Effects of Targeted Anticancer Medicines on Post-Cell Removal Surface Morphology of Cancer Cells Cultivated on 3-Aminopropyltriethoxysilane Surface

Chung-Ping Hsu1,2, You-Lin Wu2*, Wan-Yun Lee1, Li-Wen Li3 and Jing-Jenn Lin4
1Division of Thoracic Surgery, Department of Chest Surgery, Taichung Veteran General Hospital, Taichung, Taiwan, Republic of China
2School of Medicine, National Yang Ming University, Taipei, Taiwan, Republic of China
3Department of Electrical Engineering, National Chi Nan University, Puli, Nantou, Taiwan, Republic of China
4Department of Applied Materials and Optoelectronic Engineering, National Chi Nan University, Puli, Nantou, Taiwan, Republic of China

Abstract

A post-cell-removal surface morphology (PCRSM) profiling technique was used to identify the effects of targeted anticancer medicines on cancer cells. Living non-small lung cancer cells, A549 and H1299, were cultivated on a 3-aminopropyltriethoxysilane (γ-APTES) coated silicon wafer surface with and without targeted anticancer medicine added in the culture medium. Atomic force microscopy (AFM) was used to examine the surface morphology profile on the γ-APTES wafer surface after removing the cells. Two different targeted anticancer medicines, epidermal growth factor receptor (EGFR)-inhibitor Iressa (gefitinib) and protein kinase c (PKC)-inhibitor Staurosporine were examined. Our experimental results show that only the cancer cells treated with Staurosporine can have the PCRSM profiles resemble to those of normal cells, whereas those treated with Iressa reserve the PCRSM profiles of the pre-medicine treated cancer cells. This observation indicates that the PCRSM technique is able to detect the cell-traction force difference caused by EGFR-inhibitor and PKC-inhibitor, respectively and Staurosporine is more effective than Iressa in deactivating the cell-substrate interaction of the cancer cells.

Keywords: Surface morphology; Targeted anticancer medicine; Lung cancer; Cell traction force; Atomic force microscopy; Cell-substrate interaction

Introduction

Activating the apoptosis mechanism of the cancer cells has been recognized as one of the key factors in development of anticancer medicines. It has also been pointed out that manipulation of the intrinsic pathway for mitochondria-dependent apoptosis activation is an efficient way for cancer cells treatment [1]. Upon anticancer medicine treatment, the mitochondria proteins cytochrome c (cyt-c) and Smac protein/direct inhibitor of apoptosis (IAP) binding protein with low isoelectric point (pl) (DIABLO) are released into the cytosol. Here, they synergistically activate caspases through activating apoptotic protease activating factor 1 (APAF-1) and relieving the apoptotic inhibition with IAPs [2-7]. In recent years, new agents aimed at target-specific intracellular pathways, such as angiogenesis inhibitors, EGFR inhibitors, and PKC inhibitors related to the distinctive properties of cancer cells have been developed and tested [8-17]. On the other hand, in vitro testing of cancer cells treated with anticancer medicines using cell-based essays has become the usual way for development of anticancer medicines. In previous work, we reported that the cell-substrate interaction property can be determined via cultivating cells on a γ-APTES modified silicon dioxide (SiO2) surface and then measure the PCRSM profiles with atomic force microscopy technique [18]. With this method, we were able to observe the evolvement of the cell–substrate interaction over time and found that the cancer and normal cells would leave different imprints on the γ-APTES surface due to different cell traction forces that are generated through actomyosin tractions and actin polymerization. In this work, we used the PCRSM profiles technique to investigation the cancer cells treated with or without the targeted anticancer medicines, Iressa and Staurosporine. These two anticancer medicines were developed based on the aforementioned target-specific intracellular pathways related mitochondria-dependent apoptosis mechanism, and are commonly used for non-small lung cancer (NSLC) treatment. It should be noted that Iressa is an anticancer medicine that inhibits EGFR tyrosine kinase through binding to the adenosine triphosphate (ATP)-binding site of the enzyme [19,20], whereas Staurosporine is a medicine that prevents the ATP binding with the PKC to deactivate the ability of growth and transformation of the affected cells [8]. In addition to the two NSLC cells of A549 and H1299, the Madin–Darby canine kidney (MDCK) epithelial cell, one of the normal cells commonly used for biochemical analysis, was also examined for the comparison. The PCRSM profiles of those cells were obtained via cultivation of the cells on γ-APTES surface after treatment either with or without anticancer medicines and measurement of the imprints left after cells removal. The reason we cultivated the cells on γ-APTES layer is that rich NH2 bonds can be provided on its surface and give the cells better attachment [21,22].

