

# Mangiferin induces cell cycle arrest at G2/M phase through ATR-Chk1 pathway in HL-60 leukemia cells

Z.G. Peng, Y.B. Yao, J. Yang, Y.L. Tang and X. Huang

Department of Hematology, The First Affiliated Hospital, Guangxi Medical University, Nanning, China

Corresponding author: Z.G. Peng E-mail: zhigangpengcn@126.com

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**ABSTRACT.** This study aimed to determine the effect of mangiferin on the cell cycle in HL-60 leukemia cells and expression of the cell cycle-regulatory genes Wee1, Chk1 and CDC25C and to further investigate the molecular mechanisms of the antileukemic action of mangiferin. The inhibitory effect of mangiferin on HL-60 leukemia cell proliferation was determined by the MTT assay. The impact of mangiferin on the HL-60 cell cycle was evaluated by flow cytometry. After the cells were treated with different concentrations of mangiferin, the expression levels of Wee1, Chk1 and CDC25C mRNA were determined by RT-PCR, and Western blot was used to evaluate the expression levels of cdc25c, cyclin B1, and Akt proteins. The inhibition of HL-60 cell growth by mangiferin was dose- and time-dependent. After treatment for 24 h, cells in G2/M phase increased, and G2/M phase arrest appeared with increased mRNA expression of Wee1, Chk1 and CDC25C. Mangiferin inhibited Chk1 and cdc25c mRNA expression at high concentrations and induced Wee1 mRNA expression in a dose-dependent manner. It significantly inhibited ATR, Chk1, Wee1, Akt, and ERK1/2 phosphorylation but increased cdc2 and cyclin

B1 phosphorylation. Furthermore, mangiferin reduced cdc25c, cyclin B1, and Akt protein levels while inducing Wee1 protein expression. It also antagonized the phosphorylation effect of vanadate on ATR, and the phosphorylation effect of EGF on Wee1. These findings indicated that mangiferin inhibits cell cycle progression through the ATR-Chk1 stress response DNA damage pathway, leading to cell cycle arrest at G2/M phase in leukemia cells.

Key words: Mangiferin; HL-60; Cell cycle; ATR; Chk1

#### INTRODUCTION

Recent studies have found that the occurrence of leukemia is related to the dysregulation of the cell cycle (Nong et al., 2005). Due to p53 deletion, p53 mutations or G1/S check point deletion, most tumor cells are more and more dependent on the G2/M check point (Garrido et al., 2004; Muruganandan et al., 2005; Pardo-Andreu et al., 2008; Marquez et al., 2012). Therefore, the regulation of the G2/M check point is particularly important in the regulation of the cell cycle in tumor cells.

Flavonoids are a class of polyphenolic compounds widely existent in natural plants, which have been found to have significant antitumor effects (Chattopadhyay et al., 1987; Ang et al., 2011; Rivera et al., 2011). Their derivatives such as flavopiridol and genistein have been used safely in human trials (Guha et al., 1996; Huang et al., 2001) to be further applied to experimental evaluation of solid tumors in clinical phaseII (Makare et al., 2001; Garcia et al., 2003). Therefore, these compounds have potential medicinal value. Mangiferin is a flavonoid and it is extracted from the mango tree. Studies have found that mangiferin can inhibit the growth of liver cancer cells, colon cancer cells, leukemia K562 cells, and other tumor cells, and the inhibition of tumor cells is related to the induction of cell cycle arrest at the G2/M phase (Huang et al., 2002; Rajendran et al., 2008; Garcia-Rivera et al., 2011). On the basis of our previous studies, which showed that mangiferin inhibits the proliferation of HL-60 leukemia cells and induces apoptosis, we investigated the impact of mangiferin on the HL-60 cell cycle and the expression regulation of related genes to clarify the antileukemic mechanism of mangiferin, so as to provide experimental evidence for using mangiferin as a new leukemia-treatment drug.

Leukemia is a malignant blood disease that is associated with a disorder in the cell cycle and apoptosis. Cancer cells, including leukemic cells, tend to be resistant to chemotherapeutic agents and become refractory, resulting in failure of treatment. Although different types of chemotherapeutic agents are available for the treatment of leukemia, the toxicity and side effects of the drugs and the resistance of the leukemic cells to the agents remain unsolvable issues in clinical practice. Thus, to break through this bottleneck, new chemotherapeutic agents with mild side effects and multi-target efficacy with low resistance are needed and will be ideal in curing this life-threatening blood malignancy.

