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Retraction notice for: "Baicalein restrains proliferation, migration, and invasion of human malignant melanoma cells by down-regulating colon cancer associated transcript-1" [Braz J Med Biol Res (2019) 52(12): e8934]

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Retraction for: Braz J Med Biol Res | doi: 10.1590/1414-431X20198934 | PMID: 31778440 | PMCID: PMC6886380

The authors would like to retract the article "Baicalein restrains proliferation, migration, and invasion of human malignant melanoma cells by down-regulating colon cancer associated transcript-1" that was published in volume 52 no. 12 (2019) (Epub Nov 25, 2019) of the Brazilian Journal of Medical and Biological Research.

After the publication of this study, the corresponding author requested its retraction due to "the identification of unspecified data inconsistency that could lead to mistaken conclusions." The Editors agreed with and endorsed that decision.

The Brazilian Journal of Medical and Biological Research had received authorization from all authors before the publication of the paper. We regret the unprofessional behavior of the authors involved.

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Baicalein restrains proliferation, migration, and invasion of human malignant melanoma cells by down-regulating colon cancer associated transcript-

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Abstract

Baicalein (BAI) is an acknowledged flavonoids compound, which is regarded a useful therapeutic pharmaceutical for numerous cancers. However, its involvement in melanoma is largely unknow. This sture aimed to examine the anti-melanoma function of BAI and unraveled the regulatory mechanism involved. A375 at SN 28 were treated with BAI for 24 h. Then, CCK-8 assay, flow cytometry, and transwell assay were carried out to investig to ell growth, migration, and invasion. RT-qPCR was applied to detect the expression of colon cancer associated transcript- (CCAT1) in melanoma tissues and cells. The functions of CCAT1 in melanoma cells were also evaluated. Western blot was ulized to appraise Wnt/β-catenin or MEK/ERK pathways. BAI restrained cell proliferation and stimulated or apply tic capability of melanoma by suppressing cleaved-caspase-3 and cleaved-PARP. Cell migratory and invasive a ties were restrained by BAI via inhibiting MMP-2 and vimentin. CCAT1 was over-expressed in melanoma tissues and own-regulatory by BAI in melanoma cells. Overexpressed CCAT1 reversed the BAI-induced anti-growth, anti-migratory, a darti-invalue effects. Furthermore, BAI inhibited Wnt/β-catenin and MEK/ERK pathways-axis via regulating CCAT1. Our right dicated that BAI blocked Wnt/β-catenin and MEK/ERK pathways via regulating CCAT1, thereby inhibiting melanoma cells eliferation, migration, and invasion.

Key words: Malignant melanoma; Colon can a cate transcript-1; Wnt/β-catenin; MEK/ERK

Introduction

Melanoma evolves from skin mu membrane and is the most remmon cover with high metastatic potential (1). Maliginary noma, caused by the abnormal transformation of man melanocytes, is one of the fastest grant ma nant tumors with an annual growth rate 3-F $_{\circ}$ (2). To date, surgery and chemotherapy combine the most are the most common endors of thera vitic approaches to melanoma (3,4). Neverth ... the biggest disadvantage of these therapies is toxicity. Thus, there is research focused on natural products towards cell metastasis of melanoma (5). However the potential value of traditional Chinese medieatm at of melanoma has not been assessed. cine in the aicalensis Georgi is a kind of traditional a medicine containing several flavonoids. One of rigredients is baicalein (BAI), which is commonly regalied as useful adjuvant therapeutic pharmaceutical for vanous diseases (6). Thus far, a number of researchers tested the efficacy of BAI on malignant tumors, such as breast carcinoma (7), non-small-cell lung carcinoma (8), cervical carcinoma (9), and carcinoma of urinary bladder (10). Moreover, previous research indicated that BAI impeded cell proliferation and melanogenesis of B16F10 mouse melanoma cells (11,12). What is not yet clear is the functional mechanism of BAI on human malignant melanoma.

