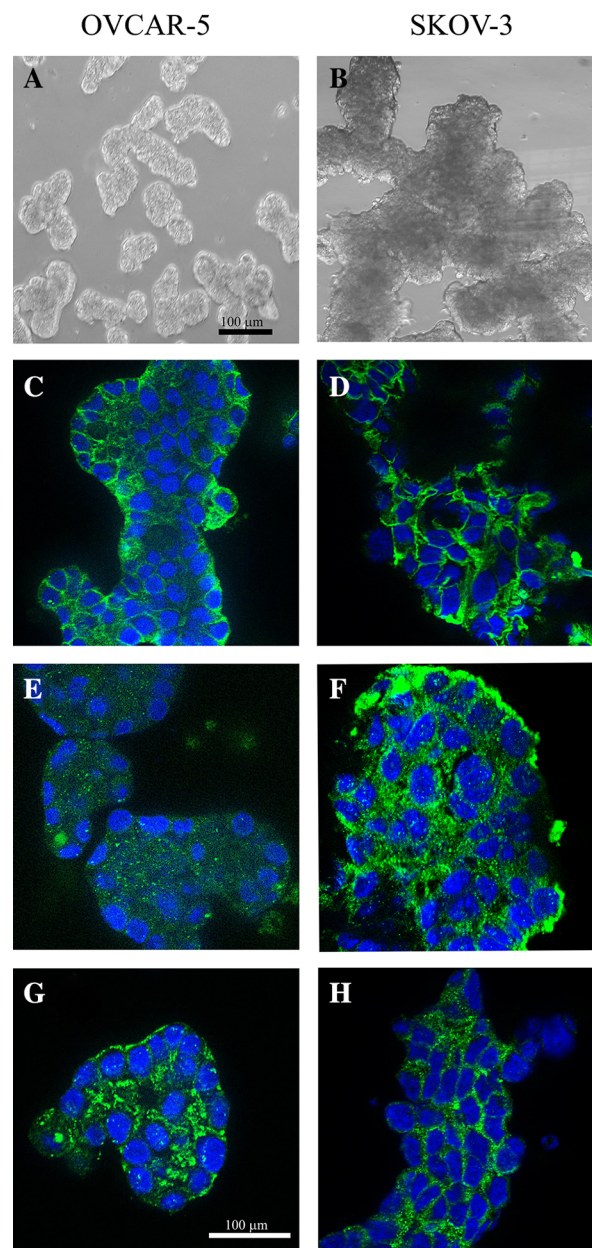
## RESULTS

### Morphology and Expression EGFR, Her-2, and c-Met

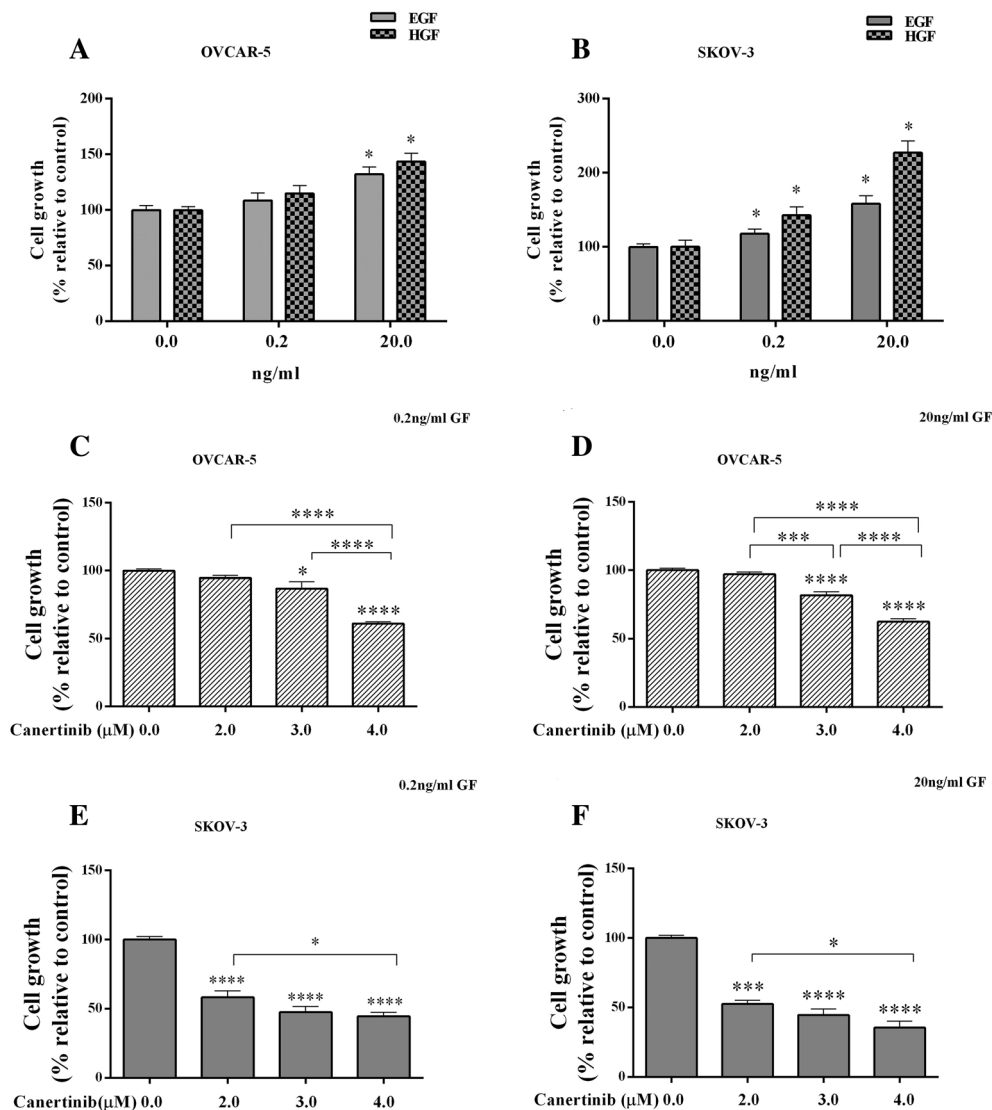
OVCAR-5 cells formed small cell clusters when cultured on a non-adherent surface (Fig. 1a). In contrast, SKOV-3 cells formed large cellular aggregates with dense multicellular structures (Fig. 1b). To determine the presence of cell receptors, we immunofluorescently labeled OVCAR-5 clusters and SKOV-3 cellular aggregates. OVCAR-5 clusters were EGFR (Fig. 1c) and c-Met positive (Fig. 1g), but had low expression of Her-2 (Fig. 1e). SKOV-3 aggregates were EGFR, Her-2 and c-Met positive (Fig. 1d, f, h).