Mangiferin, a naturally occurring flavonoid compound that can be extracted from bark and leaves of *Mangifera indica* L. (Nong et al., 2005) and other herbal species, has been demonstrated to possess antioxidative (Muruganandan et al., 2005; Pardo-Andreu et al., 2008; Marquez et al., 2012), anti-inflammatory (Garrido et al., 2004; Rivera et al., 2011), anti-bone resorption (Ang et al., 2011), and immunoregulatory effects with mild or non-existent side ef-

fects (Chattopadhyay et al., 1987; Guha et al., 1996; Huang et al., 2001; Makare et al., 2001; Garcia et al., 2003). More interestingly, this flavonoid has been further shown to possess antiproliferative and antitumor growth properties, thus making it a potential anticancer agent in recent years (Huang et al., 2001; Rajendran et al., 2008; Garcia-Rivera et al., 2011). Previous in-depth studies have found that mangiferin induces apoptosis (Huang et al., 2002; Peng et al., 2004; Sarkar et al., 2004; Cheng et al., 2007; du Plessis-Stoman et al., 2011; Shoji et al., 2011) and inhibits cell cycle progression in liver cancer and leukemic cells, leading to G2/M phase arrest (Huang et al., 2002; Yao et al., 2010); however, since the compound's antitumor properties have not been studied extensively, the mechanisms of action in cell apoptosis induction and cell cycle inhibition have not been fully elucidated. It is reported here that mangiferin inhibits cell cycle progression at G2/M phase through the ATR (ataxia telangiectasia and Rad3 related)-Chk1 (checkpoint kinase 1) pathway in HL-60 leukemia cells.

## MATERIAL AND METHODS

# Cell culture and mangiferin preparation

The HL-60 human promyelocytic leukemia cell line was purchased from Cell Bank of Shanghai Institute of Cell Biology, Chinese Academy of Science. Cells were maintained in IMDM supplemented with 10% heat-inactivated fetal bovine serum (Hyclone, Logan, UT, USA) and grown in T-25 flasks (Corning, Corning, NY, USA) at 37°C in a 5% CO<sub>2</sub> humidified incubator (Thermo Fisher Scientific, Waltham, MA, USA). The medium was replaced every other day by centrifugation. HPLC and capillary electrophoresis grade mangiferin were generously provided by Dr. Huayi Huang of Guangxi Nationalities Hospital in Guangxi, China (currently at Roswell Park Cancer Institute, Buffalo, NY, USA). The compound was dissolved in 2% NaHCO<sub>3</sub> and was prepared as a 10 mM stock solution, which was kept at -20°C in aliquoted vials away from light.

#### Cell cytotoxicity assay

The 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay (Sigma-Aldrich, St. Louis, MO, USA) was used in determining the cytotoxicity of mangiferin. Briefly,  $4 \times 10^4$  cells in 200  $\mu$ L medium were added to each well of a 96-well plate and were incubated overnight. The cells were exposed to mangiferin on the next day at final concentrations of 20, 40, 80, 160, 200, and 250  $\mu$ M for 24, 48, and 72 h. NaHCO<sub>3</sub> solution was used as the vehicle control. Each concentration and time point of treatment were tested with 5 replicate wells. At the indicated time point of treatment, 10  $\mu$ L MTT solution was added to each well and the cells incubated for 4 h in a 5% CO<sub>2</sub> incubator. The plate was then centrifuged at 2500 rpm for 10 min, and the supernatant was carefully removed by pipetting. Finally, 150  $\mu$ L DMSO was added to each well followed by incubation for 10 min at 37°C. When the purple crystals were completely dissolved, the absorbance at 490 nm was measured using a Bio-Rad microplate reader (Bio-Rad, Hercules, CA, USA). The experiment was conducted in triplicate.

# Cell cycle analysis

HL-60 cells were exposed to various concentrations of mangiferin for 24 and 48 h.

The cells were then collected by centrifugation, washed twice with cold PBS, and fixed with 5 mL 70% cold ethanol at 4°C for at least 18 h. After fixation, the cells were again washed twice with PBS, and the cell number was adjusted to 1 x  $10^6$  in a BD Falcon (BD Biosciences, Franklin Lakes, NJ, USA) centrifugation tube. These cells were kept on ice, and 1 mL (50  $\mu$ g/mL) propidium iodide (Sigma-Aldrich) was added followed by incubation for 30 min. The DNA fragments were analyzed on a Beckman Coulter Epics XL flow cytometer (Beckman Coulter, Brea, CA, USA), and the data were analyzed using the EXPO32 ADC software.

# Wright-Giemsa staining and photographing

HL-60 cells  $(5x10^4)$  were placed in each well of a 6-well plate (Corning) overnight. The cells were then exposed to 160  $\mu$ M mangiferin for 24, 48 and 72 h. Afterwards, the plate was centrifuged at 3000 rpm and 4°C for 10 min on a Beckman Coulter Allegra 6R centrifuge (Beckman Coulter). The culture medium was then removed carefully by aspiration, and the cells were air-dried for 10 min at room temperature. The cells were fixed with 1 mL methanol (slowly added along the edge of the wells) for 1 min at room temperature. The methanol solution was then aspirated and the cells were stained with Wright-Giemsa-staining solutions (gently added along the edge of the wells) for 1 min. The wells were then gently washed with PBS 3 times to remove the excess and unbound dye. The cells that were damaged by mangiferin were unable to attach to the well after centrifugation and were removed with the solutions. The cells in all wells were then observed and photographed under a ZEISS Axiovert 25 inverted microscope equipped with a digital camera system (ZEISS North America, Peabody, MA, USA). The central areas of all wells were chosen for photographing.