Materials and Methods

Substrate preparation

A p-type (100) silicon wafer covered with a 2 nm-thick SiO2 surface layer was used as the supporting substrate. A 1% ethanol solution of γ-APTES was then spin-coated onto the SiO2 surface and cured at 120°C for 5 min on a hot plate. The final thickness of γ-APTES layer was determined with ellipsometry to be almost 550 nm. After coating with γ-APTES, the Si wafer was subjected to sterilization in an autoclave sterilizer at 110°C for 90 min in vacuum. The Si wafer was then divided into 12 pieces and placed in a 12-well culture plate for cell cultivation.

*Corresponding author: You-Lin Wu, Department of Electrical Engineering, National Chi Nan University, 301 University Rd., Puli, Nanto, Taiwan, 54561, Tel: +886-49-2910960 (X-4805); Fax: +886-49-2917810; E-mail: ylwu@ncnu.edu.tw

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Cell culture and removal

Before cultivation of cells on the substrate surface, all the cell lines were immersed in a 5 mL Dulbecco modified Eagle’s medium (DMEM) supplemented with 10% fetal bovine serum containing 10 mg/mL penicillin/ streptomycin and 2 mL glucose, and cultured in an incubator with 5% CO₂ at 37°C. After cultivation of A549 and H1299 (obtained from the American Type Culture Collection, Manassas, VA, USA) for 3 days and the MDCK cells for 7 days, each cell line was seeded separately onto the γ-APTES modified silicon surface placed in the 12-well culture plate with an initial concentration of 8×10³ cell/well. The 12-well plate was then immersed in culture medium DMEM solutions either with or without anticancer medicine addition. For the anticancer medicine examinations, 10 μL Iressa and Staurosporine (Sigma-Aldrich, USA) solutions with various concentrations were prepared and added separately to the culture medium. The cells cultured on the γ-APTES modified SiO₂ surface were then removed after 24 h of cultivation, respectively through immersing the samples in an 75% ethanol solution for 10 min followed by a phosphate buffer solution (PBS) wash-off process and afterward drying at room temperature under normal atmospheric environment. The 24 h cultivation is needed to ensure that the cells are completely adhered to the substrate surface. Imprints were left on the γ-APTES surface after the removal of cells. Optical microscopy was used to make sure the removal of the cells.

PCRSM profiles measurement

For surface morphology image of the cells before they were removed, we placed each cell line cultured on the γ-APTES modified SiO₂ surface into a dish-type slot filled with culture medium and loaded it into a SEIKO 300HV AFM system for surface morphology measurement which was then carried out under a liquid environment. After the removal of the cells, each sample was again loaded into a SEIKO 300HV AFM system for surface morphology image measurement that was performed under a vacuum of 5×10⁻⁶ Torr. The surface morphology examinations were carried out under tapping mode with a resonant frequency of 130 kHz, a force constant of 15 N/m and a speed of 0.7 Hz. The PCRSM profiles were obtained by scanning a straight line across the imprint of each cell using the AFM tip. In this work, a Si tip with a radius of smaller than 7 nm was used for all the measurements. Figure 1 shows schematically the process steps of the PCRSM profiling technique.

