

RESEARCH PAPER

## Curcumin induces apoptosis in human leukemic cell lines through an IFIT2-dependent pathway

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### ABSTRACT

Curcumin, the primary bioactive component isolated from turmeric, has been shown to possess variety of biologic functions including anti-cancer activity. However, molecular mechanisms in different cancer cells are various. In the present study, we demonstrated that curcumin induced G2/M cell cycle arrest and apoptosis by increasing the expression levels of cleaved caspase-3, cleaved PARP and decreasing the expression of BCL-2 in U937 human leukemic cells but not in K562 cells. We found some interferon induced genes, especially interferon-induced protein with tetratricopeptide repeats 2 (*IFIT2*), were significantly upregulated when treated with curcumin in U937 cells by gene expression chip array, and further confirmed that the expression of *IFIT2* was obviously higher in U937 than that in K562 cells by Western blot assay. In addition, inhibiting the expression of *IFIT2* by shRNA in U937 rescued curcumin-induced apoptosis and exogenous overexpression of *IFIT2* by lentiviral transduction or treating with IFN $\gamma$  in K562 cells enhanced anti-cancer activity of curcumin. These results indicated for the first time that curcumin induced leukemic cell apoptosis via an *IFIT2*-dependent signaling pathways. The present study identified a novel mechanism underlying the antitumor effects of curcumin, and may provide a theoretical basis for curcumin combined with interferon in the cancer therapeutics.

**Abbreviations:** IFNs, Interferons; ISGs, IFN-stimulated genes; *IFIT2*, IFN-induced protein with tetratricopeptide repeats 2; TPR, tetratricopeptide repeats; IRES, internal ribosome entry site; ECMV, encephalomyocarditis virus

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Apoptosis; curcumin; *IFIT2*;  
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### Introduction

Leukemia, cancer of the haematopoietic cells, is the leading cause of cancer death in adults and children.<sup>1</sup> In 2012, leukemia developed in 352,000 people globally and caused 265,000 deaths.<sup>2</sup> Chemotherapy is the major form of treatment of leukemia, and the survival rates have improved remarkably over the past decades. However, the potential side effects of cytotoxic chemotherapy significantly impact the effect of drugs.<sup>3</sup> Thus, new therapeutic weapons, or methods of prevention are required.

Curcumin, an active ingredient derived from turmeric, has been recognized for its medicinal properties.<sup>4</sup> Multiple studies in both animals and human indicated that curcumin possesses a variety of potent properties, including antioxidant, anti-inflammation, radical-scavenging, anticancer, and so on.<sup>5,6</sup> Several studies indicated that curcumin can suppress the proliferation of leukemia and induce cell death.<sup>7-9</sup> Besides, it shows no significant side effects compared with other chemotherapeutic drugs.<sup>10</sup> Therefore, curcumin may represent a new therapeutic strategy for treatment of leukemia.<sup>11</sup>

Considerable work has been conducted to explore how curcumin works.<sup>7-9</sup> Extensive research over the last decades has revealed that curcumin exerts anticancer effect through affecting numerous molecular and biochemical cascades.

The molecular targets of curcumin include growth factors, growth factor receptors, transcription factors, cytokines, enzymes, and genes regulating apoptosis and proliferation.<sup>12</sup> However, the molecular mechanisms in different tumor cells are different.

Interferon-induced protein with tetratricopeptide repeats 2 (*IFIT2*), also known as ISG54, GARG39, and MuP54, is originally discovered as a direct response to type I IFN. Further studies proved that it was also induced as a primary stress response to infection which independent of IFNs, and might play a role in anticancer.<sup>13-16</sup> *IFIT2* regulates the functions of cell cycle, apoptosis, tumor colonization and viral replication, which confer cellular resistance to viral infections and regulates proliferation, apoptosis and migration of cancer cells.<sup>17-20</sup>

In this report, we found that curcumin induced expression of interferon regulatory genes (especially *IFIT2*) in leukemia cell U937. upregulation of *IFIT2* by exogenous expressing or treating with IFN $\gamma$  in K562 increased cells apoptosis and enhanced anticancer effect of curcumin. And shRNA-mediated *IFIT2* knockdown inhibited curcumin-induced apoptosis in U937 cells. Our results demonstrated for the first time that there is a cross-talk between curcumin and interferon signaling pathways, which provides the basis for curcumin combined with interferon in cancer therapeutics.

## Results

### Sensitivity of leukemia cell lines to curcumin

To assess anticancer effect of curcumin on leukemia, 5 leukemia cell lines (NB4, NB4-R1, K562, HL60 and U937) were treated with different concentration of curcumin for 24 h and growth viability were analyzed by MTT assay. As shown in Fig. 1A, The concentration of curcumin inhibiting 50% cell proliferation ( $IC_{50}$ ) in leukemia cell lines were:  $15.77 \pm 2.17 \mu\text{M}$  for NB4,  $16.67 \pm 1.63 \mu\text{M}$  for NB4-R1,  $27.86 \pm 2.45 \mu\text{M}$  for HL60,  $27.67 \pm 2.72 \mu\text{M}$  for K562 and  $8.63 \pm 2.34 \mu\text{M}$  for U937 respectively.