# RNA extraction and RT-PCR

HL-60 cells were treated with mangiferin for 24 h. Total RNA was extracted using Trizol reagent (Invitrogen, Carlsbad, CA, USA) following manufacturer instructions. The RNA quantity and quality were analyzed on a Beckman Coulter DU-800 spectrophotometer using the nucleic acid II software-RNA/Oligo.

cDNA was synthesized using SuperScript III First-Strand Synthesis System (Invitrogen) following manufacturer instructions. Oligo (dT) was used as primer. The resulting cDNA concentration was determined with the Beckman Coulter DU-800 spectrophotometer using the nucleic acid II software-ssDNA.

Chk1, cdc25c, and Wee1 cDNA were amplified using the primers as follows: Chk1: forward primer: 5'-ATG AAG CGT GCC GTA GAC T-3' and reverse primer: 5'-TCC AAG GGT TGA GGT ATG T-3' to generate an amplicon of 571 bp. cdc25c: forward primer: 5'-GAA CAG GCC AAG ACT GAA GC-3' and reverse primer: 5'-GCC CCT GGT TAG AAT CTT CC-3' to generate an amplicon of 179 bp. Wee1: forward primer: 5'-ATG AGC AGA ACG CTT TGA G-3' and reverse primer: 5'-CTT GTG GTA TCC GAG GTA ATC-3' to generate an amplicon of 594 bp. β-actin served as the endogenous standard with a primer set of forward: 5'-TCC AAG GGT TGA GGT ATG T-3' and reverse: 5'-CAG GTC CAG ACG CAG GAT GGC-3'. The size of the amplicon was 273 bp. After reverse transcription, we prepared a 20-μL reaction mixture containing 2 μL 10X PCR buffer, 2 μL 2.5 mM dNTP mixture, 1.2 μL 25 mM MgCl<sub>2</sub>, 1 μL (100 ng) cDNA, 1 μL forward and reverse primers, 0.1 μL (0.5 U) DNA polymerase and 11.7 μL H<sub>2</sub>O. Thermal cycling was carried out on a Bio-Rad iCycler with the

following conditions: 95°C for 3 min; 28 cycles of 94°C for 30 s, 57°C (*Wee1* and *Chk1*) or 59°C (*cdc25c* and  $\beta$ -*actin*) for 30 s and 72°C for 30 s; and further extension at 72°C for 10 min. The PCR products were analyzed by 2% agarose gel electrophoresis and were visualized and quantified by the UVP BioImaging system (UVP, Upland, CA, USA). The densities of *Chk1*, *cdc25c*, and *Wee1* over the density of  $\beta$ -*actin* were obtained for further statistical analysis.

#### Western blot

The HL-60 cells (1 x 10<sup>6</sup>) were placed in T-25 flasks (Corning). There were three different treatment groups. In group 1, cells were treated with 160 µM mangiferin alone for 24, 48 and 72 h. In group 2, cells were treated with mangiferin with the same procedures as group 1 plus treatment with epidermal growth factor (EGF, 100 ng/mL; Invitrogen) for 15 min prior to harvesting. In group 3, cells were treated with mangiferin with the same procedures as group 1 plus treatment with 1 mM sodium orthovanadate (Sigma-Aldrich) for 3 h prior to harvesting. The cells were then washed with cold PBS twice and lyzed with a lysis buffer [25 mM Tris-HCl, pH 7.5, 150 mM NaCl, 1 mM EDTA, 1% Triton X-100, and 1% protease and phosphatase inhibitor cocktails (Sigma-Aldrich)]. The crude lysates were transferred to pre-chilled centrifuge tubes and centrifuged at 13,000 rpm for 15 min at 4°C. For immunoblotting, lysate protein (30 µg) was resolved on an SDS-polyacrylamide gel followed by transfer to a PVDF membrane using the Bio-Rad Mini-Protein 3 electrophoresis and transfer apparatus (Bio-Rad). Membranes were then incubated with 5% non-fat dry milk for 1 h at room temperature to block the nonspecific protein binding sites on the membrane followed by incubation with the primary antibodies (rabbit anti-phospho-ATR, ser428; rabbit anti-phospho-Chk1, ser345 or ser317; rabbit anti-phospho-Wee1, ser642; rabbit antiphospho-cdc2, tyr15; rabbit anti-phospho-cyclin B1, ser133; rabbit anti-phospho-ERK1/2, T202/Y; mouse anti-Wee1 antibody; rabbit anti-cdc2; rabbit anti-Akt; rabbit anti-ERK1/2 (Cell Signaling Technology, Danvers, MA, USA); rabbit anti-phospho-Akt, ser473 (R & D Systems, Minneapolis, MN, USA); mouse anti-Chk1; mouse anti-cdc25c; rabbit anti-cyclin B1 (Santa Cruz Biotechnology, Santa Cruz, CA, USA); and mouse anti-actin (Sigma-Aldrich) overnight at 4°C on a shaker (VWR International, Radnor, PA, USA). This was followed by incubation with a goat anti-rabbit or goat anti-mouse HRP-conjugated secondary antibody (KPL, Gaithersburg, MD, USA) at a dilution of 1:5000 at room temperature for 1 h. Protein bands were visualized using SuperSignal West Pico Chemiluminescent Substrate (Pierce, Rockford, IL, USA) and X-ray film exposure.

