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ต่อเซลล์มะเร็งลำไส้ใหญ่ของมนุษย์

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CYTOTOXIC ACTIVITY AND MECHANISM OF GLYCOSMIS PARVA  
LEAF EXTRACT ON HUMAN COLORECTAL CANCER CELLS

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จุฬาลงกรณ์มหาวิทยาลัย

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Field of Study	Pharmacology
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ณัฐพร บุรณะบุญวงศ์ : การศึกษาความเป็นพิษและกลไกการออกฤทธิ์ของสิ่งสกัดจากใบส้มขึ้นต่อเซลล์มะเร็งลำไส้ใหญ่ของมนุษย์. (CYTOTOXIC ACTIVITY AND MECHANISM OF GLYCOSMIS PARVA LEAF EXTRACT ON HUMAN COLORECTAL CANCER CELLS) อ.ที่ปริกษาวิทยานิพนธ์หลัก: ผศ. ดร. วชิร ลิมนสิทธิกุล, อ.ที่ปริกษาวิทยานิพนธ์ร่วม: รศ. ดร. นิจศิริ เรืองรังษี, 127 หน้า.

ต้นส้มขึ้นเคยมีรายงานว่ามีความเป็นพิษและมีฤทธิ์ต้านการอักเสบโดยไปลดการแสดงออกของยีนคืออกซ์ 2 ในหลอดทดลอง การศึกษานี้มีวัตถุประสงค์เพื่อศึกษาผลของสิ่งสกัดเอทิลอะซีเตทจากใบส้มขึ้น (จีพีอี) ต่อเซลล์มะเร็งลำไส้ใหญ่ของมนุษย์ที่มีการแสดงออกของยีนคืออกซ์ 2 (เซลล์เอชที 29) และไม่มีการแสดงออกของยีนคืออกซ์ 2 (เซลล์โคโล 205) รวมไปถึงกลไกการออกฤทธิ์ ความเป็นพิษของสิ่งสกัดจีพีอีต่อเซลล์เอชที 29 และเซลล์โคโล 205 แปรผันตามความเข้มข้นและระยะเวลาที่ใช้ทดสอบ โดยมีค่าความเข้มข้นที่ทำให้เซลล์ตายร้อยละ 50 ตาย เท่ากับ  $69.49 \pm 2.04$ ,  $55.89 \pm 1.86$  และ  $48.94 \pm 2.99$  ไมโครกรัม/มิลลิลิตร หลังจากได้รับสารนาน 24, 48 และ 72 ชั่วโมง ตามลำดับในเซลล์เอชที 29 ส่วนในเซลล์โคโล 205 ค่าความเข้มข้นที่ทำให้เซลล์ตายร้อยละ 50 ตาย เท่ากับ  $59.92 \pm 4.34$ ,  $28.85 \pm 1.44$  และ  $25.42 \pm 1.65$  ไมโครกรัม/มิลลิลิตร หลังจากได้รับสารนาน 24, 48 และ 72 ชั่วโมง ตามลำดับ สิ่งสกัดจีพีอีสามารถเหนี่ยวนำให้เซลล์เกิดการตายแบบเอพอพโตซิสในเซลล์ทั้งสองชนิด โดยดูได้จากการย้อมด้วย Annexin V-FITC/Propidium Iodide ผลการศึกษาความเป็นพิษและการเหนี่ยวนำให้เซลล์เกิดการตายแบบเอพอพโตซิสนั้น พบว่าสิ่งสกัดจีพีอีมีผลต่อเซลล์โคโล 205 มากกว่าเซลล์เอชที 29 สิ่งสกัดจีพีอีที่ความเข้มข้น 25-100 ไมโครกรัม/มิลลิลิตรมีผลยับยั้งการแบ่งตัวของเซลล์มะเร็งทั้งสองชนิด ซึ่งผลการยับยั้งการแบ่งตัวของสิ่งสกัดจีพีอีนี้มีผลต่อเซลล์เอชที 29 มากกว่าเซลล์โคโล 205 การศึกษาถึงผลของสิ่งสกัดจีพีอีต่อวัฏจักรเซลล์ พบว่าทำให้เกิดการลดลงของเซลล์ในระยะ S และเพิ่มการสะสมของเซลล์ในระยะ G0/G1 และ G2/M โดยมีผลคล้ายกันทั้งสองชนิด การศึกษาในระดับกลไกของสิ่งสกัดจีพีอีพบว่าสิ่งสกัดน่าจะมกลไกการออกฤทธิ์ทั้งที่ผ่านทางคืออกซ์ 2 และไม่ผ่านคืออกซ์ 2 กลไกที่ผ่านทางคืออกซ์ 2 เกี่ยวข้องกับการลดการแสดงออกของยีนคืออกซ์ 2 ซึ่งจะไม่มีผลต่อการส่งสัญญาณภายในเซลล์ที่อยู่ระดับถัดจากคืออกซ์ 2 ลงไป ส่วนกลไกที่ไม่ผ่านคืออกซ์ 2 นั้นเกี่ยวข้องกับการเปลี่ยนแปลงการแสดงออกของยีนที่ควบคุมวัฏจักรเซลล์ และความสมดุลของยีนในกลุ่ม Bcl-2 ในเซลล์เอชที 29 พบว่าสิ่งสกัดจีพีอีมีผลลดการแสดงออกของยีน cyclin A, cyclin E, Bcl-2 และ COX-2 และเพิ่มการแสดงออกของยีน Bak ส่วนในเซลล์โคโล 205 สิ่งสกัดจีพีอีมีผลลดการแสดงออกของยีน cyclin A, Bcl-2 และ Bcl-XL และเพิ่มการแสดงออกของยีน p21 โดยสรุปสิ่งสกัดจีพีอีมีความเป็นพิษ, เหนี่ยวนำให้เกิดการตายแบบเอพอพโตซิส, ยับยั้งเพิ่มจำนวน, และหยุดวัฏจักรเซลล์ ในเซลล์มะเร็งลำไส้ใหญ่ของมนุษย์ทั้งสองชนิด ซึ่งกลไกการออกฤทธิ์ผ่านทางคืออกซ์ 2 และไม่ผ่านคืออกซ์ 2 การศึกษาในครั้งนี้จึงสามารถเป็นพื้นฐานเพื่อที่จะพัฒนาายาต้านมะเร็งจากสิ่งสกัดจีพีอีซึ่งมีสารออกฤทธิ์ที่มีความเป็นไปได้ในการวิจัยต้านมะเร็งลำไส้ใหญ่ต่อไป

สาขาวิชา เกษษวิชา

ปีการศึกษา 2556

ลายมือชื่อนิสิต .....

ลายมือชื่อ อ.ที่ปริกษาวิทยานิพนธ์หลัก .....

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KEYWORDS: GLYCOSMIS PARVA / COLORECTAL CANCER / CYTOTOXICITY / APOPTOSIS / CELL CYCLE ARREST / COX-2 / BCL-2 / CYCLINS / HT-29 / COLO-205

NATTAPORN BURANABUNWONG: CYTOTOXIC ACTIVITY AND MECHANISM OF GLYCOSMIS PARVA LEAF EXTRACT ON HUMAN COLORECTAL CANCER CELLS. ADVISOR: ASST. PROF. WACHAREE LIMPANASITHIKUL, Ph.D., ASSOC. PROF. NIJSIRI RUANGRUNGSI, Ph.D., 127 pp.

Glycosmis parva Craib (Rutaceae) was reported to have cytotoxicity and anti-inflammatory activities with the reduction of COX-2 expression, in vitro. This study aimed to investigate the effect of ethyl acetate extract from leaves of *G. parva* (GPE) on human colorectal cancer expressing COX-2, HT-29, and not express COX-2, Colo-205 and its underlying mechanisms of action. Cytotoxicity of GPE against HT-29 and Colo-205 were exhibited in both dose- and time- dependent manner. The IC<sub>50</sub> values of GPE against HT-29 were 69.49±2.04, 55.89±1.86 and 48.94±2.99 µg/ml at 24, 48 and 72 h, respectively. In Colo-205, the IC<sub>50</sub> were 59.92±4.34, 28.85±1.44 and 25.42±1.65 µg/ml at 24, 48 and 72 h, respectively. GPE significantly induced apoptosis in both cell lines as evidenced by AnnexinV/FITC and PI staining. The effect of GPE on cytotoxicity and induction of apoptosis were higher in Colo-205 than HT-29. GPE at 25-100 µg/ml significantly inhibited cell proliferation in both cells. The effect was greater in HT-29 cells. Cell cycle analysis demonstrated that GPE caused a decrease in the cells in S phase which was associated with G<sub>0</sub>/G<sub>1</sub> and G<sub>2</sub>/M accumulation. The changes in cell cycle pattern following GPE treatment were similar in both cell lines. Mechanistic studies suggested that the effects of GPE may be mediated through both COX-2 dependent and COX-2 independent pathway. The COX-2 dependent involved with reduction of COX-2 expression which affected downstream COX-2 pathway. The COX-2 independent pathway included the changes in the expression of cell cycle control genes and the alteration in the balance of Bcl-2 family gene expression. In HT-29, GPE down-regulated cyclin A and cyclin E expression and increased the expression of pro-apoptotic Bak while decreased the anti-apoptotic Bcl-2. It also significantly decreased COX-2 expression. In Colo-205, GPE decreased cyclin A and up-regulated p21 expression. It also decreased the expression of anti-apoptotic Bcl-2 and Bcl-XL. Taken together, GPE exhibited cytotoxic activity, induction of apoptosis, inhibition of cell proliferation and arrest cell cycle in both cells. The underlying mechanisms involved both COX-2 dependent and COX-2 independent pathway. These findings provide the fundamental knowledge of the anti-cancer effect of GPE which could be a potential compounds for the treatment of colorectal cancer.

Field of Study: Pharmacology

Student's Signature .....

Academic Year: 2013

Advisor's Signature .....

Co-Advisor's Signature .....

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## LIST OF ABBREVAITIONS

$\mu\text{g/ml}$	=	Microgram per milliliter
$\mu\text{l}$	=	Microliter
$\mu\text{M}$	=	Micromolar
15-PGDH	=	15-hydroxyprostaglandin dehydrogenase
5FU	=	5-fluorouracil
ADCC	=	Antibody-dependent cell-mediated cytotoxicity
ANOVA	=	Analysis of variance
AOM	=	Azoxymethane
Apaf-1	=	Apoptotic protease-activating factor-1
APC	=	Adenomatous polyposis coli
ATCC	=	American Type Culture Collection
BAD	=	BCL-2 antagonist of cell death
BAK	=	BCL-2-antagonist/killer-1
BAX	=	BCL-2-associated X protein
BCL-2	=	B-cell lymphoma-2
BID	=	BH3-interacting domain death agonist
BIK	=	BCL-2-interacting killer
BIM	=	BCL-2-like-11
BMF	=	BCL-2 modifying factor
Caspase	=	Cysteine Aspartic Acid Specific Protease
CDKI	=	Cyclin-dependent kinase inhibitor
Cdks	=	Cyclin-dependent kinases
cDNA	=	Complementary DNA
$\text{CH}_2\text{THF}$	=	5,10-methylene tetrahydrofolate
$\text{CHCl}_3$	=	Chloroform
$\text{CO}_2$	=	Carbondioxide
COX	=	Cyclooxygenases
CRC	=	Colorectal Cancer
$C_T$	=	Cycle threshold

dATP	=	Deoxyadenosine Triphosphate
DEPC	=	Diethyl pyrocarbonate
DHFU	=	Dihydrofluorouracil
DISC	=	Death inducing signalling complex
DMEM	=	Dulbecco's modified Eagle's medium
DMSO	=	Dimethylsulfoxide
DNA	=	Deoxyribonucleic acid
dNTP	=	Deoxynucleotide
DPD	=	Dihydropyrimidine dehydrogenase
dTMP	=	Deoxythymidine monophosphate
dUMP	=	Deoxyuridine monophosphate
dUTP	=	Deoxyuridine triphosphate
EC <sub>50</sub>	=	50% Effective Concentration
EDTA	=	Ethylene diamine tetraacetic acid
EGFR	=	Epidermal growth factor receptor
ER	=	Endoplasmic reticulum
ERK	=	Extracellular signal-regulated kinase
EtOAc	=	Ethyl acetate
FADD	=	Fas-associated death domain
FAP	=	Familial adenomatous polyposis
Fas	=	Fibroblast associated antigen
FBS	=	Fetal bovine serum
FDA	=	Food and drug administration
FdUMP	=	Fluorodeoxyuridine monophosphate
FdUTP	=	Fluorodeoxyuridine triphosphate
FUDP	=	Fluorouridine diphosphate
FUMP	=	Fluorouridine monophosphate
FUR	=	Fluorouridine
FUTP	=	Fluorouridine triphosphate
GPE	=	Ethyl acetate extracts from the leaves of <i>Glycosmis parva</i>

H	=	Hour
HCl	=	Hydrochloric acid
HEPES	=	4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid
HRK	=	Harakiri (death protein-5)
HSV-2	=	Herpes simplex virus type 2
IC <sub>50</sub>	=	50% Inhibition Concentration
IgG	=	Immunoglobulin G
IL	=	Interleukin
iNOS	=	Inducible nitric oxide synthase
LV	=	Leucovorin
MAPK	=	Mitogen-Activated Protein Kinase
MeOH	=	Methanol
mg/ml	=	Milligram per milliliter
MgCl <sub>2</sub>	=	Magnesium Chloride
mRNA	=	Messenger RNA
NaHCO <sub>3</sub>	=	Sodium bicarbonate
NaOH	=	Sodium hydroxide
NF- $\kappa$ B	=	Nuclear factor kappa B
nm	=	Nanometer
NSAID	=	Non-Steroidal Anti-Inflammatory Drug
°C	=	Degree centigrade
OD	=	Optical density
OPRT	=	Orotate phosphoribosyltransferase
PBS	=	Phosphate buffer saline
PG	=	Prostaglandin
pH	=	The negative logarithm of hydrogen ion concentration
PI	=	Propidium iodide
PI3K	=	Phosphoinositide 3-kinase
PKC	=	Protein kinase C
PPAR	=	Peroxisome Proliferator Activated Receptor
PRPP	=	Phosphoribosyl pyrophosphate



PUMA	=	BCL-2 binding component-3
Rb	=	Retinoblastoma protein
RNA	=	Ribonucleic acid
Rpm	=	Round per minutes
RPMI	=	Roswell Park Memorial Institute medium
RQ	=	Relative Quantitation
RR	=	Ribonucleotide reductase
RT-PCR	=	Reverse transcription polymerase chain reaction
TBE	=	Tris-Borate-EDTA buffer
tBid	=	Truncated Bid
TK	=	Thymidine kinase
TNF $\alpha$	=	Tumor necrosis factor $\alpha$
Topo I	=	Topoisomerase I
TRADD	=	TNFR1-associated death domain
TS	=	Thymidylate synthase
TXA2	=	Thromboxane A2
UDG	=	Uracil-DNA glycosylase
UK	=	Uridine kinase
UP	=	Uridine phosphorylase
VEGF	=	Vascular-endothelial growth factor
VEGFR	=	Vascular-endothelial growth factor receptor

## CHAPTER I INTRODUCTION

### Background and Rationale

Colorectal cancer (CRC) is one of the most commonly diagnosed cancers worldwide. It ranks the third in males and the second in females with over 1.2 million new cancer cases (1). In Thailand, colorectal cancer is the second most commonly diagnosed cancers in male (14.8% of all cases) and the third in females (10.4% of all cases) (2). Despite the advances in novel cancer therapeutic agents, colorectal cancer remains the leading cause of malignancy-related death with approximately 608,700 deaths estimated to have occurred in 2008 worldwide (1).

Fluoropyrimidines base, 5-fluorouracil (5-FU), remains the first line therapy in colorectal cancer for over 40 years (3). Other standard cytotoxic drugs used in treating metastatic colorectal cancer are capecitabine, irinotecan and oxaliplatin (4). Recently, new targeted therapies that have been approved to use in addition to standard chemotherapy for treating advanced colorectal cancer which are anti-vascular endothelial growth factor-bevacizumab and anti-epidermal growth factor receptor-cetuximab and panitumumab. However, these agents come with an increase in toxicity and treatment costs (5).

Over the past decade, epidemiologic studies have shown that patients taking nonsteroidal anti-inflammatory drugs (NSAIDs), such as aspirin, have approximately a 40% to 50% reduction in mortality from CRC (6). As elevated COX-2 expression was found in approximately 50% of adenomas and 85% of adenocarcinomas, it was hypothesized that NSAIDs exert some of anti-inflammatory and antitumor effects through inhibition of the COX-2 (7). Recently, cyclooxygenase-2 (COX-2) inhibitor has become a new candidate to the prevention and treatment of CRC.

The discovery of agents that regulate biological pathways related to COX-2 pathway may provide new avenues for the treatment of CRC. Major attention has been focused on identifying phytochemicals that can inhibit tumor development processes. *Glycosmis parva* CRAIB (Rutaceae) is a wild small shrub found in Thailand. Some plants in the genus *Glycosmis* have been used as Thai traditional medicines for

treatments of abscess, scabies, and snakebite. The genus *Glycosmis* is a rich source of acridone alkaloids and sulfur-containing propanamides. Many biological activities of acridone alkaloids have been reported such as antimalarial activity, anti-viral activity, inhibition of tumor cell growth, and antiproliferative effect (8). Recent works have shown that the ethyl acetate (EtOAc) extract from the leaves of *G. parva* exhibited potent cytotoxic activity in human B-cell lymphoma (9). Moreover, the hexane and EtOAc extract of *G. parva* exhibited anti-inflammatory activity with the reduction in mRNA expression of COX-2 in LPS-activated macrophage J774A.1 cells (10). Taken together, *G. parva* has interesting pharmacological properties that could be further explore in the area of chemotherapy and chemoprevention involved with COX-2 pathway. This study is, therefore, aimed to investigate the effect of ethyl acetate extract from the leaves of *Glycosmis parva* on human colorectal cancer cells and its underlying mechanism of action.

### Research Objectives

1. To examine the *in vitro* effect of ethyl acetate extract from the leaves of *Glycosmis parva* on human colorectal cancer cells.
  - 1) Cytotoxic effect
  - 2) Induction of apoptosis
  - 3) Anti-proliferative effect
  - 4) Cell cycle analysis
2. To investigate the molecular mechanism(s) of ethyl acetate extract from the leaves of *Glycosmis parva* on human colorectal cancer cells.
  - 1) Expression of cell-cycle regulatory genes.
  - 2) Expression of genes involved in intrinsic apoptotic pathway (pro-apoptotic and anti-apoptotic of BCL-2 family)
  - 3) Expression of COX-2 gene.

## Hypothesis

Ethyl acetate extract from the leaves of *Glycosmis parva* can induce apoptosis of human colorectal cancer cells by modulation of the expression of genes involved in cell-cycle regulation, induction of apoptosis and COX-2 pathway.

## Expected benefit and application

The result of anti-cancer activity of *Glycosmis parva* leaf extract on human colorectal cancer cells and its underlying mechanism involved COX-2 dependent and COX-2 independent pathway could provide the basis for further development of the compounds that target COX-2, BCL-2, and cell cycle regulation pathways which are the potential cancer therapeutic and chemoprevention targets in colorectal cancer.

## Key Words

*Glycosmis parva*, colorectal cancer, cytotoxicity, apoptosis, cell cycle arrest, COX-2, BCL-2, cyclins, HT-29, Colo-205.

## CHAPTER II

### LITERATURE REVIEWS

#### 1. Colorectal Cancer (CRC)

Colorectal cancer is the third most common cancer in men (663,000 cases, 10.0% of the total) and the second in women (570,000 cases, 9.4% of the total) worldwide. Almost 60% of the cases occur in developed regions (1; 11). However, the incidence rates are rapidly increasing in several areas historically at low risk including Eastern Asia. About 608,000 deaths from colorectal cancer are estimated worldwide, accounting for 8% of all cancer deaths, making it the fourth most common cause of death from cancer (1). In Thailand CRC is the second most commonly diagnosed cancers in male (14.8% of all cases) and third in females (10.4% of all cases) (2).

The etiology of CRC involves the complex interaction between dietary and genetic factors. Most cases (> 90%) are sporadic. Sporadic cancers arise within adenomatous polyps by the acquisition of a series of somatic genetic mutations following a stereotyped adenoma–carcinoma sequence. Carcinogenesis proceeds through the accumulation of series of genetic mutations involving several tumor-suppressor genes (*APC*, *P53*), oncogenes (*K-ras*), as well as epigenetic changes (methylation) (12). COX-2 is expressed early during these sequences, suggesting an important role in carcinogenesis as shown in Figure 1. Two major types of inherited (familial) cancer syndromes have been characterized. Familial adenomatous polyposis (FAP) (< 1% of cases) arises as a result of germ-line mutations in the *APC* gene (13). Individuals who are affected typically develop multiple colonic polyps in adolescence and, without treatment, invasive cancer in early adulthood. Hereditary non-polyposis CRC (3 – 5%) is characterized by an autosomal dominant mode of inheritance and early age of cancer diagnosis (14). Modifiable risk factors for CRC include smoking, physical inactivity, overweight and obesity, red and processed meat consumption, and excessive alcohol consumption (1).

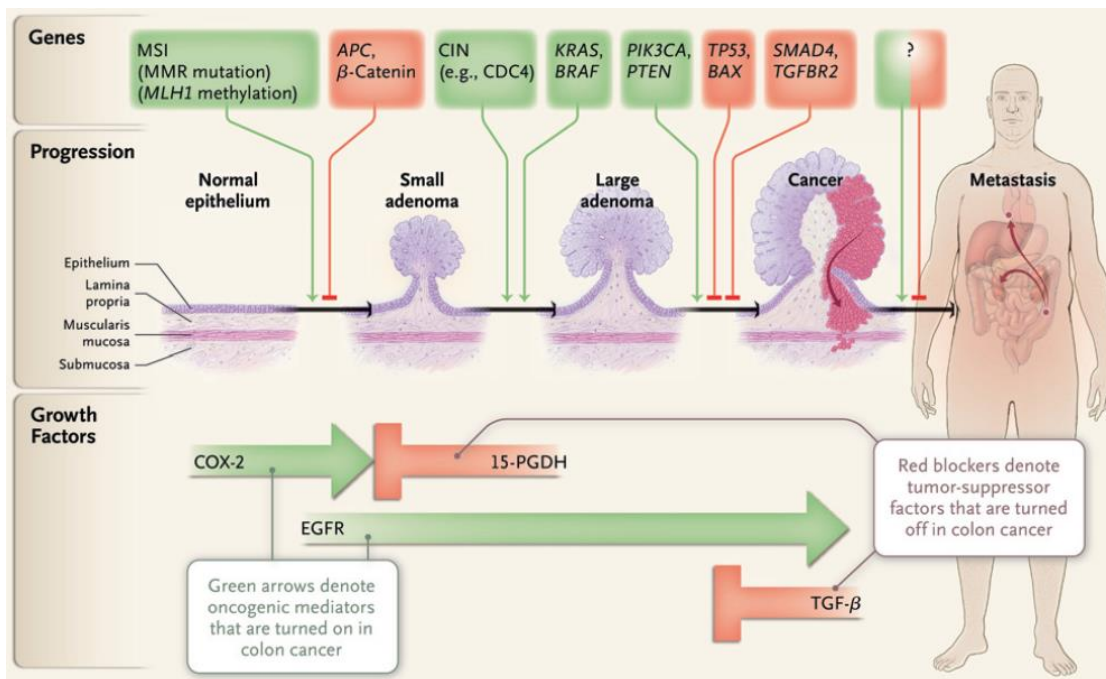


Figure 1: Genes and growth factor pathways that drive the progression of colorectal cancer (15).

The disease begins as a benign adenomatous polyp, which develops into an advanced adenoma and then progresses to an invasive cancer. Invasive cancers that are confined within the wall of the colon (stages I and II) are curable, but if untreated, they spread to regional lymph nodes (stage III) and then metastasize to distant sites (stage IV). Stage I and II tumors are curable by surgical excision, and up to 73% of cases of stage III disease are curable by surgery combined with adjuvant chemotherapy. Recent advances in chemotherapy have improved survival, but stage IV disease is usually incurable (15).

## 2. Management of colorectal cancer

Resectable colon cancer is recommended to do surgical resection with adequate lymphadenectomy. Adjuvant chemotherapy with FOLFOX, FLOX, CapeOx, 5FU/LV or capecitabine is recommended for patients with stage III disease. Adjuvant chemotherapy for patients with high-risk stage II disease is also an option with 5FU/LV with or without oxaliplatin (FOLFOX or FLOX) or capecitabine with or without oxaliplatin. Recommended initial therapy options for advanced or metastatic disease

depend on whether or not the patient is appropriate for intensive therapy. The more intensive therapy options include FOLFOX, FOLFIRI, CapeOx, and FOLFOXIRI. Addition of biological agents such as Bevacizumab, cetuximab, or panitumumab is either recommended or listed as an option in combination with some of these regimens (16).

Table 1: Chemotherapy regimen for the treatment of CRC

Name	Regimen
FOLFOX	Oxaliplatin 85 mg/m <sup>2</sup> IV over 2 hours, day 1 Leucovorin 400 mg/m <sup>2</sup> IV over 2 hours, day 1 5-FU 400 mg/m <sup>2</sup> IV bolus on day 1, then 1200 mg/m <sup>2</sup> /day x 2 days (total 2400 mg/m <sup>2</sup> over 46 – 48 hours IV continuous infusion) Repeat every 2 weeks
FLOX	5-FU 500 mg/m <sup>2</sup> IV bolus weekly x 6 + Leucovorin 400 mg/m <sup>2</sup> IV weekly x 6, each 8 week cycle x 3 with oxaliplatin 85 mg/m <sup>2</sup> IV administered on weeks 1, 3, and 5 of each 8 week cycle x 3.
CapeOx	Oxaliplatin 130 mg/m <sup>2</sup> IV over 2 hours, day 1 Capecitabine 850 – 1000 mg/m <sup>2</sup> twice daily PO for 14 days Repeat every 3 weeks
5-FU/ Leucovorin	Leucovorin 500 mg/m <sup>2</sup> IV over 2 hours and repeated weekly x 6 5-FU 500 mg/m <sup>2</sup> IV bolus given 1 hour after the start of leucovorin and repeated weekly x 6 Every 8 weeks for 4 cycles
Capecitabine	Capecitabine 1250 mg/m <sup>2</sup> twice daily PO days 1-14 every 3 weeks x 24 weeks
FOLFIRI	Irinotecan 180 mg/m <sup>2</sup> IV over 30-90 minutes, day 1 Leucovorin 400 mg/m <sup>2</sup> IV infusion to match duration of irinotecan infusion, day 1 5-FU 400 mg/m <sup>2</sup> IV bolus on day 1 then 1200 mg/m <sup>2</sup> /day x 2 days (total 2400 mg/m <sup>2</sup> over 46 – 48 hours) IV continuous infusion Repeat every 2 weeks

Table 1: Chemotherapy regimen for the treatment of CRC (Cont.)

Name	Regimen
FOLFOXIRI	Irinotecan 165 mg/m <sup>2</sup> IV day 1, Oxaliplatin 85 mg/m <sup>2</sup> IV day 1, Leucovorin 400 mg/m <sup>2</sup> IV day 1, 5-FU 3200 mg/m <sup>2</sup> over 48 hour continuous infusion starting on day 1 Repeat every 2 weeks

### 3. Chemotherapeutic drugs used in CRC

The chemotherapeutic drugs that have been the gold standard for the treatment of advanced CRC for over 40 years are the combination of 5-fluorouracil/Leucovorin (5-FU/LV). It has been proven to provide a survival benefit over supportive care. The newer cytotoxic agents that come into regular practice are irinotecan and oxaliplatin due to improved response rates and survival when combine with 5FU/LV. The newest alternatives are the addition of biological agents, bevacuzimab and cetuximab, to the standard cytotoxic drugs which have a very promising result (17).

#### 3.1 5-Fluorouracil/Leucovorin

The 5-fluorouracil (5-FU) is a fluoropyrimidine antimetabolite drugs. It acts specifically to the cell in S phase of the cell cycle. 5-FU is widely used in the treatment of a range of cancers, including colorectal and breast cancers. 5-FU-based chemotherapy improves overall and disease-free survival of patients with resected stage III colorectal cancer (18).

5-FU is an analogue of uracil with a fluorine atom at the C-5 position in place of hydrogen. Due to the structure similarity, 5-FU rapidly enters the cell by the same facilitated transport mechanism as uracil. It is converted intracellularly to several active metabolites: fluorodeoxyuridine monophosphate (FdUMP), fluorodeoxyuridine triphosphate (FdUTP) and fluorouridine triphosphate (FUTP). The anticancer mechanisms of 5-FU exert through the inhibition of thymidylate synthase (TS) and



incorporation of its metabolites into RNA and DNA therefore disrupting RNA and DNA synthesis (18).

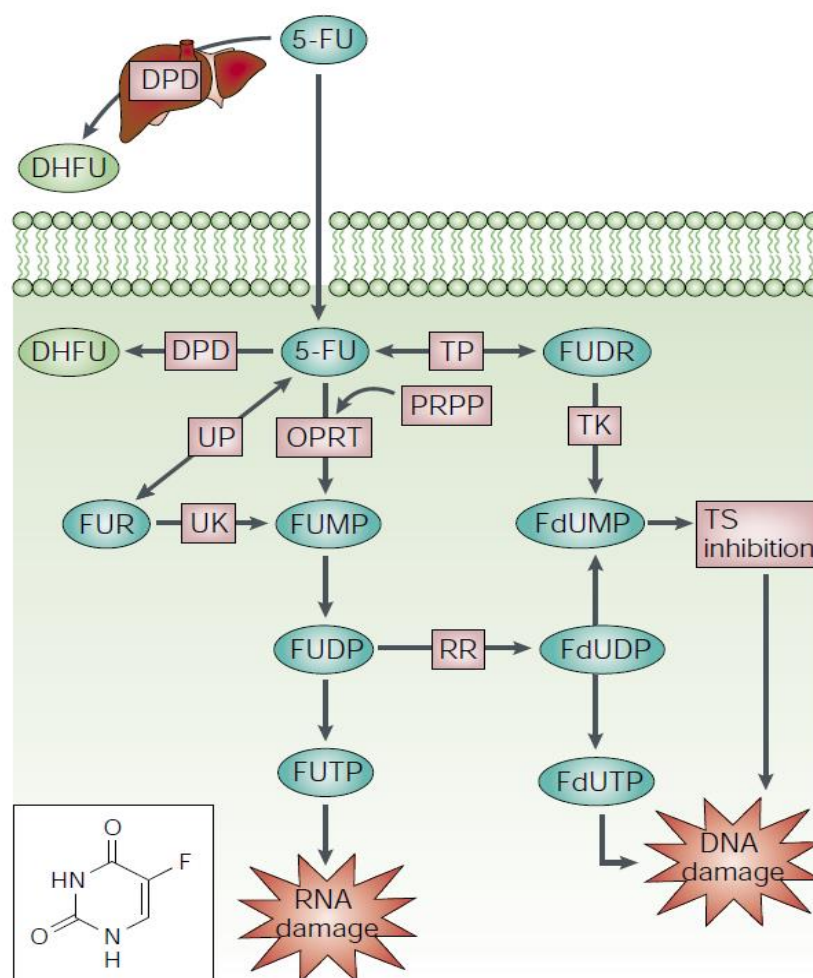


Figure 2: 5-Fluorouracil metabolism (18).

The anticancer activity of 5FU can be enhanced by combining with leucovorin (LV, 5'-formyltetrahydrofolate). LV enters the cell via the reduced folate carrier and is anabolized to reduced folate ( $\text{CH}_2\text{THF}$ ). High intracellular levels of the  $\text{CH}_2\text{THF}$  are necessary for optimal binding of FdUMP to TS. Moreover,  $\text{CH}_2\text{THF}$  is polyglutamated by foyllypolyglutamate synthetase. Polyglutamation results in increases the cellular retention of  $\text{CH}_2\text{THF}$ , and enhances the stabilization of its ternary complex with TS and FdUMP. LV has been used to increase the intracellular concentration of  $\text{CH}_2\text{THF}$  and has been shown to enhance the toxicity of 5-FU in many cancer cell lines (18).

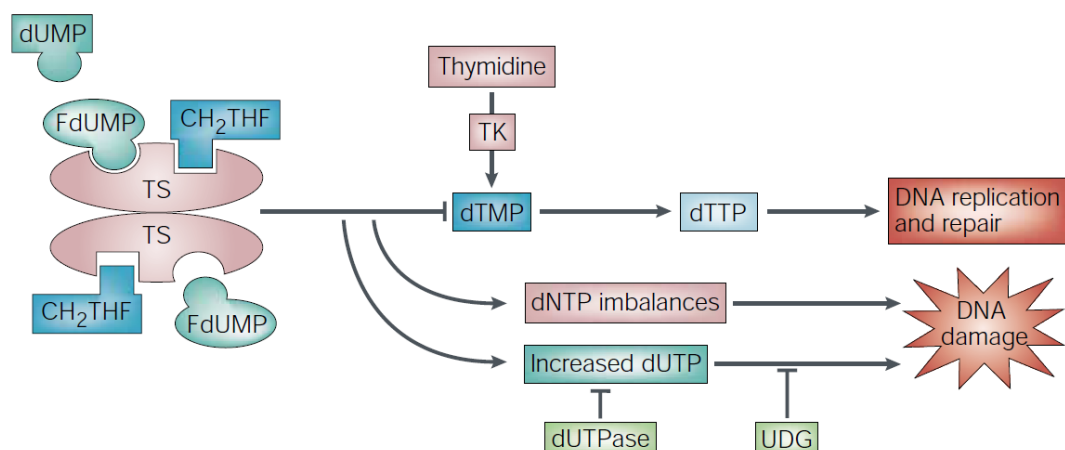


Figure 3: Mechanism of thymidylate synthase inhibition by 5-fluorouracil (18).

### 3.2 Oxaliplatin

Oxaliplatin is a third-generation platinum derivative, which, when combined with 5FU and leucovorin, is among the most effective chemotherapies for metastatic colorectal cancer (19). The mechanism of action of platinum compounds is similar to the alkylating agents. It covalently binds to nucleophilic sites on DNA to form a drug-DNA complex. The platinum compounds can form both intrastrand and interstrand cross-links with the DNA. DNA adducts inhibit DNA replication and transcription and lead to breaks and miscoding, and if recognized by p53 and other checkpoint proteins, induction of apoptosis (20).

Oxaliplatin, a diaminocyclohexane-containing platinum, has a spectrum of activity and mechanisms of action and resistance that different from those of other platinum-containing compounds (20).

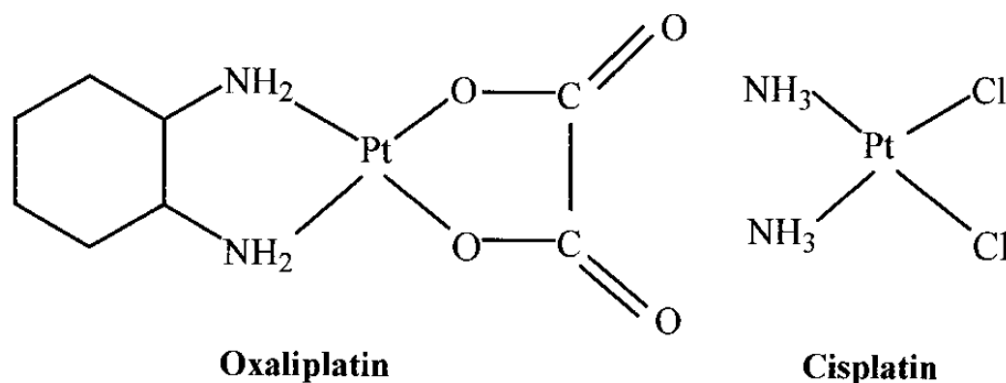


Figure 4: Chemical structures of oxaliplatin and cisplatin (20).

### 3.3 Irinotecan

Irinotecan is a semisynthetic analog of camptothecin, originally isolated from the Chinese/Tibetan ornamental tree *Camptotheca acuminata*. The prototype drug is topotecan (21). Irinotecan has shown activity against colorectal, esophageal, gastric, non-small-cell and small-cell lung cancers, leukemia and lymphomas, as well as central nervous system malignant gliomas (22).

The mechanism of action of irinotecan is by interacting with topoisomerase I (Topo I) in the cell. The topoisomerase I is the enzyme that unwind DNA strand to facilitate DNA replication. The Topo I cuts one strand of double-stranded DNA, relax the strand, and reanneal the strands. Irinotecan binds to the Topo I-DNA complex, stabilizing it and preventing re-ligation of the complex. Collision with advancing replication forks results in the formation of double-stranded DNA breaks. These breaks activate cell cycle arrest and, if unrepaired, can cause cell death. It is a chemotherapy agent that acts in the S-phase of cell cycle (23).