Results

As we reported in our previous paper, the PCRSM profiling technique enables us not only to observe the evolvement of the cell-substrate interaction over time, but also differentiate the difference of cell-substrate interaction between normal and cancer cells [18]. Prior to measuring the PCRSM profiles, we checked the surface morphology images before and after the removal of the cells. Figure 2(a) shows the AFM surface morphology images of the normal cells MDCK, Beas-2B, and MRC5 cultivated for 24 h on a γ-APTES modified SiO₂ surface before they were removed. Their corresponding PCRSM AFM images are shown in figure 2b. Figures 3a and 3b show the AFM surface morphology images of cancer cells A549 and H1299 cultivated for 24 h on a γ-APTES surface before and after they were removed, respectively. Both normal and cancer cells were observed being completely adhered to the substrate surface and imprints were left on the γ-APTES surface after the cells were removed due to cell-substrate interaction. Figure 4a shows the PCRSM profiles of the normal cells Beas-2B, MRC5, and MDCK, whereas figure 4b depicts those of the cancer cells A549 and H1299 after 24 h cultivation on γ-APTES surface, as reported by Hsu et al. [18]. The surface morphology profiles shown in figure 4 were normalized to the original γ-APTES surface before they were removed, that is, the original γ-APTES surface was taken as reference zero. As observed, the cancer cells tend to form deeper trench along the circumference and a protrusion at the center of the PCRSM profiles, whereas the normal cells exhibit flatter PCRSM profiles at the bottom of the adhesion region. That is, the PCRSM profile for the cancer cells is nearly in W-shape, but that of the normal cells is somewhat like a flat line. In this work, only the MDCK cell was used to be compared with the cancer cells since it produces the deepest PCRSM profiles, and among the three normal cells is closest to the PCRSM profiles of the cancer cells. In order to check the effect of anti-cancer medicine on the PCRSM profiles, we added anti-cancer medicines Staurosporine and Iressa into the culture medium, respectively. Figure 5a and b shows respectively the normalized PCRSM profiles of cancer cells A549 and H1299 after 24 h cultivation in culture medium with various concentrations of Staurosporine and Iressa.
concentration of anticancer medicine Staurosporine. As observed, for both the A549 and H1299, the PCRSM profiles change from W-shape to nearly flat line as the Staurosporine concentration increases. The same amount of Staurosporine (100 nM) was also added to the culture medium for the normal cell MDCK. The PCRSM profiles of the MDCK cells after treatment with and without anticancer medicines are shown in figure 6a. The PCRSM profiles of the cancer cells A549 and H1299 treated with and without 100 nM Staurosporine are given in figure 6b and 6c, respectively, for comparison. The difference of the PCRSM profiles between normal cells and cancer cells is clear. As observed, although the depth of PCRSM profile of the MDCK cell decreases after the treatment of anticancer medicine, the profile basically remains the flat line shape. For cancer cells, the PCRSM profiles also turn into nearly flat after been treated with 100 nM Staurosporine.

We also conducted an experiment through culturing the cancer cells in Staurosporine-containing medium with concentrations of 10 and 100 nM for 24 h followed by replacing the culture medium with anticancer medicine free ones. Then, their PCRSM profiles were taken after another 24 h cultivation period. The resultant PCRSM profiles for the cancer cells A549 and H1299 are shown in figure 7a and b, respectively. The PCRSM profiles of the cancer cells H1299 and A549 before and after the treatment with 100 nM Staurosporine are also included in figure 7 for comparison. As shown in figure 7a, after replacing with anticancer medicine free culture medium the PCRSM profile of A549 changes from almost flat profile to the one with deeper trench along the circumference and a protrusion at the center, regardless of whether the cancer cells had been treated with 10 or 100 nM Staurosporine. That is, the PCRSM profiles change from nearly flat-line shape to W-shape. Similar result was observed for the PCRSM profiles of the cancer cell H1299, as shown in figure 7b.