#### **Statistics analysis**

The experimental data were analyzed utilizing the SPSS11.5 statistical software with one-way ANOVA. A difference of P < 0.05 was considered to be significant. GraphPad Prism 5 was used to determine the IC $_{50}$ .

#### **RESULTS**

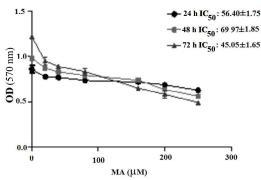
# Cell proliferation and viability

HL-60 cells were exposed to 20, 40, 80, 160, 200, and 250 μM mangiferin for 24, 48

and 72 h. The MTT cell proliferation/cytotoxicity assay results are shown in Table 1. The data represent an average of three different experiments with 5 replicate wells with the corresponding standard deviation. The  $\rm IC_{50}$  from 24-, 48- and 72-h treatments are shown in Figure 1.

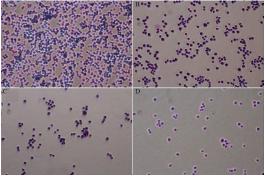
Table 1. Effect	of mangiferin (MA) on cell prolif	eration/cytotoxicity (MTT assay, OI	D:490 nm).
MA (μM)	24 h	48 h	72 h
0	$0.8583 \pm 0.0945$	$0.9800 \pm 0.0562$	$1.2150 \pm 0.0333$
20	$0.7767 \pm 0.0326*$	$0.8717 \pm 0.0671^{\#}$	$0.9500 \pm 0.1020^{\#}$
40	$0.7683 \pm 0.0402^{\#}$	$0.8300 \pm 0.0438^{\#}$	$0.8883 \pm 0.0483^{\#}$
80	$0.7350 \pm 0.0281^{\#}$	$0.7900 \pm 0.0704$ #	$0.8333 \pm 0.0814$ #
160	$0.7183 \pm 0.0591^{\#}$	$0.7383 \pm 0.0331^{\#}$	$0.6500 \pm 0.0374^{\#}$
200	$0.6850 \pm 0.0187^{\#}$	$0.6350 \pm 0.0432^{\#}$	$0.5817 \pm 0.1038^{\#}$
250	$0.6267 \pm 0.0485^{\#}$	$0.5617 \pm 0.0232^{\#}$	$0.4917 \pm 0.0147^{\#}$

<sup>\*</sup>P < 0.05,  $^{\#}$ P < 0.01 comparing with vehicle control (0  $\mu$ M).



**Figure 1.** Effect of mangiferin (MA) on cell growth and viability. HL-60 cells were treated with 20, 40, 80, 160, 200, and 250  $\mu$ M mangiferin for 24, 48, and 72 h (MTT assay). The absorbance at 570 nm of wavelength was obtained using a microplate reader. The IC<sub>50</sub> from various time points of treatment are presented.

The Wright-Giemsa cell viability assay was carried out as described above. The results are shown in Figure 2 with a magnification of 200X.

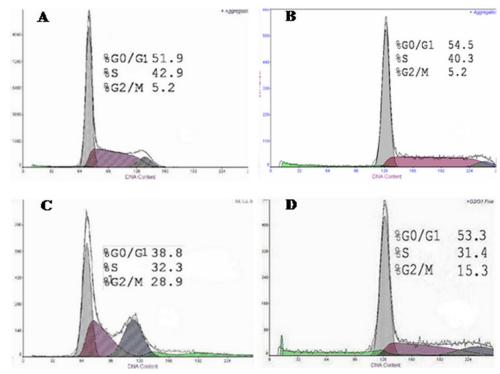


**Figure 2.** HL-60 cells (20,000) were grown on 6-well plates and treated with 160  $\mu$ M mangiferin for 24, 48, and 72 h. The cells were then centrifuged at 3,500 rpm for 10 min at 4°C followed by Wright-Giemsa staining and photographing as described above. Result exhibits that treatment with 160  $\mu$ M mangiferin reduced cell viability and attachment to the wells. **A.** No mangiferin; **B.** 24 h treatment; **C.** 48 h treatment; **D.** 72 h treatment.