### EGF and HGF Increase the Growth of OVCAR-5 Clusters and SKOV-3 Aggregates

We chose two concentrations of the growth factor EGF and HGF; a physiological concentration (0.2 ng/ml) that approximates to that present in ovarian cancer patients, and a concentration that is commonly used in in vitro studies (20 ng/ml). The growth of OVCAR-5 cell clusters was not increased in the presence of the physiological concentration of growth factor ( $p = 0.4076$ ). However, the higher



**Fig. 1** Morphology of cell clusters and the expression of EGFR, Her-2 and c-Met. Microscopic images of OVCAR-5 ovarian cancer cell clusters (a) and SKOV-3 ovarian cancer cell aggregates (b). The expression of epidermal growth factor receptor (EGFR) in OVCAR-5 (c) and SKOV-3 (d). Total expression of human epidermal growth factor receptor-2 (Her-2) in OVCAR-5 (e) and SKOV-3 (f). Total expression of mesenchymal epithelial transition factor (c-Met) receptors in OVCAR-5 (g) and SKOV-3 (h)



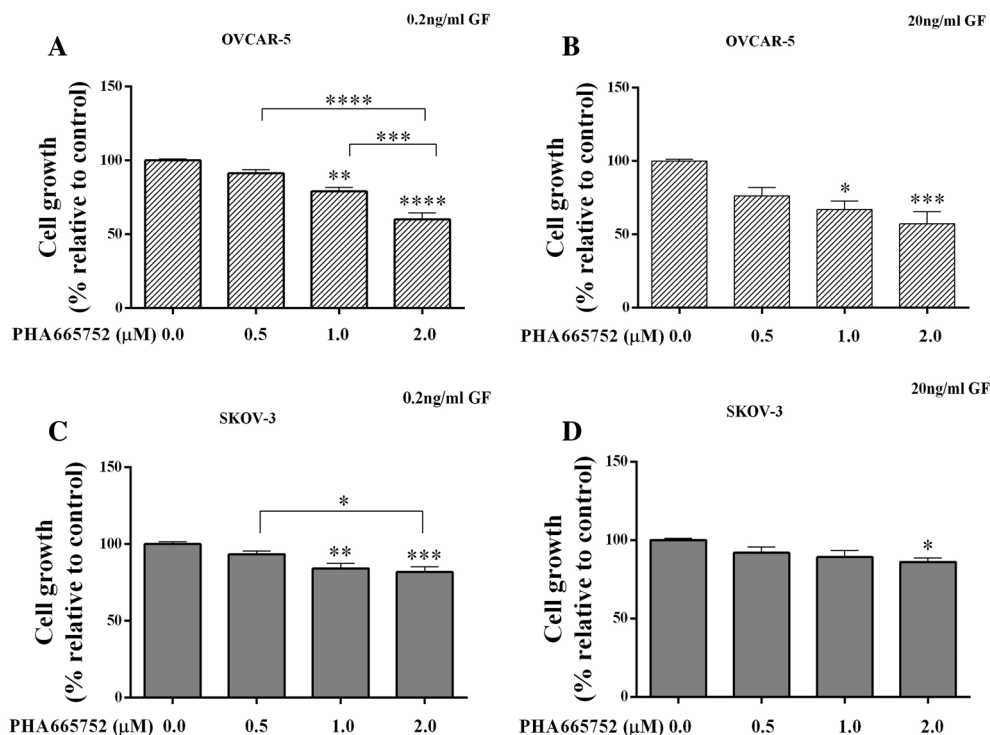
**Fig. 2** Combined growth factors induce cell growth and canertinib inhibits cell growth of ovarian cancer cells. Growth stimulation of OVCAR-5 (a), SKOV-3 (b) cells incubated with 0.2 and 20 ng/ml epidermal growth factor, EGF, or hepatocyte growth factor, HGF. Canertinib inhibited cell growth of OVCAR-5 in a dose-dependent manner in the presence of 0.2 ng/ml (c) and 20 ng/ml (d) combined growth factors. Cell growth of SKOV-3

aggregates was markedly reduced by canertinib in the presence of combined growth factors (e, f). All experiments were independently performed at least three times with triplicates. Data analyzed logarithmically using one-way ANOVA. Data considered statistically significant are indicated as  $p < 0.05$  (\*),  $p < 0.01$  (\*\*),  $p < 0.001$  (\*\*\*), and  $p < 0.0001$  (\*\*\*\*) when compared to the control or the other doses

concentration significantly increased growth ( $p > 0.0001$ ) (Fig. 2a). Both concentrations of EGF and HGF significantly increased growth ( $p > 0.0001$ ) in the SKOV-3 cell aggregates (Fig. 2b).