Discussion

The cell-substrate interaction has been recognized as an indication that cells will generate a local force via so-called cell–extracellular matrix (ECM) interactions. It has been confirmed that it is the traction force caused by actin polymerization at the cell's leading membrane edge that transmits the contractility force to the ECM via the primary mediators' focal adhesion protein integrins [23-25]. In addition, reports have shown that cell's central region also plays an important role in cell–substrate interaction [26,27]. Since cell-substrate interaction is
directly related to the traction force of the cell, the cell property can be judged from the imprint it leaves on the soft substrate surface onto which the cell is cultured. The cancer cells and the normal cells show different surface morphology profiles of the imprints, i.e., they have the PCRSM profiles due to different cell-substrate interactions. It is found that the PCRSM profile for the cancer cells is nearly in W-shape, but that of the normal cells is somewhat like a flat line, as shown in figure 4. Based on the PCRSM profiling technique, we compared the effects of anticancer medicines, Staurosporine and Iressa, on the cell-substrate interaction of NSCLC cancer cells A549 and H1299. Figure 5 shows that the PCRSM profiles of the cancer cells changed from a nearly flat-line shape to a W-shape when the Staurosporine-containing culture mediums were replaced with anticancer medicine free ones, (Figure 7). It is also noticed that, the recovered PCRSM profiles of the cancer cell H1299 are closer to the PCRSM profiles of the cancer cells subjected to no medicine treatment than that of the cancer cell A549. This is an indication that the malignancy of A549 is much more severe than that of H1299 [29-31]. Surprisingly, the effect of Iressa on the PCRSM profiles of cancer cells is not the same as that of Staurosporine. The PCRSM profiles maintained a W-shape for both the cancer cells A549 and H1299 after being subjected to a much higher concentration (50 µM) of Iressa. In figure 8 we compared the PCRSM profiles of the cancer cells subjected to 100 nM Staurosporine and 50 µM Iressa, respectively. As observed,
the effects of Staurosporine and Iressa on the cell-substrate interactions of the cancer cells were different. The PCRSM profiles adopted a flat-line shape after a 100 nM Staurosporine treatment, while those that were treated with 50 µM Iressa maintained their W-shape. From a cell-substrate interaction point of view it seems to be apparent that the PKC-inhibitor Staurosporine is more effective than the EGFR inhibitor Iressa, which is in agreement with the reports from Wu et al. [31] and Notbohm et al. [32]. It was also noticed that, although the PCRSM profiles remained in a W-shape, the central protrusion of the PCRSM profiles increased after treatment with Iressa. This observation implies that Iressa induces an increase in cell traction force toward the center region of the cell, along the circumference of the cell.

Conclusion

Based on our previously developed PCRSM profiling technique, we compared the effects of anticancer medicines, Staurosporine and Iressa, on the cell-substrate interaction of NSCLC cancer cells A549 and H1299. It was observed that the PCRSM profiles of the cancer cells change from W-shape into flat-line shape after treatment with a certain amount (100 nM) of Staurosporine, whereas the other ones remain in W-shape after treatment with the same amount of Iressa. This indicates that the PKC-inhibitor Staurosporine is more effective than the EGFR inhibitor Iressa on reducing the cell traction force. No report in the literature has addressed the differences between the EGFR-inhibitor and PKC-inhibitor on the cell-substrate interaction, although it has been reported that both Staurosporine and Iressa induce the shape and morphology change of NSCLC cells [10,32,33]. It is evident that the PCRSM profiling technique allows us to see the difference of the cell-substrate interaction between the EGFR-inhibitor and PKC-inhibitor on reducing the cell traction force. Although further investigations are needed to disclose the actual mechanism behind these observations. We believe that the PCRSM profiling technique could be used as a tool for the development of new targeted anticancer medicines in vitro.

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