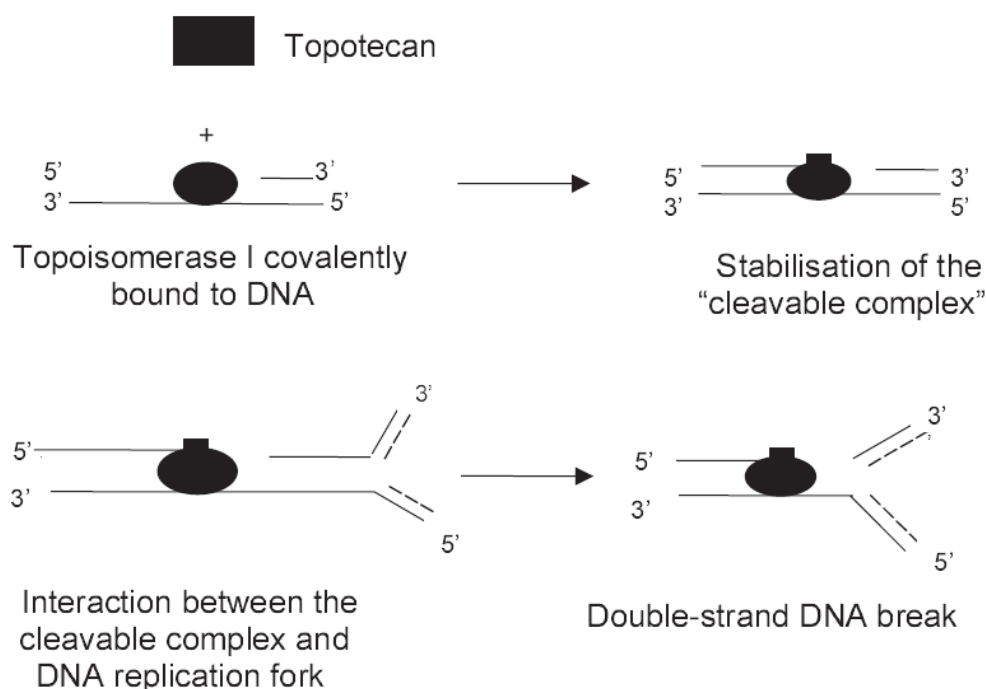


Figure 5: Mechanism of action of topotecan (24).

### 3.4 Bevacizumab

Bevacizumab is a humanized monoclonal antibody against the vascular-endothelial growth factor (VEGF). VEGF is a potent pro-angiogenic factor, the stimulator of the growth of vascular endothelial cells. Bevacizumab binds to VEGF-A, which prevents its interaction with the VEGF receptor tyrosine kinases VEGFR1 and VEGFR2 receptors (25).

VEGF expression is up-regulated in several tumor types, including breast, ovarian, non-small cell lung, and colorectal cancer. Its expression correlates with neovascularization within tumors. In colorectal cancer, micro-vessel density is associated with progression of adenomas to carcinomas and with metastatic potential and poor prognosis. Bevacizumab is FDA approved for metastatic colorectal cancer in combination with 5-FU based chemotherapy (26).

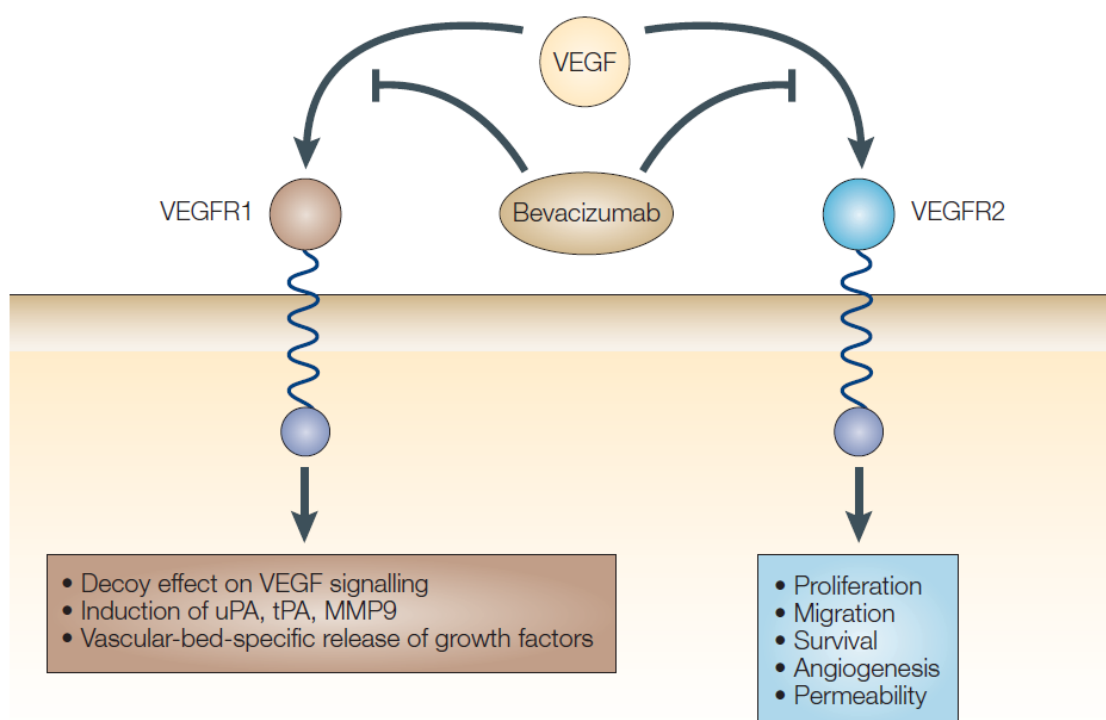


Figure 6: Mechanism of action of Bevacizumab (25).

### 3.5 Cetuximab

Cetuximab is a chimeric IgG<sub>1</sub> monoclonal antibody that recognizes the epidermal growth factor receptor (EGFR). Activation of EGFR tyrosine kinase receptor is associated with proliferation, survival, and angiogenesis. EGFR expression is found in 60–75% of colorectal cancers, where it has been linked to tumor progression and a poor prognosis. Multiple epithelial cancers, such as breast, lung, kidney, prostate, brain, pancreas, bladder, and head and neck malignancies, also express EGFR and are potential therapeutic targets of cetuximab (27).

Cetuximab binds to the extracellular domain of EGFR, blocks ligand induced receptor signaling and modulates tumor cell growth. Immune-mediated antitumor mechanisms, such as antibody-dependent cell-mediated cytotoxicity (ADCC), may also contribute to the activity of cetuximab (28).

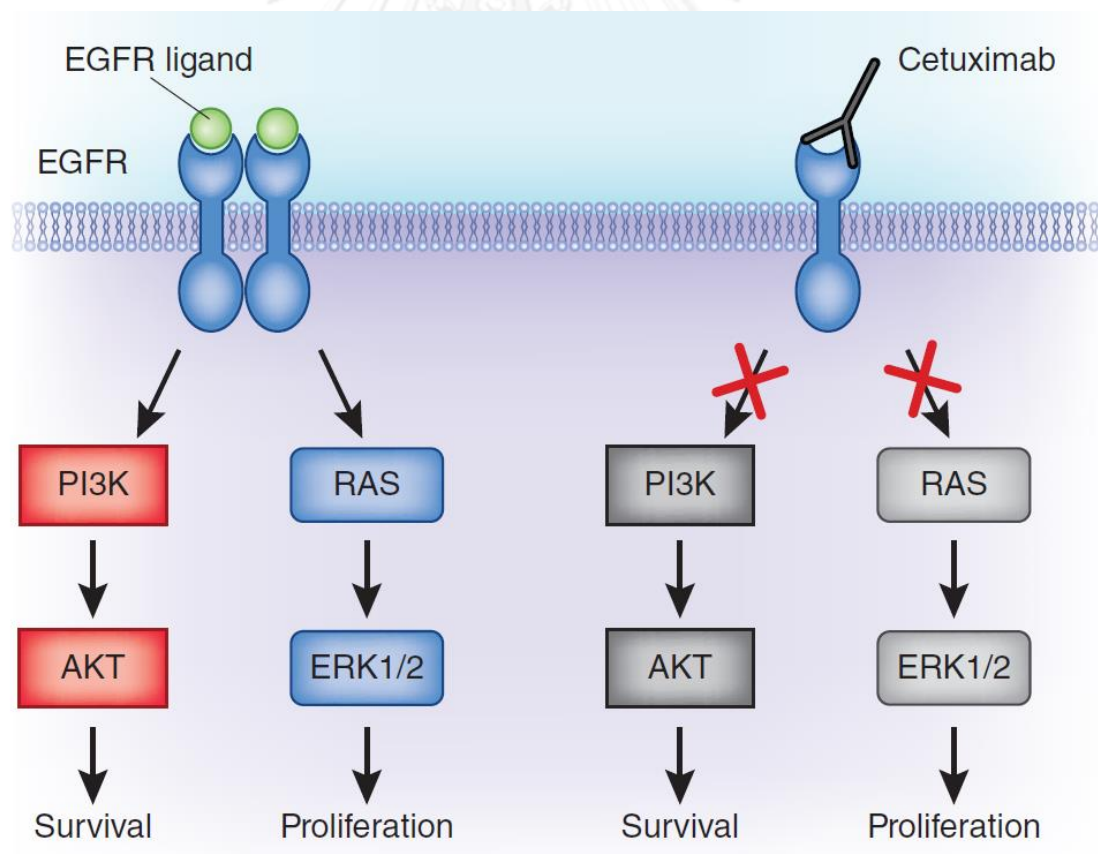


Figure 7: Mechanism of action of Cetuximab (29).

Cetuximab is FDA approved for the treatment of EGFR-positive metastatic colorectal cancer as a single agent in patients who were unable to tolerate irinotecan-based therapy, or in combination with irinotecan for refractory patients (27).

#### 4. COX-2 and CRC

##### 4.1 COX biology

Cyclooxygenases (or prostaglandin H synthases), commonly known as COXs, are a family of myeloperoxidases located at the luminal side of the endoplasmic reticulum and nuclear membrane, which catalyze the rate-limiting step of prostaglandin biosynthesis from arachidonic acid (30). These enzymes act by two coupled reactions. The first one is the conversion of arachidonic acid to prostaglandin  $G_2$  by the cyclooxygenase activity. The second reaction is the conversion of prostaglandin (PG)  $G_2$  to prostaglandin  $H_2$  which mediated by the peroxidase activity. Then, different synthases convert prostaglandin  $H_2$  to prostaglandin  $D_2$ ,  $F_{2\alpha}$ ,  $E_2$ ,  $I_2$ , and thromboxane  $A_2$ .

Prostaglandins are immediately released from the cells to act locally in an autocrine and paracrine way through different receptors and activating different intracellular pathways (31). Important physiological functions of prostaglandins are vasodilatation ( $PGD_2$ ,  $PGE_2$ ,  $PGI_2$ ), gastric cytoprotection ( $PGI_2$ ), maintenance of renal homeostasis, and platelet aggregation ( $TXA_2$ ). Besides, prostaglandins play a major role in mediating fever ( $PGE_2$ ), pain sensitivity, and inflammation. Up to now, three isoforms of COXs have been identified.

Cyclooxygenase-1 (COX-1) is constitutively expressed in different tissues. COX-1 plays a role in tissue homeostasis by modulating several cellular processes ranging from cell proliferation to angiogenesis or platelet aggregation due to thromboxane production (30). Cyclooxygenase-2 (COX-2) is the inducible isoform, which is regulated by growth factors and different cytokines such as  $IL1\beta$ ,  $IL6$ , or  $TNF\alpha$  which overexpressed during inflammation (32). The COX-2 promoter displays an NF- $\kappa$ B response element as well as other cytokine-dependent (i.e.,  $IL6$ ) response elements

(30). Lastly, COX-3 has been identified as a splice variant of COX-1, and it is present mainly in brain and spinal cord. Currently, the role of COX-3 is not known (33).

#### 4.2 COX-2 expression in CRC

The relationship between COX expression and tumor biology has shown that higher COX-2 expression is associated with larger tumor size, more advanced stage of disease, an increased likelihood of lymph node involvement and a reduced likelihood of patient survival (34). Five-year survival in the absence of COX-2 expression is 91.6% compared with a survival rate of 40.5% for those patients whose tumors express the enzyme (35). Animal models provide additional evidence of the role of COX-2 in colorectal carcinogenesis (36). The multiple intestinal neoplasia mice model resembles the familial adenomatous polyposis (FAP) syndrome in human. COX-2 levels are elevated in the polyps of these MIN mice (37). COX-2 is also detected in pre-cancerous lesions and is increased in the later stages of cancer development (38). The elevation of COX-2 early in the development of CRC, suggests a role of COX-2 in the carcinogenesis. Other evidence of the involvement of COX-2 in CRC was found by using gene knockout techniques. In the APC MIN mice model, the COX-2 gene was disrupted to investigate the role of COX-2 in the formation of polyps. The development of intestinal adenomas was less than six-fold in these COX-2 knockout MIN mice compared to the COX-2 wild-type MIN mice. Although polyp formation in COX-2<sup>-/-</sup> MIN mice was not totally prevented, this study demonstrated direct genetic evidence of the importance of COX-2 in the carcinogenesis in CRC (39). Taken together, the evidence from animal studies reinforces the concept that COX-2 plays an important role in tumor development.

Cyclooxygenase-2-derived PGs are involved in many of pathological processes include CRC. Multiple lines of evidence have demonstrated that PGE<sub>2</sub> mediates the tumor-promoting effects of COX-2 in CRC. The steady-state cellular levels of PGE<sub>2</sub> depend on the relative rates of COX-2/PGE synthase-dependent biosynthesis and 15-hydroxyprostaglandin dehydrogenase (15-PGDH)-dependent degradation (7). PGE<sub>2</sub> has a predominant role in promoting colorectal tumor growth. PGE<sub>2</sub> is the most abundant PG found in human CRC (40). In contrast, 15-PGDH is highly expressed in

normal colon mucosa but is lost in most human CRCs (41). Moreover, the loss of 15-PGDH leads to resistance of the antitumor effects of celecoxib in an azoxymethane (AOM) mouse model (42). Recent studies showed that PGE<sub>2</sub> treatment drastically increased both small and large intestinal adenoma burden in Apc<sup>Min/+</sup> mice and significantly enhanced AOM-induced colon tumor incidence and multiplicity (43). In contrast, inhibition of endogenous PGE<sub>2</sub> by genetic deletion of mPGES-1 suppressed intestinal tumorigenesis in Apc<sup>Min/+</sup> and AOM models (44).

Many reports have shown that PGE<sub>2</sub> promotes colorectal tumor growth by stimulating angiogenesis, cell invasion, cell growth and inhibiting apoptosis. These processes are regulated by PGE<sub>2</sub>-activated signaling pathways, including EGFR, PI3K-Akt, Ras-MAPK, PPAR $\delta$ , VEGF, BCL-2, chemokines and their receptors (45; 46). On the other hand, many downstream targets of PGE<sub>2</sub> can also up-regulate COX-2 expression. Such positive feedback loops may amplify the activity of the COX-2 pathway and magnify the potency of COXIBs (7).

### 4.3 NSAIDs and coxibs in colorectal cancer

NSAIDs are widely used as analgesics, anti-inflammations and anti-pyretics. Their anti-inflammatory actions result from the inhibition of COX activity and consequently PGs formation. Aspirin and many other NSAIDs inhibit non-specifically both isoforms of COX, while some newer generations, such as the coxibs, have higher selectivity on the COX-2 isoform. COX-inhibitors block the synthesis of PGs by directly competing with the substrate, arachidonic acid, for access to the active site of the COX enzyme. Aspirin, the prototype NSAID, has a distinct mechanism of action from other NSAIDs. It irreversibly inactivates COX by acetylating a serine residue within the active site of both COX-1 and COX-2 (12).

A multiple line of evidence from population-based studies, case-control studies and clinical trials indicate that regular use of NSAIDs including aspirin and COXIBs over a 10- to 15-year period reduces the relative risk of developing CRC by 40–50% (47-49). Moreover, aspirin use was associated with a risk reduction in patients whose colon tumors expressed higher levels of COX-2 (50). Besides prevention, regular aspirin use after the diagnosis of CRC at stage I-III improves overall



survival, especially among patients with tumors that overexpress COX-2 (51), suggesting the potential therapeutic use of NSAIDs in advanced CRC.

Furthermore, several large randomized double-blind placebo-controlled trials have demonstrated preventive effects of NSAIDs on patients with previous polyps by reducing the risk of colorectal (adenomatous) polyp recurrence (52; 53). These findings alerted investigators to evaluate the preventive effects and safety of COXIBs in patients with FAP or previous history of adenomas. Observational studies demonstrating that both celecoxib and rofecoxib prevented the occurrence and recurrence of CRC stimulated growing interest in the anti-neoplastic properties of this new generation of NSAIDs (54). Particularly, the evidence that treatment of FAP patients with celecoxib significantly reduced the polyp burden in a randomized controlled trial led to the Food and Drug Administration approval of celecoxib for use in patients with FAP at 400 mg twice a day (55). Initial enthusiasm for the use of selective COX-2 inhibitors waned following reports of an increased risk of cardiovascular events in patients taking these agents for long periods in CRC chemoprevention trials. The results of the APPROVe (Adenomatous Polyp Prevention on Vioxx™) trial showed a doubling in the relative risk of thrombotic events in patients receiving rofecoxib for > 18 months and led to the withdrawal of the drug by the manufacturer (56). Likewise, long-term treatment with celecoxib in the APC (adenoma prevention with celecoxib) trial was also associated with increased risk of death from cardiovascular causes (57). However, the APPROVe, APC and subsequent polyp prevention trials confirmed that both rofecoxib and celecoxib were effective in the reduction of colorectal adenoma recurrence (52; 58; 59). The mechanism of increased cardiovascular risk of COXIBs probably involves reduction in vascular production of the anti-thrombotic prostacyclin, without a change in the generation of the pro-thrombotic thromboxane (derived from COX-1 activity in platelets) (60).

## 5. Apoptosis pathway

Apoptosis or programmed cell death is an essential physiological process that plays a crucial role in controlling the number of cells in development and

throughout the life by removal of cells at the appropriate time. However, apoptosis is also involved in a broad range of pathological conditions include neurodegenerative diseases, cardiovascular diseases, immunological diseases, and cancer (61). There are 3 different pathways to initiated apoptosis: 1) extrinsic apoptotic pathway which can be trigger by binding of death ligands to death receptors and consequent caspase-8 activation; 2) intrinsic apoptotic pathway, which is initiated by cellular stress followed by activation of caspase-9; or 3) Perforin/granzyme pathway, where the cytotoxic T cell produces protease enzyme and delivers to sensitize target cells. Each of these pathways converges to a common execution step that requires proteolytic cleavage and activation of caspases-3 and/or -7 (62).

### **5.1 Intrinsic apoptotic pathway**

The intrinsic apoptotic pathway or mitochondria pathway is triggered by various extra- and intracellular stimulus, including oxidative stress, irradiation, and cytotoxic drugs (62). This pathway is involved with the permeability of the mitochondrial membrane which is regulated by the BCL-2 family proteins. When induction of apoptosis, the pro-apoptotic BCL-2 proteins, Bax and Bak, insert into mitochondrial membrane and cause cytochrome c release from the mitochondrial inter-membrane space into the cytosol. Anti-apoptotic BCL-2 family proteins, BCL-2 and BCL-XL, prevent cytochrome c release by binding and inhibition of BAX and BAK. Cytochrome c then binds to the Apaf1 and together with dATP causes recruitment of pro-caspase-9 to the complex, called apoptosome (63; 64). Caspase-9 is then activated and in turn activates caspase-3 to initiate the proteolytic cascade (65).

### **5.2 Extrinsic apoptotic pathway**

The extrinsic apoptotic pathway is initiated by extracellular ligands bind with death receptors. Typical death receptors are Fas (fibroblast associated antigen or Apo-1 or CD95) and tumour necrosis factor receptor 1 (TNFR1). Binding of ligands to death receptors induce receptor clustering and formation of a death inducing signalling complex (DISC) (66) in which the adaptor proteins FADD and/or TRADD bind

with their death domain (DD) to a death domain of the receptors in the cytoplasmic region. This leads to the recruitment of caspase-8 or -10 (initiator caspases) to the DISC (67). Activated caspase-8 then directly cleaves pro-caspase-3 or other executioner caspases, and finally leading apoptosis cell death. Caspase-8 can also cleave the BH3-only protein Bid. The resulting truncated Bid (tBid) then moves to the mitochondria and induces cytochrome c release, leading to activation of caspase-9 and caspase-3 (68).

### 5.3 Perforin/Granzyme pathway

Cytotoxic T cells can exert their cytotoxic effects on tumor cells and virus-infected cells via a pathway that involves secretion of the transmembrane pore-forming molecule perforin with a subsequent exophytic release of cytoplasmic granules through the pore and into the target cell (69). The serine proteases granzyme A and granzyme B are the most important component within the granules. Granzyme B will cleave proteins at aspartate residues and will therefore activate pro-caspase-10 (70). Moreover, granzyme B can utilize the mitochondrial pathway for amplification of the death signal by specific cleavage of Bid and induction of cytochrome c release (71). However, granzyme B can also directly activate caspase-3. In this way, the upstream signaling pathways are bypassed and there is direct induction of the execution phase of apoptosis.

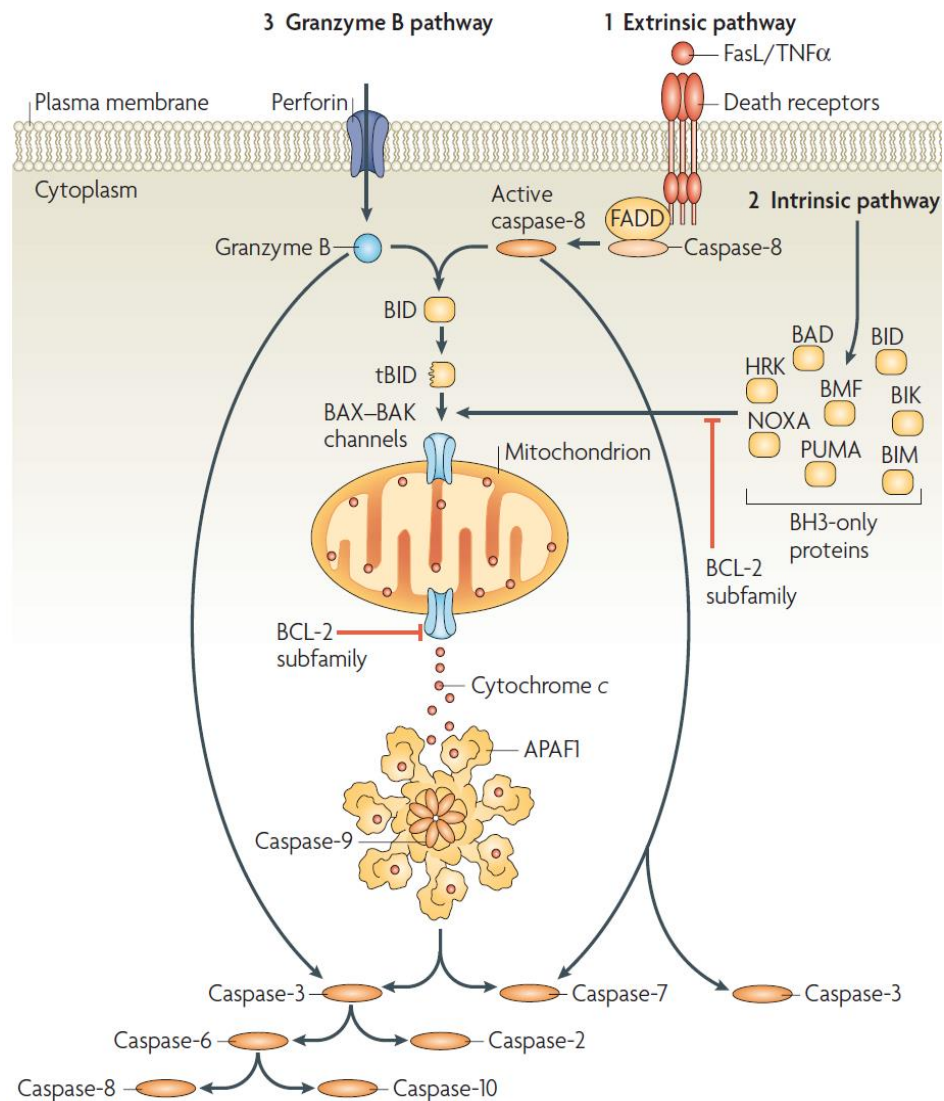


Figure 8: Apoptosis pathway (72).

## 6. BCL-2 family protein and the regulation of apoptosis

The BCL-2 family proteins are key molecules which function as anti-apoptotic or pro-apoptotic in the intrinsic apoptotic pathway. All of BCL-2 family structure shares the same conserved regions termed BCL-2 homology (BH) domains. It can be divided into 3 sub group according to the function and structure.

- 1) Pro-apoptotic BH3 only proteins - BIM, PUMA, BID, BAD, BIK, BMF, HRK, NOXA
- 2) Pro-apoptotic BAX-like proteins – BAX, BAK
- 3) Anti-apoptotic Bcl-2-like proteins - BCL-2, BCL-XL, BCL-W, MCL-1, A1

The pro-apoptotic BH3-only proteins share only the BH3 region of homology and serve as upstream sensitizer, becoming activated in response to various forms of cellular stress, whereas pro-apoptotic Bax-like proteins contain multiple BH domains (BH1, BH2 and BH3) and are required downstream of BH3-only proteins to induce apoptosis. The anti-apoptotic proteins share up to four regions of sequence homology (BH1-4) (73).

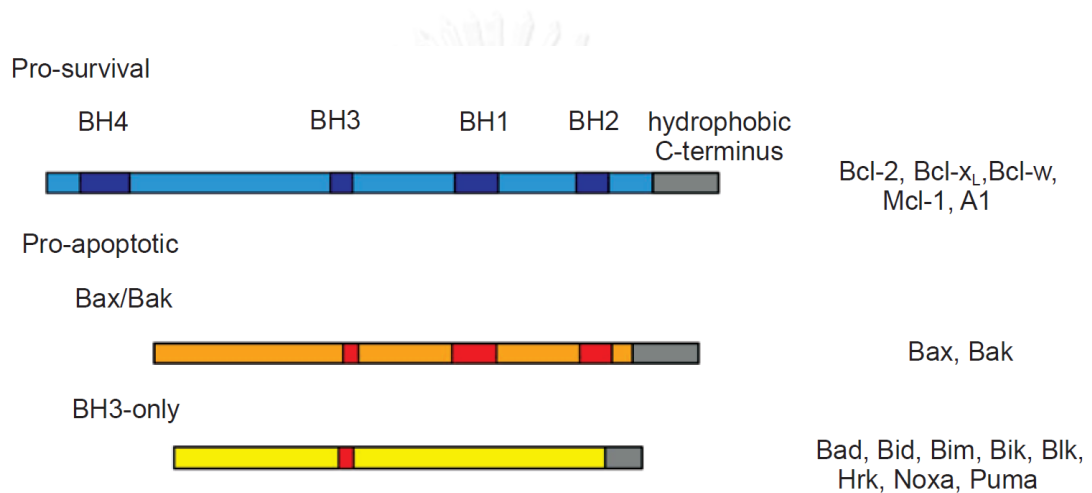


Figure 9: Sequence homology between proteins of the BCL-2 family (73).

Activating intrinsic apoptotic pathway requires both types of pro-apoptotic proteins. The BH3-only proteins act as damage sensors and direct antagonists of anti-apoptotic BCL-2-like proteins. whereas the Bax-like proteins, once activated, act further downstream, probably by permeabilizing the mitochondrial outer membrane and perhaps also by perturbing the endoplasmic reticulum (ER)/nuclear envelope (74).

Bax and Bak appear to be largely redundant in function. A biochemical difference between Bax and Bak is that, in healthy cells, Bax is largely cytosolic or loosely associated with mitochondria, whereas Bak is an integral membrane protein on the cytosolic face of the mitochondrion and ER. In response to cytotoxic signals, Bax translocates to those membranes, and both Bax and Bak change conformation and form membrane-associated homo-oligomers. These oligomers are thought to permeabilize the outer mitochondrial membrane (74).

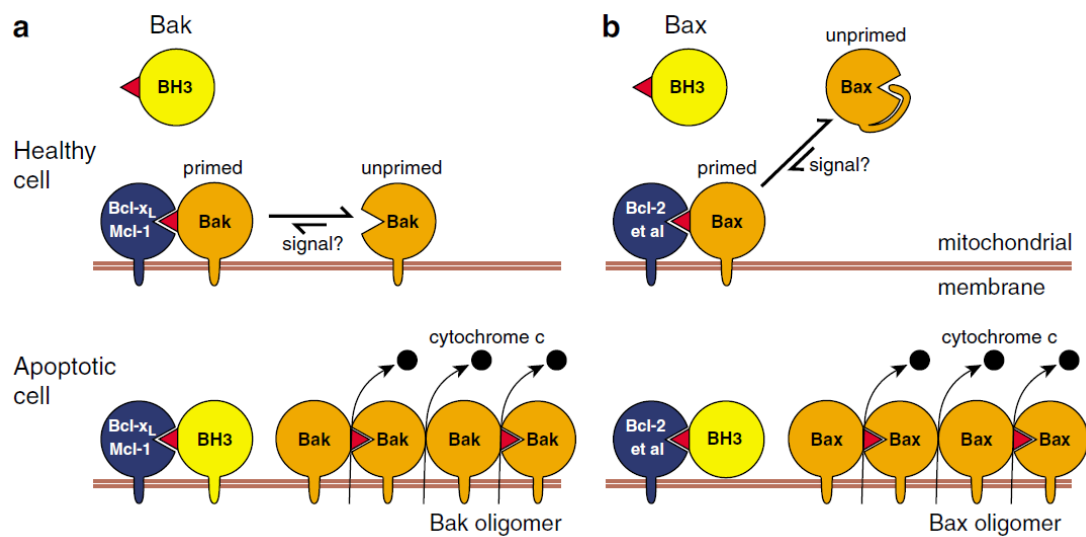


Figure 10: Model for the regulation of Bak and Bax by their pro-survival relatives (74).

The anti-apoptotic activity of BCL-2 has been demonstrated in both overexpression and gene targeting studies. Overexpression of any of the five pro-survival members protects cells against apoptosis induced by a variety of cytotoxic stimuli family (73). Anti-apoptotic BCL-2 proteins are generally integrated within the mitochondrial outer membrane, but may also be in the cytosol or ER membrane. They preserve mitochondrial outer membrane integrity by directly inhibiting the pro-apoptotic BCL-2 proteins (75).

## 7. The regulation of Cell cycle

The eucaryotic cell cycle is divided into four stages: G1, S, G2, and M. G1 is the gap phase during which cells prepare for the process of DNA replication. During the G1 phase, cell integrates mitogenic and growth inhibitory signals and makes the decision to proceed, pause, or exit the cell cycle. S phase is defined as the stage in which DNA synthesis occurs. G2 is the second gap phase during which the cell prepares for the process of division. M stands for mitosis, the phase in which the replicated chromosomes are segregated into separate nuclei and cytokinesis occurs to form two daughter cells. In addition to G1, S, G2, and M, the term G0 is used to describe cells that have exited the cell cycle and become quiescent (76).

The ability of the cells to divide is mainly attributed to the presence of two classes of molecules, cyclin-dependent kinases (CDKs), a family of serine/threonine kinases and their binding partners, cyclins, which are synthesized and destroyed at specific times during the cell cycle, thus regulating kinase activity in a timely manner.. The discovery of CDKs and cyclins has increased considerably during evolution. However, only certain CDK–cyclin complexes are thought to control cell cycle progression. According to the ‘classical’ model for the mammalian cell cycle, specific CDK–cyclin complexes are responsible for driving the various events known to take place during interphase in a sequential and orderly fashion (77).

In early G1 phase, Cdk4 and/or Cdk6 are activated by D-type cyclins and initiate phosphorylation of the retinoblastoma protein (Rb) family. This leads to the release of E2F transcription factors and results in the activation and transcription of E2F responsive genes required for cell-cycle progression. Early E2F responsive genes include E- and A-type cyclins. In the late G1 phase, Cdk2 is activated by binding to cyclin E and completes the phosphorylation of Rb leading to further activation of E2F mediated transcription. This leads to passage through the restriction point at the boundary of the G1/S phase, and to S phase initiation. Later Cdk2 plays an important role in S phase progression by complexing with cyclin A. A-type cyclins are synthesized at the onset of the S phase and phosphorylate proteins involved in DNA replication. During the G2/M transition, Cdk1/cyclin A activity is required for the initiation of prophase. Finally, Cdk1/cyclin B complexes actively participate in and complete mitosis (78).

The activities and functions of Cdk/cyclin complexes under normal as well as extreme conditions, such as stress, DNA damage, telomere dysfunction and others, are regulated by two families of Cdk inhibitors; the INK4 family (p16<sup>INK4a</sup>, p15<sup>INK4b</sup>, p18<sup>INK4c</sup>, p19<sup>INK4d</sup>) specifically bind to Cdk4 and Cdk6 and prevent D-type cyclin activity and the Cip/Kip family (p21<sup>Cip1/Waf1/Sdi1</sup>, p27<sup>Kip1</sup>, p57<sup>Kip2</sup>) inhibits Cdk2/cyclin E, Cdk2/cyclin A, Cdk1/cyclin A, as well as Cdk1/cyclin B activity (78).

The basic concept put forward by this model, that is, that each phase of the cycle is driven by specific CDKs, has been recently challenged by genetic studies in mice. Extensive analyses of these models revealed that most of the Cdk and

cyclins, originally thought essential for cell cycle regulation, are in fact largely dispensable. Cyclin A2 and B1 emerged as the most non-redundant cyclins. CDK2, CDK4 and CDK6 are not essential for the mammalian cell cycle. Instead, they are only required for the proliferation of specific cell types. By contrast, Cdk1 emerged as the master regulator of mammalian cell-cycle regulation whereas and Cdk1 alone by complexing with all the cyclins can drive the cell cycle, at least until mid-gestation (77).

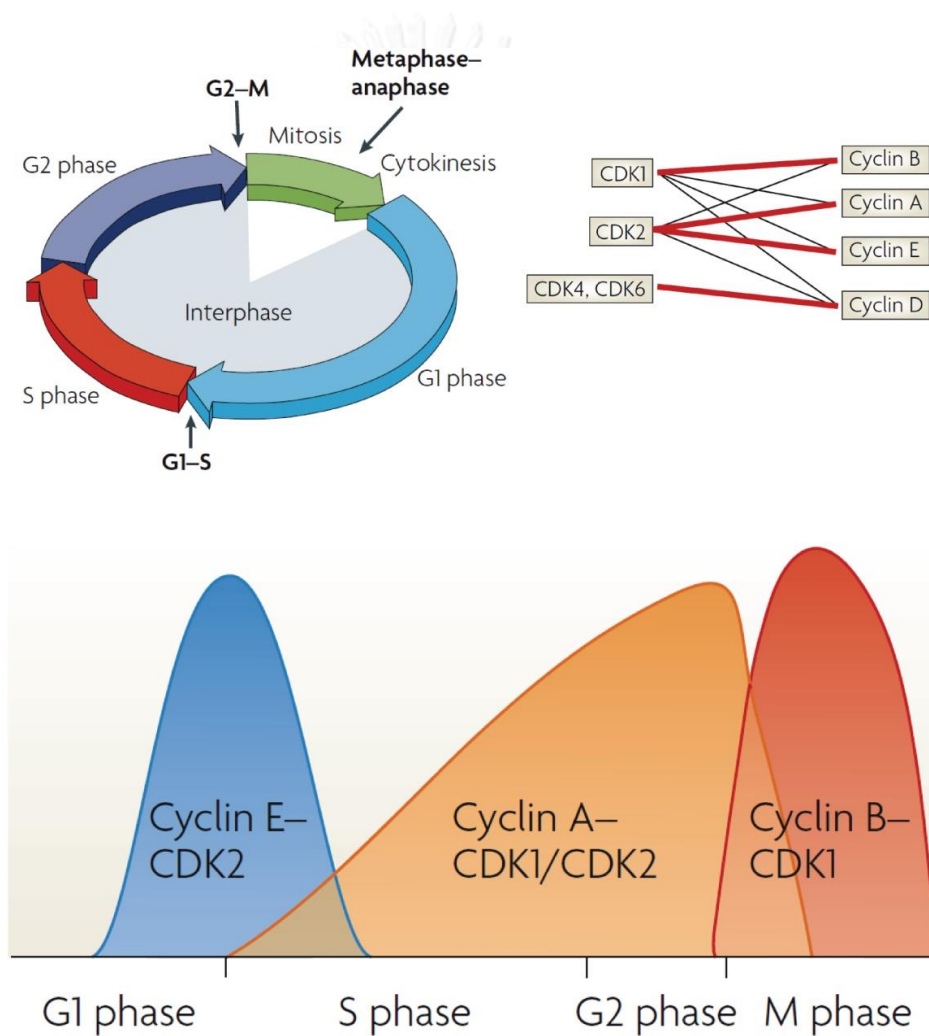


Figure 11: The control of cell cycle (79).



## 8. *Glycosmis parva*

### 8.1 Plant

*Glycosmis parva* Craib (Rutaceae) is wild small shrub distributed in Thailand with general name of Som-chuen or Prayon-Kluean. The botanical characteristics are leaves alternate, 1-5 foliate. Flowers are usually small, axillary panicles, calyx 4 or 5 partial imbricate. Petals 4 or 5 imbricate, stamen 8 to 10 free, filaments dilated below. Ovary 2 to 5 celled, the style very short, not jointed ovule 1 in each cell. Fruits globose, freshy, berry. Seed 1 to 3 oblong, testa membranous (80).