# Cell cycle

HL-60 cells were exposed to various concentrations of mangiferin for 24 and 48 h and were processed for cell cycle analysis as described above. The flow cytometry analysis results of DNA content are shown in Table 2 and Figure 3. Also in Figure 2, only the vehicle control and  $160~\mu M$  mangiferin groups are shown to represent the typical results.

MA (µM)	24 h treatment			48 h treatment		
	G0/G1	S	G2/M	G0/G1	S	G2/M
0	51.9	42.9	5.2	54.5	40.3	5.2
20	48.5	39.1	12.4	54.8	38.5	6.7
40	46.8	35.7	17.5	54.7	37.0	8.3
80	44.0	31.3	24.7	56.0	34.0	9.4
160	38.8	32.3	28.9	53.3	31.4	15.3
200	44.3	33.2	22.5	51.8	34.1	14.1
250	47.4	35.5	17.1	54.7	37.7	7.6



**Figure 3.** Effect of mangiferin on cell cycle checkpoints. HL-60 cells were treated with 20, 40, 80, 160, 200, and 250  $\mu$ M of mangiferin for 24 and 48 h. The DNA fragments were analyzed using flow cytometry propidium iodine-staining method. DNA fragments in different phases of cell cycle are represented in percentage. **A.** and **B.** Cells treated with vehicle control without mangiferin for 24 and 48 h. **C.** and **D.** Cells treated with 160  $\mu$ M mangiferin for 24 and 48 h. The DNA fragments in all phases are represented in percentage.

# Chk1, cdc25c, and Wee1 mRNA expression

RT-PCR was carried out as described above. The electrophoretic results of PCR products are shown in Figure 4A; the corresponding net densities over actin are shown in Figure 5. After 24 h of treatment, mangiferin induced Chk1 and cdc25c mRNA expression at low concentrations; however, the mRNA levels decreased at high concentrations. Nevertheless, up to as high as 250  $\mu$ M, the Chk1 mRNA level was still higher than that of control. cdc25c mRNA level was reduced to less than that of the control and displayed a flattened pattern. In contrast, mangiferin induced Wee1 mRNA expression in a dose-dependent manner up to 80  $\mu$ M. This effect plateaued starting at 160  $\mu$ M.

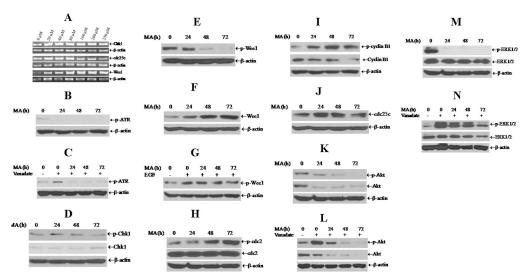


Figure 4. A. Chk1, cdc25c, and Wee1 mRNA expression, HL-60 cells were treated with 20, 40, 80, 160, 200, and 250 µM mangiferin for 24 h. The total RNA was extracted and the RT-PCR was performed according to the procedures described above. Oligo (dT) primer was used in the reverse transcription, and a specific pair of primers to each gene was used in the PCR. The β-actin was amplified and used as endogenous control. The PCR products were analyzed on 1% agarose gel, then visualized and photographed. B.-D. ATR phosphorylation. HL-60 cells were treated with 160 µM mangiferin alone for 24, 48, and 72 h (B), or with mangiferin plus epithelial growth factor (EGF) for 15 min before harvesting (C), or with mangiferin plus sodium orthovanadate for 3 h before harvesting (D). The cells were then harvested and subjected to Western blotting. Anti-phospho-ATR, anti-phospho-Chk1, or anti-total Chk1 antibody was used to detect the phosphorylation status of ATR and Chk1, the total Chk1 protein in the samples.  $\beta$ -actin was used as endogenous control. E.-G. Weel phosphorylation. HL-60 cells were treated with 160 µM mangiferin alone for 24, 48, and 72 h (E, F), or with mangiferin plus EGF for 15 min before harvesting (G). The cells were then harvested and subjected to Western blotting. Anti-phospho-Wee1 or an anti-Wee1 antibody was used to detect the phosphorylation status of Wee1 (E, G) and total Wee1 protein (F) expression in the samples. β-actin was used as endogenous control. H.-J. cdc2, cyclin B1 phosphorylation, and total cdc2, total cyclin B1, and cdc25c expression. HL-60 cells were treated with 160 µM mangiferin alone for 24, 48, and 72 h. The cells were then harvested and subjected to Western blotting. Anti-phospho-cdc2 (H), anti-phospho-cyclin B1 (I), anti-total cdc2 (H), anti-total cyclin B1 (I), or anti-total cdc25c (J) antibody was used to detect the phosphorylation status of cdc2 and cyclin B1, total cdc2, total cyclin B1, and total cdc25c expression in the samples. β-actin was used as endogenous control. K.-N. Akt, ERK1/2 phosphorylation, and total Akt, total ERK1/2 protein expression. HL-60 cells were treated with 160 µM mangiferin alone for 24, 48, and 72 h (K, M), or with mangiferin plus sodium orthovanadate for 3 h before harvesting (L, N). The cells were then harvested and subjected to Western blotting. Anti-phospho-Akt, anti-Akt, anti-phospho-ERK1/2, or anti-ERK1/2 was used to detect the phosphorylation status of Akt (K, L) and ERK1/2 (M, N), the total Akt (K, L) and total ERK1/2 (M, N) protein expression in the samples. β-actin was used as endogenous control.