Given the intriguing results, we chose K562 and U937 as research objects to study the mechanism under this condition. We treated K562 and U937 cells with different concentration of curcumin (0–10  $\mu\text{M}$ ) for 24 h and 48 h. MTT results showed that curcumin significantly inhibited U937 cell growth in time- and dose-dependent manner (Fig. 1B), while K562 cells were not sensitive to curcumin (Fig. 1C). These results indicated that sensitivity of leukemic cells to curcumin was different.

### Curcumin induces G2/M phase arrest in leukemia cells

To further define anti-tumor effect of curcumin in leukemia cells, we analyzed cell cycle progression in K562 and U937 cells treated with curcumin (0, 5 and 10  $\mu\text{M}$ ) for 24 hours by FCM assay. As the results shown in Fig. 2, although curcumin blocked the cells in the G2/M phase, there were some differences in K562 and U937 cells: a significantly increased number of cells were detected in G2/M phase and corresponding reduction in G0/G1 phase of U937 cells were observed in a dose-dependent manner. However, in K562 cells, the proportion of G2/M phase slightly increased and accompanied by a decrease in the S phase.

### Curcumin induces apoptosis in U937 but not in K562 cells

To determine apoptosis induction capacity of curcumin in leukemia cells, we treated U937 and K562 cells with curcumin (0, 5, and 10  $\mu\text{M}$ ) for 24 h, and detected the percent of apoptotic cells by Annexin-V/PI staining assay. The results showed that curcumin induced apoptosis in U937 with a dose-dependent manner but not in K562 (Fig. 3A). Furthermore, in U937 cells, 2

apoptotic proteins (cleaved caspase3 and PARP) were markedly increased and the apoptosis inhibitor protein Bcl-2 was decreased. However, the proteins in the K562 changed slightly (Fig. 3B).

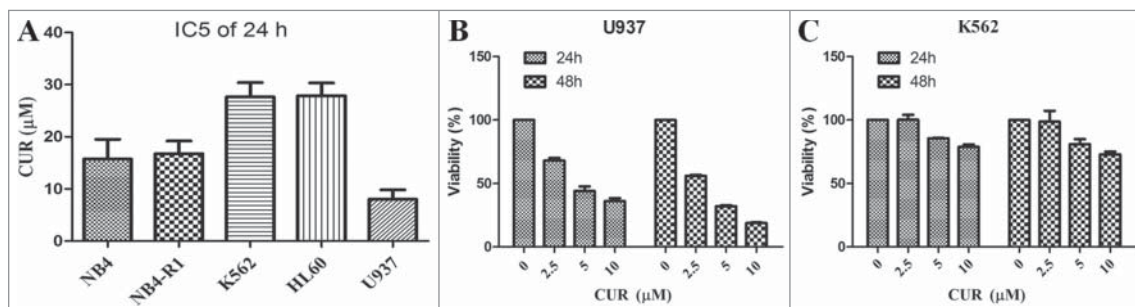
### Curcumin induces the expression of interferon regulatory genes in U937 but not in K562 cells

To explore the molecular mechanisms of different impact of curcumin on K562 and U937 cells, we analyzed change of gene expression in U937 after treated with curcumin (10  $\mu\text{M}$ ) for 24 h by gene expression chip. Chip results showed that over 1000 genes were changed and most genes have been reported previously, such as *skp2*, *myc*, and *GADD45A*. We found that some of the interferon regulatory genes, especially *IFIT2* gene expression was significantly upregulated when treated with curcumin (Fig. 4A).

