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Original Paper

The Effect of 6-Gingerol on Human AML **Cell Lines**

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Key Words

6-Gingerol • AML cell lines • Proliferation • Apoptosis • ROS •

Abstract

Background/Aims: Acute myeloid leukemia (AML) is a devastating hematological malignancy without a definitive cure. 6-Gingerol, a bioactive compound, has shown promise in treating various cancers, yet its impact on AML remains elusive. *Methods:* To elucidate the potential of 6-gingerol in AML, we conducted comprehensive experiments. Cell growth and clonogenic capacity were assessed using CCK-8 testing and colony formation assays. Flow cytometry was employed to analyze cell cycle progression and apoptosis. The invasive capability of AML cells was evaluated through the Transwell migration assay. Fluorescent probe staining was used to determine intracellular reactive oxygen species (ROS) concentration, while Western Blot was utilized to assess the expression levels of key proteins including Bcl-2, caspase3, MAPK, and p-MAPK in AML cells. Potential targets of 6-gingerol in AML were identified through various bioinformatics databases such as STP, SEA, STICH, OMIM GeneMap, and GeneCards. Enrichment analysis of Gene Ontology (GO) and Kyoto Encyclopedia of Genes and Genomes (KEGG) pathways was performed using the cluster Profiler package (v4.16.0). Results: Our findings indicated that 6-gingerol effectively inhibits proliferation, colony formation, and invasive capacity of AML cells, arresting them at the G1 phase of the cell cycle. Furthermore, 6-gingerol enhanced the expression levels of ROS, caspase 3, MAPK, and p-MAPK in AML cells. 67 overlapping targets between 6-gingerol and AML were identified, which are enriched within the MAPK signaling pathway and ROS-related pathways. Notably, NFKB1 emerged as a pivotal hub gene through which 6-gingerol exerts its influence on AML. Conclusion: 6-Gingerol could act as a promising agent sourced from Traditional Chinese Medicine (TCM) for AML treatment. © 2025 The Author(s). Published by

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Zhang et al.: the Effect of 6-Gingerol on Human AML Cell Lines

Introduction

Acute myeloid leukemia (AML), a malignancy arising from hematopoietic stem or progenitor cells within the bone marrow [1], presents a challenging clinical scenario with a median survival of approximately one year post-diagnosis [2]. Globally, AML incidence stands at 1.5 per 100, 000 individuals, exhibiting a higher rate of 2.4 per 100, 000 in Western nations [3]. The worldwide mortality rate for AML is 1.3 per 100, 000, escalating to approximately 2.2 per 100, 000 in Western Europe and North America [3]. In accordance with the 2022 classification by the European Leukemia Net, complete remission rates vary among risk groups: 73% for favorable-risk, 66% for intermediate-risk, and 45% for poor-risk AML patients. Correspondingly, five-year progression-free survival (PFS) rates are approximately 52%, 32%, and 16%, while overall survival (OS) outcomes reach 55%, 34%, and 15%, respectively [4]. The standard induction therapy for AML employs the "3+7" regimen, a combination of cytarabine and anthracyclines. The US Food and Drug Administration has approved several targeted therapies for AML in recent years, including FMS-like tyrosine kinase 3 inhibitors [5], isocitrate dehydrogenase inhibitors [6, 7], and B-cell lymphoma 2 (BCL2) inhibitors [8]. Furthermore, drugs such as TP53 are under clinical investigation, demonstrating potential to enhance complete remission rates, recurrence-free survival, and OS in select AML patients [9]. Nevertheless, the prognosis remains bleak for older patients (≥60 years) and those with relapsed/refractory AML, with five-year OS rates of merely 4-18% and 10%, respectively [10]. Therefore, there is an urgent need to develop novel treatment strategies to address this unmet medical need in AML management.

6-Gingerol, an active constituent derived from ginger (a plant renowned for its dual roles in both medicine and cuisine), exhibits a broad spectrum of biological activities, including anticancer, anti-inflammatory, anti-apoptotic, antioxidant, and anti-obesity effects [11-14]. Specifically, 6-gingerol has been shown to inhibit the progression of cervical cancer by generating reactive oxygen species (ROS), which not only damages DNA but also suppresses cell proliferation and promotes cell death [15]. Nevertheless, the efficacy of 6-gingerol in treating AML cells remains largely uncharted territory.