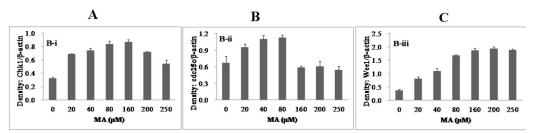


Figure 5. Net density of individual mRNA was obtained by comparing the density of each gene over correspondent density of the  $\beta$ -actin (B-i to B-iii). MA: mangiferin.

# ATR, Chk1 and Wee1 protein expression

On the basis of the cell proliferation/cytotoxicity (MTT) assay and flow cytometry cell cycle analysis, it was determined that  $160~\mu\text{M}$  was the optimal concentration of mangiferin for further study of its effect on the phosphorylation and total protein expression of several protein kinases. Briefly, HL-60 cells were exposed to  $160~\mu\text{M}$  mangiferin for 24, 48 and 72 h. Cell lysates were obtained and processed for Western blot analysis. The results showed that  $160~\mu\text{M}$  mangiferin inhibited ATR phosphorylation (Figure 4B) starting at 24 h of treatment. Mangiferin also attenuated the phosphorylation effect of orthovanadate on ATR (Figure 4C). Mangiferin treatment at first induced Chk1 phosphorylation at 24 h and then reduced it at 48 and 72 h of treatment, with 72 h being significant (Figure 4D). In addition, treatment of mangiferin for 24 h inhibited Wee1 phosphorylation. This effect was dramatic after 48 and 72 h of treatment (Figure 4E). It also antagonized, to some extent, the effect of EGF on Wee1 phosphorylation (Figure 4F); however, the total Wee1 protein was significantly induced starting at 24 h of treatment (Figure 4G).

#### cdc2, cyclin B1 and cdc25c protein expression

Treatment with 160  $\mu$ M mangiferin increased the phosphorylation of both cdc2 and cyclin B1 at 48 and 72 h, while decreasing the total cyclin B1 protein level (Figure 4H and I). In addition, mangiferin induced cdc25c at 24 and 48 h of treatment but then reduced its expression at 72 h of treatment to a level similar to that of the vehicle (Figure 4J). The total cdc2 protein level was unchanged (Figure 4H).

## Akt and ERK1/2 protein expression

Phospho-Akt and phospho-ERK1/2 were markedly inhibited by  $160 \,\mu\text{M}$  mangiferin at different treatment time points; this concentration of mangiferin also attenuated the phosphorylation effect of orthovanadate on the kinases (Figure 4K-N). Furthermore, it dramatically inhibited total Akt protein expression and slightly inhibited total ERK1/2 protein expression.

#### **DISCUSSION**

The flavonoid compound mangiferin exhibits diverse biological activities, but its anticancer property has not been extensively studied. The antitumor capability of this compound

can be traced back as far as 1987 (Chattopadhyay et al., 1987). In the later stage of studies of this compound, several significant findings relevant to its anticancer properties have been reported (Chattopadhyay et al., 1987; Huang et al., 2002; Rajendran et al., 2008; Garcia-Rivera et al., 2011). As mentioned above, this compound has a mild pharmacological action that is less toxic both *in vitro* and *in vivo* (Chattopadhyay et al., 1987; Huang et al., 2001, 2002). Chemotherapeutic agents are usually highly toxic to both cancer and normal cells, thus causing significant adverse effects. In addition, cancer cells become tolerant and subsequently resistant to the chemotherapeutic agents during the treatment course. Studies have indicated that this naturally occurring flavonoid compound is effective in inducing apoptosis and blocking cell cycle progression (Huang et al., 2002; Peng et al., 2004; Cheng et al., 2007); however, the molecular mechanisms involved in these actions are not fully understood. In this study, it was shown that the stress response DNA damage pathway, or the ATR-Chk1 pathway, is involved in the inhibition of cell cycle progression by mangiferin in HL-60 leukemia cells.