Long noncoding RNAs (IncRNAs) are RNA segments with no fewer than 200 nucleotides in length that do not encode proteins (13). IncRNAs are closely linked to miscellaneous regulations, functioning as regulators of gene transcription, RNA splicing, and miRNA regulatory systems (14,15). A number of investigators reported that IncRNAs SLNCR1 and HEIH interfered with the melanoma cell proliferative potential, migratory status, and invasive ability via regulating corresponding downstream targets (16,17). Colon cancer associated transcript-1 (CCAT1), an innovative tumor-related IncRNA, plays an essential role in tumor progression, being up-regulated

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Received June 15, 2019 | Accepted September 9, 2019

in malignancies (18). However, the extent to which CCAT1 is related to malignant melanoma remains poorly understood.

Here, we demonstrated a crucial role of BAI in inhibiting cell growth and motility by mediating CCAT1 as well as the underlying mechanism of BAI-induced signaling pathways in human melanoma cells. Our findings might provide new insights into the application of traditional Chinese medicine and feasible therapies for malignant melanoma.

Material and Methods

Clinical tissues

Twenty-two pairs of human melanoma tissues and corresponding paracancerous skin specimens were collected from patients at Qingdao Central Hospital (Qingdao, Shandong) from January 2017 to July 2018. Thirteen cases were from males and 9 were from females, who did not receive any radiation or chemotherapy before surgery. Participants signed an authorization and the Ethics Committee of Qingdao Central Hospital approved the procedures and the study.

Cell culture and treatment

The malignant melanoma cell lines A375 and SK-MEL-28, which were cultured in DMEM (Gibcr JSA enriched with 10% fetal bovine serum (FBS, Gibco), we obtained from ATCC (USA). The conditions for cell culturer 5% CO_2 and 37°C. BAI was obtained to Nanjing ZeLang Medical Technology Co. Ltd. (#ZL100708, hina). BAI was diffused in DMSO as a storage concentration and diluted using DMEM to work concentrations (4 0, 50, 20, and 10 μ M). The cells were treated van BAI for 24 h.

Cell transfection

The entire length of CCAT was concatenated into the pcDNA3.1 vector (Concatenated into the pcDNA3.1 vector (Concatenated in a pcomposition plasmid was sampled as pccAT1. The lipofectamine 2000 reagent (Concatenated in pcccate in pc

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Is (a 1 /well) were seeded into 96-well plates of which remarks a harmonic of the real way of the cultures. Then, cultures were incubated for 1 ha. 37°C. Microplate Reader (Bio-Rad, USA) was employed to evaluate the cell viability at 450 nm.

Bromodeoxyuridine (BrdU) assay

Cell proliferation was determined using BrdU (Sigma-Aldrich, USA). After treatment of BAI, BrdU (1 mg/mL)

was added to the cells for 3 h. Then, immunofluorescence assay was carried out to estimate the BrdU-tagg cells, providing the cell proliferation rate.

Cell migration and invasion assays

Cell migratory capacity and invasive potential assessed by transwell culture chamber rning C satar, USA), which consists of 8-um re 'v .cbonate membrane. Firstly, 200 μ L of 1 10^4 cells, which were cultured in DMEM without FBS, were see and into the top chamber, which had been vere with Matrigel matrix (Becton Dickinson, US/ for vasion assay or kept uncovered for migrative assay. Insequently, 800 μL medium was injected to the wer chamber. After 24 h, the migratory cells we fixed who methyl alcohol and dyed with 0.5% cryst viol liquid (Solarbio). Then, the relative migration rate. ver ated. After 48 h, the invading cells were proces of in the above same manner and the number vading Is was counted.

Apoptosis 3ay

Apoptotic cells proportion was measured utilizing PI/
-Annexin V staining kit (Invitrogen, USA). In brief, cells ×10⁶/well) were cultured into 6-well plates and starve in FBS-free medium for 12 h. Next, PI and the control of the cell cultures. Flow cytometry was performed with FACScan (Becton ckinson). The apoptosis ratio was calculated using FlowJo software (Becton Dickinson).