In this study, we delved into the impact of 6-gingerol on AML cell lines, examining its effects on proliferation, colony formation, cell cycle progression, apoptosis induction, invasive capacity, as well as the modulation of key biomarkers including ROS, caspase 3, MAPK, and p-MAPK. This investigation aims to provide novel insights into the potential therapeutic benefits of 6-gingerol in AML management.

Materials and Methods

Cells and cell culture

We obtained human AML cells (HL-60 and SKM-1) from Procell. We cultured HL-60 cells using IMDM (Procell) supplemented with 20% fetal bovine serum (FBS) and 1% penicillin-streptomycin (P/S). SKM-1 cells were grown in RPMI 1640 (Procell) containing 10% FBS and 1% P/S.

Cell proliferation assay

HL-60 and SKM-1 cells were exposed to various concentrations of 6-gingerol for 0, 24, 48, and 72 hours, respectively. Cell proliferation was assessed using the CCK-8 (BKMAM) assay, following the manufacturer's instructions. A microplate reader (Shanpu) measured the absorbance at 450 nm, enabling the calculation of cell viability.

Colony formation assay

To prepare the lower gel layer, 1.2% agar was combined with 2× medium. This mixture was added to 6-well plates and allowed to solidify at 37°C for 30 minutes. Once the lower gel layer had solidified, cell suspensions were mixed with a 0.7% agar-medium mixture to create the upper gel layer. The plates were then incubated until visible colonies formed (2-4 weeks). Subsequently, the colonies were stained with nitroblue tetrazolium (NBT) staining solution at 37°C for 1-4 hours, photographed, and counted.

920

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Zhang et al.: the Effect of 6-Gingerol on Human AML Cell Lines

Cell cycle analysis

Cells were either treated with 200 µM 6-gingerol (Med Chem Express) or left untreated for 72 hours. Following this treatment, the cells were collected and preserved in 75% ethanol. DNA staining was performed using the Cell Cycle Detection Kit (Solarbio). The proportions of each phase in the cell cycle were determined through flow cytometric analysis and processed using NovoExpress software.

Cell apoptosis assay

Cells were harvested and stained with Annexin V-phycoerythrin/7-aminoactinomycin D from the Annexin V-phycoerythrin/Apoptosis Detection Kit (Solarbio) according to the manufacturer's instructions. Apoptosis cells were analyzed using flow cytometry (Agilent NovoCyte) and processed with NovoExpress software (Agilent NovoCyte).

Transwell migration assay

After gel formation and a 72-hour incubation period, cell suspension was introduced into transwell inserts. A 24-well plate was then filled with 600 µL of 4% paraformaldehyde, and the inserts were submerged within this solution for 20-30 minutes to secure cell integrity. Subsequently, cells were stained with 0.1% crystal violet, air-dried, and observed to enumerate the number of migrated cells.

ROS detection

Cells were collected post-treatment with 6-gingerol and suspended in serum-free medium. The ROS Detection Kit (Solarbio) was employed for ROS assessment. A DCFH-DA solution was added to the cell suspension, and cells were subsequently washed. The intracellular fluorescence, an indicator of ROS levels, was visualized using a fluorescence microscope.

Western blot analysis

Cells were lysed and combined with protein phosphatase inhibitors. The protein concentration of the sample was determined using the bicinchoninic acid Protein Quantification Kit (CWBIO). Following lysis, proteins were subjected to SDS-PAGE and transferred onto PVDF membranes via electroblotting. The membranes were then blocked with 5% non-fat milk at room temperature for one hour. Subsequently, they were incubated overnight at 4°C with gentle agitation using primary antibodies targeting Bcl-2, caspase-3, MAPK, and p-MAPK (Servicebio). After thorough washing, the membranes were exposed to a secondary antibody for 1.5 hours. Following another wash with 1×Tris-buffered saline containing Tween 20 (TBST), an ECL reagent from Sangon Biotech was added for chemiluminescent detection. ACTIN (TransGen Biotech) served as an internal control in this process.

