

Original Research

80MAP17 promotes the tumorigenesis of papillary thyroid carcinoma by reducing the stability of p53

Kun Yu^{1,†}, Hongjiang Lu^{2,†}, Yanhong Chen³, Ying Xin¹, Zhuo Tan¹, Qiong Yang^{4,*}

¹Department of Head and Neck Surgery, Center of Otolaryngology-Head and Neck Surgery, Zhejiang Provincial People's Hospital, People's Hospital of Hangzhou Medical College, Key Laboratory of Endocrine Gland Diseases of Zhejiang Province, 310014 Hangzhou, Zhejiang, China, ²Department of Radiology, The 903 Hospital of the joint logistics support force of the Chinese people's Liberation Army, 310014 Hangzhou, Zhejiang, China, ³Laboratory Animal Center, Zhejiang University, 310058 Hangzhou, Zhejiang, China, ⁴Department of Breasts Surgery, Center of Otolaryngology-Head and Neck Surgery, Zhejiang Provincial People's Hospital, People's Hospital of Hangzhou Medical College, Key Laboratory of Endocrine Gland Diseases of Zhejiang Province, 310014 Hangzhou, Zhejiang, China

TABLE OF CONTENTS

1. Abstract
2. Introduction
3. Material and methods
 - 3.1 UALCAN analysis
 - 3.2 Gene expression profiling interactive analysis (GEPIA) database
 - 3.3 Collection of tissue samples and immunohistochemistry
 - 3.4 Cell culture
 - 3.5 Western blotting
 - 3.6 MAP17 overexpression and knockdown
 - 3.7 CCK8 assay
 - 3.8 Clone formation assay
 - 3.9 Terminal deoxynucleotidyl transferase deoxyuridine triphosphate (dUTP) nick end labeling (TUNEL) staining
 - 3.10 Wound healing assay
 - 3.11 Transwell assay
 - 3.12 Immunofluorescence staining
 - 3.13 Animal modeling and randomization
 - 3.14 Immunohistochemistry
 - 3.15 Statistical analysis
4. Results
 - 4.1 The expression of MAP17 was increased in patients with PTC
 - 4.2 MAP17 promoted the growth of PTC cells and inhibited cell apoptosis in vitro
 - 4.3 MAP17 promoted the migration, invasion and epithelial-mesenchymal metastasis (EMT) of PTC cells
 - 4.4 MAP17 reversed NUMB-mediated p53 stabilization
 - 4.5 MAP17 promoted the growth and metastasis of PTC cells by p53
 - 4.6 Knockdown of MAP17 inhibited tumor growth in vivo
5. Discussion
6. Conclusions
7. Author contributions
8. Ethics approval and consent to participate
9. Acknowledgment
10. Funding
11. Conflict of interest
12. References

1. Abstract

Background: Papillary thyroid cancer (PTC) is an endocrine malignancy whose incidence has increased rapidly worldwide. MAP17 (PDZKIP1) is a small protein related to tumor progression. The aim of this study was to investigate the role of MAP17 in PTC and the underlying molecular mechanism. **Methods:** Bioinformatics, Western blotting and immunohistochemistry were used to analyze the expression of MAP17 in PTC. The gene transcription was measured by qPCR. Cell viability was determined by CCK8 assay. Cell growth was measured by clonal formation assay. Cell apoptosis was measured by TUNEL. Wound healing assay and transwell assay were used to measure the mobility of cells. The expression of E-cadherin and N-cadherin was determined by immunofluorescence. The effect of MAP17 on tumor growth was determined in animal experiments. **Results:** The results showed that MAP17 was up-regulated in PTC, which significantly promoted the growth and motility of PTC cells, but inhibited cell apoptosis. Besides, overexpression of MAP17 accelerated cycloheximide (CHX, a protein synthesis inhibitor)-induced p53 degradation, while low expression of MAP17 slowed down CHX-induced p53 degradation, suggesting that MAP17 can regulate p53 stability. Notably, NUMB exhibited an opposite effect on P53 stability. Interestingly, p53 overexpression reversed the effects of MAP17 overexpression on cell viability, motility, and apoptosis, indicating that p53 was involved in the progression of PTC. *In vivo* studies have shown that tumor growth was positively correlated with MAP17 expression and negatively correlated with p53 expression. **Conclusion:** Our findings revealed that MAP17 exhibited carcinogenic effects through interacting with NUMB to reduce the stability of p53, demonstrating that MAP17 may serve as a potential prognostic biomarker for PTC treatment.

2. Introduction

Papillary thyroid cancer (PTC) is an endocrine malignancy that occurs in thyroid follicular cells [1]. The incidence of PTC has increased rapidly worldwide over the last 3 decades. In the United States, the incidence of PTC increased 3.7-fold from 3.4 per 100000 to 12.5 per 100000 [2]. Most PTC patients have a good prognosis with a low mortality rate, but cervical lymph node metastasis is relatively common [3, 4]. Although great progress has been made for detection and treatment of PTC, the diagnostic accuracy still needs to be improved [5]. Therefore, it is urgently needed to develop alternative diagnostic therapeutic markers.

