# 國立嘉義大學生命科學院

學生學術研究成果優良海報評選獲獎名單

時間:103年6月4日

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# Role of down-regulation of ERCC1 expression in metformin enhancing paclitaxel-induced cytotoxicity in human lung cancer cells

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### ABSTRACT

Metformin, a widespread used and well-tolerated biguanide class drug for treating type 2 diabetic patients, has recently discovered therapeutic effects on diversified human cancer cells. Paclitaxel (taxol), as a mitotic inhibitor isolated from the bark of the Pacific Yew tree and Taxus brevifolia, is an universal antitumor drug that can use in treatment of non-small cell lung cancer (NSCLC). Excision repair cross-complementary 1 (ERCC1) is a DNA excision repair protein. Whereas, overexpression levels of ERCC1 in cancer cells have been positively associated with the DNA repair capacity and repress cancer cells death to cause a poor prognosis in humans NSCLC treated with platinum-based chemotherapy. According to my preceding research, when paclitaxel used in human NSCLC cell lines H1703 cell, increased phosphorylation of mitogen-activated protein kinase (MAPK) kinase 3/6 (MKK3/6)-p38 MAPK as well as protein and mRNA levels of ERCC1. Moreover, down-regulation of p38 MAPK activation by either the pharmacological inhibitor SB202190 or p38 MAPK-siRNA was capable of enhancing NSCLC sensitivity to paclitaxel. In this research, we presumed that metformin raise the paclitaxel-induced cytotoxic effect by decreasing ERCC1 expression and MKK3/6-p38 MAPK activation. Together, our experiments demonstrated that down-regulation of ERCC1 expression and paclitaxel co-treated with metformin may provide varied lung cancer cells therapeutic effects in clinically useful combination.





ERCC1



# Regulatory effects of natural products on cell growth, melanin content and metastasis in melanoma cells

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#### Introduction

Malignant melanoma can resistant to radiation therapy and chemotherapy, and has characteristics like high rate of metastasis and rapid cell growth. It makes the clinical therapy of malignant melanoma with the limited success. In previous experiments, we found that B16 melanoma cells with a-melanocyte-stimulating hormone (a-MSH) stimulation showed less sensitive to STI-571 (imatinib)-mediated cytotoxicity.

In melanoma cells, there is no definitive evidence that metastasis and drug resistance are correlation with melanin content. Therefore, the aim of this study assayed the correlation of natural products, such as resveratrol and arbutin, between melanin content and metastatic potency on B16 melanoma cells. We use a-MSH to promote melanin biosynthesis, for researching the effects of melanin biosynthesis on cell growth, drug resistance and metastasis in B16 melanoma cells, and analyzing the effects on melanin-related pathways, such as c-kit and Wnt/\beta-catenin pathways, as well as protein expression, including microphthalmia-associated (MITF) transcription factor and matrix metalloproteinases (MMPs).

We also use recombinant human stem cell factor (rhSCF), which can bind c-kit receptor on cell membrane, to activate c-kit signaling pathways, to further investigate the cell viability, melanin content, ckit signaling pathway and MITF protein expression on B16 cells with the treatment of STI-571, resveratrol and arbutin.





Fig 1. Effects of a-MSH-combined treatment with STI-571, resveratrol and arbutin on cell viability in B16 melanoma cells.

#### **Melanin content:**



Fig 2. Effects of a-MSH-combined treatment with STI-571, resveratrol and arbutin on melanin content in B16 melanoma cells. β-catenin expression:

# **β**-catenin GAPDH **B**-cat 6-tubulin

Fig 3. Expression of cytoplasmic and nuclear β-catenin in B16 melanoma cells by α-MSH-combined treatment with

#### STI-571, resveratrol and arbutin. **MITF expression:**





Fig 4. Expression of cytoplasmic and nuclear MITF in B16 melanoma cells by a-MSH-combined treatment with STI-571, resveratrol and arbutin.

#### **MMP-9** expression:



Fig 5. Expression of MMP-9 in B16 melanoma cells in a-MSH-combined treatment with STI-571, resveratrol and arbutin.

· ASCT

Stem cell factor (SCF) effect"

on viability of B16 cells:

Fig 6. Effects of rhSCF-combined treatment with STI-571. resveratrol and arbutin on cell viability in B16 melanoma cells.