### Canertinib and PHA665752 Alone Inhibit the Growth of Clusters and Aggregates

We next investigated whether growth of cell clusters and aggregates could be inhibited by the



**Fig. 3** Inhibition of c-Met inhibitor, PHA665752 on cell growth of ovarian cancer cells. PHA665752 inhibited cell growth of OVCAR-5 clusters in a dose-dependent manner in the presence of 0.2 ng/ml (a) and 20 ng/ml (b) combined growth factors. Cell growth of SKOV-3 aggregates was marginally reduced by PHA665752 in the presence of combined growth factors (c, d). All

experiments were independently performed at least three times with triplicates. Data analyzed logarithmically using one-way ANOVA. Data considered statistically significant are indicated as  $p < 0.05$  (\*),  $p < 0.01$  (\*\*),  $p < 0.001$  (\*\*\*), and  $p < 0.0001$  (\*\*\*\*) when compared to the control or the other doses

EGFR/Her-2 inhibitor, canertinib. In their in vivo microenvironment, floating ovarian cancer cells are exposed to a mixture of growth factors (GF). Therefore, we stimulated the cells with the mixture of EGF and HGF. Canertinib inhibited the growth activity in OVCAR-5 cell clusters, in a concentration-dependent manner, in the presence of low and high concentrations of combined growth factors ( $p > 0.0001$ ) (Fig. 2c, d). The inhibitor was very potent against SKOV-3 cell aggregates ( $p > 0.0001$ ) (Fig. 2e, f). Interestingly, the inhibition of canertinib was growth factor independent.

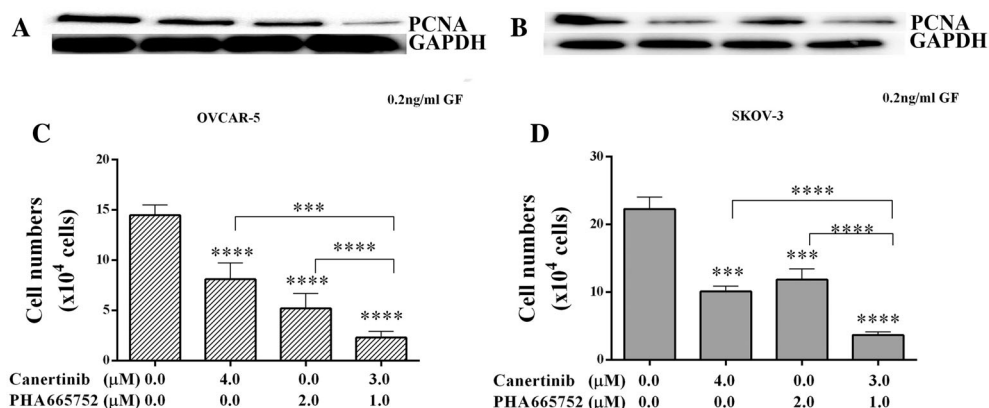
We then examined the effect of PHA665752 on growth. PHA665752 inhibited growth of OVCAR-5 clusters in a concentration-dependent

manner ( $p > 0.0001$ ) (Fig. 3a, b). Its effect on SKOV-3 aggregates was less pronounced, but there still was a decrease in growth ( $p > 0.05$ ) (Fig. 3c, d). Interestingly, growth inhibition by PHA665752 in the presence of low and high concentration mixtures of growth factors was not statistically different (data not shown).

#### Effect of a Combination of Both Canertinib and PHA665752

We investigated the effects of a combination of canertinib and PHA665752 at respective concentrations of 3 μM and 1 μM, which were the lowest effective doses from the single drug treatment experiments. The combination of





**Fig. 4** The effect of the combination of canertinib and PHA665752 on the growth of ovarian cancer cells. The expression of proliferative cell nuclear antigen (PCNA) and a reference protein (GAPDH) in ovarian cancer cells treated with single inhibitors and combination in OVCAR-5 (a) and SKOV-3 (b). Cell numbers of OVCAR-5 cells (c) and SKOV-3 (d) were reduced with single and combination of both inhibitors compared to

untreated cells in presence of 0.2 ng/ml combined growth factors. All experiments were independently performed at least three times with triplicates. Data analyzed using one-way ANOVA. Data considered statistically significant are indicated as  $p < 0.05$  (\*),  $p < 0.01$  (\*\*),  $p < 0.001$  (\*\*\*), and  $p < 0.0001$  (\*\*\*\*) when compared to the control or the other doses

these concentrations reduced total expression of the cell proliferative protein marker, PCNA in OVCAR-5 (Fig. 4a) and SKOV-3 (Fig. 4b) cells. Growth was more reduced with the inhibitor combination, than with the respective single inhibitor alone, in both OVCAR-5 clusters (Fig. 4c) and SKOV-3 aggregates (Fig. 4d) ( $p > 0.0001$ ) in the presence of 0.2 ng/ml GF. Similar patterns of inhibition were obtained with 20 ng/ml GF for OVCAR-5 and SKOV-3 (Fig. 5).

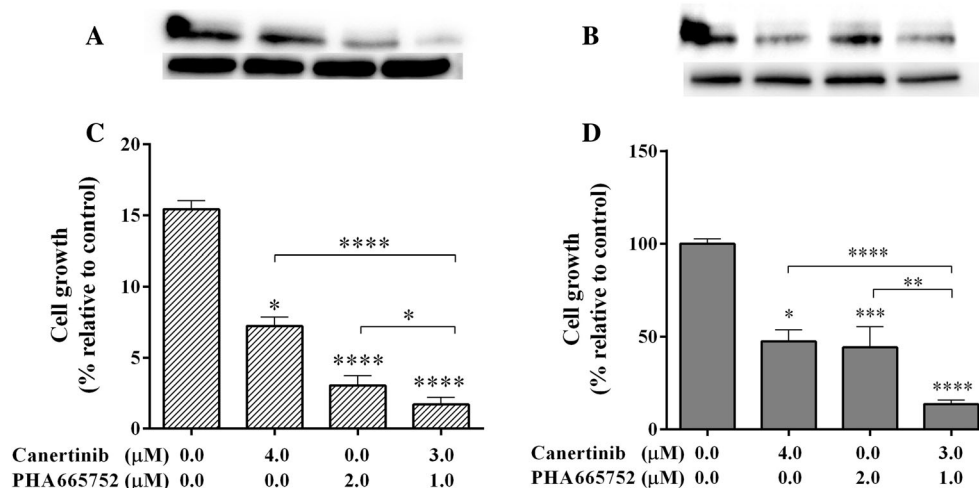
**Canertinib and PHA665752 Reduce Expression of EGFR, Her-2 c-Met, and Their Downstream Signaling Molecules in OVCAR-5 Cell Clusters**

