RESEARCH ARTICLE

Cucurbitacin B Diminishes Metastatic Behavior of Cholangiocarcinoma Cells by Suppressing Focal Adhesion Kinase

Putthaporn Kaewmeesri, Veerapol Kukongviriyapan, Auemduan Prawan, Sarinya Kongpetch, Laddawan Senggunprai*

Abstract

Objective: Cholangiocarcinoma (CCA) is a malignant tumor with aggressive metastatic property resulted from dysregulation of metastasis-regulated signaling pathways. The aim of this study was to investigate the effect of cucurbitacin B on metastatic behavior of CCA cells through modulation of focal adhesion kinase (FAK) protein. Methods: KKU-452 cells were treated with a specific FAK inhibitor, FAK inhibitor-14, or cucurbitacin B at various concentrations for 24 h. Cell viability was assessed by sulforhodamine B assay. The migratory and invasive abilities of the cells were investigated using wound healing and transwell invasion assays, respectively. The fibronectin-coated plate was used for adhesion assay. The effects of the test compounds on FAK activation and the expression of metastasis-associated proteins were determined by Western blot analysis. The amount of MMP-9 was evaluated using a commercial ELISA Kit. Results: FAK inhibitor-14 and cucurbitacin B at concentrations which minimally affected KKU-452 cell viability could suppress FAK activation, evidently by decreased level of phospho-FAK protein after exposure to the compound. At these conditions, cucurbitacin B suppressed metastatic behavior including migration, invasion and adhesion abilities of CCA cells similar to FAK inhibitor-14. Further molecular studies demonstrated that FAK inhibitor-14 and cucurbitacin B downregulated the expression of metastasis-associated proteins including MMP-9, ICAM-1 and VEGF. Consequently, exposure to cucurbitacin B inhibited the production of MMP-9 enzyme in CCA cells similar to FAK inhibitor-14 treatment. Conclusion: FAK participated in regulation of metastatic behavior of KKU-452 CCA cells. Cucurbitacin B suppressed FAK activation in the cells which was associated with inhibition of metastasis essential steps and their related metastatic proteins. The compound may be developed as a novel therapeutic agent for CCA metastasis therapy.

Keywords: FAK- cholangiocarcinoma- metastasis- cucurbitacin B- FAK inhibitor- MMP-9- ICAM-1- VEGF

Asian Pac J Cancer Prev, 22 (1), 219-225

Introduction

Cholangiocarcinoma (CCA) or bile duct cancer is a malignant tumor with extremely aggressive metastatic behavior resulting in a very poor prognosis (Doherty et al., 2017). Chronic inflammation by liver fluke infection is one of the established risk factors associated with cholangiocarcinogenesis particularly for people residing in Southeast Asia (Zheng et al., 2017). According to changes in intracellular signaling networks contributing in CCA development, targeted therapies to these pathways have been intensively studied for eventually gaining promising treatment of CCA (Fouassier et al., 2019).

Metastasis of cancer is an important clinical problem for cancer treatment. In CCA, most patients are diagnosed at an advanced or metastatic stage, where curative surgical treatment is not appropriate and chemotherapy is a remaining option. Prognosis for patients with metastatic cancer is generally poor. Recently, the association between metastasis and survival rate of CCA patients in Eastern region of Thailand has been analyzed. The results showed that metastasis was a significant clinical predictor of death among these patients (Charonponsuntorn et al., 2019). It is therefore, effective therapeutic compounds capable of suppressing CCA metastasis are still highly needed. Cancer cell metastasis depends on the ability of cells to migrate and invade in which they are influenced by the surrounding extracellular matrix (ECM) (Welch and Hurst, 2019). Thus, degradation of ECM components considered as an essential step in facilitating the metastasis process. It has been reported that matrix metalloproteinase-9 (MMP-9) is the major enzyme participating in proteolytic degradation of ECM proteins particularly type IV collagen (Curran and Murray, 2000). Besides migratory and invasive properties,
the adhesion ability of cells to ECM is also required for cancer cells with metastatic behavior to be settled at the target organ site (Paschos et al., 2009). Additionally, angiogenesis, the establishment of new blood vessels, is also a necessary component of the metastatic pathway. The spread of metastatic cells depends on an adequate supply of oxygen and nutrients from the vascular network formed (Bielenberg and Zetter, 2015). Hence, the interventions targeting on cancer cell migration, invasion, adhesion as well as angiogenesis could be the potential strategies for suppression of cancer metastasis. It has been reported that several compounds could inhibit these metastasis-associated events and molecules such as ellagic acid inhibited MMP-9 expression in human prostate cancer cells (Pitchakarn et al., 2013) and trimellitic acid had the capability of angiogenic inhibition by suppressing proangiogenic activity of fibroblast growth factor 2 and inhibiting HUVEC tube formation (Mahdavi et al., 2017).