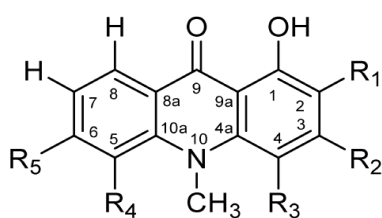
Figure 12: *Glycosmis parva* CRAIB

### 8.2 Compounds

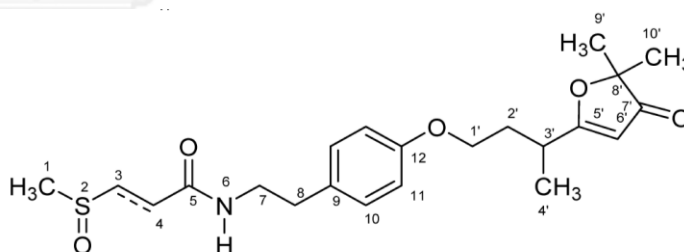
The genus *Glycosmis* is a rich source of acridone alkaloids and sulfur-containing propanamides especially in branches and leaves, respectively. Chemical examination of the branches and leaves of *G. parva* led to the identification of the following compounds (8);

Table 2: Chemical components from the branches and leaves of *G. parva*

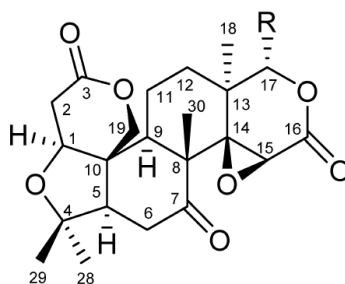
Part	Main Structure	Chemical name
Branches	Acridone alkaloids	1. Glycosparvarine 2. N-methylatalaphylline 3. N-methylcyclo-atalaphylline-A 4. Glycofolinine 5. Citramine
	Limonoids	1. Limonin 2. Limonexic acid 3. Isolimonexic acid
Leaves	Sulfur-containing propanamide derivatives	1. S-deoxydihydroglyparvin 2. S-deoxytetrahydroglyparvin 3. Glyparvin-A 4. Dihydroglyparvin
	Acridone alkaloids	1. Arborinine



Acridone alkaloids



Sulfur-containing propanamide derivatives



Limonoids

Figure 13: Structure of chemical constituents found in *G. parva* (8)

### 8.3 Pharmacological activities

Some plants in the genus *Glycosmis* have been used as Thai traditional medicines for treatments of abscess, scabies, and snakebite. Numbers of biological activities of the compound isolated from *Glycosmis* spp. have been reported. The majority of them are cytotoxicity, anti-tumor activity, antiviral activity and anti-malarial activity of acridone alkaloids. For sulfur-containing propanamides, antifungal activities were mainly reported.

#### *Antimarial effects*

Many acridone alkaloids from *Glycosmis* spp. exhibited antimarial activities that were comparable to anti-malarial drug-chloroquin. 5-hydroxynoracronycine at 10 µg/ml inhibited 90% of *Plasmodium yoelii*, *in vitro* (81).

#### *Antiviral activities*

Atalaphillidine, citracridone-I and 5-hydroxy-N-methylseverifoline possessed antiviral activity against herpes simplex virus type 2 (HSV-2) (82). Compounds isolated from leaves and branches of *G. parva* exhibited an antiviral activity against both HSV-1 and HSV-2. Glycosparvarine, glycofolinine and (+)-tetrahydroglyparvin showed moderate activities with EC<sub>50</sub> of 151 µM, 348 µM and 229 µM, respectively. While (+)-5-deoxydihydroglyparvin exhibited more potent activities with EC<sub>50</sub> of 29.8 µM and 44.6 µM, respectively, against HSV-1 and HSV-2 (8).

#### *Antifungal activities*

Many sulfur-containing propanamide compounds found in *Glycosmis* spp. exhibited antifungal activity. Methyllukumbin-A at very low concentration (10 µg/ml) showed antifungal activity against *Cladosporium cladosporioides* (83). Moreover, it could inhibit *Cladosporium herbarum* with an ED50 of 5.5 µg/ml (84).

#### *Anti-inflammation*

The hexane and EtOAc extracts from the branches and leaves of *G. parva* inhibited nitric oxide production, mRNA expression of iNOS, COX-2, and pro-inflammatory cytokines (IL-1, IL-6, TNF- $\alpha$ ) in LPS-activated macrophage J774A.1 cells (10).

#### 8.4 Anti-cancer activities

Several effects related to anti-cancer activities have been reported from the compounds found in *Glycosmis* spp.

##### *Anti-proliferative effect*

Acridone alkaloids have been reported to possess antiproliferative effect. Atalaphyllidine and des-*N*-methylnoracronycine inhibited clonal proliferation and induce cell differentiation in HL-60 (Human promyelocytic leukemia) (85). In addition, atalaphyllidine, 5-hydroxy-*N*-methylseverifoline and des-*N*-methylnoracronycine exhibited antiproliferative against several cancer cell lines include human lung carcinoma (A-549), mouse melanoma (B16-melanoma 4A5), T-cell leukemia (CCRF-HSB-2) and human gastric cancer cell and lymph-node metastasized (TGBC11TKB) (86).

Arborinine which is the main acridone alkaloids found in EtOAc extract from the leaves of *G. parva* showed potent antiproliferative effect against cervix adenocarcinoma (HeLa) with an IC<sub>50</sub> of 1.84 μM which was lower than that of cisplatin (12.43 μM) (36).

##### *Cancer prevention*

Glycocitrine-II and O-methylglycocitrine-II had potent cancer prevention properties by inhibiting Epstein-Barr virus early antigen activation with IC<sub>50</sub> of 280 and 281 mol ration/32 pmol TPA (87).

##### *Inhibit cell cycle progression and induction of apoptosis*

Glyfoline inhibited leukemic HL-60 cell growth in vitro with IC<sub>50</sub> of 1.1 μM (88). Moreover, it induced nasopharyngeal carcinoma (NPC) cell growth arrest at G<sub>2</sub>/M and induced apoptosis (89). The EtOAc extract from the leaves of *G. parva* exhibited potent cytotoxic activity (IC<sub>50</sub> 15.68 μg/ml), induction of apoptosis (decrease BCL-XL, increase BAK), and caused G<sub>1</sub>/S phase accumulation (decreased cyclin D1 and cyclin E, increase p21) in human B-cell lymphoma (9).

## CHAPTER III

### MATERIAL AND METHOD

#### 1. Material

##### 1.1 Cell cultures

Human colorectal cancer cell lines, HT-29 and Colo-205, were obtained from American Type Culture Collection-ATCC (Rockville, MD). HT-29 was maintained in Dulbecco's modified Eagle's medium (DMEM) while Colo-205 was propagated in Roswell Park Memorial Institute medium (RPMI) 1640. Medium were supplemented with 10% fetal bovine serum and penicillin (100 units/ml), streptomycin (0.1 mg/ml) at 37°C in a humidified atmosphere with 5% CO<sub>2</sub>.

##### 1.2 Tested compounds

The leaves of *G. parva* were collected at Sakaerat, Wang Nam Khieo district, Nakorn Ratchasima province, Thailand in December 2007. A voucher (NSR 510209) has been deposited at the Department of Pharmacognosy and Pharmaceutical Botany, Faculty of Pharmaceutical Sciences, Chulalongkorn University, Thailand. The GPE is a part of solvent partitioned extraction method as mentioned before (8). Ethyl acetate extract from the leaves of *Glycosmis parva* (GPE) were prepared and identified by Associate Professor Nijsiri Ruangrunsi, College of Public Health Sciences, Chulalongkorn University, Bangkok, Thailand. The extract was prepared by partitioned extraction method. Briefly, the dried leaves of *Glycosmis parva* were grounded and macerated with methanol. Afterward, the methanol extract was partitioned with hexane to obtain the hexane extract. The remaining layers were added with water and further partitioned with ethyl acetate (EtOAc) to yield the EtOAc extract (8). All solvents were removed by evaporation. The extract in solid form was stored at -20°C prior to use. The percentage yield of EtOAc extract was 2.08% of dried *G. parva* leaves. Repeated chromatographies of the GPE using the solvent systems of MeOH-CHCl<sub>3</sub>, acetone-hexane, and acetone-CHCl<sub>3</sub> led to the isolation of one acridone alkaloid and four sulfur-containing propanamide derivatives. 0.2% dimethylsulfoxide

(DMSO) was used as a negative control. Standard cytotoxic drugs including cisplatin, oxaliplatin, and 5-FU were used as positive controls.

Table 3: Chemical components found in the EtOAc extract from the leaves of *G. parva* (80)

Main Structure	Chemical name	% in GPE	% of Total
Acridone alkaloids	Arborinine	0.12	65.04
Sulfur-containing propanamide derivatives	S-deoxydihydroglyparvin	0.03	14.81
	S-deoxytetrahydroglyparvin	0.02	7.95
	Glyparvin-A	0.02	7.95
	Dihydroglyparvin	0.01	4.24

### 1.3 Chemicals

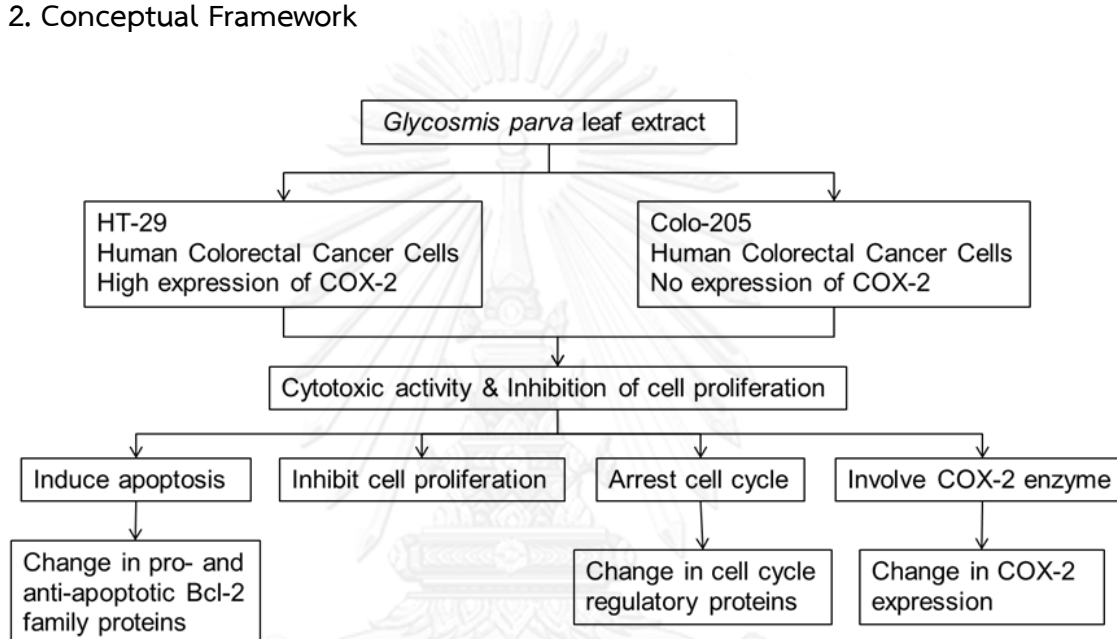
The following reagents and reagent kits were used in this study; DMEM medium (Gibco, USA), RPMI-1640 (Gibco, USA), fetal bovine serum (Gibco, USA), L-glutamine (Gibco, USA), penicillin/streptomycin (Gibco, USA), Trypsin/EDTA (Gibco, USA), 0.4% trypan blue dye (Sigma, USA), DMSO (Sigma, USA), Resazurin (Sigma, USA), annexin V apoptosis detection kit (Invitrogen, USA), Propidium Iodide (Santa Cruz Biotechnology, USA), TRIzol Reagent (Invitrogen, USA), diethyl pyrocarbonate (DEPC) (Molekula, UK), absolute ethanol (Merck, Germany), Improm-II reverse transcription system (Promega, USA), TaqMan<sup>®</sup> Gene Expression System (Applied Biosystems, USA).

### 1.4 Equipments and Instruments

The following equipments and instruments were used in this study; analytical balance (GMPH, Satorius, Germany and UMT2, Mettler Toledo, Switzerland), autopipette (Gilson, USA), biohazard laminar flow hood (Science, Germany and Labconco, USA), centrifuge (Hettich, USA and Eppendorf, Germany), autoclave (Hirayama, Japan), pH meter (Mettler Tuledo, Switzerland), refrigerator 4°C and -20°C (Sanyo, Japan), Incubator (Thermo, USA), Light microscope (Nikon, Japan), vortex mixer (Scientific Industries, USA), fluorescence flow cytometer (BD Biosciences, USA),

PCR thermal cycler (Eppendorf, Germany), StepOnePlus™ Real-Time PCR System (Applied Biosystems, USA), T25 tissue culture flasks (Corning, USA), 96-well microplates (Corning, USA), and 6-well culture plates (Corning, USA). MicroAmp Fast 8-tube strip (0.1 ml) (Applied Biosystems, USA), MicroAmp optical 8-cap strip (Applied Biosystems, USA), Scepter™ Handheld Automated Cell Counter (Millipore, USA).

## 2. Conceptual Framework



## 3. Methods

All assays were done at least three independent experiments in duplication or triplication. 0.2% DMSO was used as a negative control. Cells in the exponential growth phase with over 95% viability were used in the experiments.

### 3.1 Preparation of the stock solutions of GPE

The stock solutions were prepared at the concentration of 50 mg/ml in 100% DMSO and stored at  $-20^{\circ}\text{C}$  until use. Prior to use, the stock solutions were diluted with sterile double-distilled water to the final concentration of 1000  $\mu\text{g/ml}$  in 2% DMSO. The solutions then were diluted with 2% DMSO to the desired concentration. They were used to treat cells at the ratio of 1:10 which gave in the final solvent concentration constantly at 0.2% DMSO.

### 3.2 Determination of baseline COX-2 mRNA expression

The level of baseline COX-2 mRNA expression in human colorectal cancer cells, HT-29 and Colo-205, were determined by reverse transcription polymerase chain reaction (RT-PCR) and followed by agarose gel electrophoresis. Briefly, the total RNA of each cell lines were extracted using Trizol<sup>®</sup> reagent and reverse transcribed to cDNA by Improm II<sup>™</sup> reverse transcription system. PCR amplification of COX-2 gene were done using PCR thermal cycler and the house keeping gene, GAPDH, was used as a loading control. Then, the PCR products were analyzed by electrophoresis in 1.5% agarose gel, stained the gel with ethidium bromide and determined the band densities by gel documentation. The detailed method is the same as method described in section 3.8.

### 3.3 Determination of cytotoxic activities of GPE

The indicator dye resazurin is used to measure the metabolic capacity of cells—an indicator of cell viability. Viable cells have the ability to reduce resazurin (dark-blue, maximum absorption wavelength 605 nm) into resorufin (pink, maximum absorption wavelength 573 nm). Nonviable cells rapidly lose metabolic capacity thus do not reduce the indicator dye.

1. Human colorectal cancer cells, HT-29 and Colo-205, were harvested by trypsinization and re-suspended at a final concentration of  $5 \times 10^4$  cells/ml in fresh medium with 10% FBS.
2. Aliquots of 90  $\mu$ l cell suspension were seeded in 96-well plates and allowed to attach for 24 hours.
3. The cells were treated with a range of GPE concentrations (6.25, 12.5, 25, 50, 100  $\mu$ g/ml) using standard cytotoxic drugs and 0.2% DMSO as positive and negative control, respectively.
4. After 24, 48 and 72 hours, 5  $\mu$ l of a 1 mg/ml resazurin solution were added to each well and the plates were incubated for 6, 3, and 2 hours, respectively.
5. The absorbance of individual wells was determined at 570 and 600 nm using a microplate reader.



6. The percentages of cytotoxicity of the tested compound were calculated using the following formula

$$\% \text{ Cytotoxic} = \frac{(\Delta \text{ OD}_{(\text{negative control})} - \Delta \text{ OD}_{(\text{sample})})}{\Delta \text{ OD}_{(\text{negative control})}} \times 100$$

Where  $\Delta \text{ OD} = \text{Absorbance } 570 \text{ nm} - \text{Absorbance } 600 \text{ nm}$

The inhibition concentration 50 ( $\text{IC}_{50}$ ) was determined for each time point. The 3 concentrations of GPE were chosen for the next experiments.

### 3.4 Determination of apoptotic induction

Induction of apoptosis represents a most potent cellular mechanism against cancer. One biochemical feature of apoptotic cell death is the externalization of the normal inward-facing phosphatidylserine of the cell's lipid bilayer to the outer layers of the plasma membrane which result in the early phagocytic recognition of apoptotic cells by adjacent cells, permitting quick phagocytosis with minimal compromise to the surrounding tissue.

Annexin V is a recombinant phosphatidylserine-binding protein that interacts strongly and specifically with phosphatidylserine residues and can be used for the detection of apoptosis (90).

1. Human colorectal cancer cells, HT-29 and Colo-205, were harvested by trypsinization and re-suspended at a final concentration of  $5 \times 10^4$  cells/ml in fresh medium with 10% FBS.
2. Aliquots of 3.6 ml cell suspension were seeded in 6-well plates and allowed to attach for 24 hours.
3. The cells were treated with a range of GPE concentrations (25, 50, 100  $\mu\text{g/ml}$ ) using standard cytotoxic drugs and 0.2% DMSO as positive and negative control, respectively, for 18 and 24 hours.
4. The floated cells in the supernatant were collected. The remaining attached cells were harvested by trypsinization. The media were removed by centrifugation at 1,500 rpm, 25°C for 5 minutes.
5. Cell pellets were washed twice with 500  $\mu\text{l}$  cold PBS, and collected the cells by centrifugation at 1,500 rpm, 4°C for 5 minutes.

6. Cell pellets were re-suspended in 100  $\mu$ l assay buffer and transferred in to flow cytometer tube
7. Cells were stained with 1  $\mu$ l annexin V-FITC and 1  $\mu$ l of 0.05  $\mu$ g/ml propidium iodide (PI).
8. The stained cells were incubated in the dark at room temperature for 15 minutes.
9. The 400  $\mu$ l of assay buffer were added. Cells suspensions at  $3 \times 10^4$  cells/sample were immediately analyzed using fluorescence flow cytometer.
10. The type of cells were determined as following; viable cells is annexin V-FITC<sup>-</sup>/PI<sup>-</sup> cells, early apoptotic cells is annexin V-FITC<sup>+</sup>/PI<sup>-</sup> cells, necrotic cells is annexin V-FITC<sup>-</sup>/PI<sup>+</sup> cells, and non-apoptotic cells, late apoptotic or secondary necrotic cells are annexin V-FITC<sup>+</sup>/PI<sup>+</sup> cells.

### 3.5 Determination of anti-proliferative effect of GPE

Anti-proliferative effects of the GPE against human CRC cells were determined using automated cell counter.

1. Human colorectal cancer cells, HT-29 and Colo-205, were harvested by trypsinization and re-suspended at a final concentration of  $5 \times 10^4$  cells/ml in fresh medium with 10% FBS.
2. Aliquots of 3.6 ml cell suspension were seeded in 6-well plates and allowed to attach for 24 hours.
3. The cells were treated with a range of GPE concentrations (25, 50, 100  $\mu$ g/ml) using standard cytotoxic drugs and 0.2% DMSO as positive and negative control, respectively, for 24 hours (91-93).
4. The floated cells in the supernatant were collected. The remaining attached cells were harvested by trypsinization. The media were removed by centrifugation at 1,500 rpm, 25°C for 5 minutes.
5. Cell pellets were washed twice with 500  $\mu$ l cold PBS, and collected the cells by centrifugation at 1,500 rpm, 4°C for 5 minutes.
6. Cell pellets were re-suspended in 500  $\mu$ l cold assay buffer. The number of cells in each well were then counted using Scepter™ Handheld Automated Cell Counter.

7. The percentages of cell proliferation were calculated compared to the vehicle control using the following formula;

$$\% \text{ Proliferation} = \frac{\text{Total cell count}_{(\text{Sample})}}{\text{Total cell count}_{(\text{Negative control})}} \times 100$$

### 3.6 Determination of the effect of GPE on the expression of genes in BCL-2 family

The expressions of genes in BCL-2 family were determined by quantitative RT-PCR as following procedures. There were three main steps in this experiment.

1. Preparation of total RNA
2. Conversion of RNA to cDNA
3. Quantitative RT-PCR

#### *Preparation of total RNA*

Total RNA from HT-29 and Colo-205 treated cells were isolated using TRIzol<sup>®</sup> reagent according to the manufacturer's protocol.

1. Human colorectal cancer cells, HT-29 and Colo-205, were harvested by trypsinization and re-suspended at a final concentration of  $5 \times 10^4$  cells/ml in fresh medium with 10% FBS.
2. Aliquots of 3.6 ml cell suspension were seeded in 6-well plates and allowed to attach for 24 hours.
3. The cells were treated with a range of GPE concentrations (25, 50, 100  $\mu\text{g/ml}$ ) using standard cytotoxic drugs and 0.2% DMSO as positive and negative control, respectively, for 24 hours.
4. The floated cells in the supernatant were collected. The media were removed by centrifugation at 1,500 rpm, 25°C for 5 minutes.
5. The remaining attached cells were added with 500  $\mu\text{l}$  of TRIzol<sup>®</sup> reagent per well. Cells were lysed and homogenized by pipetting the cells up and down several times. The floated cells were also lysed with TRIzol<sup>®</sup> reagent.

6. All the cell lysate were transferred to 1.5 ml eppendorf tubes and incubate for 5 minutes at room temperature to permit complete dissociation of the nucleoprotein complex.
7. Phase separations were performed by adding 0.2 ml of chloroform per 1 ml of TRIzol<sup>®</sup> Reagent used for homogenization. The tubes must be cap securely and shaken vigorously by hand for 15 seconds. Then, the tubes were incubated for 2–3 minutes at room temperature.
8. The samples were centrifuged at 12,000 × g for 15 minutes at 4°C. The mixtures were separated into a lower red phenol-chloroform phase, an interphase, and a colorless upper aqueous phase. RNA remains exclusively in the aqueous phase. The upper aqueous phase is about 50% of the total volume.
9. The aqueous phase was removed of the sample by angling the tube at 45° and pipetting the solution out. Avoid drawing any of the interphase or organic layer into the pipette when removing the aqueous phase. The aqueous phase was placed into a new tube and proceeds to the RNA Isolation Procedure.
10. Isopropanol was used to precipitate RNA. 0.5 ml of 100% isopropanol was added to the aqueous phase, per 1 ml of TRIzol<sup>®</sup> Reagent used for homogenization. The tubes were incubated at room temperature for 10 minutes and centrifuged at 12,000 × g for 10 minutes at 4°C. The RNA is often invisible prior to centrifugation, and forms a gel-like pellet on the side and bottom of the tube.
11. RNA pellet were subjected to RNA wash. The supernatant was removed from the tube, leaving only the RNA pellet. The pellets were washed with 1 ml of 75% ethanol per 1 ml of TRIzol<sup>®</sup> Reagent used in the initial homogenization.
12. The samples were briefly vortexed and centrifuged the tube at 7500 × g for 5 minutes at 4°C. The wash was discarded.
13. The RNA pellets were air dried for 5–10 minutes. Do not allow the RNA to dry completely, because the pellet can lose solubility.
14. The RNA pellets were resuspended in 0.1% DEP-C treated water 10–20 µl by passing the solution up and down several times through a pipette tip.

15. The RNA concentration and possible contamination were determined by measuring the absorbance at 260 and 280 nm. The RNA samples should have ratio of  $A_{260/280}$  more than 1.8.
16. The total RNA samples were stored at  $-70^{\circ}\text{C}$  prior to converse to cDNA.

#### *Conversion of RNA to cDNA*

Total RNA samples were converted to cDNA using Improm-II<sup>TM</sup> reverse transcription system according to manufacturer's protocol.

1. Place sterile 0.2 ml PCR tube on ice. Thaw the experimental RNA on ice and return any unused portion to the freezer as soon as aliquots are taken.
2. On ice, combine the experimental RNA 1  $\mu\text{g}$ , 1  $\mu\text{l}$  of the Oilgo(dT)<sub>15</sub> primer 0.5  $\mu\text{g}/\mu\text{l}$  and Nuclease-Free Water for a final volume of 5 $\mu\text{l}$  per reaction. Mix and spin each tube to maintain the original volume.
3. Close each tube of RNA tightly. Place the tubes into a water bath at  $70^{\circ}\text{C}$  for 5 minutes. Immediately chill on ice for at least 5 minutes.
4. Spin each tube for 10 seconds in a microcentrifuge to collect the condensate and maintain the original volume. Keep the tubes closed and on ice until the reverse transcription reaction mix is added.
5. Prepare the reverse transcription reaction mix by combining the following components of the ImProm-II<sup>TM</sup> Reverse Transcription System in a sterile 1.5 ml microcentrifuge tube on ice. Prepare sufficient mix to allow 15 $\mu\text{l}$  for each cDNA synthesis reaction to be performed. Determine the volumes needed for each component, and combine them in the order listed. Vortex gently to mix, and keep on ice prior to dispensing into the reaction tubes. The reverse transcription reaction mix contains the following components

1) Nuclease-Free Water (to a final volume of 15 $\mu\text{l}$ )	7.3	$\mu\text{l}$
2) ImProm-II <sup>TM</sup> 5X Reaction Buffer	4.0	$\mu\text{l}$
3) 25 mM $\text{MgCl}_2$ (final concentration 1.5 mM)	1.2	$\mu\text{l}$
4) dNTP Mix (final concentration 0.5mM each dNTP)	1.0	$\mu\text{l}$
5) Recombinant RNasin <sup>®</sup> Ribonuclease Inhibitor (20 u)	0.5	$\mu\text{l}$
6) ImProm-II <sup>TM</sup> Reverse Transcriptase	1.0	$\mu\text{l}$

Final volume 15.0  $\mu\text{l}$

6. Add 15  $\mu\text{l}$  aliquots of the reverse transcription reaction mix to each reaction tube on ice for a final reaction volume of 20  $\mu\text{l}$  per tube. Be careful to prevent cross-contamination.
7. Place the reaction tube in the PCR thermal cycler machine to generate cDNA using the following conditions; annealing at 25°C for 5 minutes, extension at 42°C for 1:30 hour, and inactivate Reverse Transcriptase at 70°C for 15 minutes. The cDNA samples were stored at -20°C to use as the template for determining gene expression.

#### Quantitative RT-PCR

Quantitative RT-PCR were carried out for *BCL-2* (Hs00608023\_m1), *BCL-XL* (Hs00236329\_m1), *BAK* (Hs00832876\_g1) and *BAX* (Hs00180269\_m1) genes in each cDNA sample using TaqMan<sup>®</sup> Gene Expression System (Applied Biosystem, USA).

1. Thaw on ice, completely resuspend by gently vortexing, then briefly centrifuge to bring liquid to the bottom of the tube of 20X TaqMan<sup>®</sup> Gene Expression Assays (containing specific primers and probes for genes of interest) and cDNA samples. Mix the 2X TaqMan<sup>®</sup> Gene Expression master mix reagent (containing Taq Polymerase and reaction buffer) by gently swirling the bottle.
2. The assays were done in 10  $\mu\text{l}$  singleplex reactions. For each sample, pipet the following into a nuclease free 1.5-ml microcentrifuge tube:

1) 20X TaqMan <sup>®</sup> Gene Expression Assays	0.5	$\mu\text{l}$
2) 2X TaqMan <sup>®</sup> Gene Expression master mix	5	$\mu\text{l}$
3) cDNA template 50 ng	1	$\mu\text{l}$
4) RNase-free water	3.5	$\mu\text{l}$
Final volume	10.0	$\mu\text{l}$

3. Cap the tube and invert it several times to mix the reaction components. Centrifuge the tube briefly.
4. Load the microcentrifuge tubes into the Applied Biosystems StepOnePlus™ Real-Time PCR System.

5. Create an experiment for the run using the parameter values as following; standard run, reaction volume 10  $\mu$ l, thermal cycling conditions: pre-incubation at 95°C for 10 min, then cycling for 45 cycles of 95°C for 15 seconds and 60°C for 1 minute. Run the experiment.
6. The quantification of gene expression levels were used relative quantitation method where GAPDH was an endogenous control and 0.2% DMSO was a negative control as following equation;

$$\text{Relative Quantitation (RQ)} = 2^{-\Delta\Delta C_T}$$

Where  $C_T$  = cycle threshold

7. The ratios of pro-apoptotic/anti-apoptotic expression were calculated from the following equation;

$$\text{Ratio of Proapoptotic/Anti-apoptotic} = \frac{\text{RQ}_{\text{BAK}} + \text{RQ}_{\text{BAX}}}{\text{RQ}_{\text{BCL-2}} + \text{RQ}_{\text{BCL-XL}}}$$

### 3.7 Determination of the effect of GPE on the cell cycle

The nuclear DNA content of a cell can be quantitatively measured by flow cytometry. Initially, a fluorescent dye that binds stoichiometrically to the DNA is added to a suspension of permeabilized single cells or nuclei. The principle is that the stained material has incorporated an amount of dye proportional to the amount of DNA. The stained material is then measured in the flow cytometer and the emitted fluorescent signal yields an electronic pulse with a height (amplitude) proportional to the total fluorescence emission from the cell. Thereafter, such fluorescence data are considered a measurement of the cellular DNA content (94). This method of DNA staining utilizes ethanol to fix the cells and permeabilize the membrane, which allows the dye (Propidium Iodide) to enter the cells. Propidium Iodide (PI) is a DNA-binding fluorochrome that intercalates in the double-helix. Ribonuclease-A is used to eliminate the staining of double-stranded RNA.

1. Human colorectal cancer cells, HT-29 and Colo-205, were harvested by trypsinization and re-suspended at a final concentration of  $5 \times 10^4$  cells/ml in fresh medium with 10% FBS.

2. Aliquots of 3.6 ml cell suspension were seeded in 6-well plates and allowed to attach for 24 hours.
3. The cells were treated with a range of GPE concentrations (25, 50, 100 µg/ml) using standard cytotoxic drugs and 0.2% DMSO as positive and negative control, respectively, for 1 hour and further incubated for 24 hours in a fresh medium without GPE (95).
4. The floated cells in the supernatant were collected. The remaining attached cells were harvested by trypsinization. The media was removed by centrifugation at 1,500 rpm, 25°C for 5 minutes.
5. Cell pellets were washed twice with 500 µl cold PBS, and the cells were collected by centrifugation at 1,500 rpm, 4°C for 5 minutes.
6. The cells were re-suspended in 150 µl cold PBS and 350µl absolute ethanol was slowly added with continuously shaking to fix the cells and incubated at -20°C for 15 minutes.
7. Fixed cells were washed twice with 500 µl cold PBS to remove the remaining ethanol and the cells were collected by centrifugation at 1,500 rpm, 4°C for 5 minutes.
8. The cells were re-suspended in 500 µl assay buffer.
9. Five µl of 4 mg/ml RNase was added to eliminate RNA and incubated for 30 min at room temperature.
10. Cells were stained with 5 µl of 0.05µg/ml PI and incubated in the dark for 15 min at room temperature.
11. Cell cycle patterns of  $3 \times 10^4$  cells/sample were immediately analyzed using fluorescence flow cytometer.
12. The cell cycle analysis was evaluated using FCS Express 4 Image Cytometry software.



### 3.8 Determination of the effect of GPE on the mRNA expression of cell cycle regulatory genes

The mRNA expression of cell cycle regulatory genes including cyclin A, cyclin B1, cyclin D1, cyclin E, p21 and p53 were performed in GPE treated HT-29 and Colo-205 cells using RT-PCR. There were four main steps in this experiment.

1. Preparation of total RNA
2. Conversion of RNA to cDNA
3. RT-PCR
4. Agarose gel electrophoresis

#### *Preparation of total RNA*

1. Human colorectal cancer cells, HT-29 and Colo-205, were harvested by trypsinization and re-suspended at a final concentration of  $5 \times 10^4$  cells/ml in fresh medium with 10% FBS.
2. Aliquots of 3.6 ml cell suspension were seeded in 6-well plates and allowed to attach for 24 hours.
3. The cells were treated with a range of GPE concentrations (25, 50, 100  $\mu\text{g/ml}$ ) using standard cytotoxic drugs and 0.2% DMSO as positive and negative control, respectively, for 1 hour and further incubated for 24 hours in a fresh medium without GPE (95).
4. Repeat the *Preparation of total RNA* method in section 3.6 steps 4-16.

#### *Conversion of RNA to cDNA*

The method to converse the total RNA to cDNA was the same as section 3.6 steps 1-7 in the section *Conversion of RNA to cDNA*.

#### *RT-PCR*

1. Add the following components to a sterile 0.2-ml microcentrifuge tube. The reaction volumes were 25  $\mu\text{l}$ . Prepare a master mix of common components for multiple reactions.
  - 1) 10X PCR Buffer, Minus Mg 2.5  $\mu\text{l}$

2) 10 mM dNTP mixture (final concentration 0.2 mM each)	0.5	μl
3) 50 mM MgCl <sub>2</sub> (final concentration 1.5 mM)	0.75	μl
4) 10 μM Forward Primer (final concentration 0.2 μM)	0.5	μl
5) 10 μM Reverse Primer (final concentration 0.2 μM)	0.5	μl
6) cDNA Template DNA	1	μl
7) Platinum® Taq DNA Polymerase (1.0 unit)	0.2	μl
8) Autoclaved, distilled water	to 25	μl

2. Cap the tubes, mix, and centrifuge briefly to collect the contents.
3. Incubate tubes in a thermal cycler at 94°C for 2 minutes to completely denature the template and activate the enzyme.
4. Perform 35 cycles of PCR amplification as follows:

Denature	94°C	for 30 seconds
Anneal	T <sub>m</sub>	for 30 seconds
Extend	72°C	for 1 minute
5. Perform a final extension at 72°C for 10 minutes.
6. Maintain the reaction at 4°C after cycling. After the cycle is complete, analyze the PCR products or store the amplifications at -20°C until use.

Table 4: Primers for RT-PCR and their annealing temperature.

Gene	Primer sequences	Tm °C	PCR product (bp)
GAPDH	Forward: 5'-AAG GTC GGA GTC AAC GGA TTT GGT-3' Reverse: 5'-ATG GCA TGG ACT GTG GTC ATG AGT-3'	60	530
P53	Forward: 5'-CAT GAG CGC TGC TCA GAT AG-3' Reverse: 5'-CTG AGT CAG GCC CTT CTG TC-3'	56	643
P21	Forward: 5'-GCG ATG GAA CTT CGA CTT TGT-3' Reverse: 5'-GGG CTT CCT CTT GGA GAA GAT-3'	54	352
Cyclin A	Forward: 5'-GTC ACC ACA TAC TAT GGA CAT G-3' Reverse: 5'-AAG TTT TCC TCT CAG CAC TGA C-3'	53	300
Cyclin B1	Forward: 5'-CGG GAA GTC ACT GGA AAC AT-3' Reverse: 5'-AAA CAT GGC AGT GAC ACC AA-3'	55	177
Cyclin D1	Forward: 5'-CTG GCC ATG AAC TAC CTG GA-3' Reverse: 5'-GTC ACA CTT GAT CAC TCT GG-3'	54	482
Cyclin E	Forward: 5'-AAT AGA GAG GAA GTC TGG-3' Reverse: 5'-AGA TAT GAC ACC TGC ATG-3'	55	440
COX-2	Forward: 5'- TTC AAA TGA GAT TGT GGG AAA AT-3' Reverse: 5'- AGA TCA TCT CTG CCT GAG TAT CTT-3'	50	305

#### *Agarose gel electrophoresis*

1. Prepare 1.5% Agarose gel by weighing 1.2 gram of agarose and adding agarose to Erlenmeyer flask. Add 80 ml of 1X TBE to Erlenmeyer flask. Swirl vigorously to thoroughly mix agarose.
2. Put agarose and 1X TBE slurry into microwave. Heat on high for 30 seconds at a time. After 30 seconds, remove from microwave and swirl. Be careful as the mixture was hot. After 1 minute of heating, repeat heating and swirling procedure every 10 seconds until mixture is clear.
3. Place gel tray into casting chamber. Add casting combs into the appropriate slots.

4. Pour the gel when the agarose has cooled to about 55° C. When pouring the gel, avoid creating bubbles as this will prevent current from flowing through the gel.
5. The gel should be allowed to cool for at least 30 minutes until it has set (it will turn whitish and opaque when ready). Gels should be fairly thin, approximately 1/4 to 1/2 inch.
6. Carefully remove combs by pulling them upwards firmly and smoothly in a continuous motion.
7. Place gel and tray into gel rig, with wells on the cathode side. Then, fill gel rig with 1X TBE sufficient to cover the entire gel.
8. Cut a piece of parafilm and place a 2 µl drop of 6X loading dye onto the waxy side for each sample to be loaded.
9. Keeping PCR product samples on ice, pipette up 8 µl of a sample and add the sample to one of the drops of loading dye.
10. Mix the sample and loading dye by pipetting the mixture up and down several times then load the 9 µl of the mixture into a well.
11. Continue loading the rest of the samples, placing 3 µl of DNA ladder at the first well of each row as a marker.
12. Place the lid on the gel box, connecting the electrodes. Connect the electrode wires to the power supply, making sure the positive (red) and negative (black) are correctly connected.
13. Turn on the power supply to 100 volts. Run the gel for 50-60 minutes. Stop the run before the bromophenol blue loading dye front exits the gel.
14. Turn off the power pack, remove the gel and place it in a staining solution with 0.5 µg/ml ethidium bromide 400 ml in 1X TBE for 15 minutes. Then, destain the gel in 1X TBE for 15 minutes.
15. Determine the band densities by gel documentation using Gel Doc™ XR+ System and Image Lab™ software.