Both single and combined applications of the inhibitors significantly reduced p-EGFR (Fig. 6a, b) in OVCAR-5 cell clusters in the presence of 0.2 ng/ml GF. However, the total expression of EGFR was only reduced in canertinib-treated cells (Fig. 6c). Both c-Met and phosphorylated-c-Met proteins were significantly reduced with a single treatment of PHA665752, and this was further

decreased with the combination (Fig. 6a, d, e). The combination of inhibitors reduced the total expression of Akt (Fig. 6g) and there was a tendency for reduction of Akt phosphorylation (Fig. 6f). The phosphorylation of Erk was reduced in a single treatment alone, but there was greater reduction of p-Erk in the combined inhibitors (Fig. 6h). The total expression of Erk (Fig. 6i) was unchanged. Expression of Her-2 was undetectable in OVCAR-5 cell clusters. With 20 ng/ml combined growth factors, an overall reduction of total expression and phosphorylation of EGFR, and c-Met was observed with the inhibitor combination (Fig. 7). Reductions in p-EGFR, EGFR, p-Met, Met, p-Akt, Akt and p-Erk were observed.

**Canertinib and PHA665752 Reduce the Expression of EGFR, Her-2 c-MET, and Their Downstream Signaling Molecules in SKOV-3 Cellular Aggregates**

Single treatment of SKOV-3 cell aggregates with canertinib reduced the amount of p-EGFR



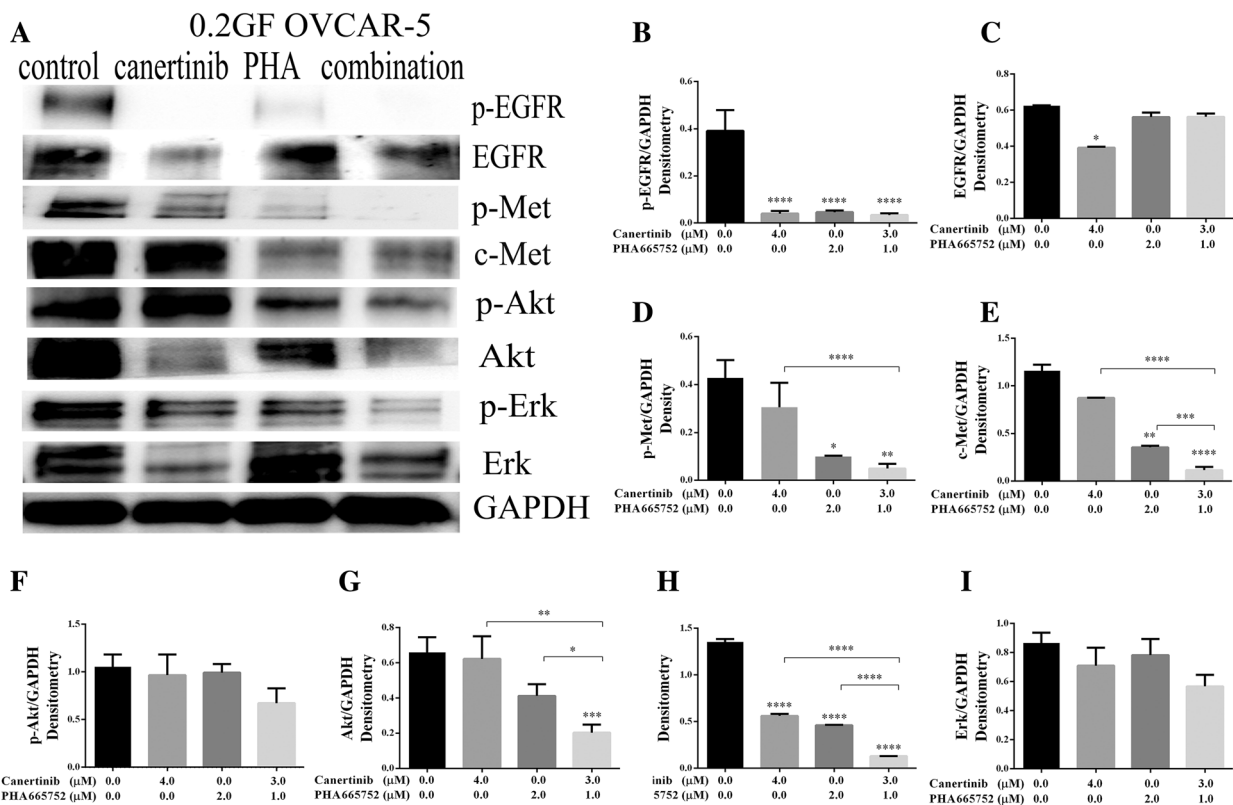
**Fig. 5** The effect of the combination of canertinib and PHA665752 on the growth of ovarian cancer cell in the presence of 20 ng/ml combined growth factors. Expression of proliferative cell nuclear antigen (PCNA) and a reference protein (GAPDH) in ovarian cancer cell clusters treated with single inhibitors and combination in OVCAR-5 (a) and SKOV-3 (b). Cell numbers of OVCAR-5 cells (c) and SKOV-3 (d) were reduced with

single and combination of both inhibitors compared to untreated cells in presence of 20 ng/ml combined growth factors. All experiments were independently performed at least three times with triplicates. Data analyzed using one-way ANOVA. Data considered statistically significant are indicated as  $p < 0.05$  (\*),  $p < 0.01$  (\*\*),  $p < 0.001$  (\*\*\*), and  $p < 0.0001$  (\*\*\*\*) when compared to the control or the other doses