Identification of 6-gingerol and AML-related potential target

To identify potential therapeutic targets for 6-gingerol in AML, we utilized several bioinformatic databases and tools. The potential targets of 6-gingerol were predicted using the SwissTargetPrediction, SEA, and STITCH databases. AML-related genes were retrieved from the OMIM GeneMap and GeneCards databases using the keyword "Acute Myeloid Leukemia." The overlapping targets between 6-gingerol and AML were considered as candidate therapeutic targets. To evaluate potential target interactions, we utilized the STRING (Search Tool for the Retrieval of Interacting Genes/Proteins) database to construct a proteinprotein interaction network. The analysis results were then imported into Cytoscape software (version 3.10.3) to construct a "component-target-pathway" network. To identify key genes within this network, we employed the CytoHubba extension in Cytoscape, utilizing four scoring models: Degree, Edge Percolated Component (EPC), Maximum Clique Centrality (MCC), and Maximum Neighborhood Component (MNC). Genes that ranked concurrently by all four criteria were designated as core hubs. The intersections among these results were illustrated through an UpSet diagram. Additionally, GO (Gene Ontology) and KEGG (Kyoto Encyclopedia of Genes and Genomes) enrichment analyses were performed using the clusterProfiler (version 4.16.0) package in R, retaining only pathways with p < 0.01 for further interpretation.

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Zhang et al.: the Effect of 6-Gingerol on Human AML Cell Lines

Statistical analysis

The data were expressed as mean ± standard deviation (x ± SD). Statistical analysis was conducted using SPSS 19.0 with one-way ANOVA. P < 0.05 was considered statistically significant. All figures were created using GraphPad Prism 9.0.

Results

6-Gingerol inhibited the proliferation and colony formation of AML cells

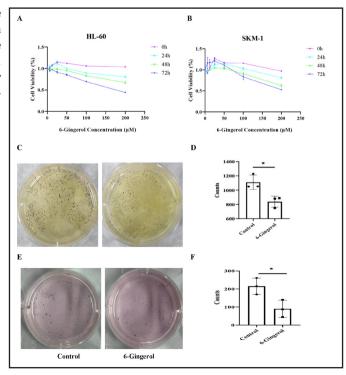
To evaluate the impact of 6-gingerol on AML cell proliferation, we employed the CCK-8 assay. As shown in Fig. 1A, treating HL-60 cells with 25 µM 6-gingerol for 0 or 24 hours resulted in a slight increase in cell viability, followed by a gradual decrease as the 6-gingerol concentration increased. After 48 and 72 hours of treatment, HL-60 cell viability decreased gradually. Notably, at a concentration of 200 µM, cell viability was reduced to 67.3% and 43.9%, respectively. For SKM-1 cells (Fig. 1B), cell viability initially increased slowly and then decreased rapidly within the 6-gingerol concentration range of 0-100 µM. When the concentration was increased from $100~\mu M$ to $200~\mu M$ for 72 hours, SKM-1 cell viability decreased from 83.1% to 52.6%.

Compared to untreated controls, exposure to 6-gingerol significantly reduced the number of colonies. Specifically, the number of HL-60 colonies decreased from 1110±102.20 to 839±78.58, and SKM-1 colonies decreased from 215±45.50 to 90±48.09 (Fig.1C-F). These findings demonstrate that 6-gingerol effectively inhibits the proliferation and colony-forming capacity of AML cells.

6-Gingerol arrested G1 phase of HL-60 and SKM-1

To further investigate the impact of 6-gingerol on cell cycle progression in HL-60 and SKM-1 cells, we employed flow cytometry to study the cell cycle progression of these two cell lines (Fig. 2). In HL-60 cells, treatment with 6-gingerol resulted in a significant increase in the G1 phase population from 33.6% to 65.1%, accompanied by a decrease in the S and G2 phases from 30.5% to 17.0% and from 18.5% to 10.6%, respectively. Similarly, in SKM-1 cells,

1. 6-Gingerol inhibited proliferation and colony formation of AML cells. 6-gingerol inhibited the proliferation of HL-60(A) and SKM-1 (B); 6-gingerol inhibited the colony formation of HL-60 (C-D) and SKM--(E-F).*P < 0.05.