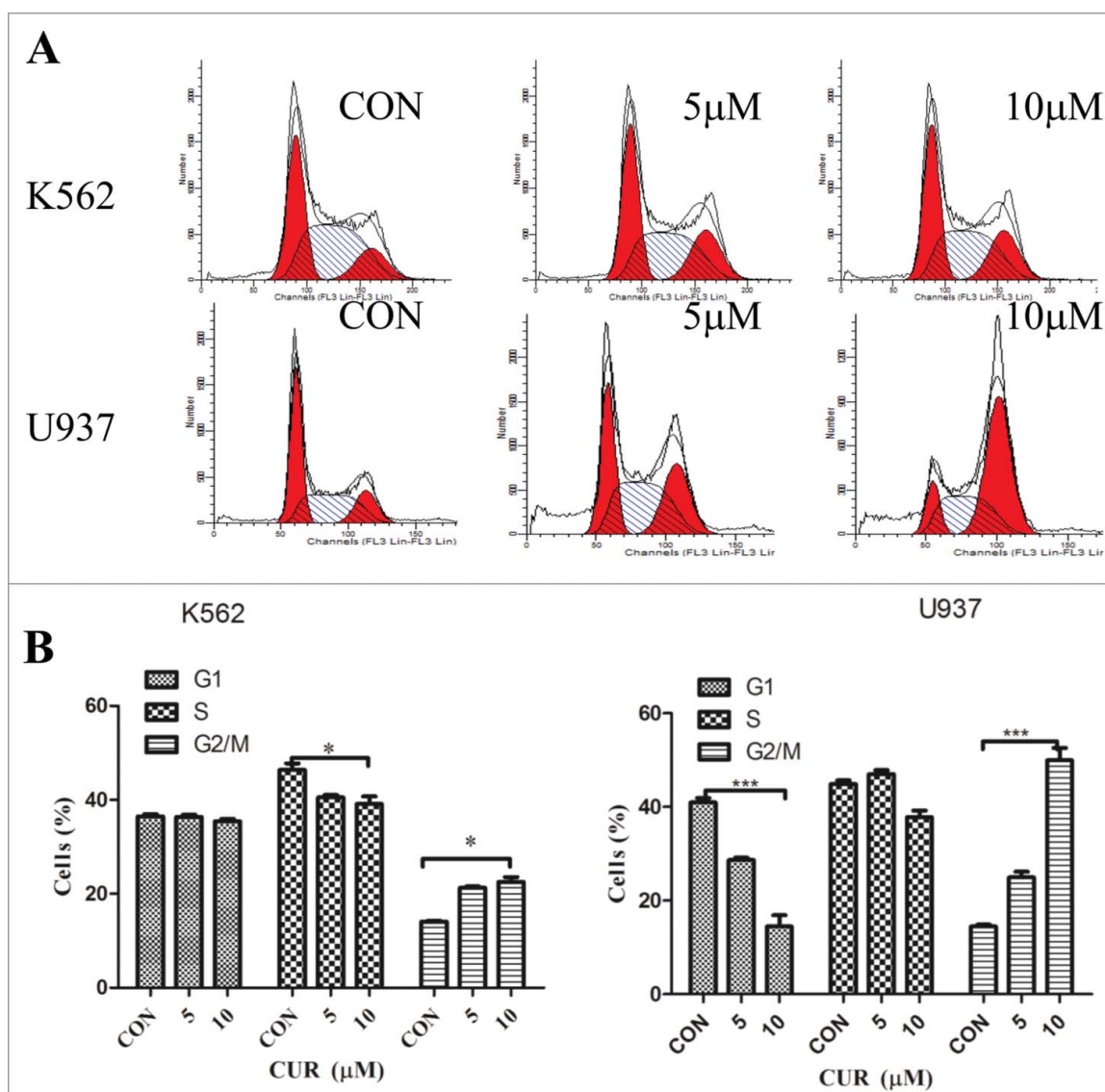
Subsequently, RT-qPCR and western-blot were performed to confirm the change of gene expression in U937 observed by gene chip. SKP2, MYC, IFIT2, IFIT3 were selected owing to their biologic function. U937 cells were treated with different concentration of curcumin (0, 5 and 10  $\mu\text{M}$ ), and genes expression of *MYC*, *IFIT2*, *IFIT3* and proteins level of *SKP2*, *IFIT2*, and *IFIT3* were determined by RT-qPCR and western blot respectively. As shown in Fig. 4B and D, the results were consistent with the previous results of gene chip assay. However, there is no difference in K562 cells with or without curcumin (Fig. 4C, D), which suggesting the gene expression induced by curcumin in U937 cells might associated with sensitivity of leukemic cells to curcumin.

### Downregulation of IFIT2 by RNA interference rescues curcumin-induced apoptosis in U937 cells

We knocked down IFIT2 in U937 so as to confirm the hypothesis that IFIT2 plays vital role in anticancer process of curcumin. After infected with IFIT2-shRNA or nonspecific shRNA lentiviral for 3 days, U937 cells were exposed with 10  $\mu\text{M}$  curcumin for 24 h, and the knockdown efficiency of IFIT2 were analyzed by using RT-qPCR to examine the expression of IFIT2 in cells. As seen in Fig. 5A, compared with nonspecific shRNA, the mRNA of IFIT2 decreased more than 65% in IFIT2-knocked down cells when treated with curcumin. The percentage of apoptotic cells was also reduced significantly in IFIT2-shRNA



**Figure 1.** Sensitivity of leukemia cell lines to curcumin (A) Leukemia cells (NB4, NB4-R1, HL60, K562 and U937) were treated with curcumin,  $IC_{50}$  were determined by MTT assay. (B-C) K562 and U937 were exposed to curcumin (0, 2.5, 5 and 10  $\mu\text{M}$ ), cell viability were measured by MTT on 24 h and 48 h; Each value represents the mean  $\pm$  SD.



**Figure 2.** Effects of curcumin on cell cycle arrest in leukemia cell lines. (A) curcumin induced the cell cycle arrest in K562 and U937 cells. (B) The percentage of cells in G1, S and G2/M of cell cycle when treated with curcumin. \* $p < 0.05$ , \*\*\* $p < 0.001$ .

infected cells after treating with curcumin (Fig. 5B and C). These data add to the growing evidence that IFIT2 plays vital role in the apoptosis-induction effect of curcumin.