### **3.9 Determination of the effect of GPE on the COX-2 mRNA expression**

The effect of GPE on the COX-2 expression was performed only in HT-29 cells which constitutively expressed COX-2. The level of COX-2 mRNA expression were

determined by reverse transcription polymerase chain reaction (RT-PCR) and followed by agarose gel electrophoresis.

#### *Preparation of total RNA*

1. Human colorectal cancer cells, HT-29, were harvested by trypsinization and re-suspended at a final concentration of  $5 \times 10^4$  cells/ml in fresh medium with 10% FBS.
2. Aliquots of 3.6 ml cell suspension were seeded in 6-well plates and allowed to attach for 24 hours.
3. The cells were treated with a range of GPE concentrations (25, 50, 100  $\mu\text{g/ml}$ ) using standard cytotoxic drugs and 0.2% DMSO as positive and negative control, respectively, for 24 hours.
4. The total RNA from HT-29 treated cells were isolated using TRIzol<sup>®</sup> reagent according to the manufacturer's protocol as previously described in section 2.6.

The total RNA samples were subjected to converse to cDNA, RT-PCR, and agarose gel electrophoresis as described in section 3.8.

#### **3.10 Statistical analysis**

All data were presented as mean  $\pm$  S.E. Statistical evaluations were determined by one-way analysis of variance (ANOVA) followed by Tukey's post hoc test. Student t test was used to compare the differences between control and the tested group in the RT-PCR experiment. The differences between the two studied cell lines were assessed by student t test. Any p-value less than 0.05 were considered statistically significant. All assays were performed in duplicate or triplicate of three independent experiments.

## CHAPTER IV

### RESULTS

#### 1. Baseline COX-2 mRNA expression

COX-2 plays a pivotal role in CRC carcinogenesis and disease progression (34; 39). Human colorectal cancer cells, HT-29 and Colo-205, the two cell lines used in this study, were subjected to the determination of baseline COX-2 mRNA expression by RT-PCR. The result has shown that HT-29 cells are constitutively expressed COX-2 while Colo-205 cells lack of COX-2 expression (Figure 14).

The further studies were done using these 2 human colorectal cancer cell lines in order to compare the effect of GPE on different cell lines which carry different COX-2 expression status. Moreover, the involvement of COX-2 on GPE action was investigated in COX-2 expressing cells, HT-29.



Figure 14: Baseline COX-2 expression in HT-29 and Colo-205 cells. HT-29 and Colo-205 cells were harvested and the total RNAs were extracted by TRIZOL<sup>®</sup> Reagent. The RNAs were converted to cDNA by Improm-II<sup>™</sup> reverse transcription system and amplified with the COX-2 primers by RT-PCR. The PCR products were identified by 1.5% agarose gel electrophoresis and analyzed by gel documentation. GAPDH was used as a loading control.

## 2. Effect of GPE on human colorectal cancer cell expressing COX-2, HT-29 cells.

The anti-cancer effects of GPE were determined on human colorectal cancer cell that constitutively expressed COX-2, HT-29 cells as following.

### 2.1 Cytotoxic activities of GPE

GPE was initially tested for its cytotoxic effects against HT-29 cells by resazurin reduction assay. The cells were exposed to 6.25 - 100 $\mu$ g/ml of GPE for 24, 48 and 72 h. Standard cytotoxic drugs recommended for the treatment of CRC, 5FU and Oxaliplatin were used as positive controls in this experiment.

As shown in Figure 15, 50 and 100 $\mu$ g/ml of the extract significantly exhibited cytotoxicity against HT-29 cells. The half maximum inhibition concentrations ( $IC_{50}$ ) of GPE against HT-29 were  $69.49 \pm 2.04$ ,  $55.89 \pm 1.86$  and  $48.94 \pm 2.99$   $\mu$ g/ml at 24, 48 and 72 h, respectively (Figure 16). The cytotoxic effect of the extract was seen in a dose-dependent and time-dependent manner. However, the dose-dependency was more pronounced. At 24 h treatment time, GPE showed higher cytotoxic activities against HT-29 when compared to that of standard cytotoxic agents, 5FU and Oxaliplatin. The  $IC_{50}$  of 5FU against HT-29 were  $>1,000$ , 45.06, and 7.22  $\mu$ M at 24, 48 and 72 h, respectively. Oxaliplatin showed the  $IC_{50}$  of 849.34, 4.44, and 2.35  $\mu$ M against HT-29 at 24, 48 and 72 h, respectively.

According to the  $IC_{50}$  of GPE, three concentrations at 25, 50 and 100 $\mu$ g/ml were chosen for the rest of the experiments.

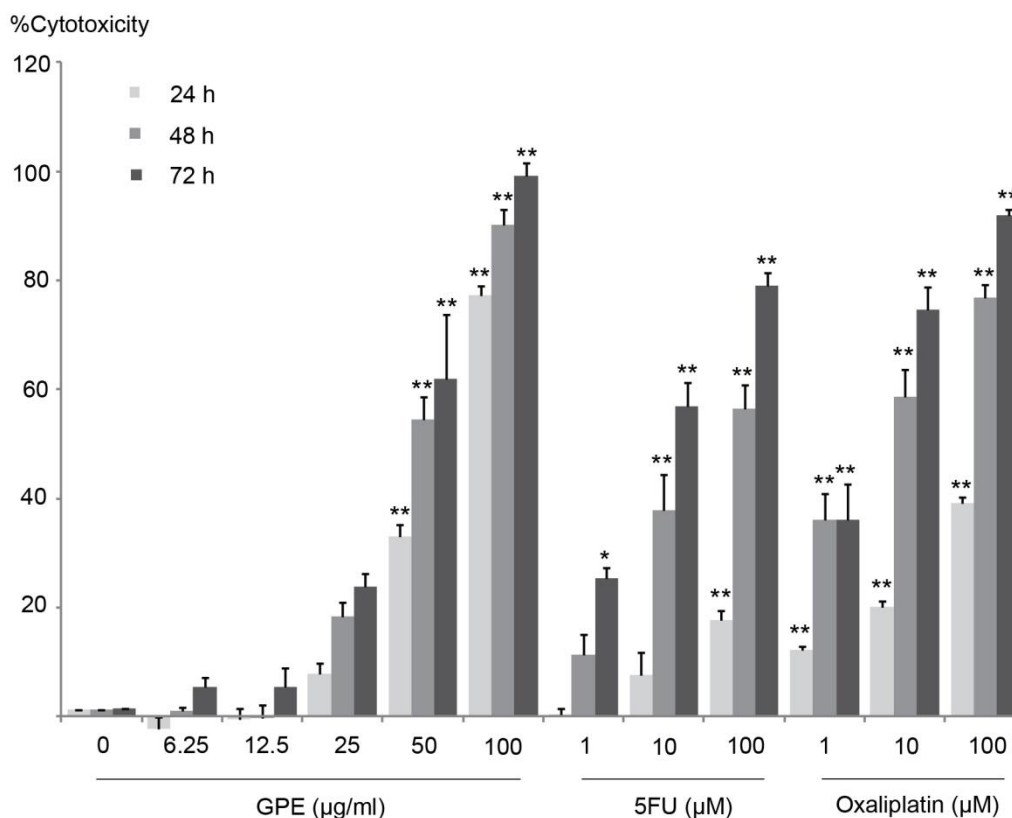


Figure 15: Cytotoxic effects of GPE against HT-29 cells. The cells were treated with GPE at 6.25 - 100 µg/ml, 5FU at 1 - 100 µM and Oxaliplatin at 1 - 100 µM for 24, 48 and 72 hours. The cytotoxic effects were determined by resazurin reduction assay. The data represent the means  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO).



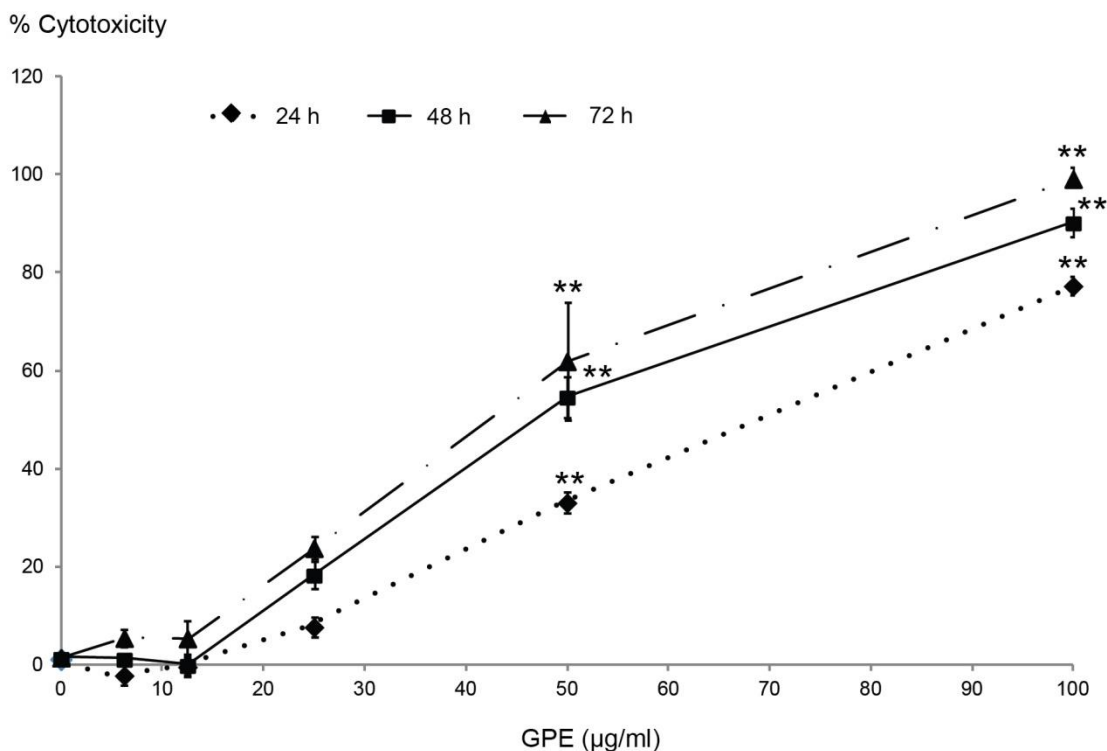


Figure 16: Line graphs show cytotoxicity of GPE against HT-29 cells. The cells were treated with GPE at 6.25 - 100 µg/ml for 24, 48 and 72 h. The cytotoxic effects were determined by resazurin reduction assay. The data represent the means  $\pm$  S.E. of three independent experiments. The percentages of cytotoxicity of GPE were plotted against GPE concentration. Linear trend lines of each treatment time were calculated. The linear equations were used to determine the IC<sub>50</sub> of GPE at each treatment time. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO).

## 2.2 Apoptotic induction effect of GPE

Induction of apoptosis is the main effect of many cytotoxic drugs to induce cancer cell death (96). The apoptotic induction effect of GPE was evaluated against HT-29. The cells were treated with 25, 50 and 100  $\mu\text{g/ml}$  of GPE for 18 and 24 h. Apoptotic cells death were determined by AnnexinV-FITC/PI staining and detected with fluorescence flow cytometer. The patterns of cell death can be categorized into viable cells (AnnexinV-FITC<sup>-</sup>/PI<sup>-</sup>), necrosis cells (AnnexinV-FITC<sup>-</sup>/PI<sup>+</sup>), late apoptosis and/or necrosis cells (AnnexinV-FITC<sup>+</sup>/PI<sup>+</sup>), and apoptosis cells (AnnexinV-FITC<sup>+</sup>/PI<sup>-</sup>). Oxaliplatin and 5FU have been reported to induce colorectal cancer cell death by apoptosis were used as positive controls in this experiment (97; 98).

GPE induced total cell death in a concentration- and time- dependent manner after 18 and 24 h exposures. GPE at 50 $\mu\text{g/ml}$  was significantly induced apoptotic cell death in HT-29 by  $14.79\pm 2.91\%$  and  $20.97\pm 0.91\%$  at 18 and 24 h, respectively. At this concentration of GPE, apoptotic cells death represented the majority type of cell death with 45.14% and 57.71% of total cell death at 18 and 24 h, respectively. On the other hand, 100 $\mu\text{g/ml}$  GPE caused majority of cell death as late apoptosis and/or necrosis stages while cells treated with 25 $\mu\text{g/ml}$  GPE remained viable (Figure 17). Oxaliplatin and 5FU showed a minimal apoptotic cells death. It was due to the short exposure time which caused the majority of cells remained viable. The results were correlated with the cytotoxicity study. The molecular mechanism of GPE on apoptotic induction was further evaluated at the same concentration at 24 h exposure.

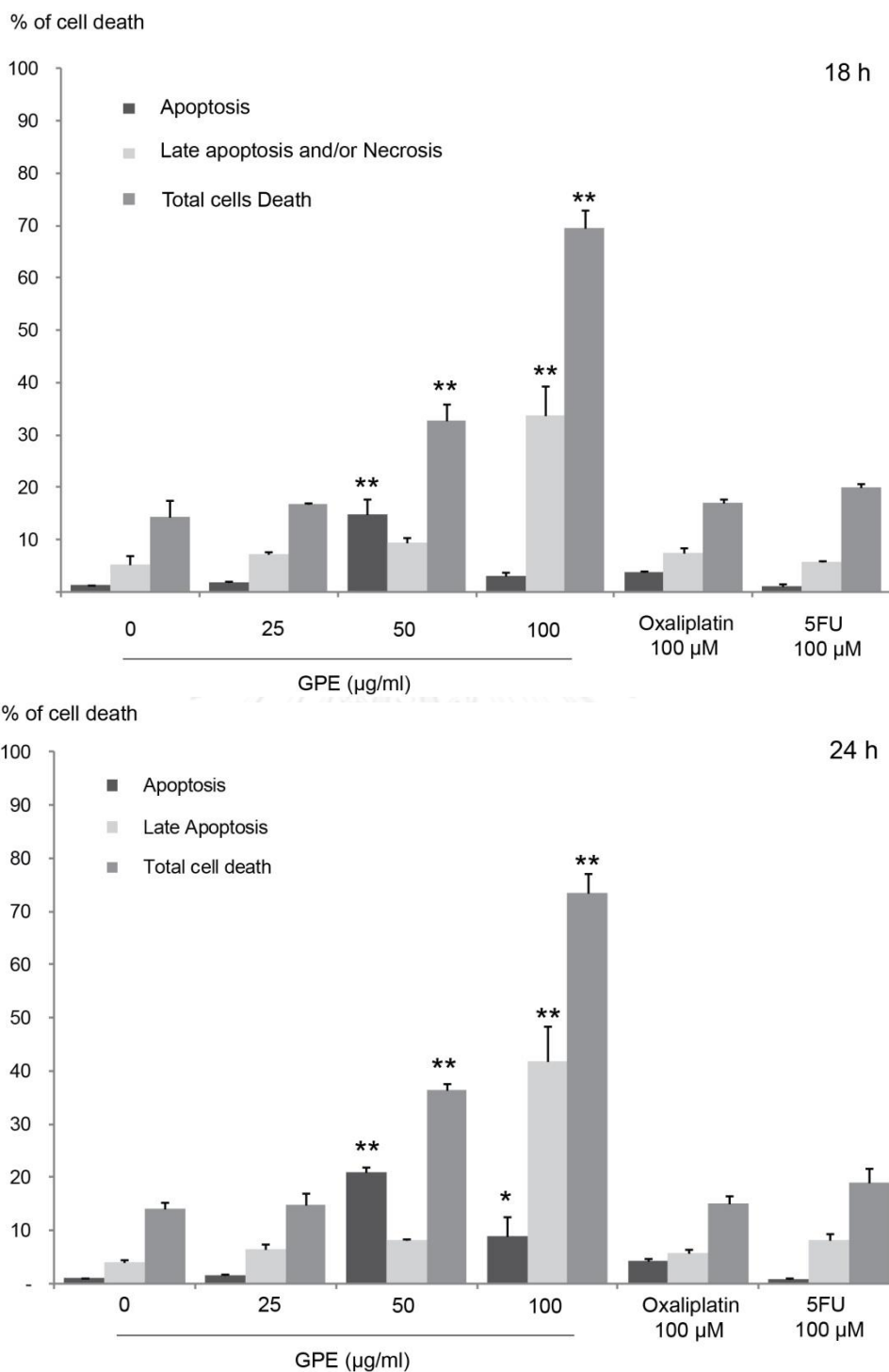


Figure 17: The apoptotic effect of GPE on HT-29 cells. The cells were treated with 25 – 100µg/ml of GPE for 18 (A) and 24 h (B). The percentage of cells labeled with annexin V+/PI-(early apoptotic cells), annexin V+/PI+ (late apoptotic cells/necrotic cells) and total cell death were plotted. Data are the mean values  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO)

### 2.3 Effect of GPE on the expression of genes in BCL-2 family

Apoptosis is regulated by a variety of pro-apoptotic and anti-apoptotic BCL-2 family proteins (90). To evaluate the mechanism by which GPE induced apoptosis in HT-29, the mRNA expression of BCL-2 family genes were assessed by quantitative RT-PCR.

HT-29 cells were treated with 25, 50 and 100  $\mu\text{g/ml}$  of GPE for 24 h, and then the cells were subjected to RNA isolation, cDNA conversion, and PCR amplification of target genes. GPE at 25 and 50  $\mu\text{g/ml}$  increased the expression of pro-apoptotic BAK to  $1.25\pm 0.16$  and  $1.79\pm 0.08$ , respectively while GPE at 100  $\mu\text{g/ml}$  tended to decrease BAK expression. The expression of anti-apoptotic BCL-2 was down-regulated to  $0.68\pm 0.2$ ,  $0.32\pm 0.13$ ,  $0.55\pm 0.28$  following GPE 25, 50, and 100  $\mu\text{g/ml}$  treatment, respectively (Figure 18). Treatment with GPE also tended to decrease both anti-apoptotic BCL-XL and pro-apoptotic BAX. However, the ratio of pro-apoptotic/anti-apoptotic (BAK and BAX/BCL-2 and BCL-XL) were favor apoptotic cell deaths (ratio > 1) across the experimental concentrations with the ratio of  $1.24\pm 0.21$ ,  $2.66\pm 0.48$ , and  $1.37\pm 0.17$  following GPE 25, 50, and 100  $\mu\text{g/ml}$  treatment, respectively (Figure 19).

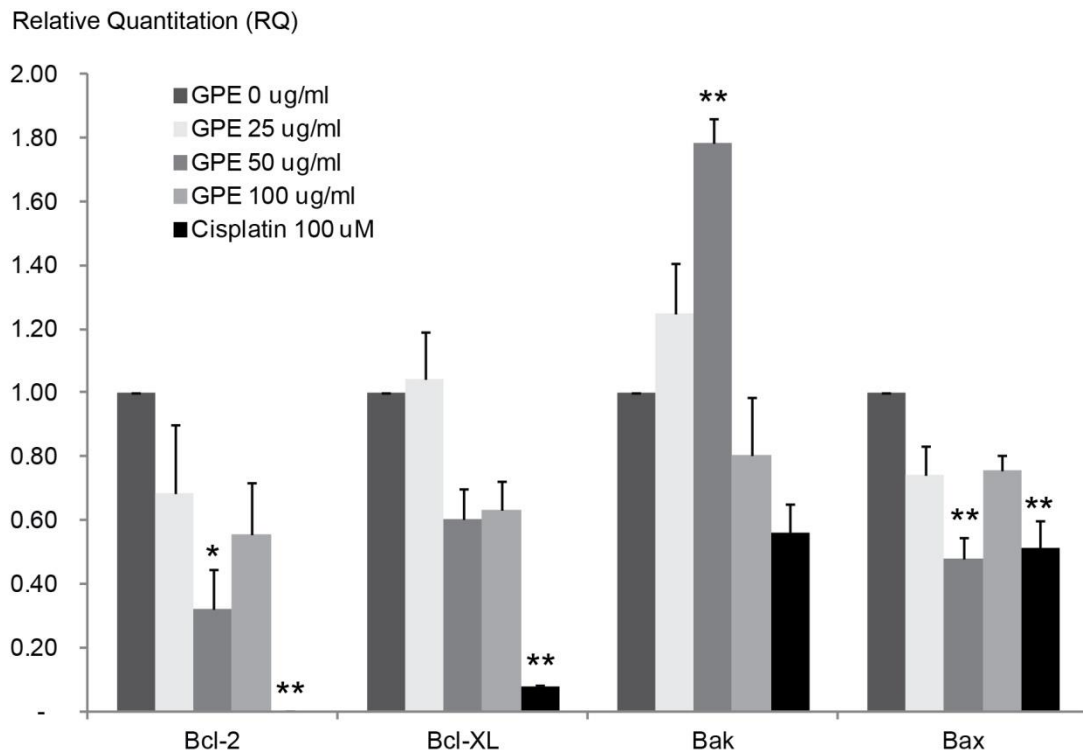


Figure 18: GPE alters the expression of BCL-2 family genes in HT-29 cells. The expressions of BCL-2 family genes were determined by quantitative RT-PCR. The cells were treated with GPE 25 - 100µg/ml for 24 h. The treated cells were subjected to RNA isolation, cDNA conversion and quantitative RT-PCR amplification of BAK, BAX, BCL-2 and BCL-XL genes. The value is given as relative copy number normalized to the endogenous control GAPDH and 0.2% DMSO solvent control. The data represent the means  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO).

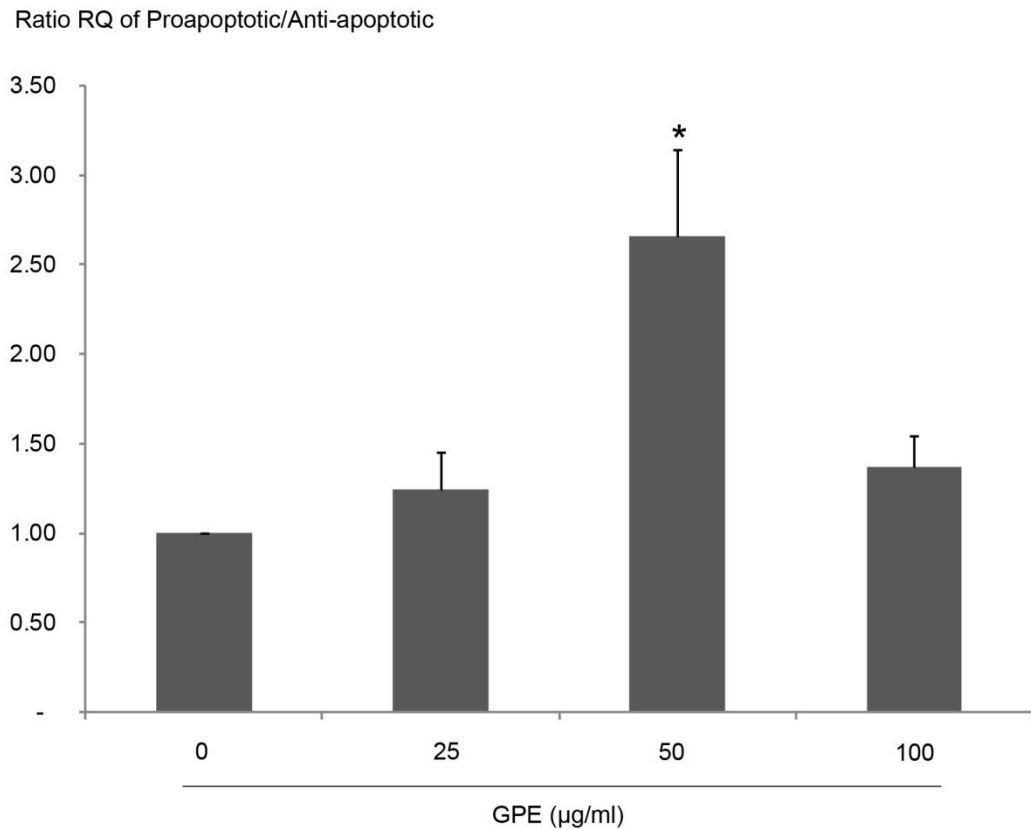


Figure 19: The ratio of pro-apoptotic/anti-apoptotic expression (BAK and BAX/BCL-2 and BCL-XL) in HT-29 treated cells. The cells were treated with GPE 25 - 100µg/ml for 24 h. The treated cells were subjected to RNA isolation, cDNA conversion and quantitative RT-PCR amplification of BAK, BAX, BCL-2 and BCL-XL genes. The ratios of pro-apoptotic/anti-apoptotic expression were calculated. The data represent the means  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO).

## 2.4 Effect of GPE on the cell proliferation

Besides the ability of GPE to induce cancer cell death, its capability to inhibit cancer cell proliferation is also contributed to its cytotoxic activity. Inhibition of HT-29 cell proliferation by GPE was determined by cell counting. Standard cytotoxic drug, 5-FU, was used as a positive control in this experiment.

HT-29 cells were treated with 25, 50 and 100  $\mu\text{g/ml}$  of GPE for 24 h afterward the total cells were counted. The results were expressed as a percentage of cells compared to the untreated control. GPE significantly inhibited HT-29 cell proliferation to  $72.40\pm 5.83\%$ ,  $52.46\pm 6.45\%$ , and  $26.68\pm 6.12\%$  at GPE 25, 50 and 100 $\mu\text{g/ml}$ , respectively when compared to the vehicle control. The effect was seen in a dose-dependent manner (Figure 20). The cell proliferation inhibition effect of GPE against HT-29 was similar to that of 5FU which 100  $\mu\text{M}$  5FU significantly inhibited cell proliferation to  $44.47\pm 0.07\%$ .

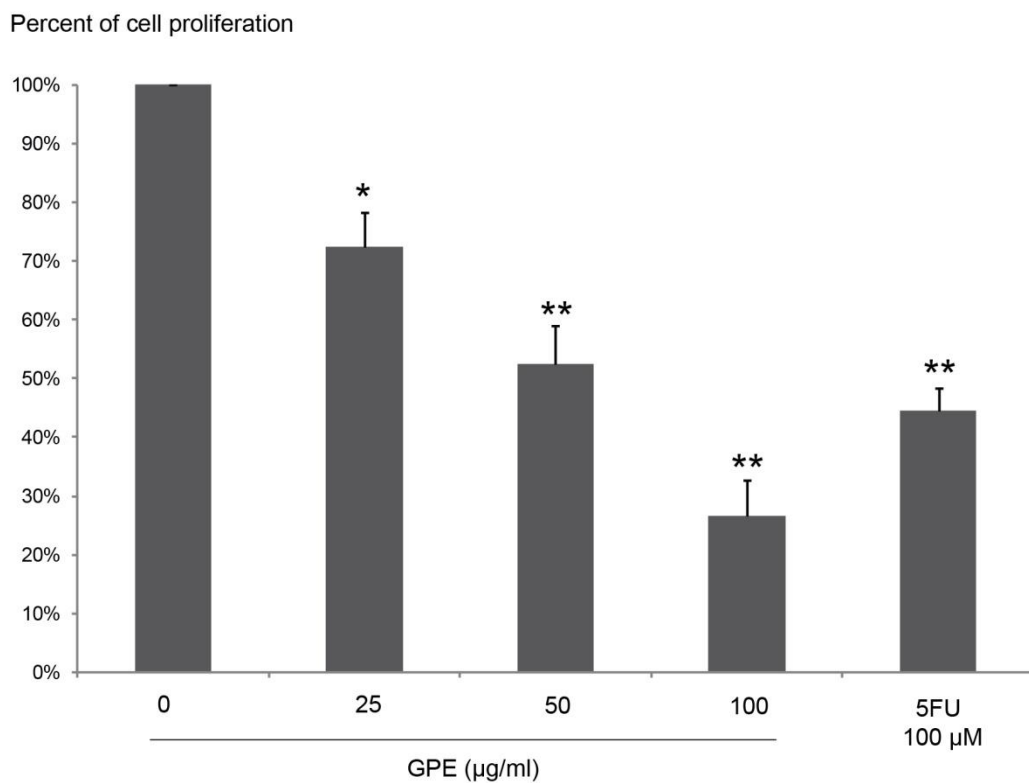


Figure 20: Anti-proliferative effects of GPE against HT-29 cells. The cells were treated with GPE 25 – 100µg/ml for 24 h. Cell proliferation was measured by cell counting using an automated cell counter. The data represent the means  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO).



## 2.5 Effect of GPE on the cell cycle progression

To determine whether the suppression of cell proliferation by GPE resulted from inhibition of cell cycle progression, HT-29 cells were treated with 25, 50 and 100 µg/ml GPE for 1 h and further incubated for 24 h in a fresh medium without GPE. The treated cells were subjected to fluorescence flow cytometric analysis for DNA content (Figure 21). 5FU which arrests the cell cycle at S phase and causes cell accumulation at G1/S phase was used as a positive control.

The effect of GPE on cell cycle pattern of HT-29 cells was difference which depended on the concentration used. GPE at 25µg/ml caused the accumulation of cells at S phase ( $32.00\pm 0.51\%$  to  $44.85\pm 2.59\%$ ) which was associated with a decrease in the cells in G0/G1 phase ( $50.13\pm 1.28\%$  to  $40.95\pm 1.33\%$ ). GPE at 50 and 100µg/ml caused a decrease in the proportion of cells in the S phase (from  $32.0\pm 0.51\%$  to  $26.78\pm 1.76\%$  and  $18.72\pm 3.55\%$ , respectively). In cells treated with 50µg/ml GPE, this decrease was associated primarily with accumulation of cells in the G0/G1 phase (from  $50.13\pm 1.28\%$  to  $60.73\pm 0.98\%$ ). At higher concentration, GPE (100µg/ml) resulted in a significant reduction in the S fraction (from  $32.0\pm 0.51\%$  to  $18.72\pm 3.55\%$ ) with a shift of cells accumulation to the G2/M phases (from  $17.86\pm 1.04\%$  to  $33.53\pm 0.85\%$ ) as shown in Figures 22. The positive control, 5FU, significantly accumulated cells population at S fraction with the reduction of the cells in G0/G1 and G2/M phases. The molecular mechanism of GPE on cell cycle pattern was further evaluated at the same concentration and treatment time.

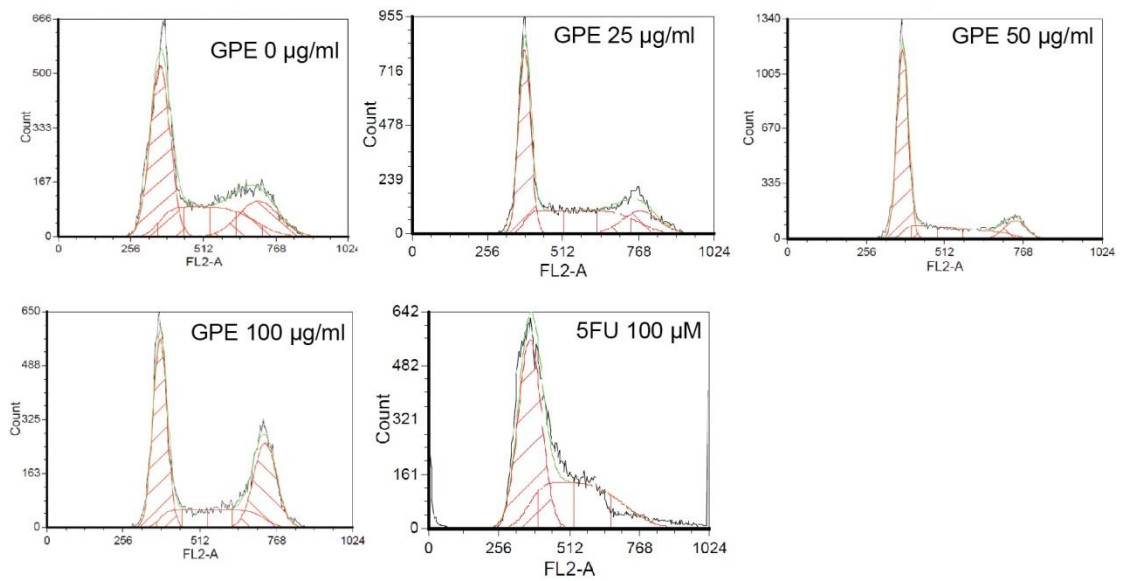


Figure 21: Representative histogram from flow cytometric analysis of HT-29 cell cycle pattern after treatment with GPE. The cells were treated with GPE at 25 - 100µg/ml for 1 h. The treated cells were washed and incubated in fresh medium for additional 24 h. Analysis of cells in the cell cycle was performed by fixing and staining the cells with PI. Cell cycle pattern of  $3 \times 10^4$  cells/sample were analyzed using fluorescence flow cytometer. The cell cycle distributions were evaluated using FCS Express 4 Image Cytometry.

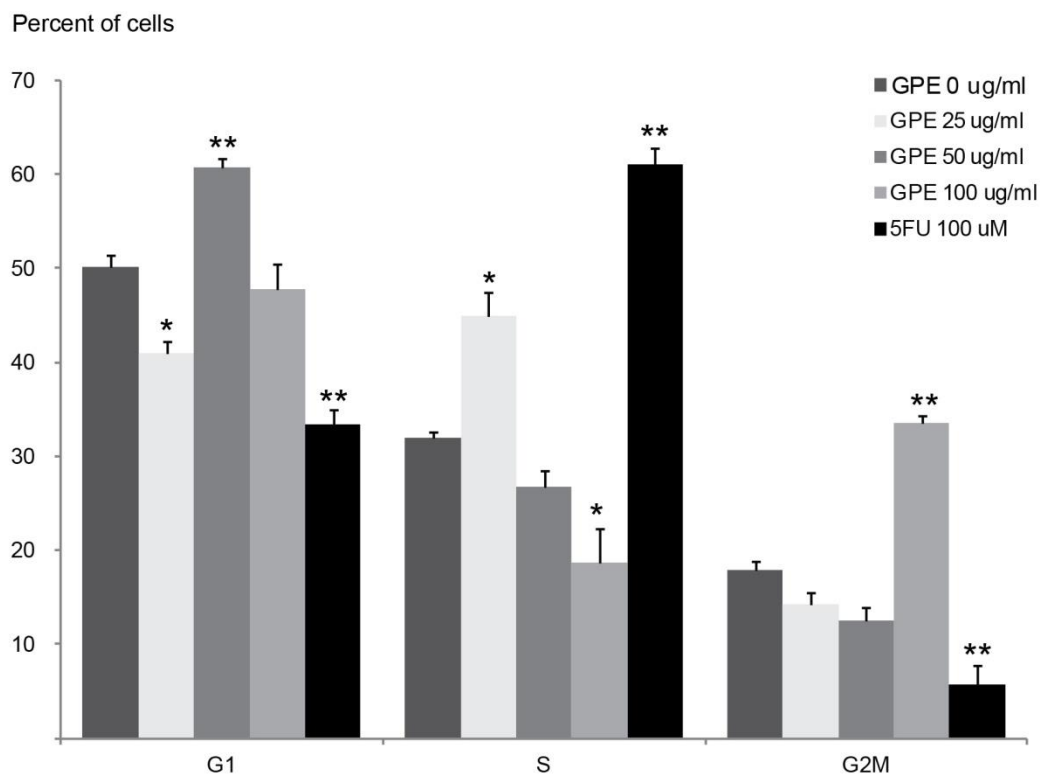


Figure 22: Distribution of HT-29 cells in the cell cycle phases after treatment with GPE. The cells were treated with GPE 25 - 100 $\mu$ g/ml for 1 h. The treated cells were washed and incubated in fresh medium for additional 24 h. Analysis of cells in the cell cycle was performed by fixing and staining the treated cells with PI followed by fluorescence flow cytometry. The percentage of cells in G1, S, and G2M phases are given as means  $\pm$  S.E. of 3 independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO).

## 2.6 Effect of GPE on the mRNA expression of cell cycle regulatory genes

Since GPE changed the cell cycle pattern of HT-29 cells, the alterations in the expression of cell cycle regulatory genes might be involved. HT-29 cells were treated with 25, 50 and 100  $\mu\text{g/ml}$  GPE for 1 h and further incubated for 24 h in a fresh medium without GPE. The treated cells were subjected to RNA isolation, cDNA conversion, PCR amplification of the genes involved in cell cycle progression including cyclin A, cyclin B1, cyclin D1, cyclin E, p21 and p53. PCR products were analyzed by agarose gel electrophoresis. The housekeeping genes, GAPDH, was used as a loading control.

As shown in Figure 23, GPE at the concentration of 50 and 100  $\mu\text{g/ml}$  significantly down-regulated cyclin A and cyclin E expression in a dose-dependent manner. However, the level of cyclin B1, cyclin D1, p21 and p53 remained unchanged.