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Zhang et al.: the Effect of 6-Gingerol on Human AML Cell Lines

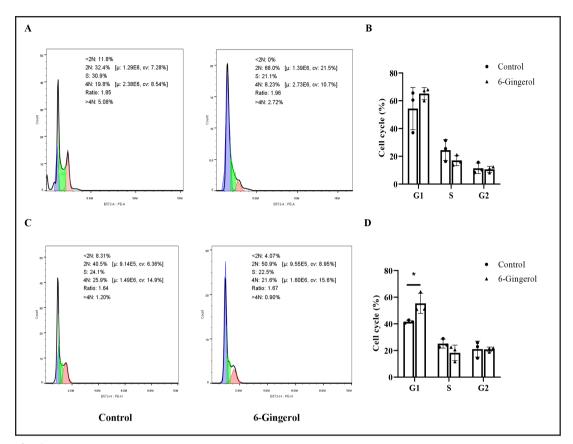


Fig. 2. 6-Gingerol arrested G1 phase of HL-60 and SKM-1. (A-B) HL-60; (C-D) SKM-1.*P<0.05.

treatment with 6-gingerol caused an accumulation of cells in the G1 phase, with concomitant reductions in both the S and G2 fractions. These findings suggest that 6-gingerol may induce DNA damage in AML cells, either triggering the apoptotic pathway or blocking cell cycle checkpoints, ultimately leading to G1 phase arrest.

6-Gingerol induced apoptosis of AML cell lines

Flow cytometry analysis revealed a substantial increase in the apoptosis rate of AML cells treated with 6-gingerol compared to the control group (Fig. 3). Specifically, the apoptosis rate of HL-60 cells increased from $1.767\pm0.18\%$ to $25.563\pm7.05\%$, and that of SKM-1 cells increased from $3.75\pm0.485\%$ to $33.287\pm10.061\%$ (P <0.01). These data unequivocally confirm that 6-gingerol effectively promotes apoptosis in AML cell lines.

6-Gingerol suppressed invasive capacity of AML cells

The invasive potential of HL-60 and SKM-1 cells, subjected to treatment with 6-gingerol, was evaluated using a transwell assay. In comparison to the control group, 6-gingerol significantly reduced the migration of HL-60 cells from 385 ± 50.587 to 130 ± 11.533 (P<0.01) and SKM-1 cells from 173.667 ± 48.645 to 62.333 ± 4.041 (P<0.05), respectively (Fig. 4). These results unequivocally demonstrate that 6-gingerol exerts a profound inhibitory effect on the invasive capabilities of both HL-60 and SKM-1 cell lines.

6-Gingerol induced the ROS levels of AML cell lines

The administration of 6-gingerol was observed to elicit a significant elevation in the intracellular ROS levels within AML cell lines, as assessed through immunofluorescence assays. As illustrated in Fig. 5, the mean fluorescent intensity (MFI) of HL-60 and SKM-1 cells subjected to 6-gingerol treatment notably rose from 8.57 ± 3.388 to 21.516 ± 6.197 (P

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923

Zhang et al.: the Effect of 6-Gingerol on Human AML Cell Lines

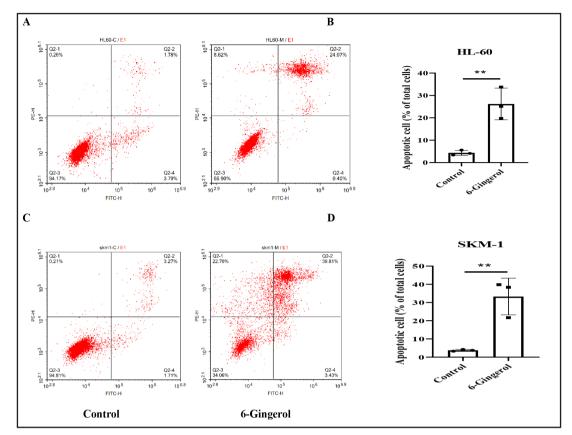


Fig. 3. 6-gingerol induced apoptosis of AML cell lines. (A-B) HL-60;(C-D) SKM-1. **P<0.01.