#### **IFIT2 enhances sensitivity of K562 cells to curcumin**

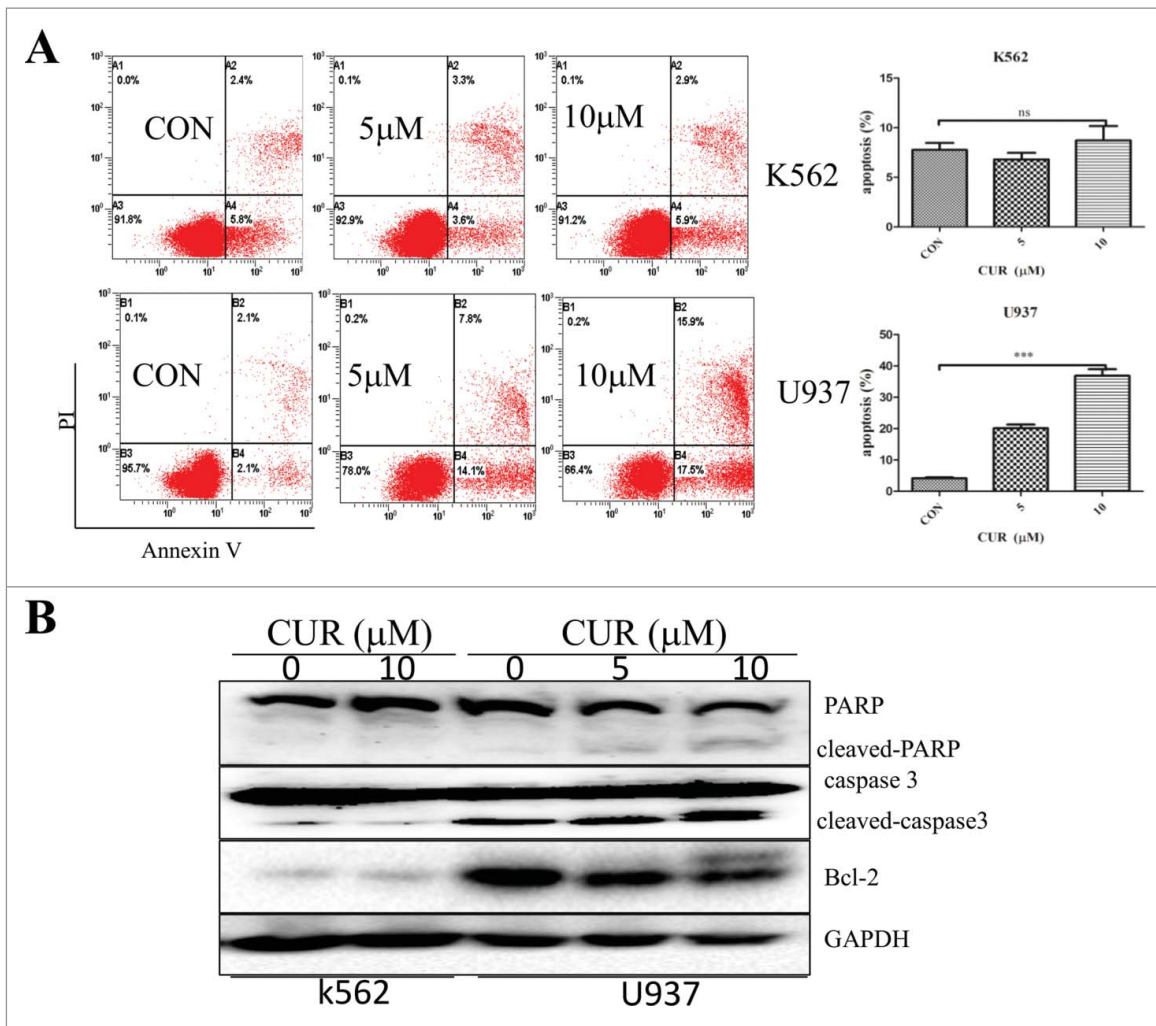
Based on effect of IFIT2 on cell cycle, apoptosis, tumor colonization and viral replication, we hypothesized that IFIT2 might make contributions to anti-tumor effect of curcumin. To test the hypothesis, IFIT2 gene-containing viruses were packaged in HEK-293T cells, and then infected K562 cells for 5 d. Expression of IFIT2 was analyzed by both RT-qPCR and western blot. As shown in Fig. 6A, mRNA and protein levels of IFIT2 were significantly increased, and expression of *SKP2* and *myc* genes was decreased at the same time (Fig. 6B), which implying IFIT2 might play a role in inhibiting proliferation in K562 cells.

To investigate the roles of IFIT2 apoptosis-induction activity of curcumin, Annexin-V/PtdIns double-staining assay was

performed to analyze apoptosis of the cells exposed to curcumin (10 μM) for 24 h. As the results seen in Fig. 6C, compared with vector group, the percentage of apoptotic cells was found to be obviously increased in the IFIT2 overexpressed cells, and curcumin further potentiated apoptosis of IFIT2 overexpressed cells. These results support the hypothesis that IFIT2 plays an important role in the apoptosis-induction activity of curcumin.

#### **IFN $\gamma$ enhances curcumin-induced apoptosis in K562 cells**

Since IFIT2 is an IFN induced protein, we supposed that IFN $\gamma$  could enhance the anticancer effect of curcumin by inducing the expression of IFIT2 protein. K562 cells were treated with IFN $\gamma$  (5000 u/ml) alone or combined with curcumin (10 μM) for 24 h. We found that IFN $\gamma$  induced the expression of IFIT2 and IFIT3 proteins (Fig. 7A), and then we detected the apoptosis of cells treated with regents by FCM analysis. As shown in Fig. 7B and C, although curcumin had little effect on the apoptosis of K562 cell, 5000u/ml IFN $\gamma$  induced apoptosis of K562 cells, and



**Figure 3.** Effect of curcumin on inducing apoptosis in leukemia cell lines. (A) The percentage of early apoptosis and late apoptosis were detected by Annexin-V/Ptdlns double-staining assay. The percentage of apoptotic cell in different concentration of curcumin. Each value were represents the mean  $\pm$  SD ns means not significant, \* $p < 0.05$ , \*\*\* $p < 0.001$ . (B) K562 and U937 cells treated with curcumin, apoptotic proteins (caspase3 and PAPR) and the apoptosis inhibitor protein Bcl-2 were detected by western blot.