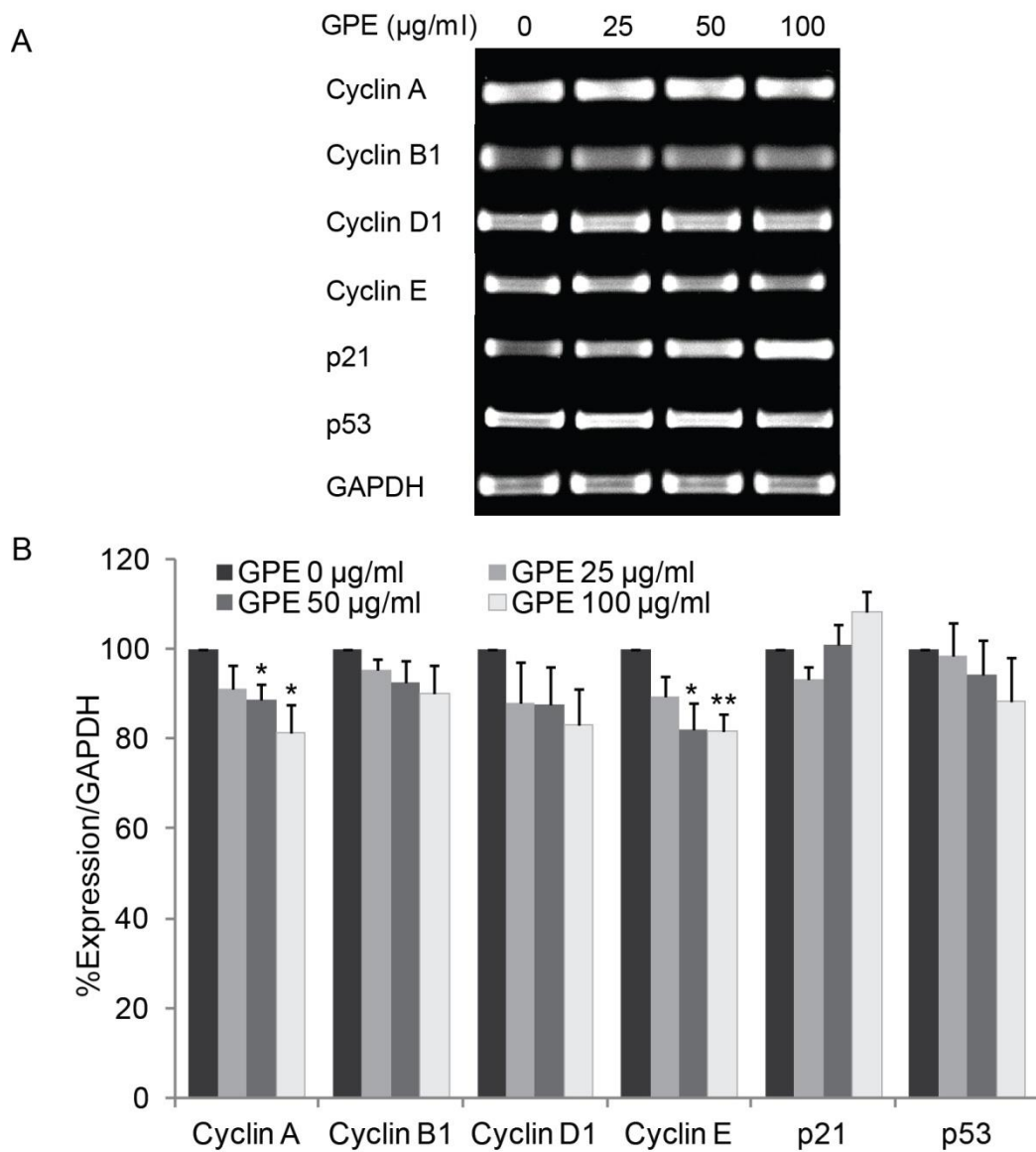


Figure 23: GPE alters the expression of cell cycle regulators genes in HT-29. The cells were treated with GPE 25 - 100 $\mu\text{g/ml}$  for 1 h and incubated with fresh medium without GPE for additional 24 h. Expression of cell cycle regulators were determined by RT-PCR. A) A representative of PCR products of the cell cycle regulatory genes. B) Densitometric analysis of the PCR products relative to GAPDH presented as % of control. The data represent the means  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO).

## 2.7 Effect of GPE on the COX-2 mRNA expression

COX-2 is an important growth factor involved the carcinogenesis of colorectal cancer (15). HT-29 cells were treated with 25, 50 and 100  $\mu\text{g/ml}$  GPE for 24 h. The treated cells were subjected to RNA isolation, cDNA conversion, PCR amplification of the COX-2 genes. PCR products were analyzed by agarose gel electrophoresis. The housing keeping genes, GAPDH, was used as a loading control.

The effect of GPE on COX-2 mRNA expression was investigated in HT-29 cells which constitutively expressed COX-2. GPE at the concentration of 25 – 100 $\mu\text{g/ml}$  significantly inhibited COX-2 mRNA expression after 24 h of treatment (Figure 24).

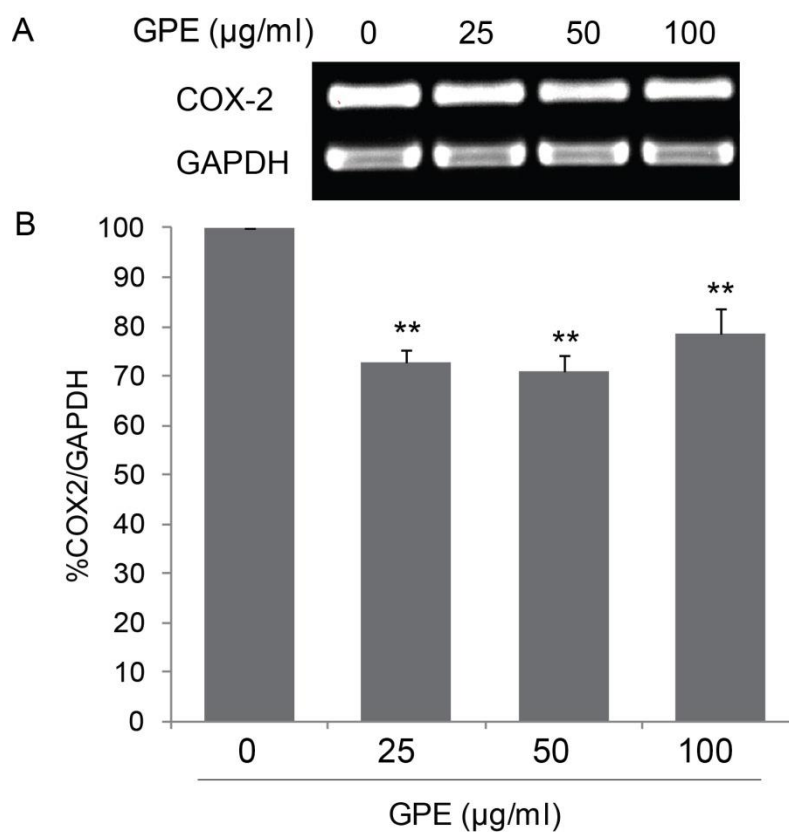


Figure 24: GPE alters the expression of COX-2 mRNA in HT-29. The cells were treated with GPE 25 - 100 $\mu\text{g/ml}$  for 24 h. Expressions of COX-2 was determined by RT-PCR. A) A representative of PCR products of COX-2 genes. B) Densitometric analysis of the PCR products relative to GAPDH presented as % of control. The data represent the means  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO).

### 3. Effect of GPE on human colorectal cancer cell not expressing COX-2, Colo-205

The anti-cancer effects of GPE were determined on human colorectal cancer cells that not expressing COX-2, Colo-205 cells as following.

#### 3.1 Cytotoxic activities of GPE

Colo-205 cells were tested for cytotoxic effects of GPE by resazurin reduction assay. The cells were exposed to 6.25 - 100 $\mu$ g/ml of GPE for 24, 48 and 72 h. Standard cytotoxic drugs recommended for the treatment of CRC, 5FU and Oxaliplatin were used as positive controls in this experiment.

As shown in Figure 25 25, 50 and 100 $\mu$ g/ml of the extract significantly exhibited cytotoxicity against Colo-205 cells. The  $IC_{50}$  values of GPE against Colo-205 were  $59.92\pm 4.34$ ,  $28.85\pm 1.44$  and  $25.42\pm 1.65$   $\mu$ g/ml at 24, 48 and 72 h, respectively (Figure 26). The cytotoxic effect of the extract was seen in a dose-dependent and time-dependent manner. However, the dose-dependency was more pronounced. At 24 h treatment time, GPE showed higher cytotoxic activities against Colo-205 when compared to that of standard cytotoxic agents, 5FU and Oxaliplatin. The  $IC_{50}$  of 5FU against Colo-205 were >1,000, 29.52, and 10.12  $\mu$ M at 24, 48 and 72 h, respectively. Oxaliplatin showed the  $IC_{50}$  of >1,000, 11.94, and 5.53  $\mu$ M against Colo-205 at 24, 48 and 72 h, respectively.

According to the  $IC_{50}$  of GPE, three concentrations at 25, 50 and 100 $\mu$ g/ml were chosen for the rest of the experiments.



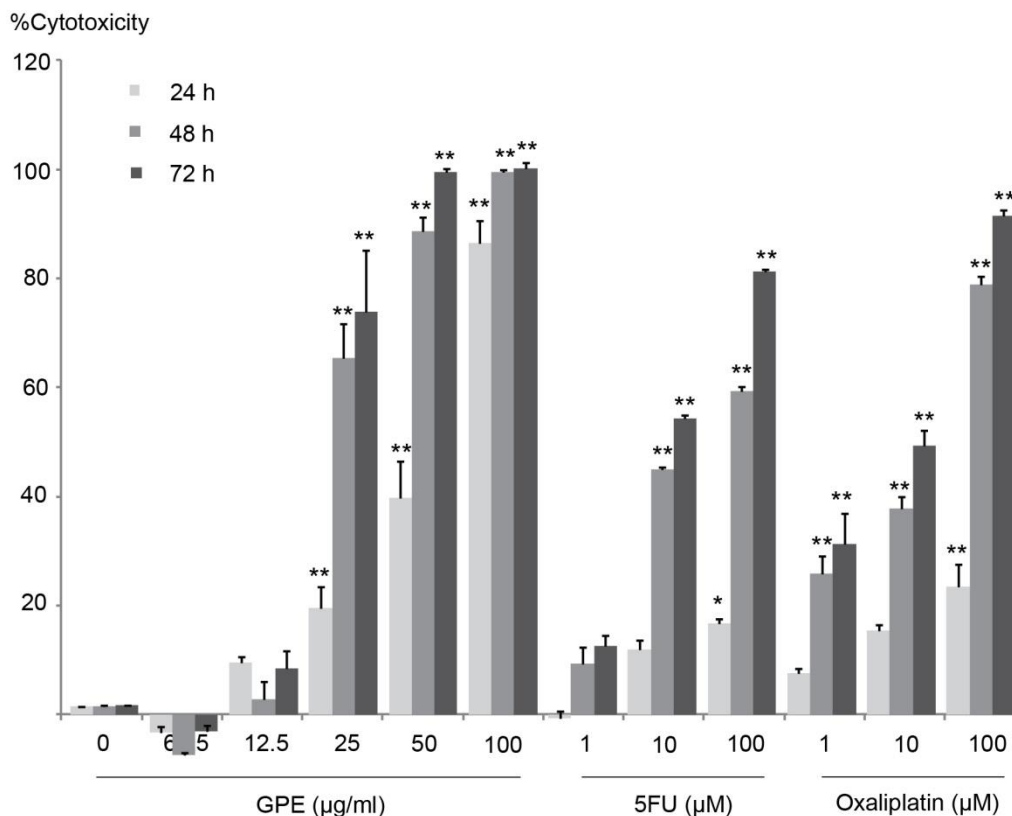


Figure 25: Cytotoxic effects of GPE against Colo-205 cells. The cells were treated with GPE at 6.25 - 100µg/ml, 5FU at 1 – 100 µM and Oxaliplatin at 1 – 100 µM for 24, 48 and 72 h. The cytotoxic effects were determined by resazurin reduction assay. The data represent the means  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO).

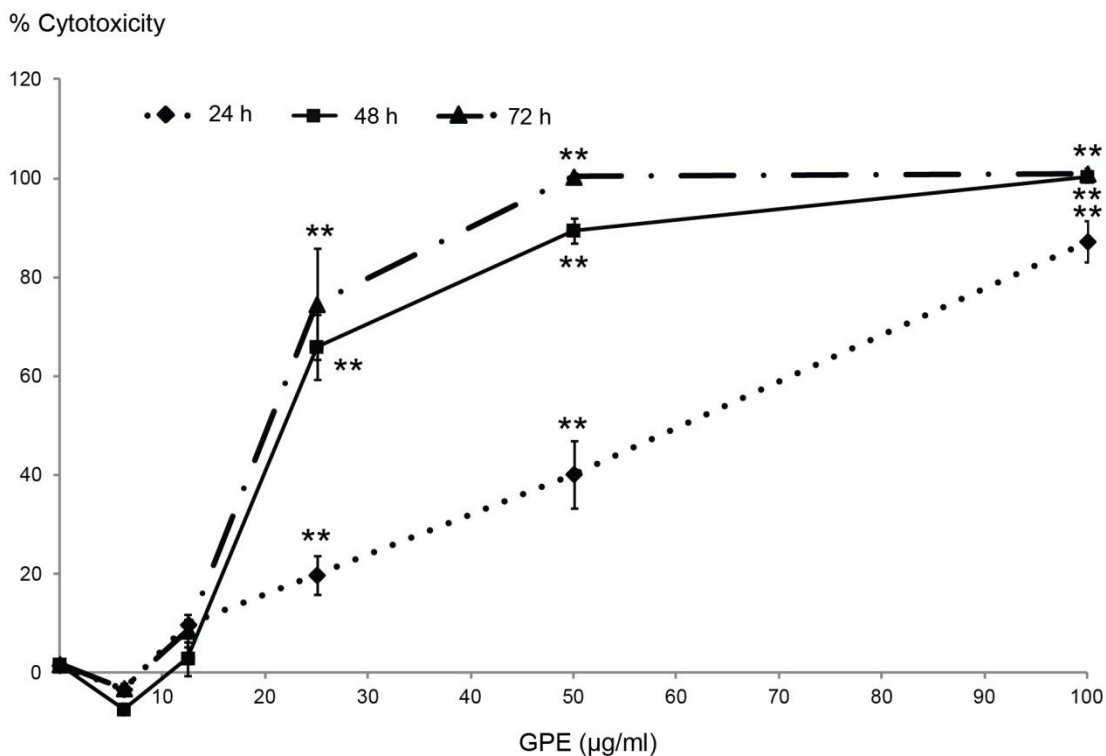


Figure 26: Line graphs show cytotoxicity of GPE against Colo-205 cells. The cells were treated with GPE at 6.25 - 100µg/ml for 24, 48 and 72 h. The cytotoxic effects were determined by resazurin reduction assay. The data represent the means + S.E. of three independent experiments. The percentages of cytotoxicity of GPE were plotted against GPE concentration. Linear trend lines of each treatment time were calculated. The linear equations were used to determine the IC<sub>50</sub> of GPE at each treatment time. \* p<0.05 and \*\* p<0.01 indicate statistically significant difference when compare to the solvent control (0.2% DMSO).

### 3.2 Apoptotic induction effect of GPE

The apoptotic induction effect of GPE was evaluated against Colo-205. The cells were treated with 25, 50 and 100  $\mu\text{g/ml}$  of GPE for 18 and 24 h. Apoptotic cells death were determined by AnnexinV-FITC/PI staining and detected with fluorescence flow cytometer. Oxaliplatin and 5FU were used as positive controls in this experiment.

GPE induced total cell death in a concentration- and time- dependent manner after 18 and 24 h exposures. GPE at 50 and 100 $\mu\text{g/ml}$  were significantly induced apoptotic cell death in Colo-205 by  $27.70\pm 5.09\%$  and  $14.42\pm 2.83\%$  at 18h and  $24.28\pm 5.72\%$  and  $18.07\pm 2.84\%$  at 24 h, respectively. At 50 $\mu\text{g/ml}$  of GPE, apoptotic cells death represented the majority type of cell death with 59.45% and 49.52% of total cell death at 18 and 24 h, respectively. On the other hand, 100 $\mu\text{g/ml}$  GPE caused majority of cell death as late apoptosis and/or necrosis stages while cells treated with 25 $\mu\text{g/ml}$  GPE remained viable (Figure 27). Oxaliplatin and 5FU showed a minimal apoptotic cells death. It was due to the short exposure time which caused the majority of cells remained viable. The results were correlated with the cytotoxicity study. The molecular mechanism of GPE on apoptotic induction was further evaluated at the same concentration at 24 h exposure.

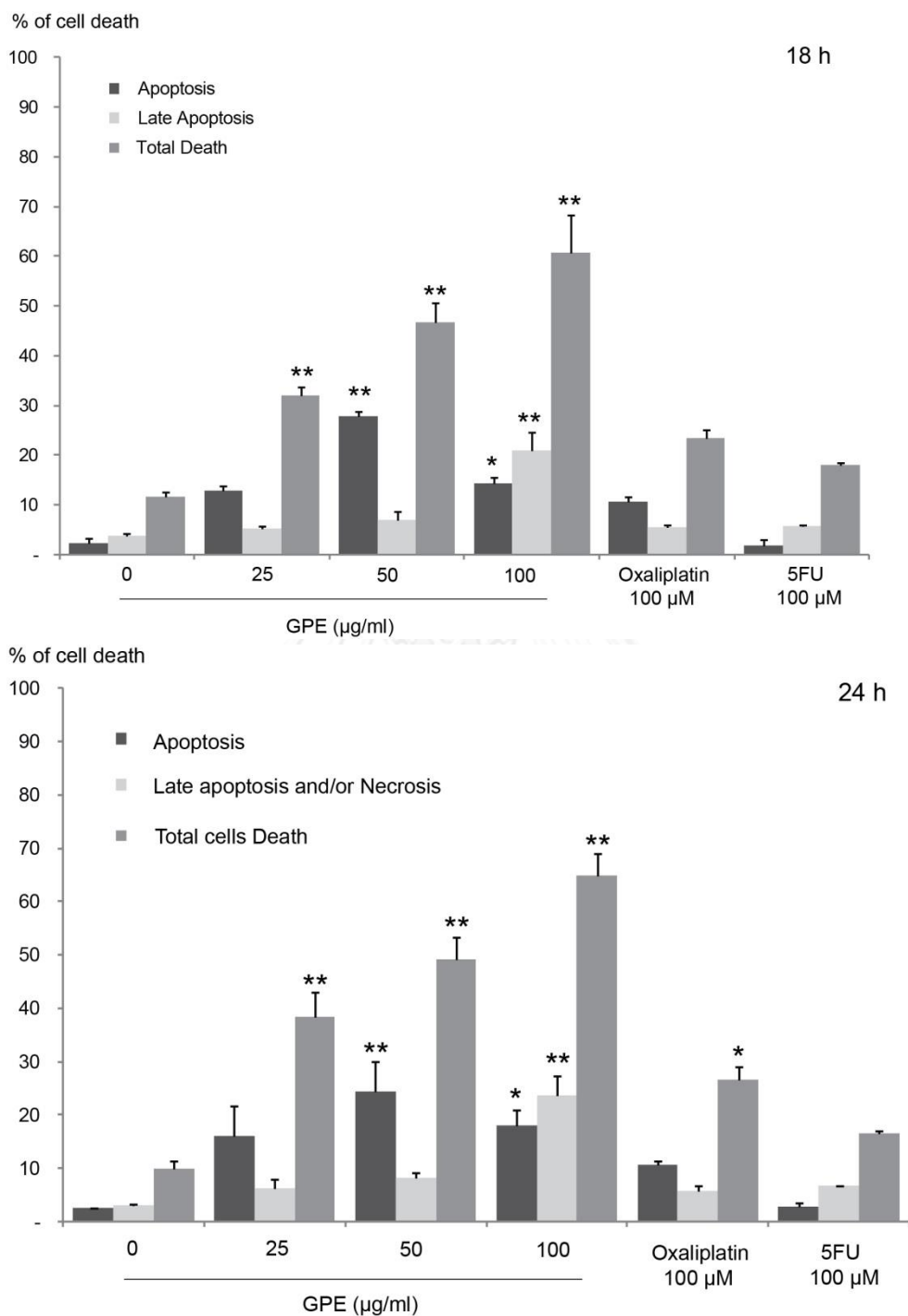


Figure 27: The apoptotic effect of GPE on Colo-205 cells. The cells were treated with 25 – 100µg/ml of GPE for 18 (A) and 24 h (B). The percentage of cells labeled with annexin V+/PI- (early apoptotic cells), annexin V+/PI+ (late apoptotic cells/necrotic cells) and total cell death were plotted. Data are the mean values  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO).

### 3.3 Effect of GPE on the expression of genes in BCL-2 family

To evaluate the mechanism by which GPE induced apoptosis in Colo-205, the mRNA expression of BCL-2 family genes were assessed by quantitative RT-PCR.

Colo-205 cells were treated with 25, 50 and 100  $\mu\text{g/ml}$  of GPE for 24 h, and then the cells were subjected to RNA isolation, cDNA conversion, and PCR amplification of target genes. GPE at 25, 50 and 100 $\mu\text{g/ml}$  significantly decreased the expression of anti-apoptotic BCL-2 to  $0.06\pm 0.01$ ,  $0.39\pm 0.07$  and  $0.09\pm 0.05$ , respectively. Moreover, the expression of anti-apoptotic BCL-XL was down-regulated to  $0.23\pm 0.00$  and  $0.42\pm 0.05$  following GPE 50 and 100 $\mu\text{g/ml}$  treatment, respectively (Figure 28). On the other hand, treatment with GPE tended to decrease pro-apoptotic BAK and BAX. However, the ratio of pro-apoptotic/anti-apoptotic (BAK and BAX/BCL-2 and BCL-XL) were favor apoptotic cell deaths (ratio > 1) with the ratio of 1.68, 1.94, and 3.09 following GPE 25, 50, and 100 $\mu\text{g/ml}$  treatment, respectively (Figure 29).

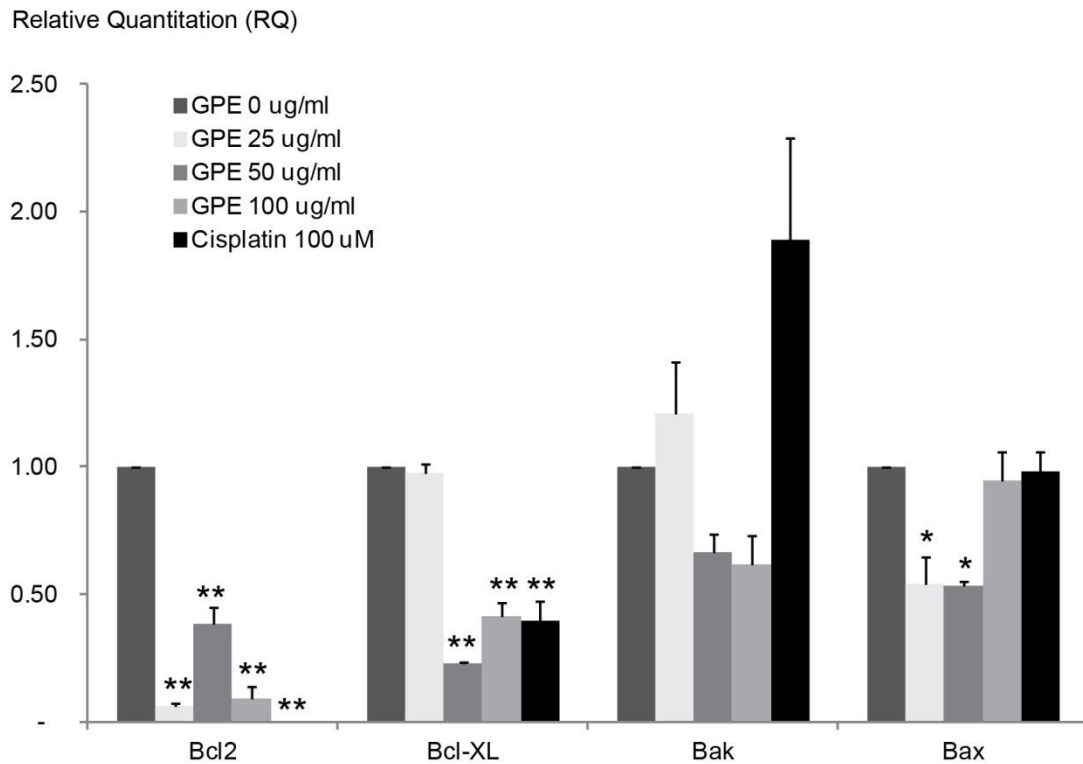


Figure 28: GPE alters the expression of BCL-2 family genes in Colo-205 cells. The expressions of BCL-2 family genes were determined by quantitative RT-PCR. The cells were treated with GPE 25 - 100µg/ml for 24 h. The treated cells were subjected to RNA isolation, cDNA conversion and quantitative RT-PCR amplification of BAK, BAX, BCL-2 and BCL-XL genes. The value is given as relative copy number normalized to the endogenous control GAPDH and 0.2% DMSO solvent control. The data represent the means  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO).

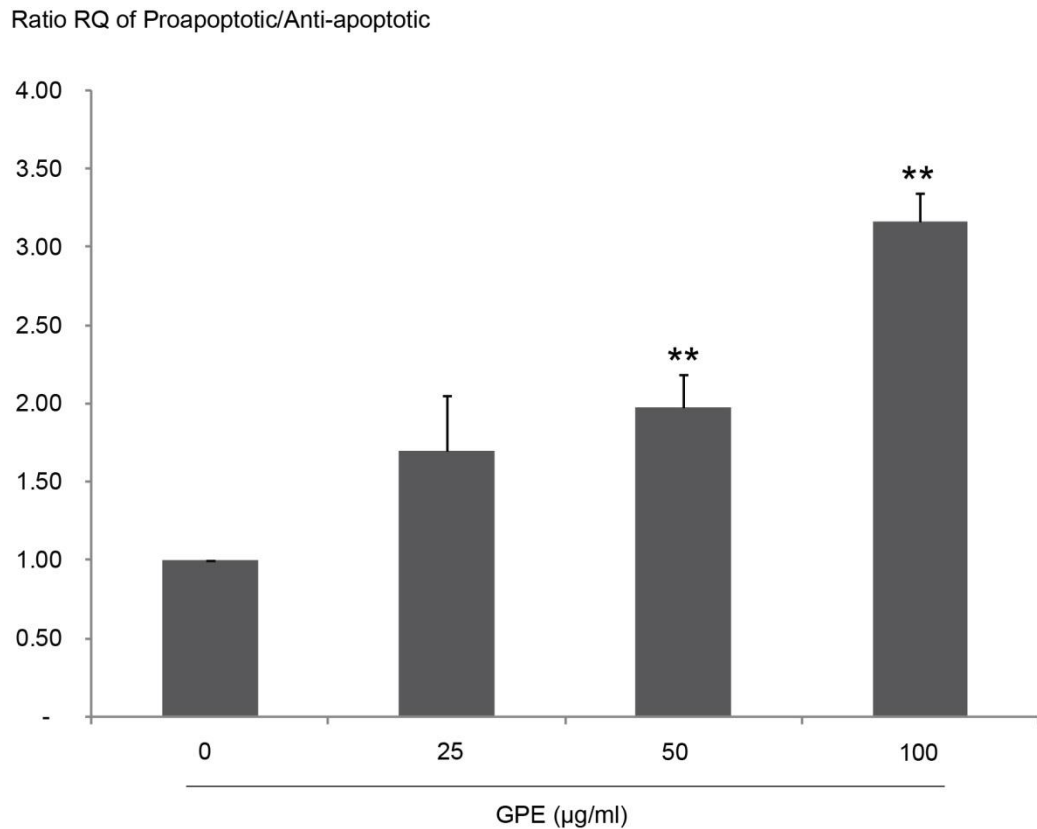


Figure 29: The ratio of pro-apoptotic/anti-apoptotic expression (BAK and BAX/BCL-2 and BCL-XL) in Colo-205 treated cells. The cells were treated with GPE 25 - 100µg/ml for 24 h. The treated cells were subjected to RNA isolation, cDNA conversion and quantitative RT-PCR amplification of BAK, BAX, BCL-2 and BCL-XL genes. The data represent the means  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO).

### 3.4 Effect of GPE on the cell proliferation

Inhibition of Colo-205 cell proliferation by GPE was determined by cell counting. Standard cytotoxic drug, 5-FU, was used as a positive control in this experiment.

Colo-205 cells were treated with 25, 50 and 100  $\mu\text{g/ml}$  of GPE for 24 h, afterward the total cells were counted. The results were expressed as a percentage of cells compared to the untreated control. GPE significantly inhibited Colo-205 cell proliferation to  $60.71\pm 5.13\%$ ,  $71.63\pm 8.84\%$ , and  $59.91\pm 2.44\%$  at GPE 25, 50 and  $100\mu\text{g/ml}$ , respectively when compared to the vehicle control. However, the effect was not seen in a dose-dependent manner (Figure 30). The cell proliferation inhibition effect of GPE against Colo-205 was slightly lower than that of 5FU which  $100\mu\text{M}$  5FU significantly inhibited cell proliferation to  $53.91\pm 5.08\%$ .



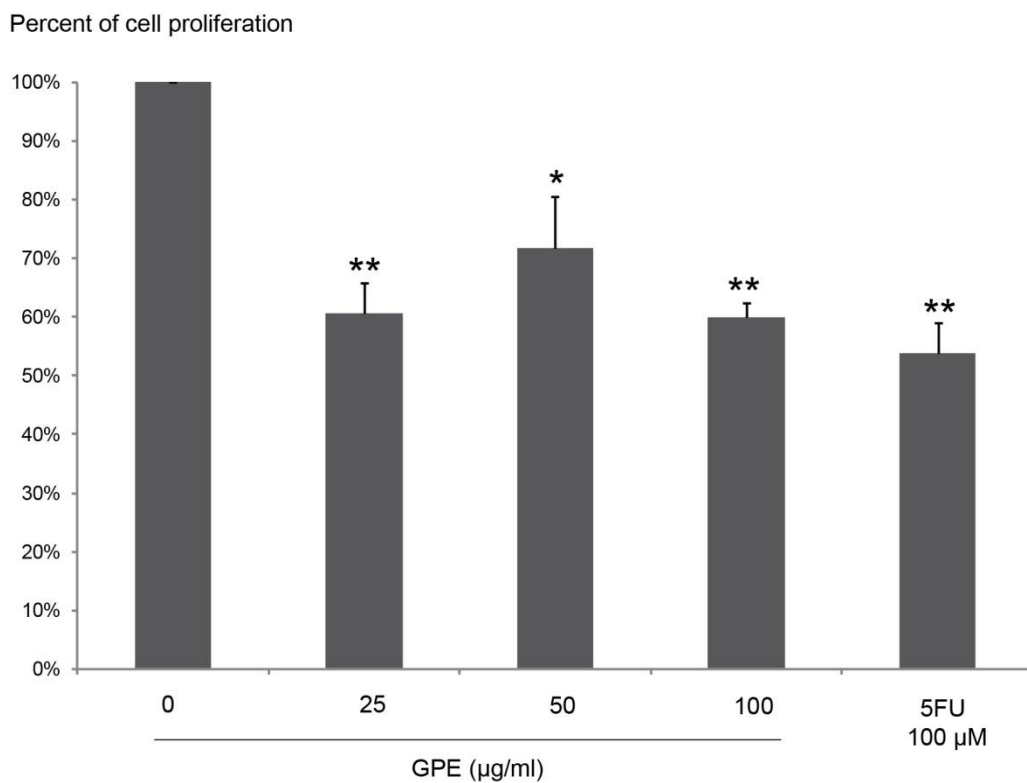


Figure 30: Anti-proliferative effects of GPE against Colo-205 cells. The cells were treated with GPE 25 – 100µg/ml for 24 h. Cell proliferation was measured by cell counting using an automated cell counter. The data represent the means  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO).

### 3.5 Effect of GPE on the cell cycle progression

Colo-205 cells were treated with 25, 50 and 100  $\mu\text{g/ml}$  GPE for 1 h and further incubated for 24 h in a fresh medium without GPE. The treated cells were subjected to fluorescence flow cytometric analysis for DNA content. 5FU which arrests the cell cycle at S phase and causes cell accumulation at G1/S phase was used as a positive control (Figure 31).

The effect of GPE on cell cycle pattern of Colo-205 cells was difference which depended on the concentration used. GPE at 25 and 50  $\mu\text{g/ml}$  caused a decrease in the proportion of cells in the S phase (from  $36.37 \pm 0.83\%$  to  $25.49 \pm 2.00\%$  and  $16.62 \pm 1.79\%$ , respectively). In cells treated with 25  $\mu\text{g/ml}$  GPE, this decrease was associated primarily with accumulation of cells in the G0/G1 phase (from  $51.49 \pm 1.73\%$  to  $65.39 \pm 1.04\%$ ). At 50  $\mu\text{g/ml}$ , GPE resulted in a significant reduction in the S fraction with a shift of cells accumulation to the G2/M phases (from  $12.14 \pm 0.95\%$  to  $30.07 \pm 1.73\%$ ) as shown in Figures 32. The positive control, 5FU, significantly accumulated cells population at S fraction with the reduction of the cells in G2/M phases. The molecular mechanism of GPE on cell cycle pattern was further evaluated at the same concentration range and treatment time.



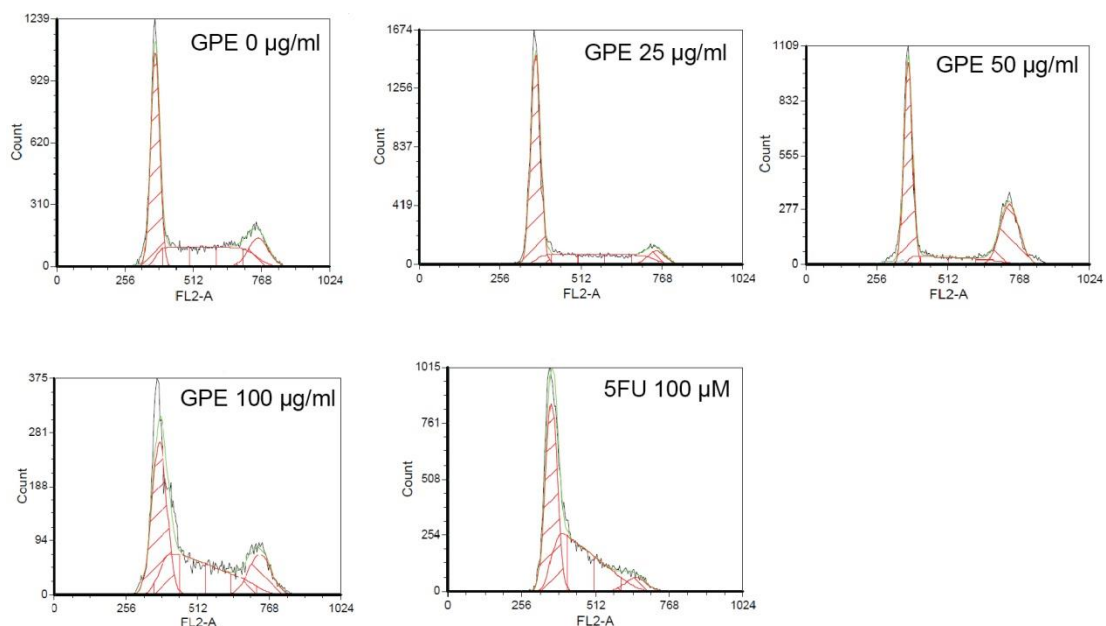


Figure 31: Representative histogram from flow cytometric analysis of Colo-205 cell cycle pattern after treatment with GPE. The cells were treated with GPE at 25 – 100 µg/ml for 1 h. The treated cells were washed and incubated in fresh medium for additional 24 h. Analysis of cells in the cell cycle was performed by fixing and staining the cells with PI. Cell cycle pattern of  $3 \times 10^4$  cells/sample were analyzed using fluorescence flow cytometer. The cell cycle distributions were evaluated using FCS Express 4 Image Cytometry.