Reverse transcription and quantitative real-time PCR (RT-qPCR)

Trizol reagent (Life Technologies Corporation, USA) was utilized to isolate total RNA of tissue samples and cell cultures. Reverse transcription of RNA was implemented utilizing SuperRT cDNA Synthesis Kit (Cwbio, China). SYBR® Green PCR Kit (Qiagen, Germany) was employed for qPCR analysis to detect CCAT1 expression. qPCR was executed on iQ5 real-time PCR Detection system (Bio-Rad). The mRNA expression of CCAT1 was normalized with β -actin. The relative quantification of CCAT1 in tumor tissues and cells was calculated using the equation: amount of target = $2^{-\Delta \Delta Ct}$ (19).

Western blot

Total proteins were extracted from cells utilizing RIPA lysis buffer (Cwbio), which contains phenylmethylsulfonyl fluoride (PMSF, Solarbio). Proteins were quantified by the Super-Bardford Pritein Assay Kit (Cwbio). The extractions were loaded into 12% polyacrylamide gel on the Bis-Tris Gel system (Bio-Rad). The products were transferred onto polyvinylidene fluoride (PVDF) membranes, which were then cultivated at 4°C overnight with primary antibodies. The primary antibodies included anti-cleaved-caspase-3 (#ab2303, Abcam, USA), anti-cleaved-PARP (#ab3246, Abcam), anti-MMP-2 (#ab37150, Abcam), anti-vimentin

(#ab92547, Abcam), anti-Wnt3a (#ab219412, Abcam), anti-β-catenin (#ab32572, Abcam), anti-t-MEK (#9126, Cell Signaling Technology, USA), anti-p-MEK (#9154, Cell Signaling Technology), anti-t-ERK (#9102, Cell Signaling Technology), anti-p-ERK (#4370, Cell Signaling Technology), and anti-β-actin (#ab179467, Abcam). Then, the PVDF membranes were rinsed and incubated with horseradish peroxidase-conjugated goat anti-rabbit IgG (#ab6721, Abcam) and goat anti-mouse IgG (#ab205719, Abcam) for 1 h at 20°C. After washing, the PVDF membranes were treated with ChemiDoc™ XRS system (Bio-Rad), and the intensity of bands was finally evaluated with ImageJ software (NIH, USA).

Statistical analysis

Each experiment was repeated three times. Graphpad 6.0 software (USA) was utilized for statistical analysis. Data are reported as means \pm SD. Analysis of variance (ANOVA) and Student's *t*-test were applied to calculate P values. A P value <0.05 was regarded as significant.

Results

BAI attenuated cell proliferation and promot d cell apoptosis of malignant melanoma cells

Figure 1A presents the inhibition of BAI on cen-Cells were sensitive to 20 µM BAI compared with the untreated group (P<0.01). Cell viability as impresed by BAI with an inhibitory concentration of s Therefore, 50 µM was consider a to be an acceptable concentration for the next progration and apoptosis assay. Figure 1B shows the BA 50 M) significantly inhibited the cell prolife tion of A375 and SK-MEL-28 cells compared to e untre d cells (P<0.001). Reversely, flow cylimetry using PI/FITC-Annexin V indicated that BAI moted compared with <0.01, Figure 1C). We analyzed the untreated grup the expression aspase-3, which acted in cell apoptosis and be icipated in the cleavage of repair ch as RP (20). The protein expression enzvme ment with the result of flow cytometry. analysis