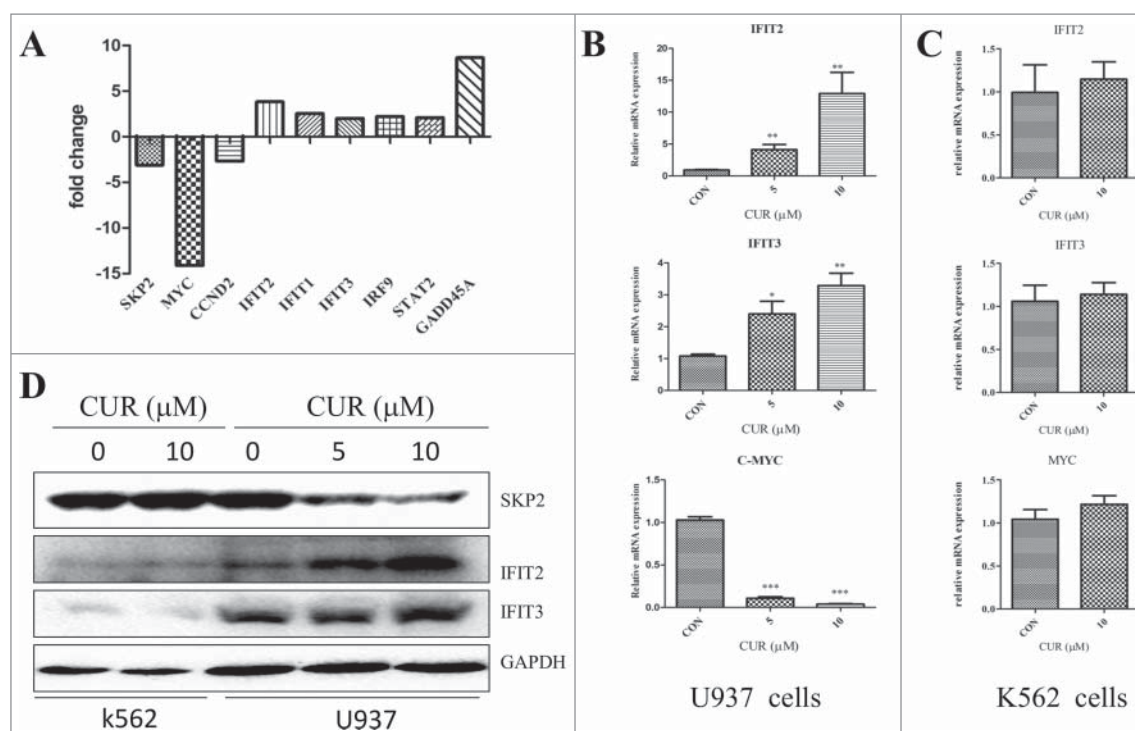
IFN $\gamma$  combined with curcumin even further strengthen apoptosis-induction effect. Depending on these results, we concluded that IFN induced IFIT2 expression and enhanced the efficacy of curcumin for the treatment of leukemia.

## Discussion

In our present study, we investigated the role of IFIT2 in anti-leukemia process of curcumin, and explored whether Interferons (IFNs) enhanced the apoptosis-induction effect of curcumin. Initially, we studied a panel of leukemia cell lines and found that sensitivity of these cell lines to curcumin were different. As curcumin is a promised anticancer drug, exploration of mechanism of curcumin can widen its clinical application and bring more benefit to patients. To clarify it, one sensitive cell line U937 and one resistant cell line K562 were chosen in the present study. We found that gene expression of IFIT2 was decreased with curcumin in both dose- and time-dependent manner in U937 cells but not in K562. To further investigate the role of IFIT2, IFIT2 was overexpressed in K562 and knocked down in U937. The results showed that IFIT2 could

increase the anticancer effect of curcumin, and when combined with IFN $\gamma$ , curcumin has a better effect on leukemia cells. These results not only demonstrate the efficacy of IFIT2, but also suggest the synergistic effects of curcumin and IFN $\gamma$ .

IFIT2, which is a member of ISRE-containing genes, have been shown could be a biomarker of cancer.<sup>19,21,22</sup> Lee TC et al. reported that IFIT2 could inhibit migration and metastasis of oral squamous cell carcinoma (OSCC) cells by activation of atypical PKC signaling.<sup>19,23,24</sup> In addition, IFIT2 also be proved to suppress development of human adenocarcinoma of gastric esophageal junction.<sup>25</sup> These results indicated that IFIT2 was related to anti-cancer effect. IFIT2 has been demonstrated involved in anticancer process of other agents, but its role in the treatment of leukemia by curcumin still unknown.<sup>26,27</sup> Several researchers have shown that IFIT2 can induce apoptosis via a mitochondrial pathway dependent on Bcl-2<sup>28,29</sup>, and curcumin has been reported inducing apoptosis via BCL-2 signaling pathway as well.<sup>30-32</sup> The previous study suggested that BCR-ABL kinase domain inhibited the expression of ISRE-containing genes,<sup>33</sup> which could in part associated with resistant of K562. All the data are consistent with our findings.



**Figure 4.** (A) Curcumin induces the expression of interferon regulatory genes in U937 but not in K562 cells. Change of partial genes expression in U937 treated with curcumin (10 μmol/L) for 24 h analyzed by gene chip; (B) Validation of genes expression in U937 and K562 when treated with curcumin for 24 h by RT-PCR. Each value represents the mean ± SD; (D) Validation of genes (*SKP2*, *IFIT2*, *IFIT3*) expression in U937 and K562 when treated with curcumin for 24 h by western blot array.