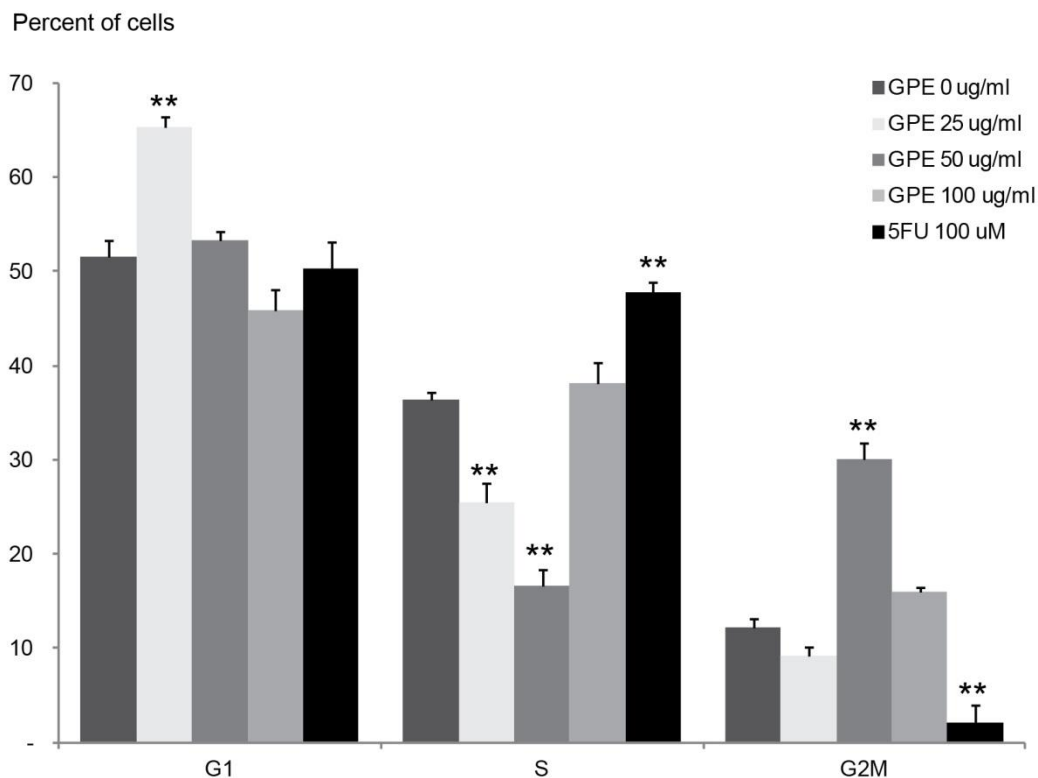


Figure 32: Distribution of Colo-205 cells in the cell cycle phases after treatment with GPE. The cells were treated with GPE 25 - 100µg/ml for 1 h. The treated cells were washed and incubated in fresh medium for additional 24 h. Analysis of cells in the cell cycle was performed by fixing and staining the treated cells with PI followed by fluorescence flow cytometry. The percentage of cells in G1, S, and G2M phases are given as means  $\pm$  S.E. of 3 independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO).

### 3.6 Effect of GPE on the mRNA expression of cell cycle regulatory genes

Since GPE changed the cell cycle pattern of Colo-205 cells, the alterations in the expression of cell cycle regulatory genes might be involved. Colo-205 cells were treated with 25, 50 and 100  $\mu\text{g/ml}$  GPE for 1 h and further incubated for 24 h in a fresh medium without GPE. The treated cells were subjected to RNA isolation, cDNA conversion, PCR amplification of the genes involved in cell cycle progression including cyclin A, cyclin B1, cyclin D1, cyclin E, p21 and p53. PCR products were analyzed by agarose gel electrophoresis. The housekeeping genes, GAPDH, was used as a loading control.

As shown in Figures 33, GPE at the concentration of 25 and 50  $\mu\text{g/ml}$  significantly down-regulated cyclin A expression in a dose-dependent manner. The cyclin E expression was likely to decrease following GPE treatment. However, the change was not statistically significant. The expression on CDKI p21 tended to decrease following GPE treatment though it reached statistically significant only at 50 $\mu\text{g/ml}$  concentration. Moreover, the level of cyclin B1, cyclin D1, and p53 remained unchanged. As the GPE at 100  $\mu\text{g/ml}$  caused majority of Colo-205 cell death which affected the expression of the housekeeping gene, GAPDH, the effects of 100  $\mu\text{g/ml}$  GPE on Colo-205 mRNA expression of cell cycle regulatory genes were excluded from the analysis.

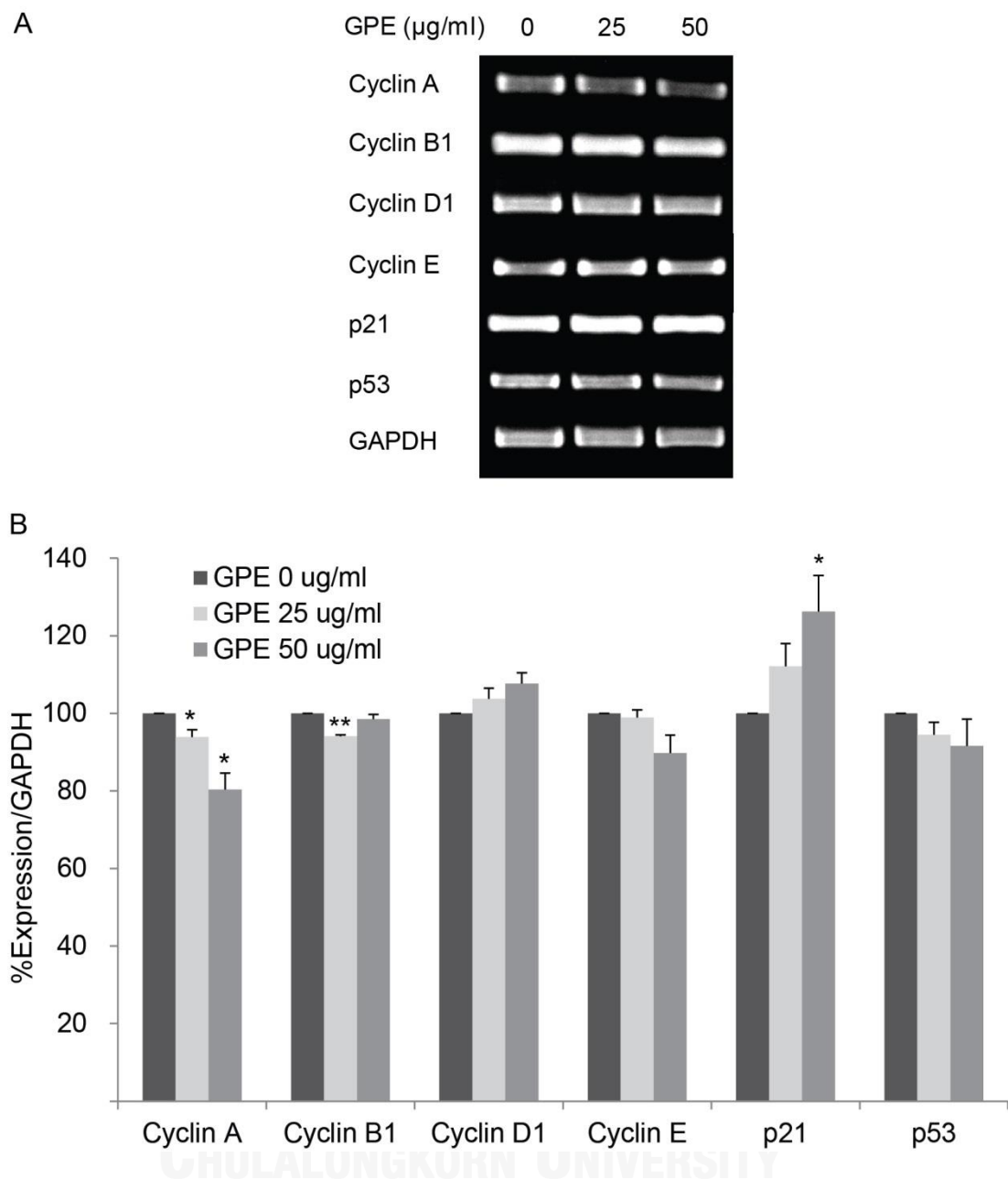


Figure 33: GPE alters the expression of cell cycle regulators genes in Colo-205. The cells were treated with GPE 25 and 50  $\mu\text{g/ml}$  for 1 h and incubated with fresh medium without GPE for additional 24 h. Expression of cell cycle regulators were determined by RT-PCR. A) A representative of PCR products of the cell cycle regulatory genes. B) Densitometric analysis of the PCR products relative to GAPDH presented as % of control. The data represent the means  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO).

4. Compare the effect of GPE on human colorectal cancer cell expressing COX-2, HT-29 cells and not expressing COX-2, Colo205 cells.

#### 4.1 Cytotoxic activities of GPE

The cytotoxic effect of GPE against HT-29 and Colo-205 was compared using independent sample student t test. At 24 h of treatment, the  $IC_{50}$  of GPE against HT-29 and Colo-205 was not significantly difference. However, with the increased treatment time to 48 h and 72 h, GPE was significantly more cytotoxic against Colo-205 than HT-29. The cytotoxic activity of GPE against both HT-29 and Colo-205 were seen at 25 – 100  $\mu\text{g/ml}$  of GPE (Figure 34).

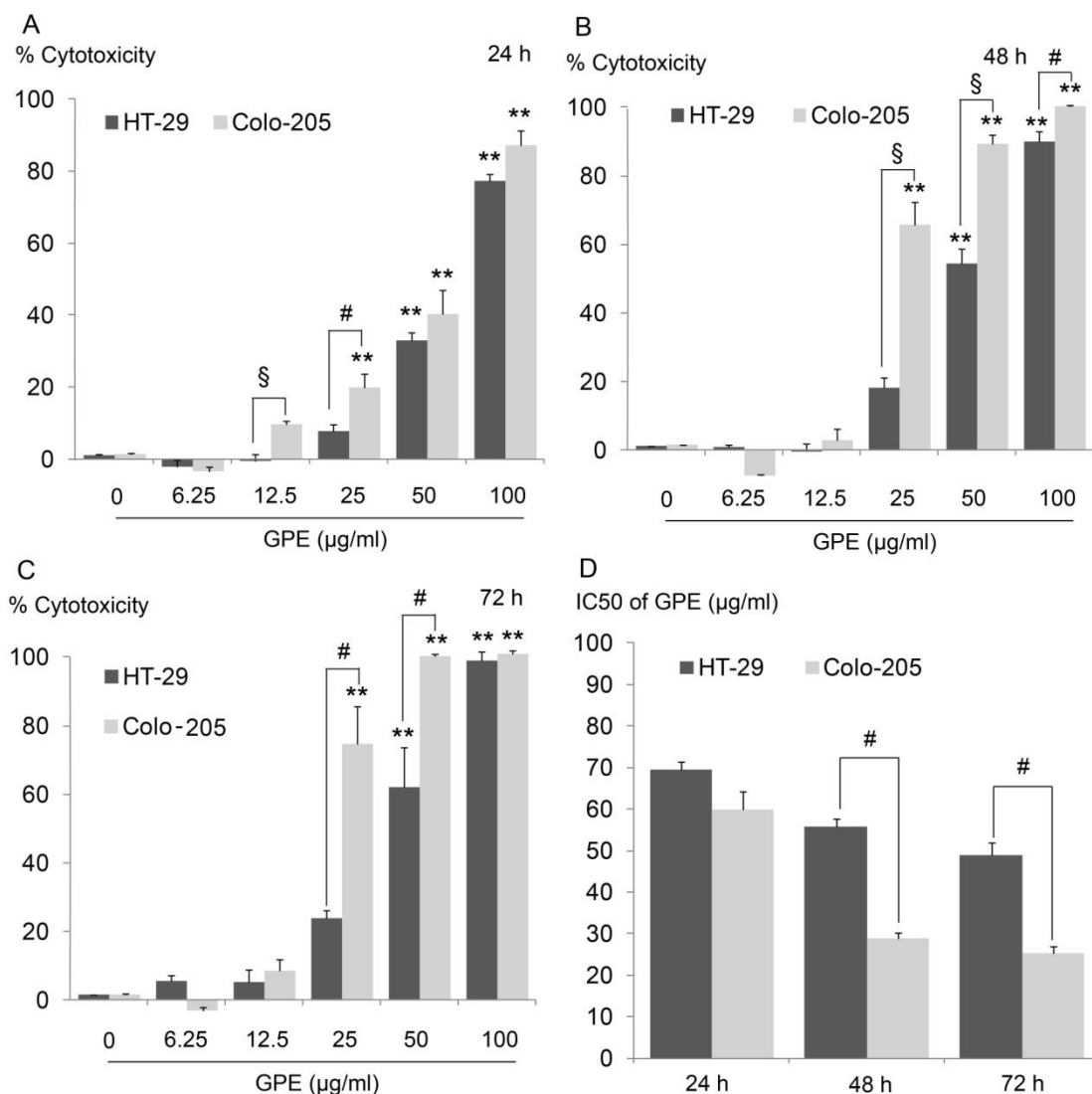


Figure 34: Cytotoxic effects of GPE against HT-29 and Colo-205 cells. The cells were treated with GPE at 6.25 - 100µg/ml for 24 (A), 48 (B) and 72 (C) h. The cytotoxic effects were determined by resazurin reduction assay. The percentages of cytotoxicity of GPE were plotted against GPE concentration. Linear trend lines of each treatment time were calculated. The linear equations were used to determine the IC<sub>50</sub> of GPE at each treatment time (D). The data represent the means ± S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO). #  $p < 0.05$  and §  $p < 0.01$  indicate statistically significant difference between the two cell lines.



## 4.2 Apoptotic induction effect of GPE

GPE can induce apoptotic cells death in both HT-29 and Colo-205 cells. At 18 h of treatment, GPE at 25 and 100  $\mu\text{g/ml}$  significantly induced apoptotic cells death in Colo-205 cells higher than HT-25 cells. At 24 h, there was no significant difference in the induction of apoptosis between the two cell lines (Figure 35). The total cells death was also significantly higher in Colo-205 treated cells.

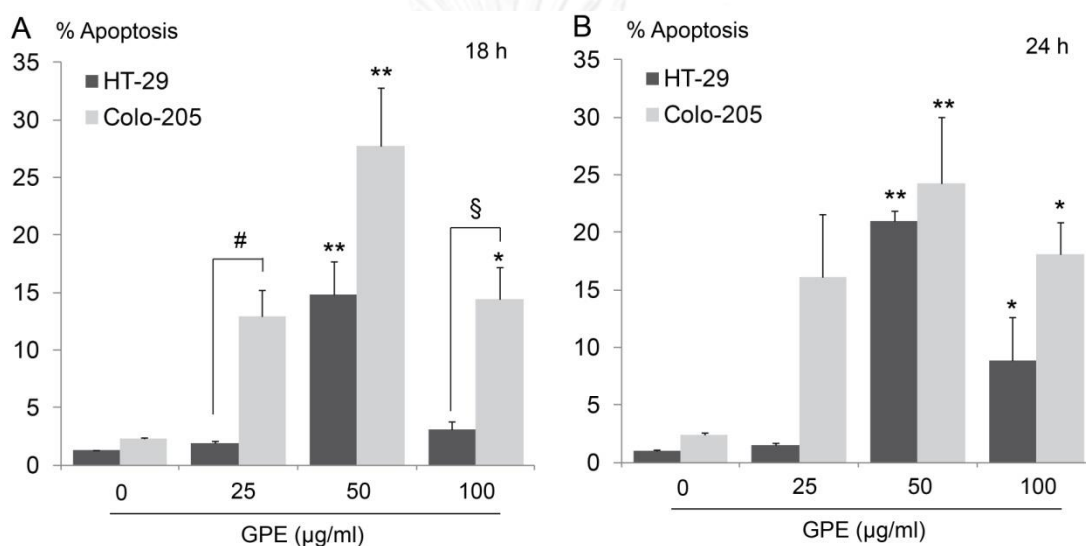


Figure 35: The apoptotic effect of GPE on HT-29 and Colo-205 cells. The cells were treated with 25 – 100  $\mu\text{g/ml}$  of GPE for 18 (A) and 24 h (B). The percentage of cells labeled with annexin  $\text{V}^+/\text{PI}^-$  (early apoptotic cells) were plotted. Data are the mean values  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO). #  $p < 0.05$  and §  $p < 0.01$  indicate statistically significant difference between the two cell lines.

### 4.3 Effect of GPE on the expression of genes in BCL-2 family

The expressions of genes in BCL-2 family in HT-29 and Colo-205 cells after treatment with GPE were both altered in the favor of apoptotic cells death. However, there were some differences in the expression of these genes in the two cell lines. Anti-apoptotic BCL-2 gene was down-regulated in both HT-29 and Colo-205 cells following GPE treatment. GPE at 25 $\mu$ g/ml significantly decreased BCL-2 gene expression in Colo-205 more than HT-29 cells. Anti-apoptotic BCL-XL gene also decreased in both cell lines after exposed to GPE. The difference in the reduction in BCL-XL expression was significantly seen at GPE 50 $\mu$ g/ml which affected Colo-205 more than HT-29. GPE tended to increase pro-apoptotic BAK in HT-29 cells. In contrast, the expression of BAK was down-regulated in Colo-205 after treated with GPE. The expression of pro-apoptotic BAX was likely to decrease following GPE treatment in both HT-29 and Colo-205 cells (Figure 36).

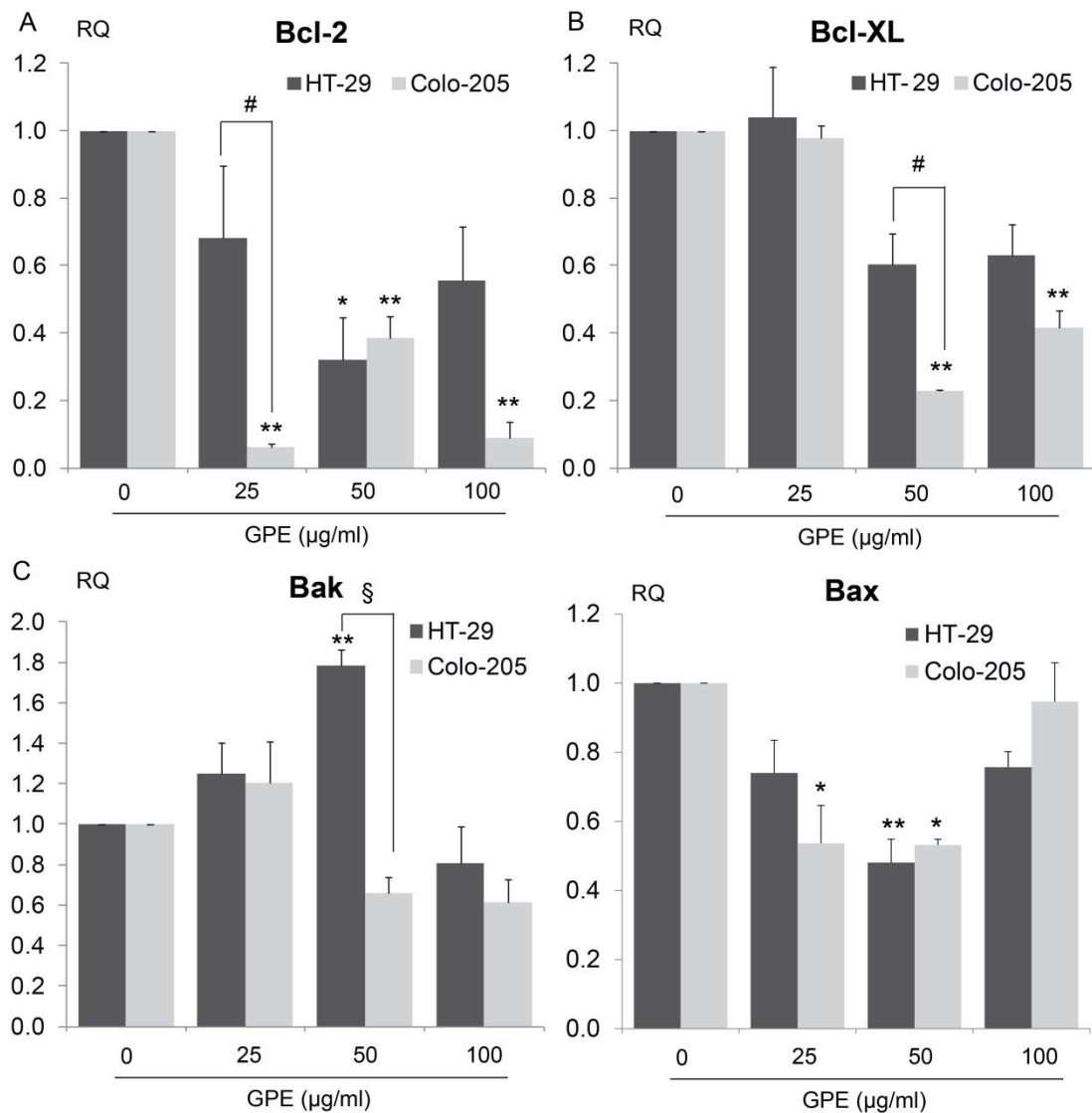


Figure 36: GPE alters the expression of BCL-2 family genes in HT-29 and Colo-205 cells. The expressions of BCL-2 family genes were determined by quantitative RT-PCR. The cells were treated with GPE 25 - 100µg/ml for 24 h. The treated cells were subjected to RNA isolation, cDNA conversion and quantitative RT-PCR amplification of BCL-2 (A), BCL-XL (B), BAK (C) and BAX (D) genes. The value is given as relative copy number normalized to the endogenous control GAPDH and 0.2% DMSO solvent control. The data represent the means  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO). #  $p < 0.05$  and §  $p < 0.01$  indicate statistically significant difference between the two cell lines.

#### 4.4 Effect of GPE on the cell proliferation

GPE at 25-100  $\mu\text{g/ml}$  significantly inhibited HT-29 and Colo-205 cells proliferation. However, the intensity was different in the two cells line. GPE at 100 $\mu\text{g/ml}$  significantly inhibit HT-29 cells proliferation more than Colo-205 cells. The concentration dependent of the cell proliferation inhibition was seen only in HT-29 treated with GPE (Figure 37).

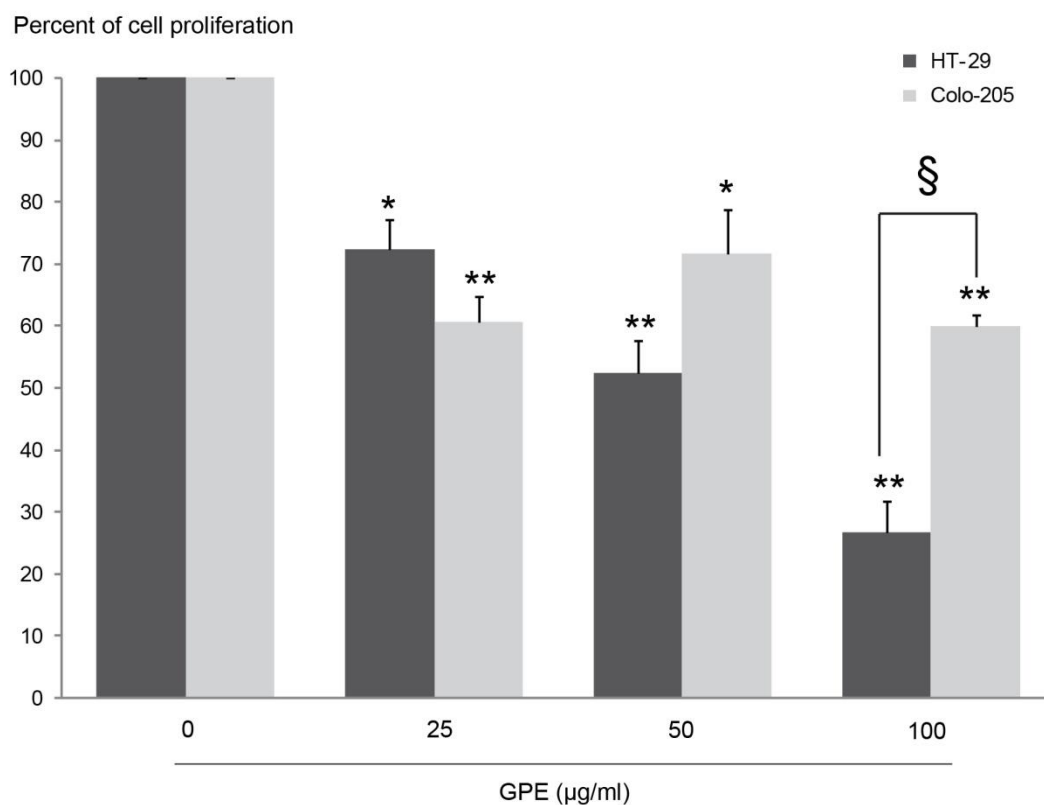


Figure 37: Anti-proliferative effects of GPE against HT-29 and Colo-205 cells. The cells were treated with GPE 25 – 100 $\mu\text{g/ml}$  for 24 h. Cell proliferations were measured by cell counting using an automated cell counter. The data represent the means  $\pm$  S.E. of three independent experiments. \*  $p < 0.05$  and \*\*  $p < 0.01$  indicate statistically significant difference when compare to the solvent control (0.2% DMSO). #  $p < 0.05$  and §  $p < 0.01$  indicate statistically significant difference between the two cell lines.

#### 4.5 Effect of GPE on the cell cycle progression

The effect of GPE on the cell cycle pattern of HT-29 and Colo-205 cells was difference depended on the concentration of GPE. Cell cycle pattern had shifted to the decrease in S phase and increase in G1 cells population following GPE 50 and 25 $\mu$ g/ml in HT-29 and Colo-205, respectively. The pattern had changed to the decrease in S phase with the cell accumulation in G2/M phase after treated with GPE 100 and 50 $\mu$ g/ml in HT-29 and Colo-205, respectively (Figure 38).

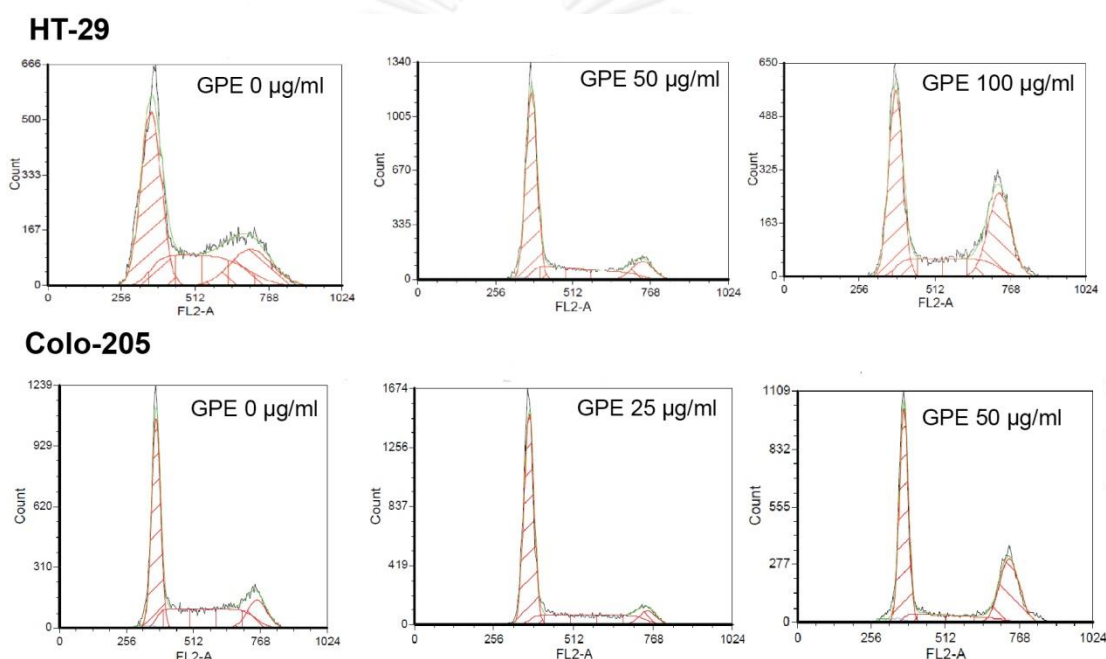


Figure 38: Representative histogram from flow cytometric analysis of HT-29 and Colo-205 cell cycle pattern after treatment with GPE. The cells were treated with GPE at 25 – 100  $\mu$ g/ml for 1 h. The treated cells were washed and incubated in fresh medium for additional 24 h. Analysis of cells in the cell cycle was performed by fixing and staining the cells with PI. Cell cycle pattern of  $3 \times 10^4$  cells/sample were analyzed using fluorescence flow cytometer. The cell cycle distributions were evaluated using FCS Express 4 Image Cytometry.

#### 4.6 Effect of GPE on the mRNA expression of cell cycle regulatory genes

GPE affected the expression of cell cycle regulatory genes in both HT-29 and Colo-205. The expression of cyclin A was down-regulated following GPE treatment in both cell lines. GPE decreased the expression of cyclin E gene significantly only in HT-29 cells. The expression of CDKI, p21, was significantly up-regulated only in Colo-205 treated cells at GPE 50 $\mu$ g/ml. The level of cyclin B1, cyclin D1, and p53 remained unchanged in both cell lines.



## CHAPTER V

### DISCUSSION AND CONCLUSION

In this study, the effects of EtOAc extract from the leaves of *G. parva* (GPE) were examined on human colorectal cancer expressing COX-2, HT-29, and non-expressing COX-2, Colo-205 cells. The GPE exhibited cytotoxic activity, inhibited cell proliferation, and induced apoptosis as well as cell cycle arrest in both cell lines but with some differences intensity. The effect of GPE on cytotoxicity and induction of apoptosis were higher in Colo-205 cells than HT-29 cells. However, the anti-proliferative effects of GPE were superior in HT-29 cells. The changes in cell cycle pattern following GPE treatment were similar in both cell lines with the reduction of cells in S phases which accompanied with the accumulation of cells in G0/G1 and G2M phases. Further studies suggested that the effects of GPE may be mediated through both COX-2 dependent and COX-2 independent pathway. The COX-2 independent pathway included the changes in the expression of cell cycle control genes and the alteration in the balance of BCL-2 family gene expression. While the COX-2 dependent involved with reduction of COX-2 expression which could affect downstream COX-2 pathway. However, these two pathways may have overlapping or crosstalk mechanisms.

Throughout medical history, plant products have been shown to be valuable sources of novel anti-cancer drugs discovery (99). *Glycosmis parva* is a wild small shrub found in Thailand. Chemical examination of the EtOAc extract from the leaves of *G. parva* led to the identification of the acridone alkaloid and sulfur-containing propanamides as its major active compounds. Arborinine, a main acridone alkaloid found in EtOAc of the leaves of *G. parva*, has been reported to have potent anti-proliferative effect against cervix adenocarcinoma HeLa cells with an  $IC_{50}$  of 1.84  $\mu$ M, which was lower than that of cisplatin (12.43  $\mu$ M) (36), indicating arborinine possesses higher anti-proliferative activity than cisplatin. Additionally, antifungal activities of sulfur-containing propanamides derivative, S-deoxydihydroglyparvin, S-deoxytetrahydroglyparvin, glyparvin-A, and dihydroglyparvin, (8) were reported (83). In the present study, the results showed that GPE possessed moderate cytotoxic

activity against HT-29 and Colo-205 with the  $IC_{50}$  of 69.49 and 59.92 $\mu$ g/ml, respectively, after 24 h exposure. The cytotoxicity of the extract was in a dose- and time-dependent manner in both cell lines. GPE was more cytotoxic to Colo-205 than HT-29 cells especially with higher exposure time. The cytotoxicity study by resazurin reduction assay demonstrated both the ability of the tested compound to kill the cells and the effect of cell proliferation inhibition. Therefore, the effects of the GPE on cell death and cell proliferation would further clarify the different effects on the two cell lines.

Treatment of cancer with ionizing radiation and several anticancer drugs kills target cells primarily by induction of apoptosis (100). The results showed that GPE at 50 $\mu$ g/ml significantly induced cell death mainly by apoptosis in both HT-29 and Colo-205 cells. The proportion of apoptotic cell death to total cell death was higher than that of currently used cytotoxic drugs, oxaliplatin and 5FU which were used as positive controls in this study. However, at higher concentration, 100 $\mu$ g/ml of GPE, the late apoptotic and/or necrotic cell death was more prominent than early apoptotic cells. The apoptotic induction effects of GPE were higher in Colo-205 than HT-29 cells.

Activation of apoptosis signaling following treatment with cytotoxic drugs has been shown to lead to activation of the mitochondrial (intrinsic) pathway of apoptosis (96). BCL-2 family proteins play a pivotal role in the regulation of the mitochondrial pathway of apoptosis. They comprise both anti-apoptotic members- e.g., BCL-2, BCL-XL, and MCL-1- as well as pro-apoptotic molecules such as BAX, BAK, and BAD (101). GPE decreased the expression of anti-apoptotic BCL-2 significantly in both HT-29 and Colo-205 cells while anti-apoptotic BCL-XL was significantly decreased only in Colo-205. However, GPE at 50 $\mu$ g/ml significantly increased the expression of pro-apoptotic BAK and BAX only in HT-29. Although, some concentrations of GPE decreased pro-apoptotic BAK and BAX expression in both cells, the ratios of pro-apoptotic/anti-apoptotic (BAK and BAX/BCL-2 and BCL-XL) were designated cancer cell death.

Cell proliferation is a key character of cancer progression (102). The results showed that GPE significantly inhibited both HT-29 and Colo-205 cells proliferation.



The dose dependence was only seen in HT-29 cells. The effects were found significantly across the experimental dose-range (25-100 $\mu$ g/ml) in both cell lines but the proliferation inhibition effects were higher in HT-29 cells.

The ability of anticancer agents to suppress the growth of cancer cells can also be associated with blocking the cell cycle progression (103). Based on the significant inhibition of HT-29 and Colo-205 cell proliferation by GPE, it was postulated that the extract might modulate the cell cycle progression of these cancer cells. Cell cycle distribution analysis showed that GPE caused a reduction in S phase population which was associated with accumulation of cells in the G<sub>0</sub>/G<sub>1</sub> phase. At higher concentration of GPE, the treated cells accumulated at the G<sub>2</sub>/M phases. The cell cycle pattern changes were similar in both cell lines but the effects were seen in different concentrations of GPE. This result may in part be due to more than one active compound presented in the extract affecting the cell cycle but at different phases of the cycle. Additional studies to identify active compounds presented in the EtOAc extract are warranted.

Cyclin dependent kinases (CDKs) and cyclins are important determinants in the regulation of cell cycle progression. As the cells progress from G<sub>1</sub> to mitosis, a series of CDKs is expressed along with that of cyclins (104). Cell cycle arrest involves down-regulation of cyclins and CDKs as well as up-regulation of cyclin dependent kinase inhibitors (CDKIs) (105). Cyclin E/CDK2 and cyclin A/CDK2 complexes play an important role in G<sub>1</sub>-S transition and DNA replication, respectively. Moreover, cyclin A can form a complex with CDC2, which is essential for mitosis (104). The CDKI p21 mediates p53-dependent cell cycle arrest through inhibition of CDK2–cyclin E (progression into and through S phase), CDK2–cyclin A (progression through S phase) and CDK1–cyclin A (progression into G<sub>2</sub>), and CDK1–cyclin B1 (progression through G<sub>2</sub> and G<sub>2</sub>/M) which lead to growth arrest at specific stages in the cell cycle (106). The results indicated that GPE down-regulated mRNA expression of cyclin A and cyclin E in a dose-dependent manner in HT-29 cells. It also decreased cyclin A expression and up-regulated p21 expression in Colo-205 cells. This may explain why GPE could induce cell cycle arrest and the changes in the cell cycle pattern in both cell lines.

Cyclooxygenase enzymes are required for the conversion of arachidonic acid to prostaglandins. COX-2 mediates the inflammatory effects of COX activity. It is induced by a wide spectrum of growth factors and pro-inflammatory cytokines and overexpressed in numerous premalignant and malignant lesions, including CRC. Treatment with the selective COX-2 inhibitor has shown promising results in the prevention of CRC (6). However, the dose-related cardiovascular toxicity of selective COX-2 inhibitors remains a concern (56). Discovery of agents that regulate biological pathways related to COX-2 pathway may provide new avenues for the treatment of CRC. Previous study has shown that GPE inhibited mRNA expressions of COX-2 in LPS stimulated-macrophage J774A.1 cells (10). Similarly, the inhibitory effect of GPE on COX-2 expression was detected in human colorectal cancer cells constitutively expressed COX-2, HT-29 cells.

The overall effects of GPE on HT-29 and Colo-205 cells showed that regardless of the COX-2 expression status, GPE possessed an anti-cancer activity against human colorectal cancer cells. The mechanisms engaged both COX-2 dependent and COX-2 independent pathway. The COX-2 dependent mechanisms mainly involved the action of PGE<sub>2</sub>, the most abundant COX-2-derived prostaglandins found in human CRC (7). PGE<sub>2</sub> stimulates proliferation and suppresses apoptotic cell death of cancer cells (45). PGE<sub>2</sub> activates multiple intracellular signaling pathways, including epidermal growth factor receptor (EGFR), protein kinase C (PKC)-ERK-c-Myc, and  $\beta$ -catenin/TCF pathways (107). However, further mechanistic study on the action of GPE needed to be done in order to confirm its action. The COX-2 independent mechanisms may involve the alteration of the expression of cell cycle regulators, arrested cell cycle, and induced apoptosis in the colon cancer cells that do not express COX-2 (108). These were in line with our findings that GPE arrested cell cycle, changed the expression of cell cycle regulators, and induced apoptosis in Colo-205 which did not express COX-2.