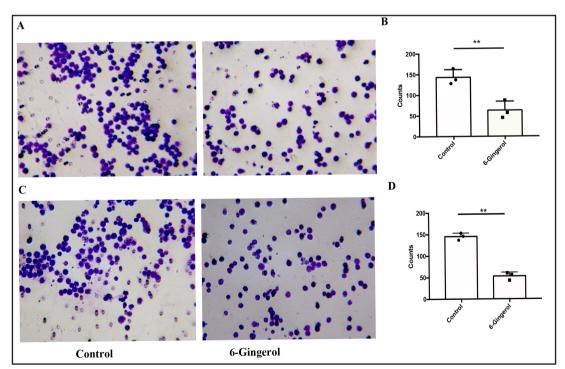


Fig. 4. 6-Gingerol suppressed invasive capacity of AML cells. (A-B) HL-60;(C-D) SKM-1.**P<0.01.

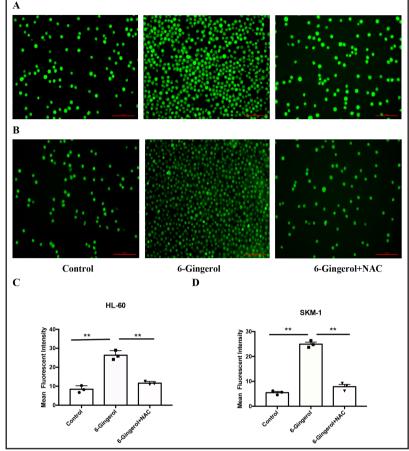
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Zhang et al.: the Effect of 6-Gingerol on Human AML Cell Lines

Fig. 5. 6-Gingerol induced the ROS levels of AML cell lines. (A, C) HL-60;(B, D) SKM-1.**P < 0.01.



< 0.05) and from 3.588 \pm 2.605 to 20.87 \pm 5.723, respectively (P < 0.01). Notably, the MFI of both cell lines decreased when the ROS inhibitor (NAC, Solarbio) was introduced, suggesting that 6-gingerol is a stimulant of ROS levels in AML cell lines.

6-Gingerol improved the protein expression levels of caspase3, mitogen-activated protein kinase (MAPK), p-MAPK of AML cells

To further elucidate the influence of 6-gingerol on AML cells, particularly in relation to Bcl-2, caspase3, MAPK, and p-MAPK, Western blotting was conducted (Fig. 6). In comparison to the control group, 6-gingerol decreased the level of Bcl-2 in HL-60 cells (from 0.548 ± 0.013 to 0.24 ± 0.048) (P < 0.001) and SKM-1 cells (from 0.401 ± 0.058 to 0.122 ± 0.008) (P < 0.01) (Fig. 6). Simultaneously, it augmented the expression of caspase 3 (from 0.123 ± 0.054 to 0.615 ± 0.013 in HL-60 cells and from 0.21 ± 0.03 to 0.52 ± 0.17 in SKM-1 cells), MAPK (from 0.138 ± 0.015 to 0.399 ± 0.023 in HL-60 cells and from 0.441 ± 0.157 to 0.921 ± 0.295 in SKM-1 cells), and p-MAPK (from 0.193 ± 0.045 to 0.356 ± 0.068 in HL-60 cells and from 0.259 ± 0.084 to 0.971 ± 0.227 in SKM-1 cells) in both cell lines. These findings collectively indicate that 6-gingerol promotes the protein expression levels of caspase3, MAPK, and p-MAPK in AML cells.

Analysis of 6-gingerol and AML-related potential targets

194 6-gingerol-related potential targets were predicted through the SwissTarget Prediction, SEA, and STICH databases while 2930 AML-related targets were retrieved from the GeneCards and OMIM databases. Notably, there was an overlap of 67 targets pertinent to both 6-gingerol and AML (illustrated in Fig. 7A). These shared targets underwent PPI analysis and were visualized using Cytoscape software (as shown in Fig. 7B).