SCF effect on melanin -HSCT content:

Fig 7. Effects of rhSCF-combined treatment with STI-571, resveratrol and arbutin on melanin content in B16 melanoma cells.



Fig 8. Expression of phospho-c-kit in B16 melanoma cells by rhSCF-combined treatment with STI-571, resveratrol and arbutin. SCF effect on MITF expression: + rbSCF





melanoma cells by rhSCF-combined treatment with STI-571, resveratrol and arbutin.

#### Conclusions

- > STI-571, resveratrol, and arbutin inhibited cell viability and melanin content, but didn't have significant effect on MMP-9 expression in B16 melanoma cells.
- STI-571 has inhibitory effects on nuclear β-catenin/MITF protein expression of B16 melanoma cells, suggesting that STI-571 may downregulate β-catenin-related pathways and inhibit downstream gene regulation, such as MITF.
- > Resveratrol and arbutin have inhibitory effects on phospho-c-kit protein level in rhSCF-treated B16 melanoma cells, but have no significant effect on MITF protein level. It could be hypothesized that resveratrol and arbutin could downregulate rhSCF-activated c-kit signal pathways, but such effect may not regulate the expression of MITE.

## 探討菌桿孢芽粉澱化液對水稻紋病枯的防治潛力

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為,一之害病要主最稻水灣臺為病枯紋稻水探討菌桿孢芽粉澱化液 稻水治防為作紋枯之劑製物生病潛力,本研究針對自臺灣本土分離所 得之液化澱粉芽孢桿菌 Bacillus amyloliquefaciens BACY1 及 A1 菌株 ,测試其對水稻紋枯病菌的生長抑制作用。利用對峙培養測試其對紋 枯病菌Rhizoctonia solani之菌絲生長抑制效果,結果發現使用500 ml 脂胜肽萃取液皆有很明顯的抑制效果。另外利用掃描式電子顯微鏡觀 察發現經過脂胜肽萃取液處理過的紋枯病菌,其菌絲或菌核皆呈現嚴 重蜷曲萎縮狀,進而使得紋枯病菌無法正常生長。而液化澱粉芽孢桿 菌是否能夠在田間施用對水稻紋枯病菌也有一樣的抑制效果,仍需進 一步的研究試驗。

### 液化澱粉芽孢桿菌脂胜肽之萃取

將液化澱粉芽泡桿菌保存管以接菌環沽取菌液於LB (Luria-Bertani Broth) 瓊脂培養基劃取單一菌落,於 28℃ 培養24小時。再挑出單一菌 落至100 ml LB液態培養基,於150 rpm,28℃ 下培養至菌液濃度A<sub>600</sub>達 到0.4時,取500 ml菌液至Best培養基 (pH 7.0,1% tryptone 和 2.5% glucose) 於150 rpm,30℃ 震盪培養48小時,將培養菌液以12,000 rpm, 4℃ 離心20分鐘,取出上清液以10 N HCI 至整調 pH 2.0液清上使,pH 於置,出析而點電等之肽胜脂於低值4℃下30以,後鐘分12,000 rpm, 4℃ 心離20再,液清上除移,部底管心離至心離肽胜脂之出析將,鐘分 以3 ml 50% 酒精以並,溶回 1 N NaOH至整調pH 7.0以後最,0.22 mm filter (Millex-GV Filter Unit, Millipore) 過濾。



圖一、Bacillus amyloliquefaciens A1及 BACY1 菌株脂胜肽 (處理量為 80 ml) 對紋枯病菌生長之抑制情形。



圖二、Bacillus amyloliquefaciens A1及 BACY1 菌株脂胜肽 (處理量為 500 ml) 對紋枯病菌生長之抑制情形。



圖三、Bacillus amyloliquefaciens A1及 BACY1 菌株脂胜肽 (處理量為 500 ml) 對不同紋枯病菌菌株生長之抑制情形 (A) 紋枯病菌RS-R2; (B) 紋枯病菌RS-R3; (C) 紋枯病菌RS-R4; (D) 紋枯病菌RS-R5; (E) 紋枯病菌RS-R6; (F) 紋枯病菌RS-R7。