Nevertheless, there are some limitations from this study. Although HT-29 and Colo-205 are both human colorectal cancer cells which have different COX-2 expression status, they came from different origin. HT-29 is a colorectal adenocarcinoma from primary site while Colo-205 is a colorectal adenocarcinoma

derived from ascites metastatic site. They have comparable population doubling time but different growth pattern. HT-29 is an adherent cell while Colo-205 grows as mixed adherent and suspension cells. They have similar oncogenes expression including c-myc, H-ras, K-ras, N-ras, myb, fos, and p53 (109). The mutational statuses of the two cell lines are somewhat different. HT-29 carries APC, BRAF, and PIK3CA mutations while Colo-205 has BRAF mutation and APC wild type. Both cells carry SMAD4 and TP53 wild type. These differences in the two cell lines needed to be taking into consideration when interpret the result from the study.

The promising anti-cancer action of GPE against human colorectal cancer cells with the interesting mechanism of actions involved both COX-2 dependent and COX-2 independent pathway deserve a further investigation of its possibility to be a candidate for colorectal cancer treatment. With the understanding of chemical constituents found in GPE, together with the confirmed anti-cancer activity, this could be a valuable source of information that benefits researchers in the anti-cancer drug development field.

## Conclusions

The anti-cancer activity of the EtOAc extract from leave of *Glycosmis parva* on human colorectal cancer cells and its underlying mechanism could provide the basis for further development of the compounds from this plant which could be a potential compounds for cancer therapeutics and chemoprevention in colorectal cancer.

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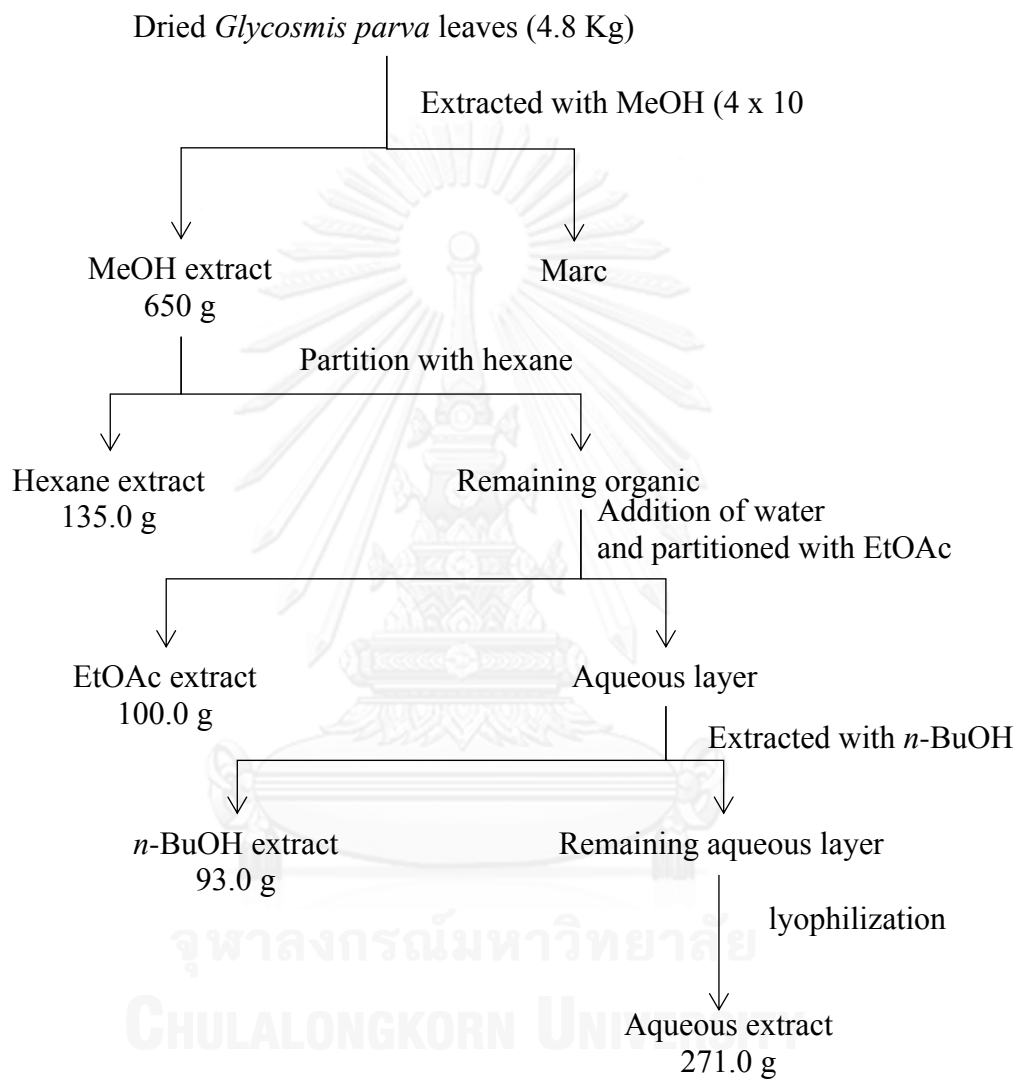
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APPENDIX

จุฬาลงกรณ์มหาวิทยาลัย  
**CHULALONGKORN UNIVERSITY**

## APPENDIX A

APPENDIX A-1: Extraction of *Glycosmis parva* leaves

## APPENDIX B

## Buffers and Reagents

## 1. RPMI 1640 stock solution 1 liter

RPMI powder	10.4	g
NaHCO <sub>3</sub>	1.5	g
Glucose	4.5	g
Sodium pyruvate	0.11	g
HEPES (1M)	10	ml
Penicillin/Streptomycin	10	ml
ddH <sub>2</sub> O	900	ml

Adjust pH to 7.2 with 1 N HCl and 1 N NaOH

Add ddH<sub>2</sub>O to 1 liter and sterilized by filtering through a 0.2 sterile membrane filter

## 2. DMEM stock solution 1 liter

DMEM powder	10.4	g
Sodium bicarbonate	3.7	g
Penicillin/Streptomycin	10	ml
ddH <sub>2</sub> O	900	ml

Adjust pH to 7.4 with 1 N HCl and 1 N NaOH

Add ddH<sub>2</sub>O to 1 liter and sterilized by filtering through a 0.2 sterile membrane filter

## 3. 10x Phosphate Buffered Saline (PBS) 1 liter

NaCl	80.65	g
KCl	2	g
KH <sub>2</sub> PO <sub>4</sub>	2	g
Na <sub>2</sub> HPO <sub>4</sub>	11.5	g
ddH <sub>2</sub> O	900	ml

Adjust pH to 7.4 with 1M HCl

Add ddH<sub>2</sub>O to 1 liter and sterilize by autoclaving

## 4. 10x Assay Buffer for Flow Cytometer 100 ml

HEPES (1M)	10	ml
CaCl <sub>2</sub> (0.1M)	28	ml
NaCl (5M)	25	ml
ddH <sub>2</sub> O	37	ml

## 5. EDTA 0.5M pH 8.0 100 ml

EDTA	18.612	g
ddH <sub>2</sub> O	80	ml

Adjust pH to 8.0 with NaOH

Add ddH<sub>2</sub>O to 100 ml and sterilize by autoclaving

## 6. 5x Tris-Borate-EDTA (TBE) Buffer 1 liter

Tris-base	54	g
Boric acid	27.5	g
EDTA 0.5M pH 8.0	20	ml

Sterilize by autoclaving

## APPENDIX C

## Results

Appendix C-1: Cytotoxic effects of GPE against HT-29 cells at 24, 48 and 72 h.

HT29 Cytotoxicity Study at 24 h	N	Mean	Std. Error	95% Confidence Interval for Mean		Min	Max
				Lower Bound	Upper Bound		
0.2% DMSO	3	1.22	0.13	-0.65	1.78	1.05	1.48
GPE 6.25 µg/ml	3	-2.12	2.07	-11.03	6.80	-5.32	1.76
GPE 12.5 µg/ml	3	-0.38	1.83	-8.24	7.48	-3.87	2.31
GPE 25 µg/ml	3	7.75	1.98	-0.75	16.26	4.07	10.85
GPE 50 µg/ml	3	33.10	2.10	24.07	42.14	29.59	36.85
GPE 100 µg/ml	3	77.31	1.81	69.50	85.11	73.80	79.87
5FU 1 µM	3	0.41	1.05	-4.10	4.91	-1.04	2.44
5FU 10 µM	3	7.57	4.18	-10.43	25.57	0.14	14.61
5FU 100 µM	3	17.70	1.78	10.03	25.38	14.32	20.38
Oxaliplatin 1 µM	3	12.14	0.77	8.84	15.44	10.94	13.57
Oxaliplatin 10 µM	3	20.23	1.00	15.94	24.52	18.91	22.19
Oxaliplatin 100 µM	3	39.23	0.94	35.21	43.26	37.59	40.82



Appendix C-1: Cytotoxic effects of GPE against HT-29 cells at 24, 48 and 72 h. (Cont.)

HT29 Cytotoxicity Study at 48 h	N	Mean	Std. Error	95% Confidence Interval for Mean		Min	Max
				Lower Bound	Upper Bound		
0.2% DMSO	3	1.29	0.06	1.04	1.53	1.20	1.40
GPE 6.25 µg/ml	3	1.12	0.59	-1.42	3.65	0.11	2.15
GPE 12.5 µg/ml	3	-0.15	2.20	-9.63	9.33	-3.87	3.76
GPE 25 µg/ml	3	18.27	2.82	6.12	30.42	14.60	23.82
GPE 50 µg/ml	3	54.57	4.12	36.82	72.31	46.52	60.16
GPE 100 µg/ml	3	90.11	2.91	77.58	102.64	84.29	93.21
5FU 1 µM	3	11.40	3.62	-4.20	26.99	4.16	15.27
5FU 10 µM	3	37.83	6.67	9.15	66.52	24.55	45.45
5FU 100 µM	3	56.52	4.36	37.77	75.27	48.21	62.94
Oxaliplatin 1 µM	3	36.14	4.79	15.51	56.76	26.62	41.92
Oxaliplatin 10 µM	3	58.61	5.08	36.75	80.47	48.45	63.82
Oxaliplatin 100 µM	3	76.78	2.55	65.81	87.74	72.02	80.73

HT29 Cytotoxicity Study at 72 h	N	Mean	Std. Error	95% Confidence Interval for Mean		Min	Max
				Lower Bound	Upper Bound		
0.2% DMSO	3	1.52	0.01	1.46	1.58	1.50	1.55
GPE 6.25 µg/ml	3	5.51	1.77	-2.09	13.10	2.27	8.34
GPE 12.5 µg/ml	3	5.42	3.48	-9.53	20.37	-0.22	11.76
GPE 25 µg/ml	3	23.85	2.31	13.89	33.81	19.29	26.83
GPE 50 µg/ml	3	61.94	11.91	10.69	113.19	39.02	79.03
GPE 100 µg/ml	3	99.11	2.45	88.58	109.65	94.22	101.78
5FU 1 µM	3	25.48	1.98	16.99	33.98	22.72	29.31
5FU 10 µM	3	56.87	4.37	38.06	75.69	48.18	62.04
5FU 100 µM	3	78.99	2.55	68.03	89.94	74.11	82.69
Oxaliplatin 1 µM	3	36.14	6.52	8.10	64.19	23.11	42.84
Oxaliplatin 10 µM	3	74.58	4.34	55.92	93.23	65.94	79.59
Oxaliplatin 100 µM	3	91.97	1.10	87.23	96.71	89.78	93.30

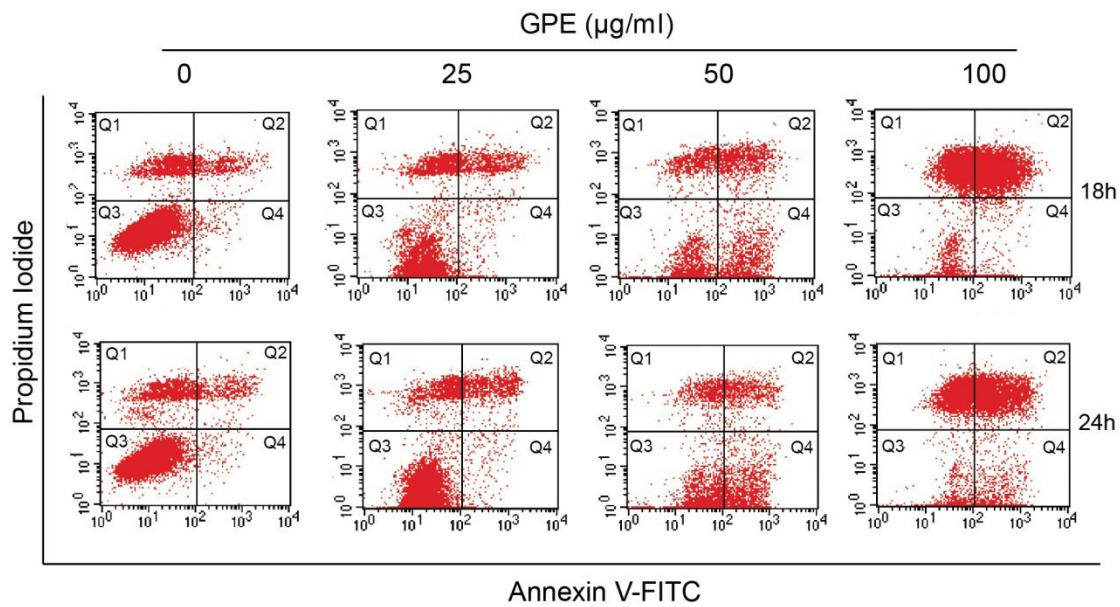
Appendix C-2: The apoptotic effect of GPE on HT-29 cells at 18 and 24.

HT-29 Apoptosis Study at 18 h		N	Mean	Std. Error	95% Confidence Interval for Mean		Min	Max
					Lower Bound	Upper Bound		
Viable	Untreated, non-stained	3	99.86	0.03	99.75	99.98	99.81	99.89
	DMEM	3	83.03	1.85	75.06	91.00	80.91	86.72
	0.2% DMSO	3	87.28	1.69	80.00	94.56	84.71	90.47
	GPE 25 µg/ml	3	83.29	0.42	81.50	85.08	82.47	83.82
	GPE 50 µg/ml	3	67.24	3.16	53.63	80.85	61.22	71.94
	GPE 100 µg/ml	3	30.39	3.39	15.79	44.99	23.86	35.25
	Oxaliplatin 100 µM	3	83.01	0.72	79.92	86.10	82.20	84.44
	5FU 100 µM	3	80.10	0.85	76.43	83.77	78.77	81.69
Necrosis	Untreated, non-stained	3	0.08	0.04	-0.09	0.26	0.01	0.15
	DMEM	3	12.47	1.76	4.90	20.03	9.03	14.83
	0.2% DMSO	3	7.98	1.69	0.72	15.24	4.68	10.25
	GPE 25 µg/ml	3	7.60	0.44	5.71	9.49	6.73	8.13
	GPE 50 µg/ml	3	8.51	0.59	5.95	11.06	7.33	9.24
	GPE 100 µg/ml	3	32.79	9.38	-7.58	73.16	22.31	51.51
	Oxaliplatin 100 µM	3	5.74	1.08	1.08	10.40	4.18	7.82
	5FU 100 µM	3	13.05	1.29	7.50	18.59	10.88	15.34
Late apoptosis/ Necrosis	Untreated, non-stained	3	0.03	0.01	-0.01	0.06	0.01	0.04
	DMEM	3	3.49	0.19	2.68	4.29	3.16	3.81
	0.2% DMSO	3	5.14	1.88	-2.94	13.21	2.94	8.87
	GPE 25 µg/ml	3	7.19	0.41	5.43	8.95	6.38	7.70
	GPE 50 µg/ml	3	9.47	0.92	5.51	13.43	7.76	10.92
	GPE 100 µg/ml	3	33.74	5.65	9.44	58.04	22.87	41.83
	Oxaliplatin 100 µM	3	7.49	1.03	3.07	11.91	5.50	8.92
	5FU 100 µM	3	5.71	0.22	4.75	6.66	5.27	5.99
Early apoptosis	Untreated, non-stained	3	0.00	0.00	0.00	0.00	0.00	0.00
	DMEM	3	1.02	0.13	0.47	1.57	0.77	1.19
	0.2% DMSO	3	1.27	0.06	1.02	1.52	1.18	1.38
	GPE 25 µg/ml	3	1.92	0.15	1.27	2.56	1.69	2.20
	GPE 50 µg/ml	3	14.79	2.91	2.26	27.32	11.07	20.53
	GPE 100 µg/ml	3	3.08	0.66	0.22	5.94	1.75	3.81
	Cisplatin 100 µM	3	4.24	0.39	2.54	5.94	3.71	5.01
	5FU 100 µM	3	1.15	0.28	-0.07	2.36	0.61	1.57
Total death	Untreated, non-stained	3	0.11	0.05	-0.10	0.32	0.02	0.19
	DMEM	3	16.97	1.85	9.02	24.93	13.29	19.09
	0.2% DMSO	3	14.39	3.15	0.82	27.96	9.53	20.30
	GPE 25 µg/ml	3	16.71	0.41	14.93	18.48	16.18	17.52
	GPE 50 µg/ml	3	32.77	3.16	19.17	46.37	28.07	38.78
	GPE 100 µg/ml	3	69.61	3.39	55.03	84.19	64.75	76.13
	Oxaliplatin 100 µM	3	16.99	0.72	13.89	20.09	15.55	17.80
	5FU 100 µM	3	19.90	0.85	16.24	23.56	18.31	21.22

Appendix C-2: The apoptotic effect of GPE on HT-29 cells at 18 and 24. (Cont.)

HT-29 Apoptosis Study at 24 h		N	Mean	Std. Error	95% Confidence Interval for Mean		Min	Max
					Lower Bound	Upper Bound		
Viable	Untreated, non-stained	3	99.90	0.02	99.81	100.00	99.86	99.93
	DMEM	3	86.05	3.64	70.39	101.70	81.50	93.24
	0.2% DMSO	3	86.05	1.42	79.93	92.17	83.43	88.32
	GPE 25 µg/ml	3	85.16	2.25	75.47	94.85	80.74	88.11
	GPE 50 µg/ml	3	63.67	1.31	58.04	69.29	61.66	66.12
	GPE 100 µg/ml	3	24.31	5.78	-0.56	49.18	13.90	33.87
	Oxaliplatin 100 µM	3	84.80	1.44	78.62	90.98	81.94	86.44
	5FU 100 µM	3	79.78	3.16	66.17	93.40	76.00	86.07
Necrosis	Untreated, non-stained	3	0.07	0.01	0.01	0.14	0.05	0.10
	DMEM	3	8.63	2.83	-3.54	20.81	3.20	12.72
	0.2% DMSO	3	9.03	1.79	1.32	16.73	5.65	11.75
	GPE 25 µg/ml	3	6.85	1.26	1.43	12.28	5.48	9.37
	GPE 50 µg/ml	3	7.21	0.19	6.37	8.04	6.93	7.58
	GPE 100 µg/ml	3	22.56	3.17	8.92	36.20	16.22	25.78
	Oxaliplatin 100 µM	3	5.07	0.39	3.39	6.75	4.43	5.78
	5FU 100 µM	3	9.97	1.23	4.69	15.25	7.52	11.34
Late apoptosis/ Necrosis	Untreated, non-stained	3	0.02	0.01	-0.01	0.06	0.01	0.04
	DMEM	3	4.52	0.92	0.55	8.49	2.82	5.99
	0.2% DMSO	3	3.94	0.50	1.77	6.10	3.16	4.88
	GPE 25 µg/ml	3	6.45	1.12	1.64	11.25	4.42	8.27
	GPE 50 µg/ml	3	8.16	0.23	7.15	9.17	7.76	8.57
	GPE 100 µg/ml	3	41.89	6.54	13.74	70.04	33.28	54.73
	Oxaliplatin 100 µM	3	5.81	0.55	3.44	8.18	5.04	6.88
	5FU 100 µM	3	8.20	1.28	2.69	13.72	5.97	10.41
Early apoptosis	Untreated, non-stained	3	0.00	0.00	0.00	0.00	0.00	0.00
	DMEM	3	0.80	0.11	0.32	1.29	0.64	1.02
	0.2% DMSO	3	0.99	0.11	0.50	1.48	0.77	1.15
	GPE 25 µg/ml	3	1.54	0.16	0.85	2.23	1.23	1.76
	GPE 50 µg/ml	3	20.97	0.91	17.05	24.89	19.19	22.20
	GPE 100 µg/ml	3	8.89	3.77	-7.34	25.13	2.79	15.79
	Oxaliplatin 100 µM	3	4.18	0.67	1.29	7.06	3.09	5.40
	5FU 100 µM	3	0.79	0.17	0.06	1.52	0.45	0.97
Total death	Untreated, non-stained	3	0.10	0.02	0.00	0.19	0.07	0.14
	DMEM	3	13.96	3.64	-1.69	29.61	6.77	18.50
	0.2% DMSO	3	13.95	1.42	7.83	20.07	11.68	16.57
	GPE 25 µg/ml	3	14.84	2.26	5.14	24.54	11.89	19.27
	GPE 50 µg/ml	3	36.34	1.31	30.70	41.97	33.88	38.35
	GPE 100 µg/ml	3	73.34	3.81	56.97	89.71	66.13	79.05
	Oxaliplatin 100 µM	3	15.06	1.52	8.51	21.60	13.13	18.06
	5FU 100 µM	3	18.96	2.61	7.72	30.20	13.94	22.72

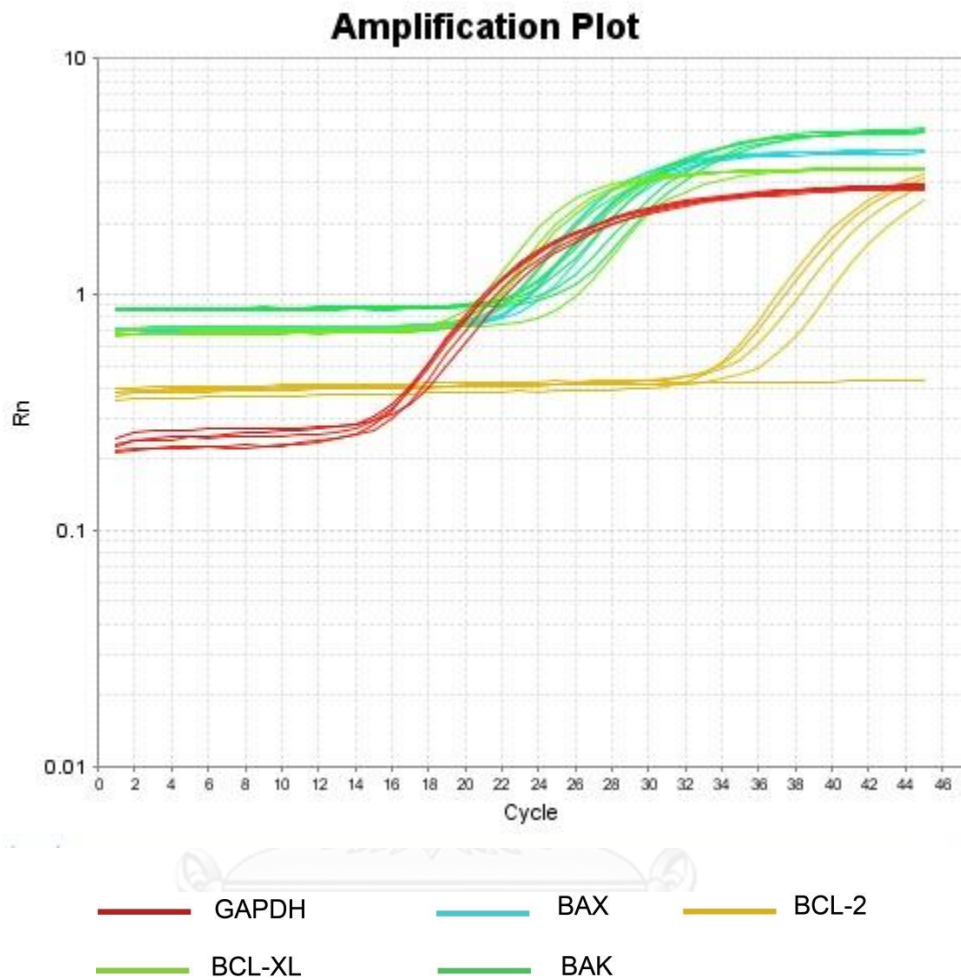
Appendix C-3: Representative flow cytometric analysis of GPE induces apoptosis in HT-29 cells.



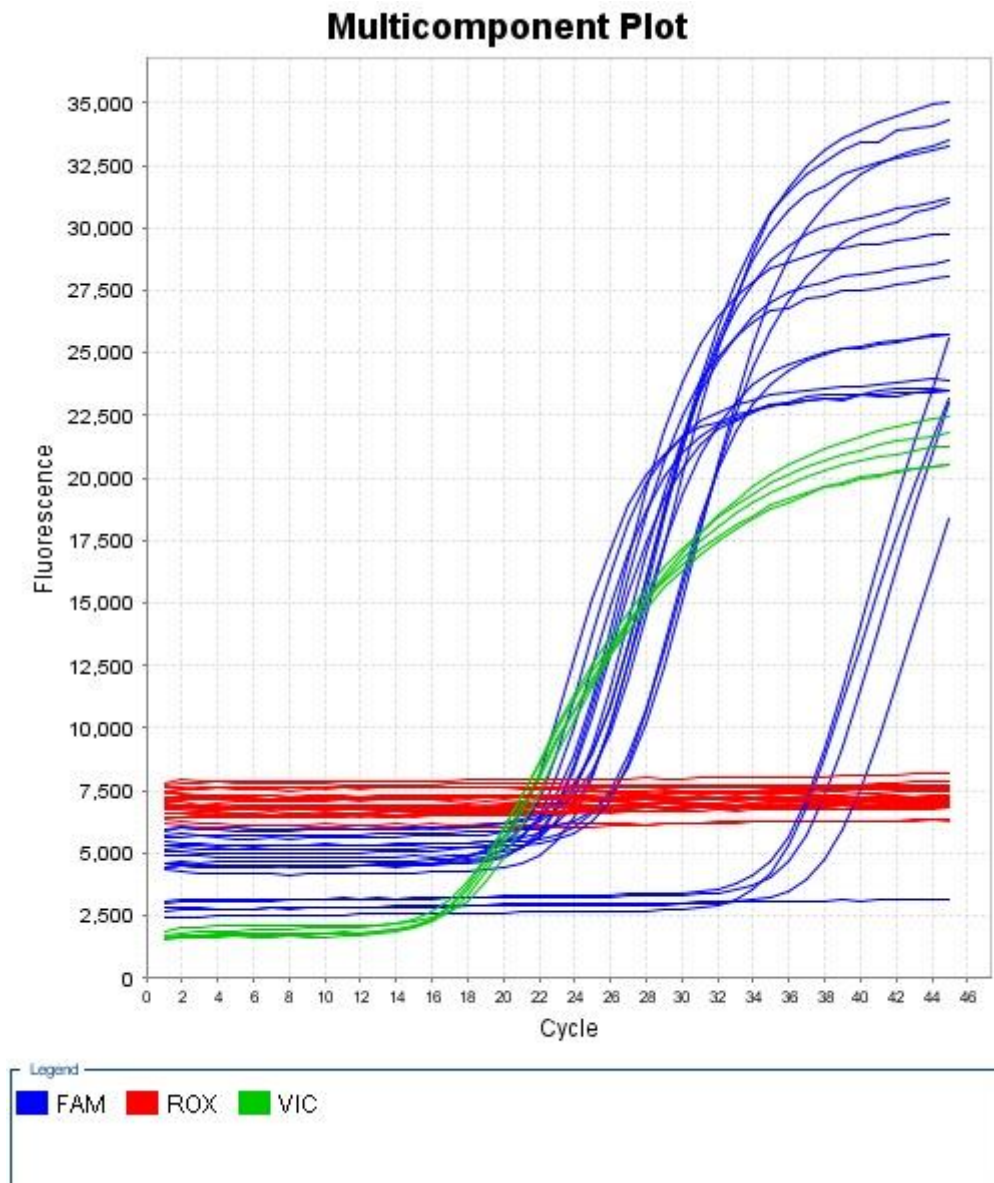
Appendix C-4: Effect of GPE on the expression of BCL-2 family genes in HT-29 cells.

BCL-2 family gene expression in HT-29		N	Mean	Std. Error	95% Confidence Interval for Mean		Min	Max
					Lower Bound	Upper Bound		
BCL-2	0.2% DMSO	3	1.00	0.00	1.00	1.00	1.00	1.00
	Cisplatin 100 $\mu$ M	3	0.00	0.00	0.00	0.00	0.00	0.00
	GPE 25 $\mu$ g/ml	3	0.68	0.21	-0.24	1.61	0.47	1.11
	GPE 50 $\mu$ g/ml	3	0.32	0.13	-0.23	0.87	0.12	0.56
	GPE 100 $\mu$ g/ml	3	0.55	0.16	-0.14	1.25	0.26	0.81
BCL-XL	0.2% DMSO	3	1.00	0.00	1.00	1.00	1.00	1.00
	Cisplatin 100 $\mu$ M	3	0.08	0.00	0.06	0.10	0.07	0.09
	GPE 25 $\mu$ g/ml	3	1.04	0.15	0.40	1.68	0.85	1.34
	GPE 50 $\mu$ g/ml	3	0.60	0.09	0.21	1.00	0.44	0.76
	GPE 100 $\mu$ g/ml	3	0.63	0.09	0.24	1.03	0.51	0.81
BAK	0.2% DMSO	3	1.00	0.00	1.00	1.00	1.00	1.00
	Cisplatin 100 $\mu$ M	3	0.56	0.09	0.16	0.95	0.38	0.67
	GPE 25 $\mu$ g/ml	3	1.25	0.16	0.57	1.92	1.06	1.56
	GPE 50 $\mu$ g/ml	3	1.79	0.08	1.46	2.11	1.64	1.90
	GPE 100 $\mu$ g/ml	3	0.81	0.18	0.02	1.59	0.61	1.17
BAX	0.2% DMSO	3	1.00	0.00	1.00	1.00	1.00	1.00
	Cisplatin 100 $\mu$ M	3	0.51	0.09	0.14	0.88	0.37	0.67
	GPE 25 $\mu$ g/ml	3	0.74	0.09	0.33	1.15	0.58	0.90
	GPE 50 $\mu$ g/ml	3	0.48	0.07	0.19	0.77	0.38	0.61
	GPE 100 $\mu$ g/ml	3	0.76	0.04	0.56	0.95	0.67	0.83

Appendix C-5: Representative amplification plot and from real-time RT-PCR experiment on the expression of genes in BCL-2 family in HT-29 cells.



Appendix C-6: Representative multicomponent plot from real-time RT-PCR experiment on the expression of genes in BCL-2 family in HT-29 cells.



FAM = Fluorescence dye labeled with studied genes.  
(BCL-2, BCL-XL, BAK, BAX)

ROX = Passive reference dye functioned as loading control.

VIC = Fluorescence dye labeled with housekeeping genes (GAPDH).

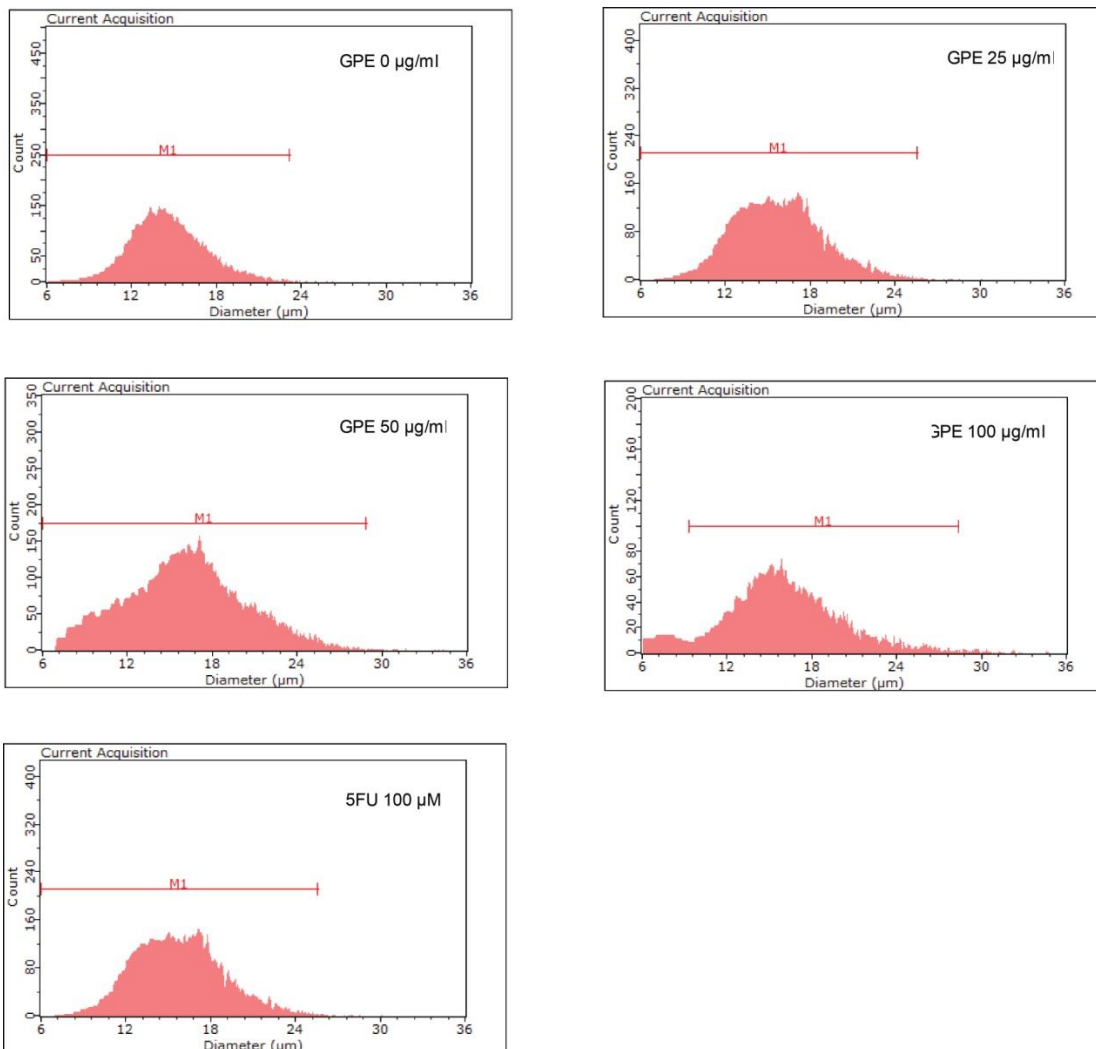
Appendix C-7: Anti-proliferative effect of GPE in HT-29 cells.

Cell Proliferation in HT-29	N	Mean	Std. Error	95% Confidence Interval for Mean		Min	Max
				Lower Bound	Upper Bound		
0.2% DMSO	3	100.00%	0.00%	100.00%	100.00%	100.00%	100.00%
GPE 25 µg/ml	3	72.40%	5.83%	47.30%	97.50%	65.28%	83.96%
GPE 50 µg/ml	3	52.46%	6.45%	24.72%	80.20%	39.58%	59.47%
GPE 100 µg/ml	3	26.68%	6.12%	0.36%	53.00%	16.07%	37.26%
5FU 100 µM	3	44.47%	3.83%	27.98%	60.95%	37.74%	51.01%





Appendix C-8: Representative histogram of cell counting from Scepter™ automate cell counter for determination of anti-proliferative effect of GPE in HT-29 cells.



Appendix C-9: Distribution of HT-29 cells in the cell cycle phases after treatment with GPE.

Cell cycle analysis in HT-29		N	Mean	Std. Error	95% Confidence Interval for Mean		Min	Max
					Lower Bound	Upper Bound		
G1	0.2% DMSO	3	50.13	1.28	44.61	55.66	47.73	52.12
	5FU 100 $\mu$ M	3	33.32	1.67	26.15	40.49	30.80	36.47
	GPE 25 $\mu$ g/ml	3	40.95	1.33	35.22	46.67	38.74	43.34
	GPE 50 $\mu$ g/ml	3	60.73	0.98	56.52	64.94	59.04	62.43
	GPE 100 $\mu$ g/ml	3	47.75	2.71	36.09	59.41	43.56	52.82
S	0.2% DMSO	3	32.00	0.51	29.80	34.21	31.00	32.68
	5FU 100 $\mu$ M	3	61.00	1.81	53.24	68.77	58.85	64.59
	GPE 25 $\mu$ g/ml	3	44.85	2.59	33.71	55.98	40.83	49.68
	GPE 50 $\mu$ g/ml	3	26.78	1.76	19.22	34.33	24.02	30.04
	GPE 100 $\mu$ g/ml	3	18.72	3.55	3.44	34.00	12.16	24.36
G2M	0.2% DMSO	3	17.86	1.04	13.39	22.33	16.78	19.94
	5FU 100 $\mu$ M	3	5.68	2.05	-3.16	14.52	2.73	9.63
	GPE 25 $\mu$ g/ml	3	14.21	1.33	8.50	19.92	11.58	15.84
	GPE 50 $\mu$ g/ml	3	12.49	1.38	6.54	18.44	10.92	15.25
	GPE 100 $\mu$ g/ml	3	33.53	0.85	29.88	37.18	32.08	35.02

Appendix C-10: Effect of GPE on the expression of cell cycle regulatory genes in HT-29 cells.