Metastasis process is regulated by several intracellular signaling cascades (Inan and Hayran, 2019). Cumulative evidence suggested that FAK acts as a prominent regulator molecule for transmitting signals to activate downstream targets associated with metastasis process (Tai et al., 2015). Elevation of FAK expression was strongly associated with tumor metastasis. The activation of FAK increased the expression of metastasis-associated molecules such as MMPs and intercellular adhesion molecule-1 (ICAM-1) in many cancer cells (Hauck et al., 2001; Schlaepfer et al., 2004; Taglia et al., 2007). Additionally, FAK has been implicated in the regulation of tumor angiogenesis through promoting vascular endothelial growth factor (VEGF) expression and activity (Peng et al., 2004). Therefore, suppression of FAK has been one of therapeutic interest strategies for treatment or inhibition of cancer metastasis.

Bioactive phytochemical compounds isolated from natural products have been focused as a potential option for cancer treatment (Pitchakarn et al., 2013; Zakaria et al., 2019). Cucurbitacin B is a tetracyclic triterpenoid derived from edible cucurbitaceous plants (Patlolla and Rao, 2012). Our previous studies demonstrated that this compound exerted antiproliferative and apoptosis-inducing effects on KKU-100 CCA cells via suppression of FAK-mediated signaling pathways (Klungsaeng et al., 2019; Klungsaeang et al., 2020). In this study, we aimed to investigate the effect of cucurbitacin B on metastatic behavior of CCA cells through modulation of FAK protein. KKU-452 CCA cell line, a newly established CCA cells derived from CCA patients with liver fluke infection with remarkably high migration and invasion properties (Saensa-Ard et al., 2017), was employed in the study.

Materials and Methods

Materials
Reagents for cell culture were purchased from Gibco-BRL Life Technologies (Grand Island, New York). Selective FAK inhibitor (FAK inhibitor-14), cucurbitacin B and sulforhodamine B were obtained from Sigma Chemical (St. Louis, MO). The primary antibodies against phospho-FAK was purchased from Cell Signaling Technology (Dancer, MA, USA). The primary antibodies against FAK, MMP-9, ICAM-1, VEGF and β-actin and the secondary horseradish peroxidase (HRP)-linked antibodies were obtained from Santa Cruz Biotechnology, Inc. (California, USA).

Cell line and cell culture
A human CCA cell line, KKU-452, was kindly provided by the Cholangiocarcinoma Research Institute, Khon Kaen University. The cells were grown in Ham’s F12 media containing 10 mM HEPES (pH 7.3), 10% fetal calf serum and antibiotics (100 U/mL penicillin and 100 μg/mL gentamicin). Cells were grown in a humidified atmosphere of 5% CO₂ at 37°C and subcultured using 0.25% trypsin-EDTA every 2 days.

Cytotoxicity assay
The effects of test compounds on the viability of KKU-452 CCA cells were assessed using sulforhodamine B colorimetric assay. Briefly, the cells were exposed to various concentrations of test compounds for 24 h. Thereafter, the cells were fixed with 10% trichloroacetic acid for 1 h at 4°C and subsequently stained with 0.4% sulforhodamine B in 1% acetic acid for 30 min. Unbound dye was discarded by rinsing with 1% acetic acid, and the protein-bound dye was then dissolved in 10 mM Tris solution. The color intensity was quantified at 540 nm using a microplate spectrophotometer.

Western blot analysis
KKU-452 cells were treated with different concentrations of test compounds for indicated time points. After treatment, whole cell lysates were prepared using ice-cold RIPA lysis buffer (Thermo Scientific, USA) following the manufacturer’s instructions. The 30 μg of proteins were then subjected to 10% SDS-PAGE and electrophoretically transferred onto a PVDF membrane. Non-specific sites on the membranes were blocked with 5% bovine serum albumin in TBS buffer containing 0.1% Tween-20 for 1 h. Thereafter, the membranes were incubated with specific primary antibodies at 4°C for an overnight. After washing, the membranes were subsequently incubated with HRP-conjugated secondary antibody at room temperature for 2 h. The target proteins were detected using chemiluminescence detection system (LuminataTM Forte Western HRP Substrate, Merek Millipore, Watford, UK) and visualized by ChemiDocTM MP Imaging system.