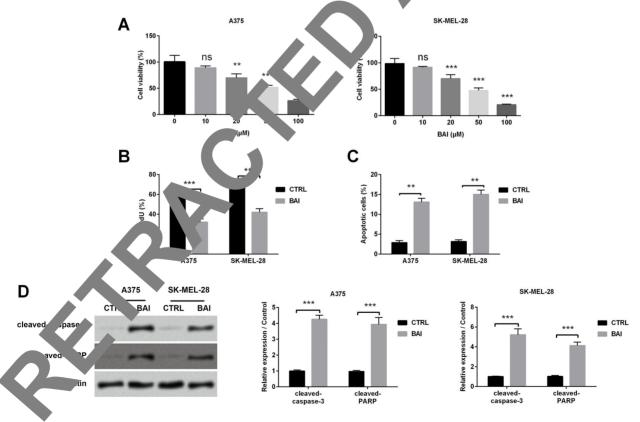


Figure 1. Baicalein (BAI) attenuated cell proliferation and strengthened cell apoptotic capacity of malignant melanoma cells. **A**, Cell viability of A375 and SK-MEL-28 cells followed by 24-h treatment with BAI (0, 10, 20, 50, and 100 μM) was assessed by CCK-8. **B**, Cell proliferation of melanoma cells was examined by bromodeoxyuridine (BrdU) assay. **C**, Flow cytometry was utilized to assess the apoptotic rate of melanoma cells. **D**, Expression of cleaved-caspase-3 and cleaved-PARP was tested by western blot assay. The relative expression of protein was normalized by β-actin. Data are reported as mean \pm SD. **P < 0.01, ***P < 0.001 (ANOVA). ns: not significant; CTRL: control.

BAI treatment accelerated cleaved-caspase-3 and cleaved-PARP expression compared with the untreated cells (P<0.001, Figure 1D). The experiments detected some evidence for the inhibitory effect of BAI on the growth of malignant melanoma cells.

BAI inhibited cell migratory capacity and invasive potential of melanoma cells

As indicated in Figure 2A, BAI significantly suppressed cell migration compared to control group (P<0.001). Figure 2B shows the results obtained from the preliminary analysis of Matrigel invasion assay. When melanoma cells were stimulated with BAI, there was an obvious decline in the relative invasive rate compared to the control group (P<0.001). It is well known that the activation of angiogenesis depends on MMP-2 and vimentin, which are known to participate in the epithelial-mesenchymal transition (21,22). As shown in Figure 2C, MMP-2 and vimentin expression was inhibited due to the treatment of BAI compared with the control (P<0.001). Overall, these results indicated that BAI apparently impaired the motility of malignant melanoma cells.

CCAT1 was up-regulated in melanoma tissues and down-regulated by BAI in melanoma cells

As can be seen in Figure 3A, CCAT1 expression vas increased in melanoma specimens compared v , the non-tumor tissues (P<0.001). Furthermore, BAL-in variables of the compared v , the non-tumor tissues (P<0.001).

down-regulation of CCAT1 (P<0.001) in malignant melanoma cells was also confirmed by RT-qPCR (Fig. e 3B). Thus, there might be an association between BAI at CCAT1.

BAI inhibited growth of malignant melanoma regulating CCAT1

To better understand the underlying n. cular pachanisms, including a possible role for CA √aenous overexpression of CCAT1 was im emented by cansfection of pCCAT1 into A375 and SK-N L-28 ce s (Figure 4A). After stable transfection with property and the stable transfection with $50~\mu\text{M}$ BAI. Proliferation as ay so wed that BAI suppressed the proliferation capar of main ant melanoma cells compared to control < 1, Figure 4B). However, the suppression was a rsed by e CCAT1 overexpression (P<0.001). In realle the promotion of BAI on cell apoptosis was also ed A375 and SK-MEL-28 cells. Furthermore, wes. blot analysis revealed that the protein levels of wed-cas, se-3 and cleaved-PARP were marksponse to the combined influence of edly decreased (P<0.001, Figure 4D). In summary, above BAI and Co data indicate that BAI inhibited cell growth and promoted pooptotic potential via weakening CCAT1.