Expression of cell cycle regulators in HT-29		N	Mean	Std. Error	95% Confidence Interval for Mean		Min	Max
					Lower Bound	Upper Bound		
CyclinA	0.2% DMSO	3	100.00	0.00	100.00	100.00	100.00	100.00
	5FU 100 $\mu$ M	3	90.58	3.73	74.55	106.61	84.25	97.15
	GPE 25 $\mu$ g/ml	3	91.05	5.29	68.31	113.79	85.54	101.62
	GPE 50 $\mu$ g/ml	3	88.54	3.75	72.40	104.69	81.07	92.87
	GPE 100 $\mu$ g/ml	3	81.44	6.03	55.47	107.40	70.33	91.08
CyclinB1	0.2% DMSO	3	100.00	0.00	100.00	100.00	100.00	100.00
	5FU 100 $\mu$ M	3	94.25	3.66	78.50	110.01	87.20	99.49
	GPE 25 $\mu$ g/ml	3	95.13	2.55	84.17	106.09	91.86	100.15
	GPE 50 $\mu$ g/ml	3	92.64	4.70	72.42	112.87	83.28	98.05
	GPE 100 $\mu$ g/ml	3	90.14	6.29	63.09	117.19	77.82	98.47
CyclinD1	0.2% DMSO	3	100.00	0.00	100.00	100.00	100.00	100.00
	5FU 100 $\mu$ M	3	95.93	3.75	79.80	112.07	89.09	102.01
	GPE 25 $\mu$ g/ml	2	88.07	9.11	-27.68	203.82	78.96	97.18
	GPE 50 $\mu$ g/ml	3	87.60	8.43	51.32	123.89	70.90	97.98
	GPE 100 $\mu$ g/ml	3	83.15	7.90	49.16	117.14	67.35	91.21
CyclinE	0.2% DMSO	3	100.00	0.00	100.00	100.00	100.00	100.00
	5FU 100 $\mu$ M	3	96.24	0.63	93.53	98.95	95.01	97.10
	GPE 25 $\mu$ g/ml	3	89.46	4.34	70.77	108.14	80.80	94.40
	GPE 50 $\mu$ g/ml	3	82.02	5.93	56.50	107.54	70.21	88.86
	GPE 100 $\mu$ g/ml	3	81.55	3.87	64.91	98.18	75.38	88.67
p21	0.2% DMSO	3	100.00	0.00	100.00	100.00	100.00	100.00
	5FU 100 $\mu$ M	3	107.94	1.80	100.20	115.68	104.47	110.49
	GPE 25 $\mu$ g/ml	3	93.36	2.71	81.72	105.00	89.07	98.36
	GPE 50 $\mu$ g/ml	3	100.79	4.72	80.48	121.09	92.85	109.18
	GPE 100 $\mu$ g/ml	3	108.24	4.64	88.30	128.18	99.63	115.52
p53	0.2% DMSO	3	100.00	0.00	100.00	100.00	100.00	100.00
	5FU 100 $\mu$ M	3	101.32	6.14	74.92	127.73	92.23	113.01
	GPE 25 $\mu$ g/ml	3	98.38	7.26	67.13	129.64	85.73	110.89
	GPE 50 $\mu$ g/ml	3	94.14	7.85	60.37	127.91	78.58	103.70
	GPE 100 $\mu$ g/ml	3	88.30	9.83	46.00	130.60	69.28	102.13

Appendix C-11: Cytotoxic effects of GPE against Colo-205 cells at 24, 48 and 72 h.

Colo-205 Cytotoxicity Study at 24 h	N	Mean	Std. Error	95% Confidence Interval for Mean		Minimum	Maximum
				Lower Bound	Upper Bound		
				0.2% DMSO	3		
GPE 6.25 µg/ml	3	-3.29	1.17	-8.31	1.73	-5.18	-1.16
GPE 12.5 µg/ml	3	9.79	1.02	5.41	14.17	8.09	11.61
GPE 25 µg/ml	3	19.81	3.82	3.36	36.26	15.08	27.38
GPE 50 µg/ml	3	40.22	6.80	10.97	69.46	31.52	53.62
GPE 100 µg/ml	3	87.25	4.09	69.65	104.84	80.28	94.44
5FU 1 µM	3	-0.51	1.18	-5.60	4.57	-1.83	1.85
5FU 10 µM	3	12.18	1.74	4.68	19.68	9.74	15.56
5FU 100 µM	3	16.97	0.84	13.36	20.58	15.37	18.21
Oxaliplatin 1 µM	3	7.76	0.79	4.36	11.16	6.66	9.29
Oxaliplatin 10 µM	3	15.62	1.10	10.89	20.36	13.55	17.29
Oxaliplatin 100 µM	3	23.67	4.16	5.77	41.58	18.61	31.92



## Appendix C-11: Cytotoxic effects of GPE against Colo-205 cells at 24, 48 and 72 h.

(Cont.)

Colo-205 Cytotoxicity Study at 48 h	N	Mean	Std. Error	95% Confidence Interval for Mean		Minimum	Maximum
				Lower Bound	Upper Bound		
0.2% DMSO	3	1.68	0.08	1.35	2.00	1.53	1.77
GPE 6.25 µg/ml	3	-7.42	0.49	-9.51	-5.33	-7.99	-6.46
GPE 12.5 µg/ml	3	2.90	3.34	-11.48	17.27	-3.40	7.98
GPE 25 µg/ml	3	65.88	6.50	37.92	93.85	56.34	78.30
GPE 50 µg/ml	3	89.40	2.57	78.33	100.47	85.61	94.31
GPE 100 µg/ml	3	100.25	0.44	98.36	102.15	99.39	100.83
5FU 1 µM	3	9.55	3.02	-3.43	22.52	6.38	15.57
5FU 10 µM	3	45.32	0.48	43.26	47.39	44.52	46.18
5FU 100 µM	3	59.74	0.92	55.77	63.70	57.91	60.84
Oxaliplatin 1 µM	3	26.13	3.29	11.96	40.30	22.20	32.67
Oxaliplatin 10 µM	3	38.15	2.11	29.06	47.25	35.64	42.36
Oxaliplatin 100 µM	3	79.54	1.60	72.64	86.44	76.49	81.92

Colo-205 Cytotoxicity Study at 72 h	N	Mean	Std. Error	95% Confidence Interval for Mean		Minimum	Maximum
				Lower Bound	Upper Bound		
0.2% DMSO	3	1.77	0.05	1.54	2.00	1.66	1.83
GPE 6.25 µg/ml	3	-3.09	1.15	-8.05	1.88	-4.60	-0.82
GPE 12.5 µg/ml	3	8.57	3.24	-5.37	22.51	4.58	14.99
GPE 25 µg/ml	3	74.58	11.22	26.30	122.85	58.30	96.09
GPE 50 µg/ml	3	100.33	0.64	97.55	103.10	99.16	101.38
GPE 100 µg/ml	3	100.95	1.17	95.92	105.97	98.69	102.59
5FU 1 µM	3	12.69	1.96	4.24	21.14	10.43	16.60
5FU 10 µM	3	54.74	0.80	51.29	58.19	53.29	56.06
5FU 100 µM	3	82.02	0.31	80.68	83.36	81.65	82.64
Oxaliplatin 1 µM	3	31.54	5.77	6.72	56.36	25.74	43.08
Oxaliplatin 10 µM	3	49.71	2.93	37.10	62.31	46.05	55.50
Oxaliplatin 100 µM	3	92.19	1.22	86.94	97.43	89.78	93.73

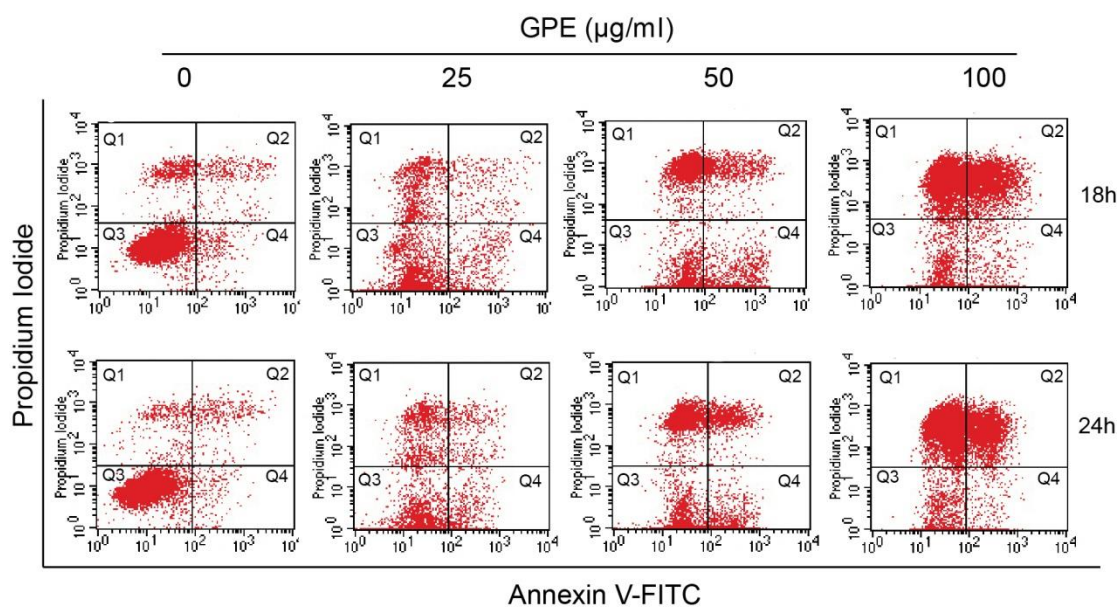
Appendix C-12: The apoptotic effect of GPE on Colo-205 cells at 18 and 24.

Colo-205 Apoptosis Study 18 h		N	Mean	Std. Error	95% Confidence Interval for Mean		Minimum	Maximum
					Lower Bound	Upper Bound		
Viable	Untreated, non-stained	3	99.68	0.12	99.18	100.17	99.45	99.83
	DMEM	3	87.82	0.43	85.97	89.68	86.97	88.36
	0.2% DMSO	3	88.45	0.96	84.32	92.59	86.82	90.15
	GPE 25 µg/ml	3	67.96	1.54	61.33	74.59	65.10	70.38
	GPE 50 µg/ml	3	53.40	4.15	35.57	71.24	48.72	61.67
	GPE 100 µg/ml	3	39.27	7.63	6.46	72.09	26.35	52.75
	Oxaliplatin 100 µM	3	76.63	1.69	69.38	83.88	73.33	78.88
	5FU 100 µM	3	82.04	0.45	80.09	83.99	81.25	82.82
Necrosis	Untreated, non-stained	3	0.29	0.12	-0.22	0.81	0.14	0.53
	DMEM	3	6.61	0.32	5.24	7.98	6.07	7.17
	0.2% DMSO	3	5.57	0.61	2.95	8.20	4.88	6.79
	GPE 25 µg/ml	3	13.81	1.13	8.96	18.66	11.59	15.25
	GPE 50 µg/ml	3	11.93	4.22	-6.23	30.09	6.81	20.30
	GPE 100 µg/ml	3	25.47	7.03	-4.77	55.72	16.45	39.32
	Oxaliplatin 100 µM	3	7.34	0.59	4.80	9.87	6.23	8.24
	5FU 100 µM	3	10.21	0.36	8.68	11.74	9.57	10.80
Late apoptosis/ Necrosis	Untreated, non-stained	3	0.03	0.01	0.01	0.05	0.02	0.04
	DMEM	3	3.47	0.21	2.55	4.39	3.10	3.84
	0.2% DMSO	3	3.71	0.47	1.67	5.74	2.81	4.42
	GPE 25 µg/ml	3	5.35	0.35	3.84	6.85	4.91	6.04
	GPE 50 µg/ml	3	6.97	1.80	-0.76	14.70	3.75	9.96
	GPE 100 µg/ml	3	20.83	3.70	4.92	36.75	13.54	25.56
	Oxaliplatin 100 µM	3	5.36	0.57	2.91	7.81	4.58	6.47
	5FU 100 µM	3	5.84	0.02	5.73	5.94	5.79	5.87
Early apoptosis	Untreated, non-stained	3	0.00	0.00	0.00	0.00	0.00	0.00
	DMEM	3	2.09	0.04	1.91	2.28	2.02	2.17
	0.2% DMSO	3	2.27	0.12	1.75	2.79	2.14	2.51
	GPE 25 µg/ml	3	12.88	2.30	2.97	22.80	9.47	17.27
	GPE 50 µg/ml	3	27.70	5.09	5.82	49.58	19.93	37.27
	GPE 100 µg/ml	3	14.42	2.83	2.26	26.58	8.77	17.26
	Oxaliplatin 100 µM	3	10.68	1.79	2.99	18.36	7.85	13.98
	5FU 100 µM	3	1.92	0.08	1.60	2.24	1.83	2.07
Total death	Untreated, non-stained	3	0.32	0.12	-0.17	0.82	0.17	0.55
	DMEM	3	12.18	0.43	10.32	14.03	11.64	13.03
	0.2% DMSO	3	11.55	0.96	7.40	15.70	9.85	13.19
	GPE 25 µg/ml	3	32.04	1.54	25.43	38.66	29.63	34.90
	GPE 50 µg/ml	3	46.60	4.15	28.76	64.44	38.33	51.28
	GPE 100 µg/ml	3	60.73	7.63	27.91	93.54	47.25	73.65
	Oxaliplatin 100 µM	3	23.37	1.69	16.11	30.64	21.12	26.68
	5FU 100 µM	3	17.96	0.45	16.04	19.89	17.19	18.74

Appendix C-12: The apoptotic effect of GPE on Colo-205 cells at 18 and 24. (Cont.)

Colo-205 Apoptosis Study 24 h		N	Mean	Std. Error	95% Confidence Interval for Mean		Minimum	Maximum
					Lower Bound	Upper Bound		
Viable	Untreated, non-stained	3	99.77	0.10	99.35	100.19	99.58	99.91
	DMEM	3	89.32	0.82	85.77	92.86	87.81	90.65
	0.2% DMSO	3	90.03	1.40	84.02	96.04	88.28	92.79
	GPE 25 µg/ml	3	61.57	4.60	41.78	81.36	53.64	69.57
	GPE 50 µg/ml	3	50.96	4.32	32.38	69.54	46.21	59.58
	GPE 100 µg/ml	3	35.20	4.22	17.05	53.35	27.38	41.85
	Oxaliplatin 100 µM	3	73.31	2.37	63.12	83.49	68.70	76.55
	5FU 100 µM	3	83.46	0.59	80.93	85.99	82.35	84.35
Necrosis	Untreated, non-stained	3	0.21	0.09	-0.17	0.59	0.08	0.38
	DMEM	3	4.67	0.69	1.71	7.64	3.96	6.05
	0.2% DMSO	3	4.54	1.33	-1.17	10.24	1.89	6.01
	GPE 25 µg/ml	3	16.22	2.69	4.66	27.78	11.37	20.65
	GPE 50 µg/ml	3	16.64	2.43	6.20	27.08	13.58	21.43
	GPE 100 µg/ml	3	23.05	3.29	8.90	37.19	17.68	29.02
	Oxaliplatin 100 µM	3	10.32	1.37	4.43	16.21	8.09	12.81
	5FU 100 µM	3	7.25	0.40	5.51	8.99	6.57	7.97
Late apoptosis/ Necrosis	Untreated, non-stained	3	0.02	0.01	-0.01	0.05	0.01	0.03
	DMEM	3	3.99	0.28	2.79	5.18	3.61	4.53
	0.2% DMSO	3	2.99	0.20	2.14	3.84	2.69	3.36
	GPE 25 µg/ml	3	6.14	1.77	-1.48	13.76	3.28	9.38
	GPE 50 µg/ml	3	8.12	1.12	3.29	12.95	6.14	10.03
	GPE 100 µg/ml	3	23.68	3.64	8.00	39.35	16.80	29.20
	Oxaliplatin 100 µM	3	5.69	0.99	1.43	9.96	3.85	7.25
	5FU 100 µM	3	6.62	0.08	6.28	6.95	6.48	6.75
Early apoptosis	Untreated, non-stained	3	0.00	0.00	0.00	0.00	0.00	0.00
	DMEM	3	2.02	0.28	0.83	3.22	1.57	2.53
	0.2% DMSO	3	2.45	0.11	1.99	2.90	2.29	2.65
	GPE 25 µg/ml	3	16.07	5.52	-7.66	39.81	6.50	25.61
	GPE 50 µg/ml	3	24.28	5.72	-0.32	48.88	12.85	30.18
	GPE 100 µg/ml	3	18.07	2.84	5.84	30.31	14.40	23.67
	Oxaliplatin 100 µM	3	10.68	0.65	7.87	13.50	9.38	11.43
	5FU 100 µM	3	2.66	0.85	-0.99	6.32	1.06	3.95
Total death	Untreated, non-stained	3	0.23	0.10	-0.18	0.64	0.09	0.41
	DMEM	3	10.68	0.82	7.14	14.23	9.35	12.19
	0.2% DMSO	3	9.97	1.40	3.96	15.99	7.21	11.72
	GPE 25 µg/ml	3	38.43	4.60	18.65	58.22	30.43	46.36
	GPE 50 µg/ml	3	49.04	4.32	30.46	67.62	40.42	53.79
	GPE 100 µg/ml	3	64.80	4.22	46.65	82.95	58.15	72.62
	Oxaliplatin 100 µM	3	26.69	2.37	16.51	36.88	23.45	31.30
	5FU 100 µM	3	16.53	0.59	14.00	19.07	15.65	17.65

Appendix C-13: Representative flow cytometric analysis of GPE induces apoptosis in Colo-205 cells.

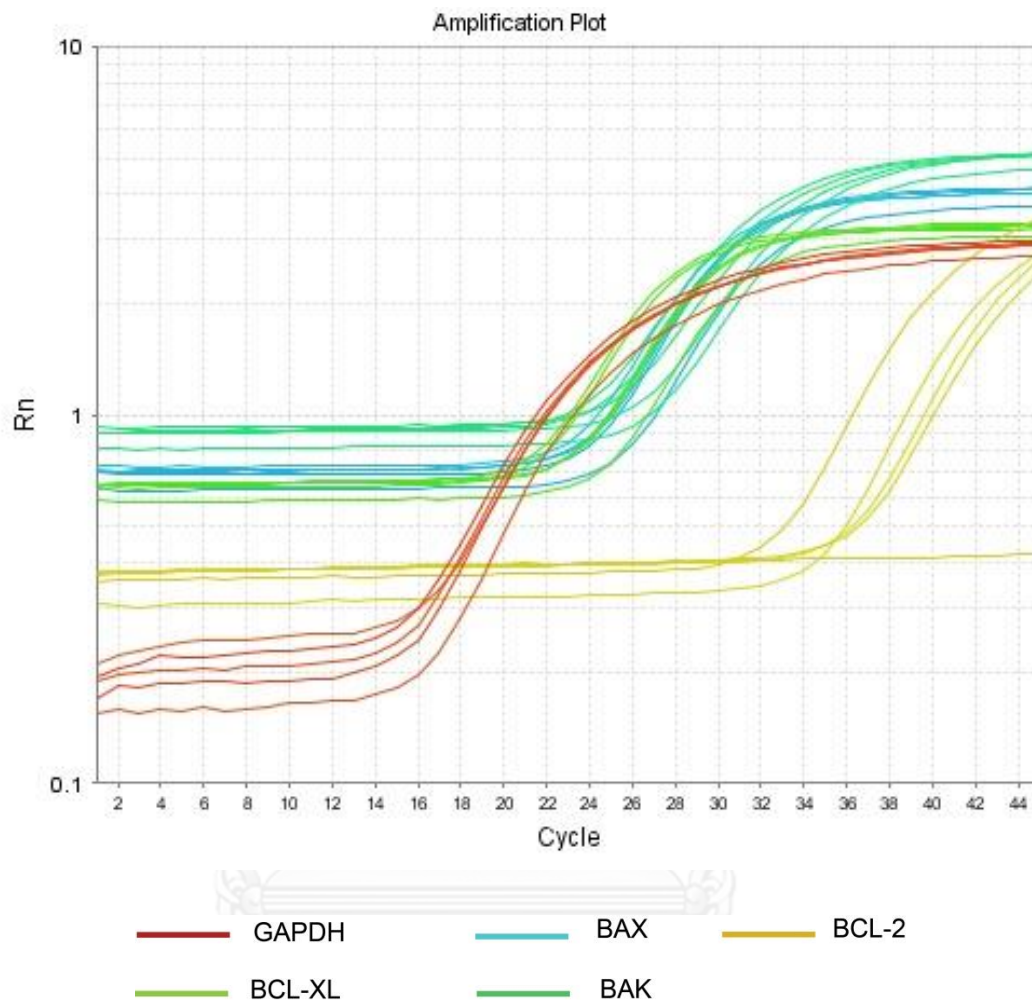




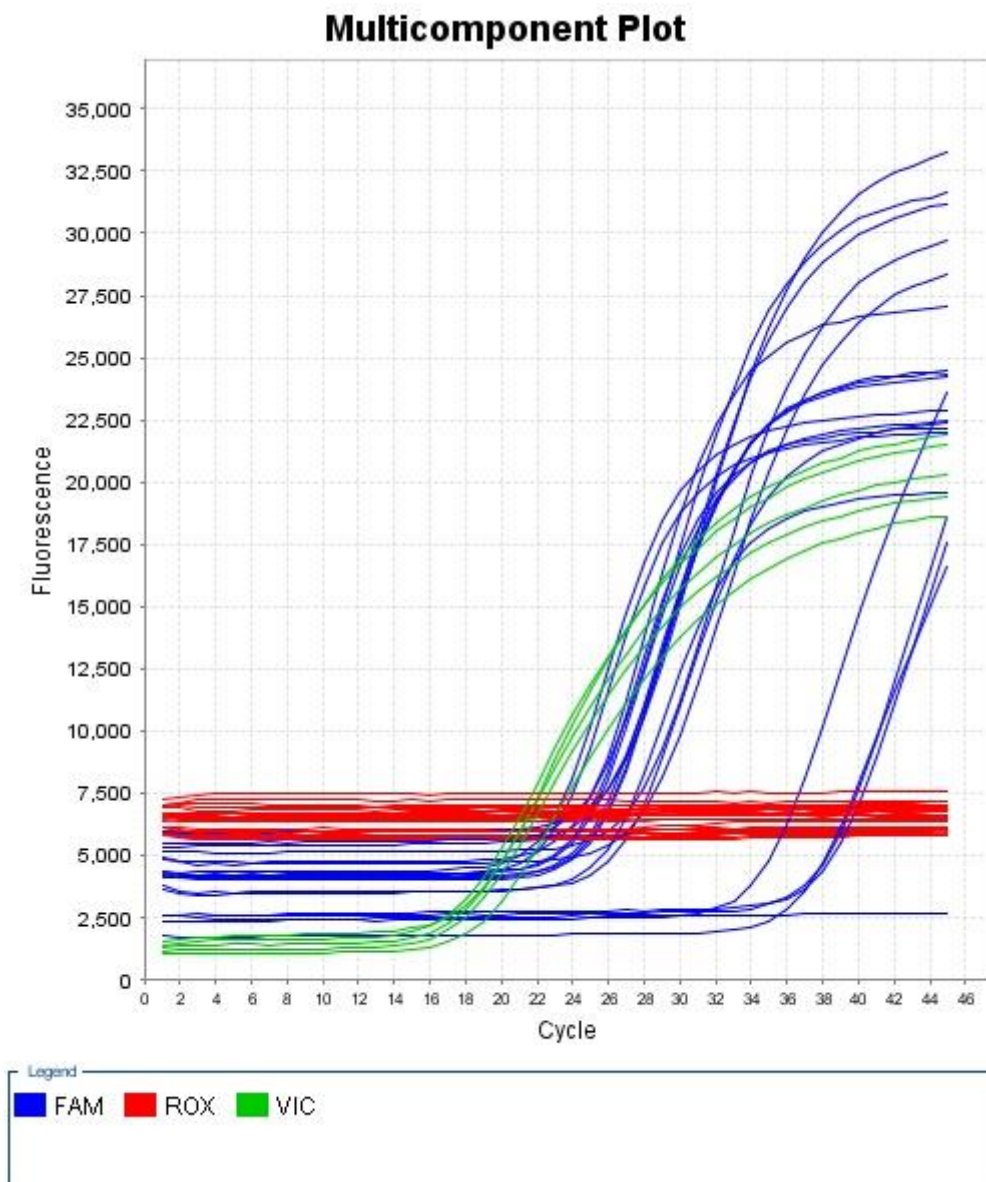
Appendix C-14: Effect of GPE on the expression of BCL-2 family genes in Colo-205 cells.

BCL-2 family gene expression in Colo-205		N	Mean	Std. Error	95% Confidence Interval for Mean		Min	Max
					Lower Bound	Upper Bound		
BCL-2	0.2% DMSO	3	1.00	0.00	1.00	1.00	1.00	1.00
	Cisplatin 100 $\mu$ M	3	0.00	0.00	0.00	0.00	0.00	0.00
	GPE 25 $\mu$ g/ml	3	0.06	0.01	0.03	0.10	0.05	0.07
	GPE 50 $\mu$ g/ml	3	0.39	0.07	0.10	0.67	0.25	0.46
	GPE 100 $\mu$ g/ml	3	0.09	0.05	-0.11	0.29	0.00	0.16
BCL-XL	0.2% DMSO	3	1.00	0.00	1.00	1.00	1.00	1.00
	Cisplatin 100 $\mu$ M	3	0.39	0.08	0.06	0.73	0.30	0.55
	GPE 25 $\mu$ g/ml	3	0.98	0.04	0.82	1.14	0.91	1.04
	GPE 50 $\mu$ g/ml	3	0.23	0.00	0.21	0.24	0.22	0.23
	GPE 100 $\mu$ g/ml	3	0.42	0.05	0.20	0.63	0.34	0.51
BAK	0.2% DMSO	3	1.00	0.00	1.00	1.00	1.00	1.00
	Cisplatin 100 $\mu$ M	3	1.89	0.40	0.18	3.60	1.45	2.68
	GPE 25 $\mu$ g/ml	3	1.21	0.20	0.32	2.09	0.98	1.62
	GPE 50 $\mu$ g/ml	3	0.66	0.08	0.34	0.98	0.58	0.81
	GPE 100 $\mu$ g/ml	3	0.62	0.11	0.13	1.10	0.47	0.84
BAX	0.2% DMSO	3	1.00	0.00	1.00	1.00	1.00	1.00
	Cisplatin 100 $\mu$ M	3	0.98	0.08	0.65	1.32	0.85	1.11
	GPE 25 $\mu$ g/ml	3	0.54	0.11	0.07	1.01	0.37	0.74
	GPE 50 $\mu$ g/ml	3	0.53	0.02	0.46	0.60	0.51	0.57
	GPE 100 $\mu$ g/ml	3	0.95	0.11	0.46	1.43	0.72	1.09

Appendix C-15: Representative amplification plot from real-time RT-PCR experiment on the expression of genes in BCL-2 family in Colo-205 cells.



Appendix C-16: Representative multicomponent plot from real-time RT-PCR experiment on the expression of genes in BCL-2 family in Colo-205 cells.



FAM = Fluorescence dye labeled with studied genes.  
(BCL-2, BCL-XL, BAK, BAX)

ROX = Passive reference dye functioned as loading control.

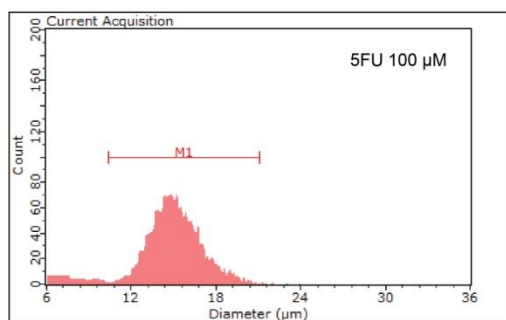
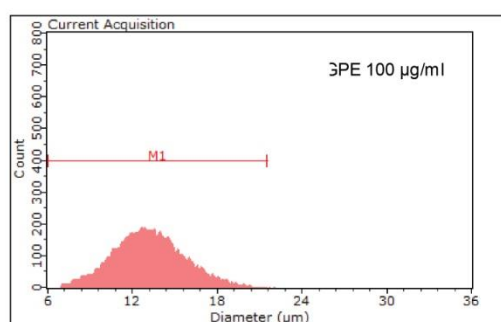
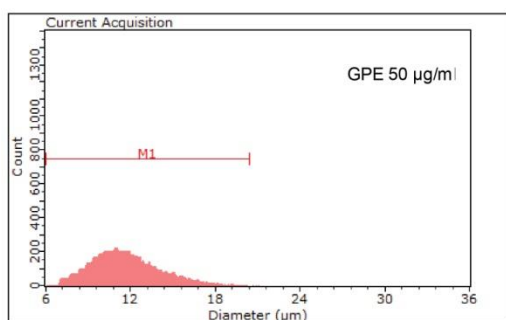
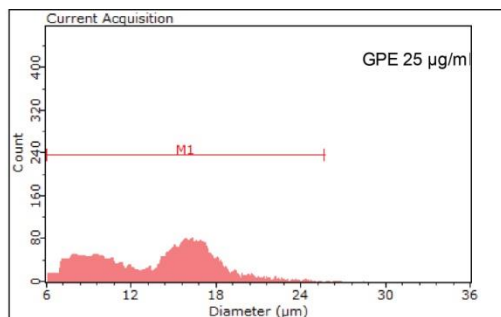
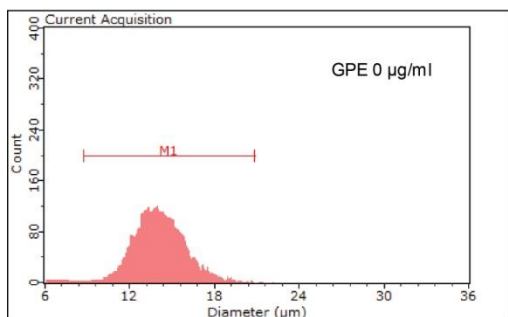
VIC = Fluorescence dye labeled with housekeeping genes (GAPDH).

Appendix C-17: Anti-proliferative effect of GPE in Colo-205 cells.

Cell Proliferation in Colo-205	N	Mean	Std. Error	95% Confidence Interval for Mean		Min	Max
				Lower Bound	Upper Bound		
0.2% DMSO	3	100.00%	0.00%	100.00%	100.00%	100.00%	100.00%
GPE 25 µg/ml	3	60.71%	5.13%	38.64%	82.78%	51.75%	69.52%
GPE 50 µg/ml	3	71.63%	8.84%	33.59%	109.66%	62.01%	89.29%
GPE 100 µg/ml	3	59.91%	2.44%	49.42%	70.40%	57.07%	64.76%
5FU 100 µM	3	53.91%	5.08%	32.04%	75.78%	43.80%	59.90%



Appendix C-18: Representative histogram of cell counting from Scepter™ automate cell counter for determination of anti-proliferative effect of GPE in Colo-205 cells.



Appendix C-19: Distribution of Colo-205 cells in the cell cycle phases after treatment with GPE.

Cell cycle analysis in Colo-205		N	Mean	Std. Error	95% Confidence Interval for Mean		Min	Max
					Lower Bound	Upper Bound		
G1	0.2% DMSO	3	51.49	1.73	44.03	58.95	48.03	53.39
	5FU 100 $\mu$ M	3	50.24	2.86	37.95	62.52	44.79	54.45
	GPE 25 $\mu$ g/ml	3	65.39	1.04	60.91	69.87	64.04	67.44
	GPE 50 $\mu$ g/ml	3	53.31	0.89	49.50	57.12	52.36	55.08
	GPE 100 $\mu$ g/ml	3	45.85	2.22	36.30	55.40	41.42	48.34
S	0.2% DMSO	3	36.37	0.83	32.81	39.93	35.18	37.96
	5FU 100 $\mu$ M	3	47.75	1.12	42.94	52.56	45.55	49.18
	GPE 25 $\mu$ g/ml	3	25.49	2.00	16.89	34.10	21.51	27.80
	GPE 50 $\mu$ g/ml	3	16.62	1.79	8.91	24.32	14.35	20.15
	GPE 100 $\mu$ g/ml	3	38.15	2.15	28.91	47.40	35.52	42.41
G2M	0.2% DMSO	3	12.14	0.95	8.07	16.22	10.98	14.02
	5FU 100 $\mu$ M	3	2.01	2.01	-6.65	10.68	0.00	6.04
	GPE 25 $\mu$ g/ml	3	9.12	1.03	4.66	13.57	7.51	11.05
	GPE 50 $\mu$ g/ml	3	30.07	1.73	22.63	37.52	27.36	33.29
	GPE 100 $\mu$ g/ml	3	15.99	0.46	14.02	17.96	15.13	16.69

Appendix C-20: Effect of GPE on the expression of cell cycle regulatory genes in Colo-205 cells.

Expression of cell cycle regulators in Colo-205		N	Mean	Std. Error	95% Confidence Interval for Mean		Min	Max
					Lower Bound	Upper Bound		
CyclinA	0.2% DMSO	3	100.00	0.00	100.00	100.00	100.00	100.00
	5FU 100 $\mu$ M	3	100.73	0.87	96.98	104.48	99.42	102.38
	GPE 25 $\mu$ g/ml	3	93.90	1.92	85.62	102.18	90.08	96.23
	GPE 50 $\mu$ g/ml	3	80.35	4.28	61.91	98.78	72.60	87.39
	GPE 100 $\mu$ g/ml	3	66.45	4.18	48.45	84.45	59.13	73.62
CyclinB1	0.2% DMSO	3	100.00	0.00	100.00	100.00	100.00	100.00
	5FU 100 $\mu$ M	3	96.31	2.12	87.21	105.41	92.08	98.58
	GPE 25 $\mu$ g/ml	3	94.09	0.34	92.62	95.56	93.51	94.69
	GPE 50 $\mu$ g/ml	3	98.51	1.20	93.34	103.68	97.02	100.89
	GPE 100 $\mu$ g/ml	3	95.49	4.80	74.83	116.16	88.19	104.55
CyclinD1	0.2% DMSO	3	100.00	0.00	100.00	100.00	100.00	100.00
	5FU 100 $\mu$ M	3	100.73	2.30	90.83	110.63	96.53	104.46
	GPE 25 $\mu$ g/ml	3	103.77	2.73	92.03	115.51	99.01	108.46
	GPE 50 $\mu$ g/ml	3	107.66	2.80	95.60	119.72	102.40	111.97
	GPE 100 $\mu$ g/ml	3	99.73	3.37	85.25	114.22	95.86	106.44
CyclinE	0.2% DMSO	3	100.00	0.00	100.00	100.00	100.00	100.00
	5FU 100 $\mu$ M	3	125.59	1.92	117.35	133.84	121.81	128.01
	GPE 25 $\mu$ g/ml	2	98.93	1.98	73.77	124.09	96.95	100.91
	GPE 50 $\mu$ g/ml	3	89.81	4.56	70.17	109.44	83.69	98.73
	GPE 100 $\mu$ g/ml	3	96.89	10.01	53.83	139.96	77.86	111.78
p21	0.2% DMSO	3	100.00	0.00	100.00	100.00	100.00	100.00
	5FU 100 $\mu$ M	3	125.34	9.03	86.47	164.21	112.23	142.66
	GPE 25 $\mu$ g/ml	3	112.12	5.88	86.82	137.41	100.36	118.07
	GPE 50 $\mu$ g/ml	3	126.29	9.25	86.49	166.10	110.85	142.84
	GPE 100 $\mu$ g/ml	3	109.72	9.49	68.89	150.56	97.10	128.31
p53	0.2% DMSO	3	100.00	0.00	100.00	100.00	100.00	100.00
	5FU 100 $\mu$ M	3	105.85	5.55	81.97	129.74	95.00	113.31
	GPE 25 $\mu$ g/ml	2	94.48	3.20	53.88	135.07	91.28	97.67
	GPE 50 $\mu$ g/ml	3	91.62	6.93	61.81	121.43	79.29	103.26
	GPE 100 $\mu$ g/ml	3	49.62	9.61	8.25	90.99	36.65	68.40

## VITA

Mrs. Nattaporn Buranabunwong was born on November 19, 1979 in Bangkok, Thailand. She received her Bachelor's degree of Pharmaceutical Science (2nd class honor) from the Faculty of Pharmaceutical Sciences, Chulalongkorn University, Bangkok, Thailand in 2002. She achieved her Master's degree Program in Marketing (International Program) from the Faculty of Commerce and Accountancy, Thammasat University, Bangkok, Thailand in 2007.

### Publication

1. Buranabunwong N., Ruangrunsi N., and Limpanasithikul W. 2013. Anti-cancer activity of *Glycosmis parva* leaf extract on human colorectal cancer HT29 cell. *Acta Pharmacologica Sinica*. 34 supp:4-4.

### Poster Presentation

1. Buranabunwong N., Ruangrunsi N., and Limpanasithikul W. 2013. Anti-cancer activity of *Glycosmis parva* leaf extract on human colorectal cancer HT29 cell. The 12th Meeting of the Asia Pacific Federation of Pharmacologists. July 9-13, 2013, Shanghai, China.