圖五、利用掃描式電子顯微鏡觀察液 化澱粉芽孢桿菌脂胜肽對紋枯病菌絲 之影響(A)未處理;(B)脂胜肽萃取液 處理;(C)菌液處理。 圖六、利用掃描式電子顯微鏡觀察 液化澱粉芽孢桿菌脂胜肽對紋枯病 菌核之影響 (A) 未處理;(B) 脂胜 肽萃取液處理;(C) 菌液處理。

結語

本研究結果顯示液化澱粉芽孢桿菌BACY1 及 A1 脂胜肽對紋枯病菌株皆 有明顯的抑制效果,其抑制效果推論是藉由使菌絲嚴重蜷曲萎縮,破壞內 部構造,進而使得無法正常生長。而液化澱粉芽孢桿菌是否能夠在土壤中 對水稻紋枯病菌也有一樣的抑制效果,仍需進一步的研究試驗,若能成功 有抑制效果,則可供日後開發水稻紋枯病菌病害防治用生物製劑的參考依 據。

## Study on removal of plant browning products and purification of a toxic protein with antineoplastic activity from *Bidens pilosa* var. *radiata*



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### Abstract

Based on our previous study, extracts of *Bidens pilosa* var. *radiata* contain anti-proliferation activity to colorectal cancer cells. Due to a lot of browning products interfering subsequent experimental analysis and purification processes, it is necessary to overcame this issue by adding various chemicals to remove browning products. The cytotoxicity of different preparation of purified proteins from *Bidens pilosa* var. *radiate* is monitored with the SRB (Sulforhodamine B) assay. The protein extract was precipitated with 35%~65% saturated ammonium sulfate, and decolored with polyvinyl- polypyrrolidone (PVPP), aluminum sulfate, polyethyleneimine (PEI), and polyethylene glycol (PEG) 4,000 plus ammonium sulfate aqueous two phase system, respectively. PEI shows the best de-colored effect on protein sample solution. The PEI is subsequent removed by a DEAE anion exchanger. The purified protein fraction still show high cytotoxicity. Therefore, dissolved ammonium sulfate precipitate solution is de-colored by PEI, and purified with DEAE Sepharose<sup>TM</sup> and Q Sepharose<sup>TM</sup> anion-exchanger for further characterization of toxin, named "bidenin".

Bidenin is separated and detected by the native or SDS polyacrylamide gel electrophoresis with the gel slicing technique. The estimated molecular weight of bidenin is identified by SDS-PAGE is about 88 kDa. The identification of bidenin will be completed by mass spectrometry in the coming future.



|       | Remove of Brow      | ning Products |           |
|-------|---------------------|---------------|-----------|
| PVPP) | Aluminum<br>sulfate | PEI           | PEG 4,000 |
|       |                     | 7             |           |
|       | Purification of 7   | Toxic Protein |           |
|       |                     | 0 51 1        |           |

Anion exchanger chromatography (DEAE & Q) Electrophoresis (Native & SDS)

### Conclusion

The result of remove for browning products, PEI show the best de-colored effect on protein sample solution and clearly analysis band on gel.

Toxic protein is successfully purified by DEAE Sepharose<sup>TM</sup> and Q Sepharose<sup>TM</sup> anion-exchanger then separated and located by Native and SDS PAGE with gel slicing technique remained its cytotoxicity, named "bidenin". The molecular weight of bidenin showed about 88 kDa on SDS-PAGE.

#### Rosults



Fig.1. Analysis of the protein extracts precipitated with different range of ammonium sulfate on the cell viability of human colon adenocarcinoma cell (DLD-1).



Fig.2. Effect of different agents on the color of ammonium sulfate precipitate dissolved protein solution.



8% Native-PAGE 8% SDS-PAGE Fig.4. Native-PAGE (A) and SDS-PAGE (B) analysis of ammonium sulfate precipitate from each treating agents.



Fig.5.DEAE-Sepharose<sup>™</sup> anion exchange chromatography of decolored protein solution.



Fig.3. Use different agents as adsorbent to treating ammonium sulfate precipitate dissolved protein solution that effect on the cell viability of human colon adenocarcinoma cell (DLD-1).



Fig.6.Q-Sepharose<sup>TM</sup> anion exchange chromatography of DEAE separated protein sample (D6).



8% Native-PAGE Fig.7.Native-PAGE (A) and SDS-PAGE (B) analysis of Q separated protein solution (Q4) purified by Native-PAGE extraction with gel slicing. Q4 : Q separated protein sample, N1–N6 : gel extracted protein solutions, M : L.M.W. protein marker.