BAI suppressed migration and invasion of malignant come via regulating CCAT1

Subsequently, we detected the cell migratory capacity and invasive potential in response to the treatment of BAI

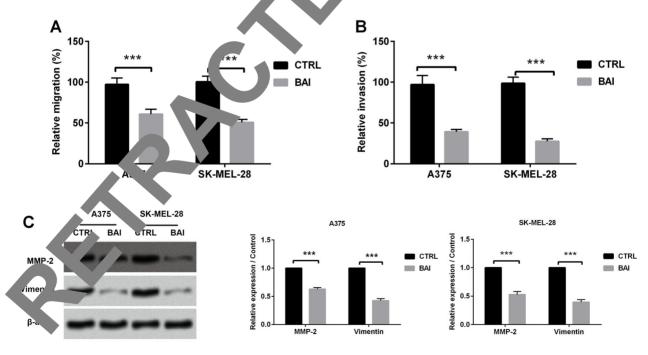


Figure 2. Baicalein (BAI) inhibited the cell migratory capacity and invasive potential of melanoma cells. **A**, Migration of A375 and SK-MEL-28 cells, treated or not with BAI, was examined by transwell assay. **B**, Invasion of A375 and SK-MEL-28 cells was detected by transwell assay with Matrigel matrix. **C**, Protein expression of MMP-2 and vimentin was tested by western blot assay. The relative expression of protein was normalized by β-actin. Data are reported as mean \pm SD. ***P < 0.001 (*t*-test). CTRL: control.

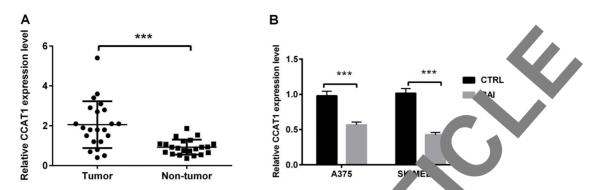


Figure 3. CCAT1 was up-regulated in melanoma tissues and down-regulated by baicalein ([, i) in relanoma cells. **A**, Expression of CCAT-1 in malignant melanoma tissues (n=22) and non-tumor skin specimens (n=22) was analyzed RT-qPCR. **B**, Expression of CCAT-1 in malignant melanoma cells after treating with BAI or not was determined by CR. Data are reported as mean ± SD. ***P < 0.001 (t-test). CTRL: control.

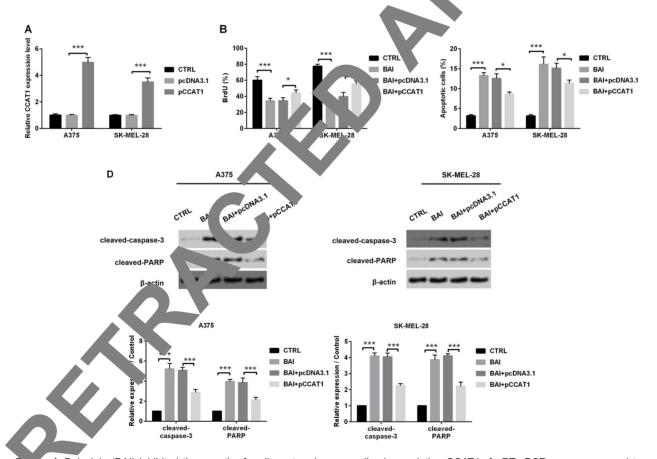


Fig. 2.4. Baicalein (BAI) inhibited the growth of malignant melanoma cells via regulating CCAT1. **A**, RT-qPCR assay was used to estimal CCAT1 expression in A375 and SK-MEL-28 after transfection with pCCAT1. **B** and **C**, BradU assay and flow cytometry assays were utilized to evaluate overexpression of CCAT1 and BAI on cell proliferation and apoptosis. **D**, Western blot assay evaluated the relative expression levels of cleaved-caspase-3 and cleaved-PARP. The relative expression of protein was normalized by β-actin. Data are reported as mean \pm SD. *P<0.05, ***P<0.001 (ANOVA). CTRL: control.