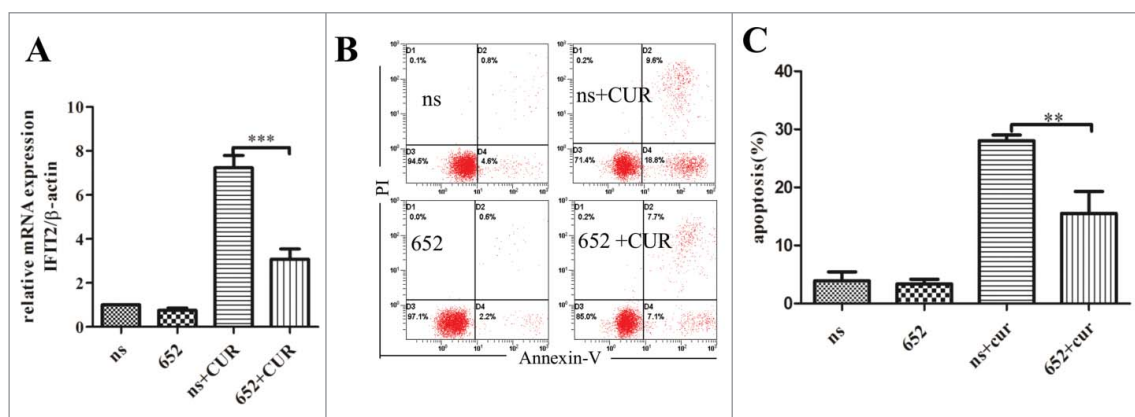
IFN $\gamma$  used to be a therapeutic agent against chronic myeloid leukemia with a moderate efficiency.<sup>34</sup> In this study, we found that combination of curcumin and IFN $\gamma$  enhanced curcumin-mediated anti-cancer activity, and showed for the first time that curcumin-induced leukemic cell apoptosis is an IFIT2-dependent signaling pathway.

Taken up, our results suggested for the first time that IFIT2 enhance the effect of curcumin, and supported that combinational of curcumin and IFN $\gamma$  would shed a new light on leukemia therapy. Because of great strides could be made in the use of curcumin, further mechanistic exploration of it is warranted.

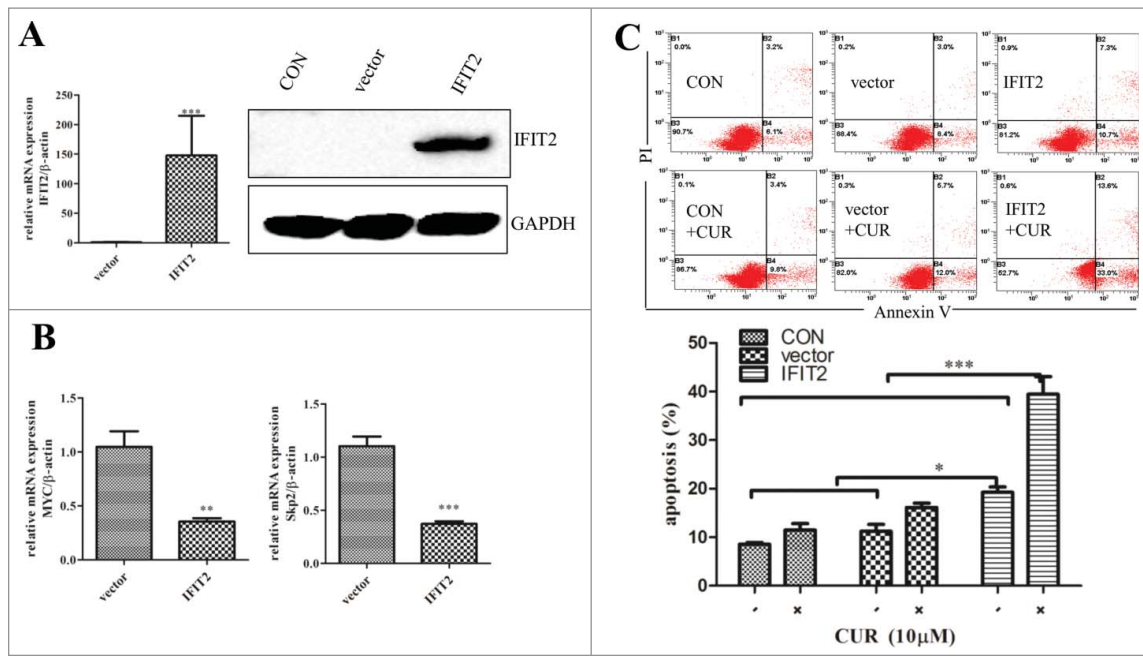
## Materials and methods

### Cells and reagents

U937, K562, NB4, NB4-R1, HL60 cells were maintained in RPMI-1640 containing 10% fetal bovine serum under 37 °C with 5% CO<sub>2</sub>. Curcumin (Sigma, St. Louis, MO) was dissolved in DMSO as a stock solution at 5 mM. IFN $\gamma$  was available from R&D systems and dissolved in PBS buffer at 10<sup>6</sup> u/ml. The antibodies against GAPDH, PARP, caspase 3, Bcl<sup>-2</sup>, Skp2, IFIT2 and IFIT3 were purchased from Santa Cruz Biotechnology (Santa Cruz, CA). Annexin-V-FITC/PtdIns kit was purchased from Bestbio Biotechnology (Bestbio, China). Cell cycle