(Fig. 8a), while the application of PHA665752 alone did not reduce p-EGFR. The combination treatment gave a similar result to canertinib alone (Fig. 8b). The total expression of EGFR itself was also reduced with canertinib, and with the combination treatment (Fig. 8c). Phosphorylation and total expression of Her-2 was significantly reduced in both the canertinib and the combination treatment (Fig. 8d, e). Canertinib alone and the combination significantly reduced the phosphorylation of c-Met (Fig. 8f). The single inhibitor treatment reduced the total expression of c-Met, but a greater reduction was observed in the combination (Fig. 8g). The reduction of p-Akt, and p-Erk were also notable in the combination treatment. The total expression of Erk was slightly reduced in the canertinib and combination treatment. Similar results were obtained when cells were activated with 20 ng/ml combined GF (Fig. 9).

## DISCUSSION

A subset of advanced ovarian cancer cells have previously been shown to express high levels of receptor tyrosine kinases that are associated with rapid growth and proliferation. EGFR and Her-2 were upregulated in 53% and 25% of patients with advanced ovarian cancers, respectively, and these were thought to be associated with resistance to chemotherapy and a poor prognosis [8]. Likewise, Swada et al. have shown that c-Met expression increased in 11% of ovarian cancer patients [17], and Davies et al. report that 96% of ovarian cancer patients have positive expression of c-Met in their cancerous tissue [10]. Thus, it is very plausible that cells may have an ability to activate multiple receptors, and this rationale has been described by other studies [2, 17, 18]. Co-activation of multiple receptor tyrosine kinases may reduce the

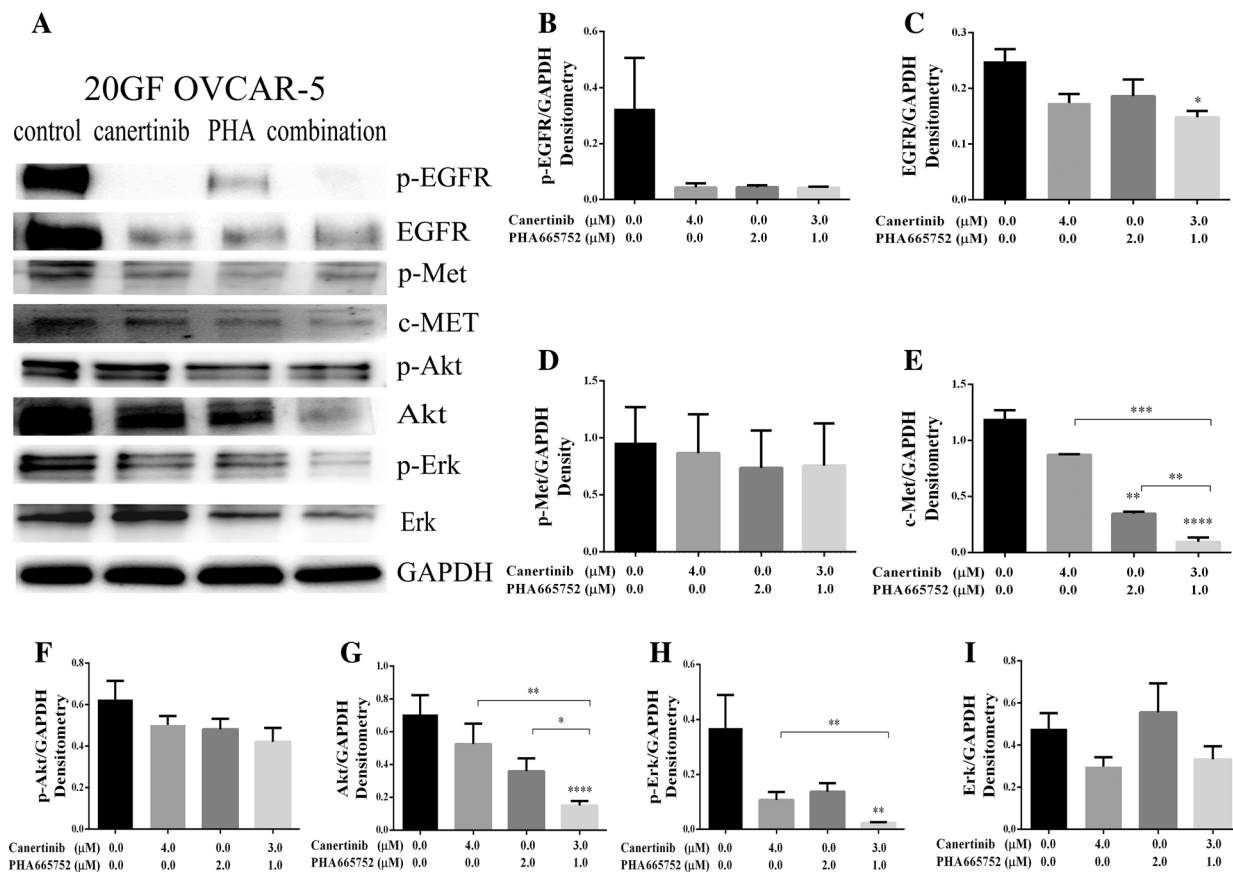


**Fig. 6** Inhibition of selective protein molecules in the presence of canertinib and PHA665752 in OVCAR-5 cell clusters with 0.2 ng/ml combined growth factors. Total protein expression and phosphorylation of EGFR, Her-2, c-Met, Akt and Erk in single and combination of inhibitors. All experiments were independently performed

at least three times with triplicates. Data analyzed logarithmically using one-way ANOVA. Data considered statistically significant are indicated as  $p < 0.05$  (\*),  $p < 0.01$  (\*\*),  $p < 0.001$  (\*\*\*), and  $p < 0.0001$  (\*\*\*\*) when compared to the control or other doses

clinical efficacy of any single drug that targets one of the receptors [19, 20]. Therefore, targeting multiple receptor tyrosine kinases may improve clinical efficacy. Our study attempts to employ such a combined approach by targeting the concurrent activation of EGFR, Her-2 and c-Met. Our study has used a 3D cell culture, in which the generation of cell clusters and aggregates replicates the phenotypic and cellular heterogeneity of floating ovarian malignant cells that are present in ascitic fluid. As such these experiments may be a better representation of the possible response to the combination of drugs in vivo [21, 22].