The cell proliferation/toxicity assay (MTT) results again showed that mangiferin inhibited the human promyelocytic leukemia cells HL-60 growth in a dose- and time-dependent manner (Table 1) with IC<sub>50</sub> of  $56.40 \pm 1.75$  (24 h),  $69.97 \pm 1.85$  (48 h), and  $45.05 \pm 1.65$   $\mu M$ (72 h) (Figure 1), with the  $IC_{50}$  at 72 h being the lowest. This means that in this cell line, 72 h of treatment is optimal for obtaining the best growth inhibitory effect on the basis of the time points tested in this study. Interestingly, Wright-Giemsa staining of the cells after exposure to 160 µM mangiferin at the indicated times showed that mangiferin significantly inhibited the viability and the adhesion/attachment ability of HL-60 cells, with 72 h of treatment being the most potent, which is consistent with the toxicity assay results (Figure 2). This result also indicates that mangiferin may have impaired or interfered with the assembly or functioning of microtubule filaments or cellular matrix components, thus disrupting the cells' adhesion/ attachment ability, although this hypothesis needs to be further investigated. In addition, this result further confirmed the growth inhibition results from previous studies on leukemia and other types of cancer cells (Huang et al., 2002; Peng et al., 2004; Cheng et al., 2007; du Plessis-Stoman et al., 2011; Shoji et al., 2011). At 72 h, with concentrations as high as 160 uM, mangiferin inhibited approximately half of the cells' growth (Table 1). This result also indicates that the growth-inhibitory effect of mangiferin is mild, which is consistent with previous reports in other cancers (Chattopadhyay et al., 1987; Huang et al., 2001). It is believed that chemotherapeutic agents such as naturally occurring flavonoid compounds causing mild growth inhibition rather than strong cell kill/toxicity may have lesser adverse side effects. Also, development of resistance of cancer cells to the agents tends to be slower. Similar studies with another type of flavonoid compound, eupatilin, showed that it had relatively low toxicity to normal cells compared to cisplatin (Cho et al., 2011). In this case, other mechanisms such as immunomodulatory efficacy may be involved in cell growth inhibition caused by mangiferin, as hinted in a previous study (Guha et al., 1996; Makare et al., 2001). This hypothesis would be of interest in future pharmacological and pharmaceutical studies.

The flow cytometry cell cycle analysis indicated that mangiferin, at various concentrations and time points, inhibited cell cycle progression in HL-60 cells. At the concentrations indicated, cell cycle arrest at G2/M phase occurred after exposure to mangiferin for 24 and 48 h (Table 2). The peak of DNA content in G2/M phase obtained with 160  $\mu M$  after 24 and 48 h of treatment and the percentage of DNA content in G2/M phase were both increased, with the 24-h treatment being the highest. Interestingly, G2/M DNA content decreased above 160  $\mu M$  with S phase being slightly increased. This phenomenon suggests that high concentrations of

mangiferin also inhibit DNA synthesis in addition to inhibiting mitosis, and a switch in checkpoint effect from low to high concentrations was observed. Figure 3 illustrates the typical flow cytometry histograms of the results at  $160 \mu M$ .

Various concentrations of mangiferin ranging from 20 to 250  $\mu$ M were tested for 24 h in this experiment. RT-PCR results of *Chk1*, *cdc25c*, and *Wee1* after mangiferin treatment indicated that lower concentrations ranging from 20 to 160  $\mu$ M induced Chk1 mRNA expression; however, the mRNA level declined at 200  $\mu$ M. Similar results were observed in the cdc25c mRNA level, with slightly higher potencies compared to that of Chk1. Contrary to Chk1 and cdc25c mRNA expression levels, mangiferin treatment for 24 h increased Wee1 mRNA expression in dose-dependent manner (Figures 4A and 5). On the basis of these results, it was hypothesized that mangiferin is more potent in targeting cdc25c and Wee1 expression than that of Chk1, even though there was insufficient time to perform the time course experiments beyond 24 h for evaluation of mRNA expression.

ATR is one of the kinases belonging to the phosphoinositide-3-like kinase (PI3K) family and shares similar domain architecture and several modes of regulation (Mordes and Cortez, 2008). ATR is activated when DNA damage occurs and preferentially phosphorylates other downstream proteins, including Chk1. ATR is essential in replicating cells, as it responds to replication stress to restore the stalled cell cycle checkpoint and ensures the re-entry of the cell cycle (Cimprich and Cortez, 2008). Thus, ATR and Chk1 are currently considered cancer chemotherapeutic targets, and inhibitors of these kinases have emerged and are being developed as new anticancer agents (Dent et al., 2011). In this study, it was found that mangiferin at  $160 \mu M$  inhibited the autophosphorylation of ATR. Exposure to mangiferin for various lengths of time completely inhibited ATR phosphorylation starting at 24 h of treatment (Figure 4B).