**Figure 5.** Knockdown of IFIT2 by shRNA decreases curcumin-induced apoptosis. (A) IFIT2 specific shRNA or non-specific (*ns*) shRNA lentivirals infected U937 cells for 72 hours, and cells stimulated with curcumin (10 μM) for 24 hours. The knockdown efficiency of endogenous IFIT2 were analyzed by RT-qPCR. (B and C) Cells expressing IFIT2 specific shRNA or non-specific (*ns*) shRNA were evaluated for their apoptotic response to curcumin by Annexin-V/PtdIns double-staining assay and flow cytometry. \*\**p* < 0.01, \*\*\**p* < 0.001.



**Figure 6.** Exogenous overexpression of IFIT2 enhances sensitivity of K562 cells to curcumin (A) FIT2 gene-containing lentiviral infected K562 cells and detected the expression of IFIT2 by RT-PCR and western blot; (B) the expression of c-myc and skp2 were analyzed by RT-PCR after overexpression of IFIT2; (C) K562, K562-Vector and K562-IFIT2 were treated with or without 10  $\mu$ mol/L for 24 h, Annexin-V/PtdIns double-staining assay was used to detect percentage of apoptotic cells. \* $P < 0.05$ ; \*\*\* $P < 0.001$ .

detection kit (COULTER DNA PREP reagent kit) was from Beckman coulter, Inc. Reverse transcription reagent and Super-Real qPCR PreMix (SYBR Green) reagent kit were purchased from TIANGEN Biotech (Beijing) Co., Ltd. The lentiviral expression vector pLVX-Puro, pLVX-shRNA1 vector and the lentiviral packaging plasmids were provided by Clontech Laboratories, Inc. The pseudoviral particle producer cell line 293T was cultured with DMEM supplemented with 10% fetal bovine serum.

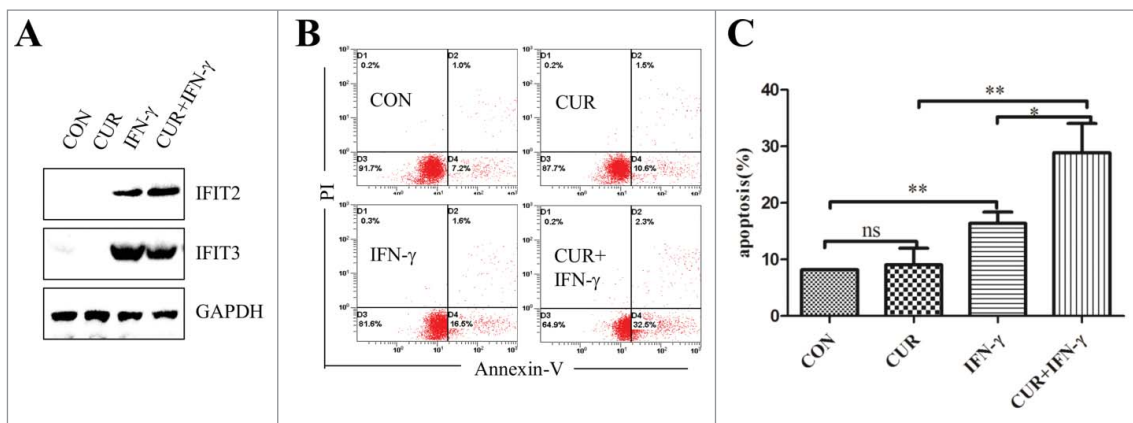
### MTT assays

Two  $\times 10^4$  cells per well were seeded into 96-well plates in 100  $\mu$ l volume and cultivated in RPMI1640 medium with 10% fetal bovine serum at 37  $^{\circ}$ C. Cells were grown for 24 h and

48 h, respectively. 10  $\mu$ l of MTT reagent was added at indicated time. Then the cells were incubated in 37  $^{\circ}$ C after covered in aluminum foil. 2 hours later, centrifuged and removed the supernatant. 100  $\mu$ l DMSO was added to each well and read the results by Elisa reader after an additional 15-minutes incubation.

### Cell cycle detection by PtdIns single-staining assay

To detect effect of curcumin on cell cycle arrest, PtdIns single-staining assay was conducted. Cells were treated with the drug and cultivated under 37 $^{\circ}$ C saturated humidity and 5% CO<sub>2</sub> for 24 h.  $10^6$  cells were harvested in the appropriate manner (centrifuged at 2,000 rpm for 5 min) and removed the supernatant. Add 50  $\mu$ l DNA PREP LPR reagent for 1 min according to the



**Figure 7.** Effect of curcumin combined with IFN on inducing apoptosis of K562. (A) Expression of IFIT2 and IFIT3 in K562 cells was measured by western blot after treating with curcumin, IFN $\gamma$ , or both IFN $\gamma$  and curcumin; (B) The apoptosis were analyzed in K562 cells following treatment of curcumin, IFN $\gamma$ , or IFN $\gamma$  combined with curcumin by FMC assay; (C) The percentage of apoptotic cells in K562 treated with curcumin, IFN $\gamma$ , or both curcumin and IFN $\gamma$ . Each value were represents the mean  $\pm$  SD (n = 3). ns means not significant; \* $P < 0.05$ ; \*\* $P < 0.01$ .

instructions, and then added 300  $\mu$ l DNA PREPStain reagent and place it at room temperature for 30 min. and then detected by CYTOMICS FC500 ( Beckman coulter, inc ) and analyzed cell cycle by MODFIT2 software.