Consistent with previous reports, we found that the OVCAR-5 cells formed small clusters while the SKOV-3 cells formed aggregates [23, 24]. We also show that OVCAR-5 cell clusters express high levels of EGFR and c-Met, but low levels of Her-2 and that the SKOV-3 aggregates express high levels of all three receptors. These observations are consistent with a previous study [25]. In previous studies, a variety of EGF and HGF concentrations have been used ranging from 10 to 200 ng/ml [26, 27]. However, these concentrations are not physiologically relevant in women with advanced ovarian cancer [28]. We show that a concentration of 0.2 ng/ml, which is a more



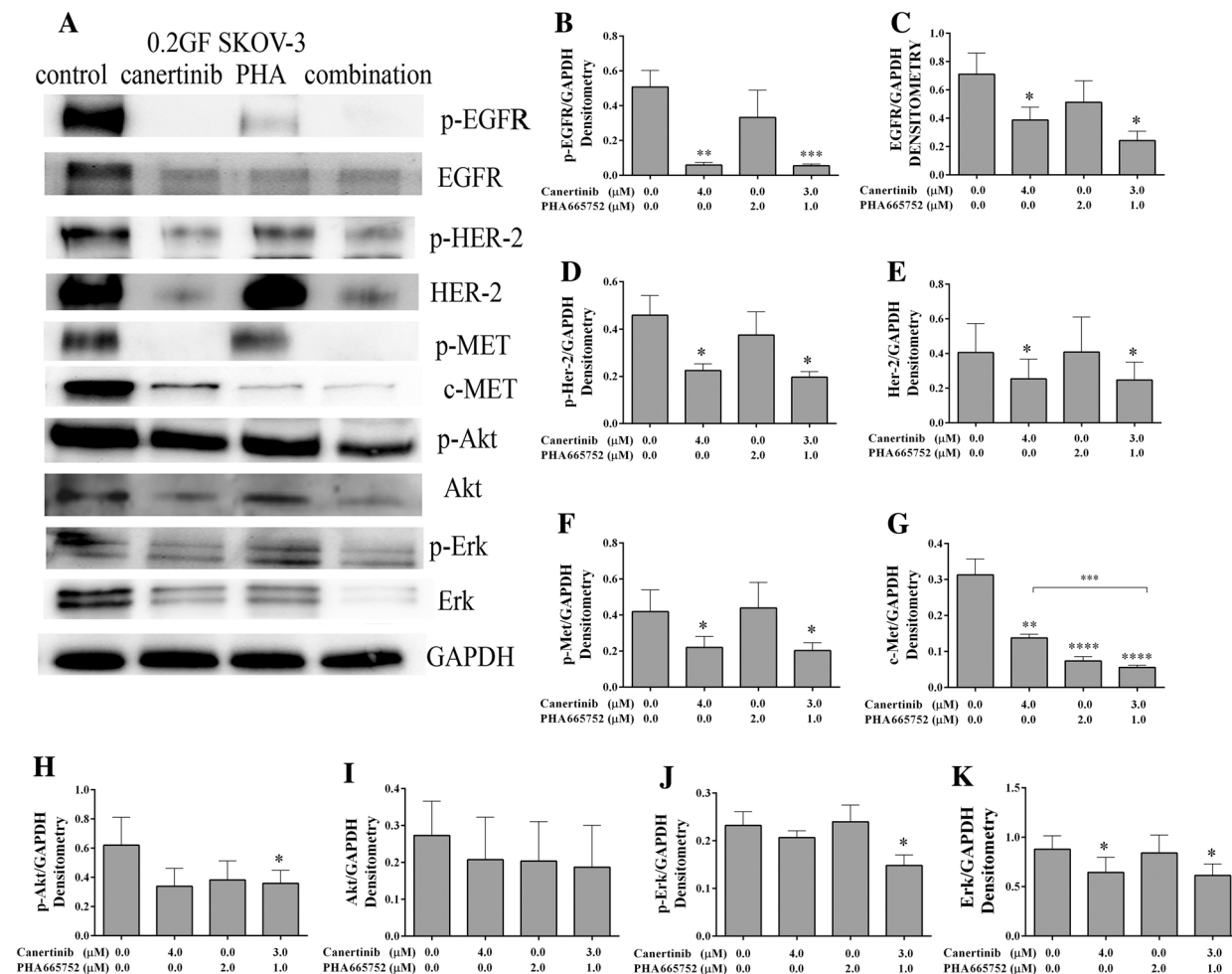
**Fig. 7** Inhibition of selective protein molecules in the presence of canertinib and PHA665752 in OVCAR-5 cell clusters with 20 ng/ml combined growth factors. Total protein expression and phosphorylation of EGFR, Her-2, c-Met, Akt and Erk in single and combination of inhibitors. All experiments were independently performed

at least three times with triplicates. Data analyzed logarithmically using one-way ANOVA. Data considered statistically significant are indicated as  $p < 0.05$  (\*),  $p < 0.01$  (\*\*),  $p < 0.001$  (\*\*\*), and  $p < 0.0001$  (\*\*\*\*) when compared to the control or the other doses

physiologically relevant level, elicits no significant change in the growth of OVCAR-5 line, compared to a concentration of 20 ng/ml. However, the growth of SKOV-3 cell line was growth factor dependent. It is unclear therefore to what extent EGF and HGF determine ovarian cell growth in physiological conditions.