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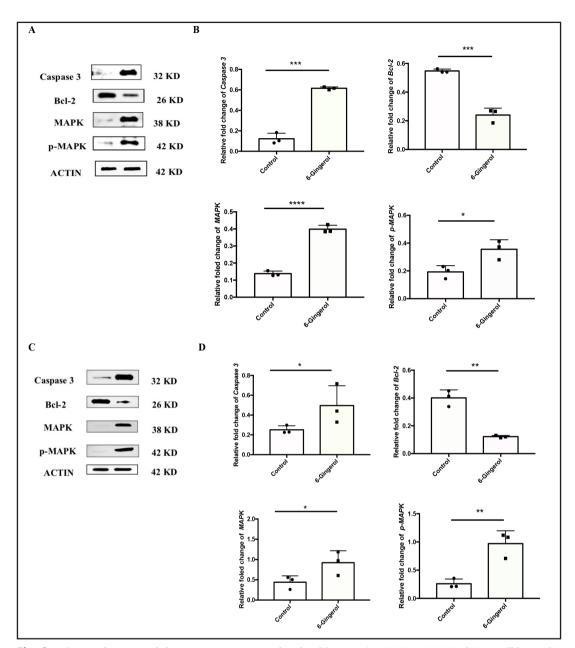


Fig. 6. 6-Gingerol improved the protein expression levels of Caspase3, MAPK, p-MAPK of AML cell lines. (A-B) HL-60; (C-D) SKM-1. *P < 0.05, **P < 0.01, ***P < 0.001, ****P < 0.0001.

In GO enrichment analysis, the overlapping targets were primarily enriched in the biological process (BP) "response to peptide hormone," the cellular component (CC) "extrinsic component of membrane," and the molecular function (MF) "histone kinase activity" (depicted in Fig. 7C). Furthermore, KEGG pathway enrichment analysis revealed their association with tumor-related pathways, including the PI3K-Akt signaling pathway, the MAPK signaling pathway, and ROS-related pathways (presented in Fig. 7D).

To identify key genes, the CytoHubba plugin (which employs four algorithms: Degree, EPC, MCC, and MNC) was employed to rank the top 10 genes based on their scores (as shown in Fig. 7E). Importantly, six hub genes consistently identified by all four algorithms were selected (displayed in Fig. 7F). The constructed "component - target - pathway" network (illustrated in Fig. 7G) underscores that these targets are predominantly enriched within the MAPK signaling pathway and ROS-related pathways. Notably, NFKB1 emerges as a pivotal hub gene through which 6-gingerol exerts its influence on AML.

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926

Zhang et al.: the Effect of 6-Gingerol on Human AML Cell Lines

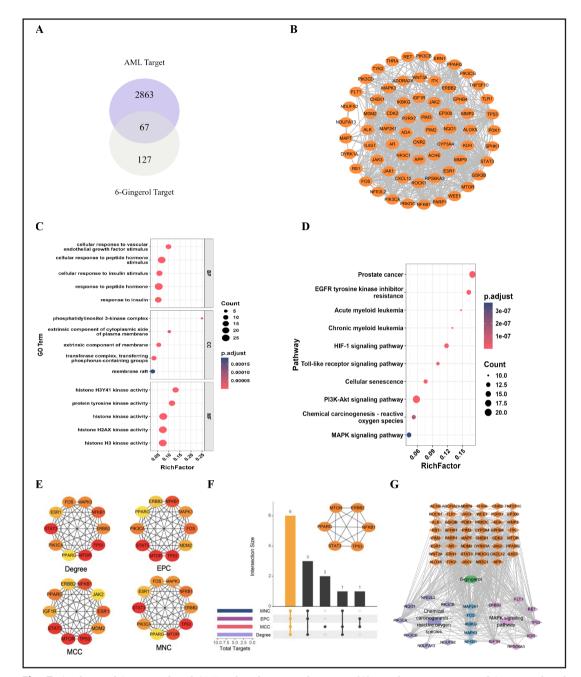


Fig. 7. Analysis of 6-gingerol and AML-related potential targets. (A) overlapping targets of 6-gingerol and AML;(B) Potential target network diagram of 6-gingerol for AML treatment;(C) GO enrichment bubble plot;(D) KEGG enrichment bubble plot;(E) the top 10 genes;(F) 6 hub genes;(G) "Component-target-pathway" network analysis plot.

Discussion

This study was the first to demonstrate that 6-gingerol effectively inhibits the proliferation, colony formation, and invasive capacity of AML cells, culminating in their arrest at the G1 phase of the cell cycle. Furthermore, 6-gingerol enhanced the expression levels of ROS, caspase 3, MAPK, and p-MAPK in AML cells.