### Apoptosis analysis by FMC

Cells were treated with or without drug and cultured under 37°C for 24 h. 10<sup>6</sup> cells were harvested in the appropriate manner( centrifuged at 2,000 rpm for 5 min), 5  $\mu$ l Annexin-V were added after adding 300 $\mu$ l Annexin-V binding solution and placed the mixture at 4°C for 15min. Added 10  $\mu$ l PtdIns at indicated time, then analyzed the results by cytometer. Annexin V+ and/or PtdIns+ cells are apoptosis cells.

### RT-qPCR

Cells ( 2  $\times$  10<sup>6</sup> ) were grown in cell culture bottles and then treated with the indicated concentrations of drug for 24 h. Total RNA were extracted by TRIzol reagent (Invitrogen) and reverse-transcribed to the cDNA synthesis and the following qPCR with the specific primers were as follows: IFIT2: 5'-ctgcaacatgagtgagaac-3' and antisense 5'-cagtgaccagactctgat-3'; IFIT3: sense 5'-tcatgagtgaggcaccag-3', antisense 5'-cctcgtgttaccatctagg-3'; MYC: sense 5'-gctgcttagacgcgattt-3' and antisense 5'-caccgagtcgtagtcgaggt-3'; Skp2: sense 5'-atgccc-caatctgtccatct-3'and antisense 5'-caccgactgagtgataggtgt-3';  $\beta$ -actin: sense 5'-catgtactgtgctatccaggc-3' and antisense 5'-ctcctaatgtcacgcacgat-3'.

The relative quantity of target gene expression was analyzed using the comparative CT (2- $\Delta\Delta$ CT) method.

### Western blot analysis

Proteins were extracted by lysis buffer (1% Triton X-100, 50 mM Tris (pH 8.0), 150 mM NaCl, 1 mM PMSF, 1 mM Na<sub>3</sub>VO<sub>4</sub>, and protease inhibitor cocktail). Proteins from cell lysates (20 $\mu$ g) were electrotransferred to nitrocellulose membranes after separated on 10% SDS-PAGE. Before blotted with primary antibody overnight at 4°C, membranes were sealed for 1 h at room temperature in Tris-buffered saline-0.05% Tween-20 (TBST) containing 5% non-fat dry milk. After 3  $\times$  10 min washes in PBS, membranes were incubated with peroxidase-conjugated secondary antibody for 1 hr. Following 3 additional 10 min washes with TBST, and proteins were imaged by enhanced chemi-luminescence detection reagent and detected with Bio-Rad ChemiDoc XRS+ chemiluminescence imaging system (Bio-rad laboratories Inc.).

### Gene chip expression

Total RNA were isolated from cells using TRIzol reagent (Invitrogen) and sent to Shanghai Genechem Co.,Ltd where further analyses were conducted. RNA integrity number for all samples should be greater than 7.0 (Agilent 2100 Bioanalyzer, Agilent, Santa Clara, CA). Hybridization and labeling (GeneChip Hybridization Oven 645) was used where aRNA was synthesized by GeneChip 3'IVT Express Kit. Posthybridization and washes were performed by GeneChip Fluidics Station 450

(Affymetrix,Santa Clara, CA).The results scanned by GeneChip Scanner 3000 (Affymetrix,Santa Clara, CA) was analyzed by GenePix Pro 4.0 software.

### Lentiviral Transduction

The lentiviral expression vector and the lentiviral packaging plasmids were transfected into 293T cells by EndoFectin<sup>TM</sup> (GeneCopoeia, Inc. US) according to the manufacturer's procedure. Lentivirus were harvested and concentrated at ultrahigh speed for 1 h and stored at -80°C.Cells were incubated 20 hours in 16-well plate at 37°C in a humidified incubator in an atmosphere of 5% CO<sub>2</sub>. Then 1 ml medium and Hexadimethrine bromide were added after removal of old medium. Before adding 200 $\mu$ l of lentiviral particles, gently swirl the plate to mix. Then the mixture was incubated at 37°C in a humidified incubator in an atmosphere of 5-7% CO<sub>2</sub>. 24 h later, the media containing lentiviral particles was removed from wells and fresh media was added. Later, the cells were cultivated for another 48 h.

### shRNA Knockdown

Nucleotide sequences (IFIT2-652) used for IFIT2 gene silencing was got from document.<sup>28</sup> Double-stranded oligonucleotides targeting human IFIT2 cDNA was cloned into pLVX-shRNA1 vector. Virus packaging and infection were seen above.

### Statistical analysis

All data were repeated 3 times and presented as mean  $\pm$  SD. (n  $\geq$  3) SPSS13.0 software (SPSS Inc., Chicago, IL) was used for all analyses.

### Disclosure of potential conflicts of interest

No potential conflicts of interest were disclosed.

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### Author contributions

Zhanglin Zhang contributed to the study design., Yonglu Zhang, Yunyuan Kong, Shuyuan Liu, Lingbing Zeng, Lagen Wan, Zhanglin Zhang performed the research and conducted the data analysis. Zhanglin Zhang and Yonglu Zhang wrote the manuscript.

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