We have assessed the effect of canertinib, an irreversible dual inhibitor for EGFR family of receptor tyrosine kinases, and PHA665752, a reversible, specific inhibitor for c-MET protein, on both cell lines in the presence of EGF and HGF which are the respective ligands for the

receptors [29]. In our study, the selective concentrations of single drug treatments were chosen on the basis of concentrations in a mouse model previously described [30]. Canertinib binds irreversibly to the ATP pocket of the intracellular domain of ErbB family receptor tyrosine kinases, forming a covalent bond that irreversibly blocks EGFR, Her-2 and Her-3 signal transduction. Reversible small molecule inhibitors of EGFR such as Gefitinib (ZD1839, Iressa) and Erlotinib (OSI-774, Tarceva) have been approved by the FDA for treatment of NSCLC [31]. The anti-growth



**Fig. 8** Inhibition of selective protein molecules in the presence of canertinib and PHA665752 in SKOV-3 cell clusters with 0.2 ng/ml combined growth factors. Total protein expression and phosphorylation of EGFR, Her-2, c-Met, Akt and Erk in single and combination of inhibitors (a). All experiments were independently

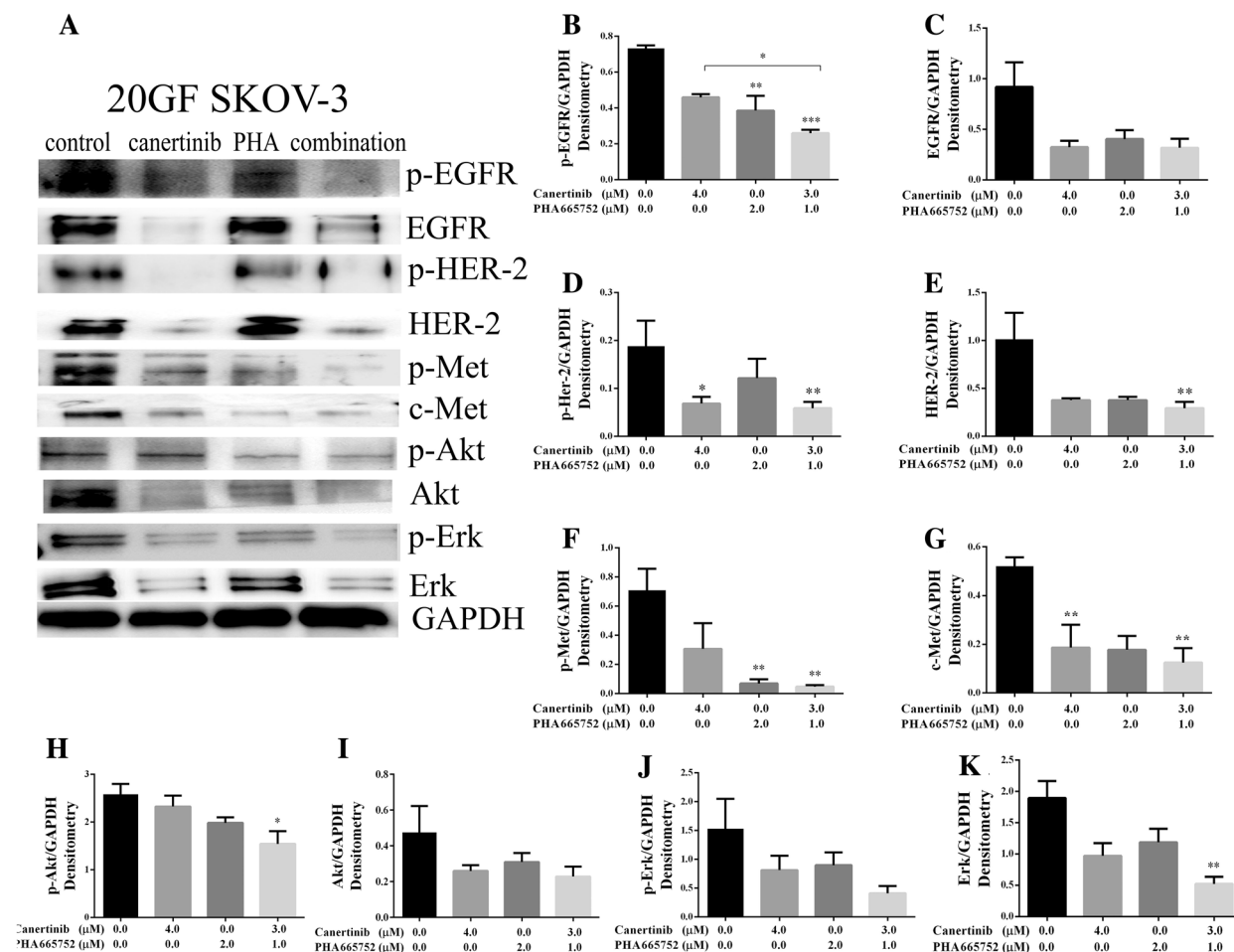
performed at least three times with triplicates. Data analyzed logarithmically using one-way ANOVA. Data considered statistically significant are indicated as  $p < 0.05$  (\*),  $p < 0.01$  (\*\*),  $p < 0.001$  (\*\*\*), and  $p < 0.0001$  (\*\*\*\*) when compared to the control or the other doses

properties of these drugs have been explored in other EGFR-positive malignancies, and unfortunately the majority of these trials have failed to exhibit prolonged clinical effects [32, 33]. The difference between canertinib, and gefitinib and erlotinib [33], is that canertinib irreversibly binds to the active site of ATP, completely blocking the ATP-binding process and thereby decreasing the activity of EGFR and Her-2 permanently. Therefore, there

is a good rationale for utilizing an irreversible inhibitor, in order to prolong the suppression of EGFR family activity and their downstream molecules [34].