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Zhang et al.: the Effect of 6-Gingerol on Human AML Cell Lines

AML presents with a dismal prognosis, and disease recurrence following shortterm clinical remission remains a significant challenge in its treatment. The persistence of leukemia stem cells (LSCs) is primarily responsible for AML recurrence. 6-gingerol, a commonly used herb in TCM [16], exhibits multiple pharmacological activities including antioxidation [17], anti-inflammation [18], anti-obesity, anti-cancer [12, 13], anti-hyperglycemia and immune regulation [14, 19]. Previous evidence indicated that 6-gingerol suppresses tumor cell proliferation in oral cancer by inducing apoptosis and halting progression at the G2/M checkpoint [20]. In contrast, our data revealed that in AML cells, 6-gingerol inhibits proliferation and colony formation, induces G1 phase arrest, and substantially diminishes their invasive capacity.

AML cells primarily rely on oxidative phosphorylation (OXPHOS) as their energy source, with ROS as a by-product. ROS plays a pivotal role in AML pathogenesis and therapeutic targeting [21]. Elevated ROS levels can damage cells and intracellular components, leading to DNA damage, protein denaturation, and tissue injury, ultimately resulting in G2/M phase arrest, apoptosis, senescence, or cell death. ROS are also implicated in mitochondria-, death receptor-, and endoplasmic reticulum-mediated apoptosis [22]. Consistently with these reports, our study demonstrated that 6-gingerol significantly increases ROS levels in AML cell lines. Furthermore, ROS levels decreased when an inhibitor of ROS was added.

It is known that MAPK are crucial for cancer cell survival [23] and involved in proliferation, differentiation, apoptosis, angiogenesis, inflammation, stress responses, and immune defense [24, 25]. Caspase 3 is a cysteinyl aspartate-specific proteinase that serves as an executioner caspase in apoptosis and mediates the anti-cancer effect of cytotoxic drugs [26]. Earlier research study reported that 6-gingerol downregulated anti-apoptotic protein BCL-2 and survivin, while enhancing Bax expression and activating caspase 3 and caspase 9, thereby inducing apoptosis in bladder cancer via MAPK- and ROS-dependent signaling cascades [27]. Our study indicated that 6-gingerol treatment also results in a reduction in Bcl-2 expression and an increase in the levels of caspase 3, MAPK, and p-MAPK in AML cell lines. When inhibitors of ROS, caspase 3, and MAPK were added, ROS levels, caspase 3 activity, and apoptosis rates decreased, improving cell viability in both cell lines (data not shown). Our findings suggested that the impact of 6-gingerol on AML cell lines may be mediated through caspase 3, ROS, and MAPK. However, the observed changes in protein expression represent molecular observations in the absence of pathway inhibition or knockdown, leaving mechanistic validation as a future direction for study. There are 67 overlapping targets between 6-gingerol and AML, which are enriched within the MAPK signaling pathway and ROS-related pathways. Furthermore, NFKB1 emerges as a central node through which 6-gingerol exerted its influence on AML cells. This represented a hypothesis-generating exercise rather than an experimentally validated target or pathway.

In conclusion, we have successfully demonstrated a previously uncharacterized fact that the effect of 6-gingerol on AML cell lines is apparent. This implies that 6-gingerol could potentially serve as a promising therapeutic agent derived from TCM for the treatment of AML.

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The authors have no acknowledgements to report.

Author contributions

M.Wu analyzed data, prepared figures and write the manuscript.

T.T. Zhang cultured cells and write manuscript and prepare figures.

C.F. Kong conducted cell proliferation and colony formation assay.

A.N. Li contributed to cell cycle and apoptosis assay.

H.B. Cheng participated in the design and data analysis.

W.R. Ding contributed to ROS assay.

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Zhang et al.: the Effect of 6-Gingerol on Human AML Cell Lines

B. Ke conducted Western Blot experiments.

C.Chen Analyzed of 6-gingerol and AML-related potential targets.

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Statement of Ethics

Cell lines in this study were purchased from Procell. It does not require ethics committee approval.

Disclosure Statement

The authors declare no conflicts of interest.

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