Wound healing assay
The migratory ability of KKU-452 cells was evaluated using wound healing assay. Briefly, the cells were cultured in a 24-well plate to achieve 90-100% confluency. A linear wound then was made using a sterile 200 μL pipette tip. The unattached cells were removed by washing with PBS. The cells were subsequently treated with various concentrations of test compounds and the wounds were photographed at 0 and 24 h of treatment using a phase-contrast microscope. The migration ability was assessed from the closing of the scratched wound which was determined using Image-Pro Plus software (Media Cybernetics, LP, USA).
Cucurbitacin B Suppresses Cholangiocarcinoma Cells Metastasis

Determination of MMP-9 level

The KKU-452 cells were treated with various concentrations of test compounds for 24 h. Thereafter, the cultured cells were trypsinized, washed with PBS and adjusted to 1 x 10^6 cells for each reaction. The cells were then completely lysed using an ultrasonic cell disruptor and centrifuged to pellet the cellular debris. The supernatants were collected to quantify the amount of MMP-9 using Human MMP-9 ELISA Kit (Elabscience, Houston, Texas) according to the manufacturer’s instructions.

Statistical analysis

The data were analyzed using One-way analysis of variance with post hoc Student–Newman–Keuls (SNK) test. Results were considered to be statistically significant with a P value less than 0.05. The statistical analyses were employed using SigmaStat v.3.5 Program (Systat Software Inc., San Jose, California).

Results

The effects of FAK inhibitor and cucurbitacin B on CCA cell viability

The effects of FAK inhibitor-14 and cucurbitacin B on KKU-452 CCA cell viability were determined by the sulforhodamine B assay. The cells were exposed to various concentrations of the test compounds for 24 h. The results
showed that IC50 values of cytotoxicity effects on KKU-452 cells of FAK inhibitor-14 and cucurbitacin B were 3.64 ± 0.09 and 0.66 ± 0.02 μM, respectively (Figures 1A & 1B). Low concentrations of these compounds which negligibility or minimally affected the cell viability were used for further studies.

Cucurbitacin B suppressed FAK activation similar to FAK inhibitor

Given that FAK acts as a regulator molecule for transmitting signals to activate downstream targets associated with metastasis processes (Tai et al., 2015). We firstly evaluated the effects of FAK inhibitor and cucurbitacin B on FAK activation in KKU-452 cells. The results showed that FAK inhibitor-14 suppressed the phosphorylation of FAK, although FAK protein level was not significantly affected (Figure 1C). Similarly, cucurbitacin B could significantly inhibit FAK activation, evidently by decreased level of phospho-FAK protein after exposure to the compound. Moreover, cucurbitacin B also suppressed the expression of FAK protein (Figure 1D).

Cucurbitacin B diminished metastatic behavior of CCA cells similar to FAK inhibitor

It has been reported that FAK is a key regulator of metastasis processes in several cancer cells. In this study, we investigated whether FAK participates in regulation of metastatic behavior of KKU-452 CCA cells. The effects of FAK inhibitor-14 on migratory, invasive and adhesion abilities of KKU-452 cells were then examined. The results clearly demonstrated that FAK inhibitor-14 significantly suppressed migration, invasion as well as adhesion of CCA cells in a concentration-dependent manner (Figure 2). Similar to FAK inhibitor, cucurbitacin B could exert anti-migratory, -invasive and -adhesive activities against CCA cells in a dose-dependent manner (Figure 2).

Cucurbitacin B suppressed MMP-9 similar to FAK inhibitor

MMP-9 is a main proteolytic enzymes involving in degradation of basement membranes. We subsequently determined the effect of FAK inhibitor-14 on MMP-9
Cucurbitacin B Suppresses Cholangiocarcinoma Cells Metastasis

Expression in KKU-452 cells. Additionally, quantitative determination of MMP-9 concentration was also performed. The results showed that FAK inhibitor-14 significantly suppressed MMP-9 expression in a dose-dependent manner, which led to a decrease of MMP-9 level in CCA cells (Figure 3A). Likewise, exposure to cucurbitacin B also inhibited the expression of MMP-9 protein. Consequently, the amount of MMP-9 was substantially decreased by cucurbitacin B treatment (Figure 3B).