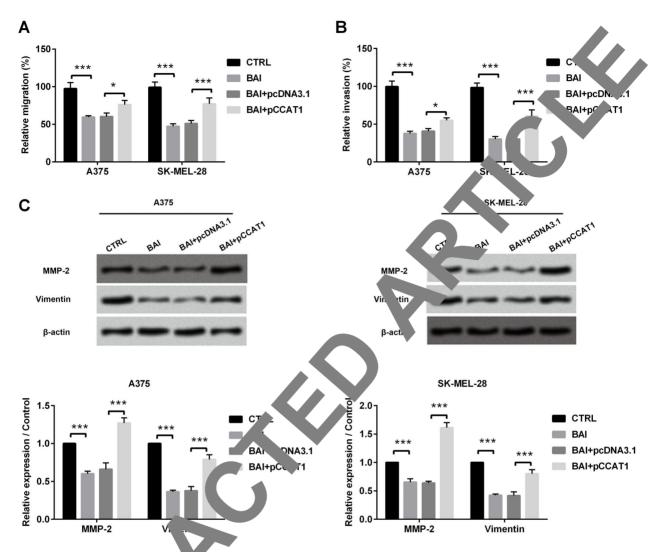


Figure 5. Baicalein (BAI) researche migration and invasion of malignant melanoma via regulating CCAT1. **A** and **B**, Transwell assay was utilized to termine active. It migratory and invasive rates. **C**, Western blot was utilized to analyze MMP-2 and vimentine expression levels in monomic acted with BAI or BAI + pCCAT1. The relative expression of proteins was normalized by β-actin. Data are reported a mean SD. *P < 0.05, ***P < 0.001 (ANOVA). CTRL: control.

and overexpress. CCAT1. Melanoma cells featured a decrea of relative ligratory rate and invasion rate with B/L tree ment compared with the untreated group (P<0.0 ligure 5A,B). We also discovered that the group translate with pCCAT1 and then treated with BAI whibind increased cell migration and invasion rate makes the group transfected with pcDNA3.1 and the treated with BAI (P<0.001). Concomitantly, we examined whether BAI could negatively regulate MMP-2 and vimentin expression through regulating CCAT1. As shown in Figure 5C, the protein expression of MMP-2 and vimentin were remarkably increased in CCAT1 overexpressed cells, which were not treated with BAI. Together, these results demonstrated that BAI exerted its

negative function of cell metastasis via regulating CCAT1 expression in malignant melanoma cells.

BAI suppressed CCAT1 to block Wnt/β-catenin and MEK/ERK signaling pathway-axis

The well-known tumor factor regulator, CCAT1, has been shown to have an overwhelming association with tumor proliferation and apoptosis by activating Wnt/ β -catenin and MEK/ERK signaling pathways (23,24). To address whether the above signaling pathways were involved in the function of BAI, protein expression was detected by western blot. Compared with the untreated group, the protein levels of Wnt3a and β -catenin were decreased in BAI-treated cells and were reversed by

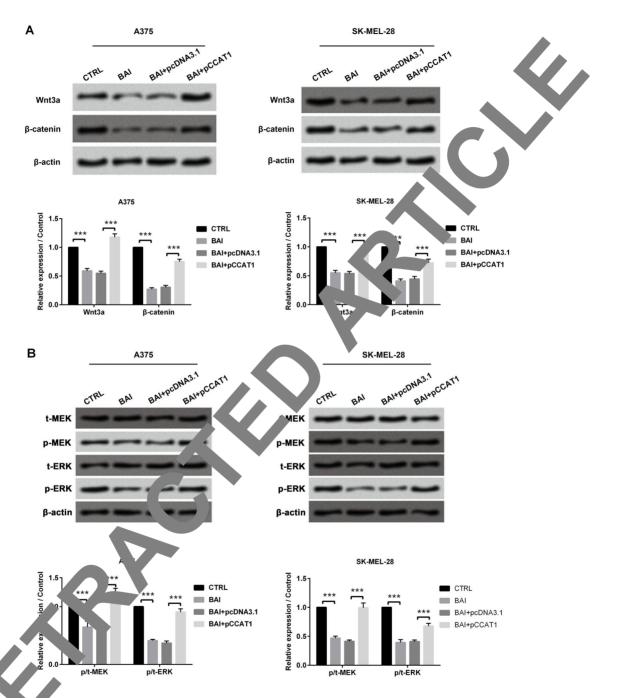


Fig. 3. Ba. (BAI) suppressed CCAT1 to block Wnt/β-catenin and MEK/ERK signaling pathway. A, The expression of Wnt3a and cate in and, is, of p-MEK, p-ERK, t-MEK, and t-ERK were examined by western blot. The relative expression of proteins was β-actin. Data are reported as mean ± SD. ***P < 0.001 (ANOVA). CTRL: control.