We show that canertinib inhibits growth of SKOV-3 cell aggregates in a dose-dependent fashion. This effect may be due to the abundant expression of EGFR and Her-2. In contrast, OVCAR-5 cell clusters exhibited less sensitivity to canertinib, possibly because they





**Fig. 9** Inhibition of selective protein molecules in the presence of canertinib and PHA665752 in SKOV-3 cell aggregates with 20 ng/ml combined growth factors. Total protein expression and phosphorylation of EGFR, Her-2, c-Met, Akt and Erk in single and combination of inhibitors. All experiments were independently performed

at least three times with triplicates. Data analyzed logarithmically using one-way ANOVA. Data considered statistically significant are indicated as  $p < 0.05$  (\*),  $p < 0.01$  (\*\*),  $p < 0.001$  (\*\*\*), and  $p < 0.0001$  (\*\*\*\*) when compared to the control or the other doses

might have other downstream mutations that facilitate cell survival [35]. Irwin, et al. showed that canertinib at 0.1 and 5  $\mu\text{M}$  inhibit the proliferative capacity of acute lymphoblastic leukemia cells [36]. Canertinib used as a single agent showed clinical responsiveness in Her-2 positive tumors of breast cancer patients [37].

In our study, we also demonstrate that c-Met inhibitor PHA665752, when used as a single agent, inhibits SKOV-3 and OVCAR-5 growth in a dose-dependent manner. However, SKOV-3

cell aggregates seem to be less responsive to the c-Met inhibitor, compared to OVCAR-5 clusters. One possible explanation is that SKOV-3 cells have a high level of Her-2 and that the SKOV-3 cells may be able to resist the inhibitory action of the c-Met inhibitor. As a single agent, PHA665752 has been shown to induce apoptosis, cease cell proliferation and spread in lung adenocarcinoma cells with *k-Ras* gene mutations [38]. Ma et al. demonstrated that PHA665752 inhibited the growth of

BaF3.TPR-Met cells in a dose-dependent manner [39] and Christensen et al. showed that PHA665752 at a concentration range of 0.1–5.0  $\mu\text{M}$  inhibited growth of gastric, pancreatic, and lung carcinoma cell lines [14, 40]. Other studies have shown NSCLC and gastric cancer cells are sensitive to concentrations of PHA665752 between 0.01 and 10  $\mu\text{M}$  [9].

In our study, we demonstrate that the combined inhibitors (using the lowest effective single treatment concentration of each inhibitor) elicit a greater reduction of cell growth than a single agent alone. This finding may suggest that the combination of EGFR/Her-2 and c-Met inhibitors is sufficient to block multiple cellular pathways and that can compromise the growth and survival of cancer cells. The rationale of combined targeting with multiple drugs has been highlighted in recent years. Engelman et al. demonstrated that a combination of gefitinib and PHA665752 reduced cell proliferation more than a single treatment alone, in NSCLC and gastric cancer cell lines [9], while Crosswell et al. indicated that combining PHA665752 with rosiglitazone strongly inhibited cell growth of a neuroblastoma cell line [41]. In addition, Tang et al. demonstrated that the inhibition of both EGFR and c-Met receptors strongly compromised the growth activity of EGFR-mutation mediated erlotinib-resistant lung cancer cells [42].

In clinical trials, the combination of cytotoxic agents with small molecule receptor tyrosine kinase inhibitors or monoclonal antibodies has been also investigated in ovarian and other cancers. The combination of cetuximab, a monoclonal antibody to EGFR, and paclitaxel and carboplatin is adequately tolerated as a primary therapy for ovarian cancer but does not demonstrate the increase

of progression-free survival [43]. The combination of lapatinib plus topotecan for the treatment of platinum-resistant ovarian cancer showed limited clinical benefit [44]. Chun et al. showed that a combination of gemcitabine and gefitinib synergistically reduced cell growth of head and neck cancer cells in an in vitro and an in vivo animal model, and the phosphorylation of EGFR and the downstream molecule Erk was also down regulated [45]. Furthermore, breast cancer patients treated with the combination of capecitabine with the Her-2 antibody, trastuzumab, or with lapatinib showed clinical activity, but increased side effects were noticed [46, 47]. Certainly, the present study is limited given the lack of any preclinical animal model study. This would be essential to confirm the efficacy of our combination inhibitors and to identify potential adverse side effects.

## CONCLUSION

Studying the SKOV-3 and OVCAR-5 ovarian cancer cell lines in vitro, we demonstrate a significant effect on growth of a combination of two tyrosine kinase inhibitors that block three receptor tyrosine kinases. Further work is required to elaborate these effects in animal and human studies. SKOV-3 and OVCAR-5 cell lines do not entirely represent in vivo ovarian cancer cells. Therefore, we need to know if these effects are seen in other cell lines, including primary ovarian cancer cells from patients, and how the effects correlate with the receptor status. The combination of receptor tyrosine kinase inhibitors at effective concentrations may, however, have the effect of increasing toxicity. Animal studies are essential to assess any adverse side effects prior to any clinical trials.

## ACKNOWLEDGMENTS

The work was supported by a Grant from the Ovarian Cancer Research Foundation (OCRF) Melbourne, Australia and School of Biological Sciences, University of Canterbury, Christchurch, New Zealand. No funding or sponsorship was received for publication of this article. All named authors meet the International Committee of Medical Journal Editors (ICMJE) criteria for authorship for this manuscript, take responsibility for the integrity of the work as a whole, and have given final approval for the version to be published. The Ministry of Education, Iraqi Government, supports the scholarship for WH. WH performed most experiments. KC, PS and AG designed the study and prepared a manuscript. All authors read and approved the final manuscript.

**Disclosures.** Wafaa Hassan, Kenny Chitcholtan, Peter Sykes and Ashley Garrill have nothing to disclose.

**Compliance with Ethics Guidelines.** This article does not contain any studies with human or animal subjects performed by any of the authors.

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