Cucurbitacin B downregulated metastasis-associated molecules similar to FAK inhibitor

Apart from MMP-9, the effect of FAK inhibitor-14 on the expression of other metastasis-associated proteins, ICAM-1 and VEGF, was also determined. It was found that the expression of these two proteins was downregulated by the inhibitor treatment (Figure 3A). Similar to FAK inhibitor, cucurbitacin B significantly suppressed the expression of ICAM-1 and VEGF in a dose-dependent manner (Figure 3B).

Discussion

The signaling pathways regulating metastasis process and the metastatic-associated proteins have been targeted for metastasis inhibition. For decades, phytochemical from fruits and vegetables have gained increased attention for development as anti-metastatic agents (Pitchakarn et al., 2013; Tuponchai et al., 2019). In the present study, it was demonstrated that cucurbitacin B suppressed FAK activation in KKU-452 CCA cells which was associated with inhibition of metastasis essential steps and their related metastatic proteins. Thus, this phytochemical compound may be a prospective candidate for the treatment of CCA metastasis.

Given the critical role of FAK in metastatic cascade, suppression of FAK activation could be a potential therapeutic strategy for prevention or inhibition of cancer metastasis (Lee et al., 2015; Tai et al., 2015). In CCA, enhanced FAK expression has been detected which was well correlated with cancer aggressiveness and poor clinical outcome of patients (Khansaard et al., 2014). Herein, the role of FAK contributing in CCA metastasis was confirmed using a specific FAK inhibitor, FAK inhibitor-14. Exposure of the inhibitor to high metastatic potential CCA cells, KKU-452, resulted in suppression of metastatic behavior of the cells. Moreover, FAK inhibitor also diminished the expression of metastasis- and angiogenesis-associated proteins including MMP-9, ICAM-1 and VEGF. These data highlighted that FAK protein positively regulates CCA metastasis.

Several natural active ingredients exerted their anti-metastasis activities against cancer cells via inactivation of FAK activity (Sun et al., 2015; Masraksa et al., 2020). Previous studies have demonstrated that the anti-metastasis effect of cucurbitacin B against breast cancer cells was mediated through FAK suppression (Luo et al., 2018; Liang et al., 2019). In the present study, the results showed that cucurbitacin B inhibited FAK activation in KKU-452
cells by alleviating the expression of both phospho-FAK and FAK proteins. The data suggested that cucurbitacin B might decrease expression or decrease phosphorylation of FAK in KKU-452 cells, which subsequently led to FAK inhibition. It is noted that cucurbitacin B had a more potent inhibitory effect on FAK protein than that of the specific FAK inhibitor-14. FAK activation has an impact on the metastatic behavior of cancer cells (Schlaepfer et al., 2004). As expected, after suppression of FAK in CCA cells by cucurbitacin B, the essential events of cancer metastasis, including migration, invasion and adhesion, were consequently inhibited. Furthermore, activation of FAK is an important phenomenon for conducting intracellular signals to various metastasis-related molecules (Tai et al., 2015). It was found that suppression of FAK in CCA cells by cucurbitacin B was accompanied with down-regulation of MMP-9 which subsequently resulted in decrease MMP-9 level. Additionally, ICAM-1 and VEGF were also down-regulated after cucurbitacin B exposure. Collectively, these findings indicated that anti-metastasis potential of cucurbitacin B against CCA cells was mediated, at least in part, through inhibition of FAK protein. Besides FAK, it has been reported that cucurbitacin B also exerted its anti-metastasis effect via modulating other signaling pathways including Hippo-YAP signaling pathway in colorectal cancer cells (Chai et al., 2018) and JAK2/STAT3 and MAPK pathways in osteosarcoma cells (Zhang et al., 2017). It is therefore likely that other intracellular signaling cascades might also be participating in the anti-metastasis activity of cucurbitacin B on CCA cells, which needs further investigation study.

In conclusion, cucurbitacin B possesses anti-metastasis property against CCA cells by blocking of FAK activation, suppressing metastatic phenotypes and down-regulating metastasis-associated molecules. These findings support the potential of cucurbitacin B to be developed as a novel therapeutic agent for CCA metastasis therapy.

Acknowledgements

This research was supported by the grant from Khon Kaen University, Thailand (6200011003), the Invitation Research grant from Faculty of Medicine, Khon Kaen University, Thailand (IN63252), and partially funded by a grant from the Office of the Higher Education Commission of Thailand. Putthaporn Kaewmeesri was supported by a Commission on Higher Education Ph.D. Scholarship. The authors would like to thank Mr. Gurdeep Singh for editing the manuscript via the English Editing Publication Clinic, Faculty of Medicine, Khon Kaen University, Thailand.

References