**Fig.8.** Native-PAGE (A) and SDS-PAGE (B) analysis Q separated protein solution (Q4) purified by SDS-PAGE extraction with gel slicing. D6 : DEAE separated protein sample, Q4 : Q separated protein sample, S1-S6 : gel extracted sample solutions, M : L.M.W. protein marker, B : bovine serum albumin. (A) (B)



Fig.9. Native-PAGE and SDS-PAGE analysis of purification samples from each purification steps. O ammonium sulfate precipitated protein solution : lane 1 : PEI decolorized sample solution : lane 2 : DEAE purified toxic sample : lane 3 : Q purified toxic sample : lane 4 : Native-PAGE gel slicing toxic protein sample (N2) : lane 5 : SDS-PAGE gel slicing toxic protein sample (S2) : M : L.M.W. protein marker.



# Role of p38 MAPK signal mediated MutS homologue-2 (MSH2) expression in regulating gefitinib and metformininduced cytotoxicity in human lung squamous cancer cells

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### ABSTRACT

Gefitinib, a quinazoline-derived tyrosine kinase inhibitor, has anti-tumor activity *in vivo* and in *vitro*. Human MutS homologue-2 (MSH2) plays a central role in promoting genetic stability by correcting DNA replication errors. The present study investigated the effects of p38 mitogen-activated protein kinase (MAPK) signal on gefitinib-induced MSH2 expression in human non-small cell lung squamous cancer cell lines. Exposure of gefitinib increased MSH2 protein and mRNA levels, which was accompanied by MKK3/6-p38 MAPK activation in H520 cells. Moreover, knockdown of p38 expression by SB202190 and specific small interfering RNA (siRNA) significantly decreased gefitinib-induced MSH2 expression by increasing mRNA and protein instability. Enhancing p38 activation using constitutively active MKK6 (MKK6E) increased MSH2 protein and mRNA levels. Specific inhibition of MSH2 by siRNA significantly enhanced gefitinib-induced cytotoxicity. Metformin, an anti-diabetic drug, might reduce cancer risk. In human lung squamous cancer cells, metformin decreased gefitinib-induced MSH2 expression and augmented the cytotoxic effect and growth inhibition by gefitinib. Transient expression of MKK6E or HA-p38 MAPK vector could abrogate metformin and gefitinib-induced synergistic cytotoxic effect in H520 cells. Together, metformin can down-regulate p38 MAPK-mediated MSH2 expression and enhance the cytotoxicity of gefitinib to human squamous lung cancer cells.





## Investigation of the molecular mechanism of tumorigenesis in 5-Fu- and Oxaliplatin-resistant human colorectal cancer stem cells

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### INTRODUCTION

Colorectal cancer in Taiwan is already the highest cancer incidence rate, and third cancer mortality rate. Chemotherapy treatments have made great contributions to the inhibition of cancer recurrence, and metastasis. But, Chemotherapy is often resistant in about half of the patients, resulting in cancer recurrence and progression. A lot of evidence are finding to suggest that cancer stem cells will lead chemoresistance and cause tumor sustained growth. However, colorectal cancer stem cells(CSCs) for drug resistance mechanisms remain unclear.

### MOTIVE

In this research, we use the mainstay chemotherapeutic drug, 5-Fluorouracil (5-FU) and Oaxliplatin (OxR) were selected with the colon cancer cell line HCT-116. The cancer stem cell lines were screened for colorectal cancer by the use of chemical resistant cloning method to analysis of whether a cancer stem cell characteristics. Study of the effect of SDF-1 on cancer stem cells signaling and uPA expression and the regulation of the Bcl-2/Bax and P-gp proteins may be related to resistant chemotherapy drug resistance mechanism.

Study Design

into the molecular mechanism and suggest promising biomarkers for

clinic therapy of colorectal cancer patients outcome of patients with

lymph node metastasis.

REFERENCE





SDF-1 Bcl-2 Fig.6 ComparedSDF-1 and Bcl-2 expressed of the CRC and CSC cells in treatment of all kind inhibitors.

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Fig. 5 Compared drug resistance differentially expressed of the CRC and CSC cells

0.5

in vitro/in vivo