Vanadate is a broad-spectrum phosphatase inhibitor, although its action is more potent on tyrosine phosphatases than on serine and threonine phosphatases (Crans et al., 1989; Huyer et al., 1997). To confirm whether mangiferin inhibits ATR phosphorylation, HL-60 cells were exposed to mangiferin for the indicated times. Prior to harvesting, cells were treated with 1 mM sodium orthovanadate for 3 h. The results showed that mangiferin inhibited the phosphorylation of ATR enhanced by vanadate (Figure 4C).

As described above, the cell cycle checkpoint kinase Chk1 is one of ATR's downstream kinases. Inhibition of the phosphorylation of ATR resulted in first inducing Chk1 phosphorylation at 24 h of treatment and then reducing it at 48 and 72 h (Figure 4D). To interpret this phenomenon, it was assumed that Chk1 activity underwent a compensating reaction to maintain the entry of G2/M under stress at 24 h of treatment due to inhibition of its upstream kinase, ATR activity; however, this compensating reaction was diminished with longer treatment time.

Wee1 is a protein kinase that controls the entry of the mitotic phase of the cell cycle through phosphorylation of cdc2 at Tyr15 and Thr14 (McGowan and Russell, 1993). Wee1 also regulates the DNA damage checkpoint that controls G2/M transition (Rowley et al., 1992). Wee1 kinase activity is deactivated when it is phosphorylated or degraded (Watanabe et al., 1995). Studies have found that inhibition of Wee1 results in enhanced cell proliferation (Chen and Gardner, 2004). In this study, the results indicated that mangiferin induced Wee1 expression at both mRNA and protein levels while decreasing autophosphorylation (Figure 4A, E, F). In addition, mangiferin, to some extent, attenuated the EGF-induced phosphorylation of Wee1 (Figure 4G). Thus, it is likely that mangiferin behaves in a dual-action manner by inhibiting ATR and Chk1 kinase activities while enhancing Wee1 activity.

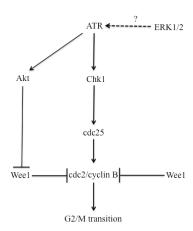
As mentioned above, cdc2 (cyclin-dependent kinase-1, CDK1) functions as serine/

threonine kinase and pairs with cyclin B to form a CDK-cyclin B complex that regulates the cell cycle including G2/M checkpoint. The cdc2/cyclin B complex is deactivated when phosphorylated by Wee1 and re-activated by cdc25 (A/B/C) phosphatase via dephosphorylation (Qiao et al., 2006; Hu and Moscinski, 2011). The cell division cycle 25s (cdc25) are dual-specificity phosphatases that are considered a subclass of protein tyrosine phosphatases. The cdc25 proteins control entry into the cell cycle and progression through various phases of the cell cycle, including the M and S phases during normal cell division, and in the event of DNA damage, they are key targets of the checkpoint machinery that ensures genetic stability (Boutros et al., 2007). In this study, 160  $\mu$ M mangiferin increased both cdc2 and cyclin B1 phosphorylation starting at 24 h of treatment but reduced the total cyclin B1 protein level at 72 h of treatment; total cdc2 showed no change (Figure 4H and I).

cdc25c was first induced by mangiferin at 24 and 48 h but was then reduced at 72 h of treatment (Figure 4J). This result is consistent with the mRNA level, which indicated that cdc25c was induced by low concentrations (20-80  $\mu$ M) of mangiferin at 24 h of treatment (Figure 4A). The phosphorylation effects of mangiferin on cdc2 and cyclin B1 indicated that this compound may behave as a cdc25c phosphatase inhibitor and a Wee1 kinase stimulator.

It is already known that PI3K/Akt is one of the downstream pathways and substrates of ATR kinase. Studies have found that Akt is hyperactivated in many human cancers, as it promotes proliferation and cancer cell survival in response to DNA-damaging agents (Fraser et al., 2011). In addition, the extracellular signal-regulated kinase 1/2 (ERK1/2) is also involved in ATM/ATR-mediated G2/M checkpoint activation (Yan et al., 2007, 2008). It was found that 160 µM mangiferin dramatically inhibited phospho-Akt and phospho-ERK1/2 as well as total Akt protein in a time-dependent manner; total ERK1/2 protein level was slightly reduced as well after treatment at all time points (Figure 4K and M). Mangiferin also diminished the effect of phosphatase inhibitor vanadate on the phosphorylation of Akt and ERK1/2, with the effect on Akt being more potent (Figure 4L and N). These results indicated that Akt and ERK1/2 are involved in ATR-mediated G2/M checkpoint in the stress response DNA damage pathways; however, the precise relationship between ATR and ERK1/2 is still unknown.

In conclusion, this study hypothesizes that mangeferin inhibits cell cycle progression at G2/M phase through the ATR-Chk1-cdc25c and cdc2/cyclin B pathway as illustrated in the following schematic chart:



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