exogenous CCAT1 (P<0.001, Figure 6A). Similarly, the protein levels of p-MEK and p-ERK were also suppressed by BAI (P<0.001, Figure 6B). The results indicated that BAI blocked Wnt/ β -catenin or MEK/ERK pathways by negatively regulating CCAT1.

Discussion

Numerous active components extracted from traditional Chinese medicinal plants exert multiple pharmacological effects (25). Among these, perhaps the most

unexpected finding was that BAI induced growth of HeLa cells via mitochondrial and death receptor pathways (9). Although it has been reported that BAI could act as an essential anti-tumor modulator, leading to ameliorated biological processes, such as programmed cell death and angiogenesis in the B16F10 cells (26), the underlying molecular mechanisms remained to be fully demonstrated. Our study found that there were intricate regulating effects between BAI and the progression of malignant melanoma.

BAI is a vigorous herbal medicine that exerts indispensable functions towards the cardiovascular system and hepatoma (27,28). The function of BAI mainly displays as two aspects: anti-oxidative and inhibitory action on cell growth. Chou et al. (11) showed that BAI caused a reduction in cellular viability of melanoma cells through generating ROS scavengers. Existing research recognized that BAI inhibited tumor growth via activation of cleaved-caspase-3 (26). The results of our study were in line with the above previous experiments. We found that BAI alleviated cell growth, and migratory and invasive ability in malignant melanoma. Our findings indicated that BAI exerted indispensable functions as tumor suppressor.

It was reported that abnormal expression of IncRNAs might be related to a wide spectrum of tumor biological processes (29). Reports such as that conducted by V et al. (30) show that overexpression of CCAT1 significantly elicit cell proliferation and invasion and inhibit cell concluder cell renal cell carcinoma. Beyond that, V et al. (a verified that CCAT1 served as an oncoming factor in melanoma genesis, accumulating cell proliferation, aigration, and invasion abilities. However, the is a relative paucity of literature concerning CC T1 invariement in regulating the effects of BAI on in Janoma biological

processes. In this study, we measured CCAT1 expression level and found that CCAT1 was up-regulated imelanoma. We showed for the first time that BAI in bit officell proliferation, migration, and invasion of maligable networks and regulating CCAT1.

Wnt3a is a key activator of Wnt pathway, generally triggering the acknowledged Wnt/β-conin simaling pathway (32) and is related to ersing processes, such as cell growth and micration (33.34). The MAPK/ERK signaling path v regules cell proliferation and differentiation in any acroslls (35,36) and is associated with melan' syn sis (37). Debates have been raised about the 'naling p' way involved in the progression of malignant in 'anoma. Results from earlier studies demonstrate that B. inhibited melanogenesis through activation of the ERK signaling pathway but did not induce AK time 2). Recent investigators found that BAI impede he migratory and invasive potential of B16F alls through the suppression of PI3K /AKT 8). The present experiments uncovsignaling 1-regulated protein expression of Wnt3a, ered the MEK, and p-ERK in malignant melanoma β-catenin, p treated with BAI. The restraint was reversed by exog ous expressed CCAT1. Therefore, we speculated that F I blocked Wnt/β-catenin or MEK/ERK signaling ays by regulating CCAT1.

Overall, our study indicated that BAI hindered Wnt/ β -utenin and MEK/ERK signaling pathways by regulating CCAT1, thereby inhibiting proliferation, migration, and invasion of melanoma cells. The present study demonstrated a pivotal role of BAI in tumor regulation, which might provide new light on the development of therapeutic strategies against malignant melanoma. Comprehensive *in vivo* experiments are crucial for future research.

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