MAP17 (PDZKIP1) is an insoluble and non-glycosylated membrane-associated protein [6], which contains two transmembrane regions and a hydrophobic c-terminal encoding a postsynaptic density-95 (PSD-95),

discs-large, zona occludens 1 (ZO-1) (PDZ) binding domain. MAP17 can interact with multiple proteins containing PDZ domains (Such as PDZK1) [7, 8]. MAP17 is overexpressed in various human cancers [9, 10], which is closely related to tumor progression [8, 9, 11], while MAP17 is rarely expressed in adenomas, benign tumors, and normal tissues, and a high proportion (50%–90%) of advanced or metastatic tumors always express high levels of MAP17, which is associated with a more dedifferentiated phenotype [8–10]. DNA methylation arrays showed that overexpression of MAP17 related to abnormal promoter hypomethylation may promote tumor growth in thyroid cancer. However, the specific role and underlying molecular mechanism of MAP17 in thyroid cancer are still unclear.

P53 is one of the most important tumor suppressors in cells, which has been found to occur deletion or mutation in more than 50% of human tumor cells. A large number of studies have shown that p53 also plays an important role in cell metabolism, especially glucose metabolism. In thyroid cancer, mutations of p53 occurred in anaplastic carcinoma (AC), which is a rare but highly aggressive form of thyroid cancer [12]. NUMB can increase the stability of p53 and reduce p53 ubiquitination degradation [13–15]. Interestingly, MAP17 can interact with NUMB to inhibit its function [16]. However, whether MAP17 can regulate PTC tumor progression through interacting with NUMB/p53 remains to be investigated.

This study investigated the expression of MAP17 in PTC and the influence of abnormal expression of MAP17 on the progression of PTC. Our results showed that MAP17 was highly expressed in PTC, which was closely related to cell growth, motility and apoptosis. *In vivo* studies have confirmed that MAP17 overexpression significantly promotes tumor growth. Our findings suggest that MAP17 may serve as a potential target for PTC treatment.

3. Material and methods

3.1 UALCAN analysis

A tool named UALCAN (<http://ualcan.path.uab.edu/index.html>) is generally used for TCGA data mining analysis to retrieve the expression of multiple genes in tumors as well as the relationship of the expression with prognosis [17]. In this study, the transcriptional expression of MAP17 (also known as PDZK1IP1) in PTC and normal thyroid tissues were analyzed using UALCAN, with a cut-off *p* value of 0.01.

3.2 Gene expression profiling interactive analysis (GEPIA) database

GEPIA (<http://gepia.cancer-pku.cn/index.html>), developed by Peking University, is an interactive analysis website containing abundant RNA sequencing expression data from TCGA and GTEx. GEPIA has various specific roles, such as tumor/normal differential expression analy-

sis, and analysis based on cancer type or pathological stage [18]. In this study, MAP17 (also known as PDZK1P1) was analyzed by multi-gene comparison module. In addition, the single gene analysis module of GEPIA was used to explore the relationship between MAP17 expression and pathological stage.

3.3 Collection of tissue samples and immunohistochemistry

A total of 50 PTC tissues and para-cancerous normal tissues from patients with PTC were collected during surgery procured from People's Hospital, People's Hospital of Hangzhou Medical College. The expression of MAP17 in PTC tissues and para-cancerous normal tissue was assessed by immunohistochemistry. In brief, the tissues were embedded in paraffin, and incubated with anti-MAP17 antibody (ab156014, 1:250, Abcam, Cambridge, UK). MAP17 staining was observed under a microscope. All analyses were conducted in triplicate.

3.4 Cell culture

PTC cell lines (HTh-7, KHM-5M, CAL62, KTC-1 and TPC-1) and normal human normal thyroid HTori-3 cell line were purchased from the Cell Bank of the Shanghai Chinese Academy of Sciences (Shanghai, China). Cells were cultured in high-glucose DMEM (Hyclone; Thermo Fisher Scientific, Waltham, MA, USA) containing 10% FBS and 1% antibiotic-antimycotic in an incubator at 37 °C with 5% CO₂.

3.5 Western blotting

The levels of MAP17, p53, β -actin, and NUMB were assessed by western blotting. Briefly, cells or tissues were lysed, and then protein concentration was determined by BCA assay. 20 μ g of each sample was separated by SDS-PAGE, and then transferred to PVDF membranes. The membranes were then probed with primary antibodies MAP17 (ab156014, 1:1000, Abcam, Cambridge, UK), p53 (ab26, 1–5 μ g/mL, Abcam, Cambridge, UK), β -actin (#4970, 1:1000, Cell Signaling Technology, Danvers, MA, USA), and NUMB (#2756, 1:1000, Cell Signaling Technology, Danvers, MA, USA) at 4 °C overnight, and then incubated with the appropriate secondary antibodies at room temperature for 1 h. After exposing to DAB, the membranes were visualized. All analyses were conducted in triplicate.

3.6 MAP17 overexpression and knockdown

For overexpression, MAP17 was overexpressed using adenovirus system. Briefly, constructs (MAP17-overexpressing (MAP17) and negative control (NC)) were inserted into an adenovirus-packaging plasmid. After that, the above adenovirus were packaged in HEK293A cells [19]. CAL62 and TPC-1 cells were seeded into 24-well plates, achieving to a cell density of about 2×10^5 , and then 5×10^{10} PFU/mL adenovirus (2 μ L) was added into

the 24-well plate, and the fresh complete culture medium was replaced after 4–8 h. WB analysis was performed 48 h later. For knockdown, MAP17 was knocked down with shRNAs. In brief, shRNAs specific MAP17 (shMAP17#1 and shMAP17#2) and negative control shRNA (shNC) were designed and synthesized by Bersinbio (Guangzhou, China). Then, shRNAs were transfected into CAL62 and TPC-1 cells using Lipofectamine 2000. After 48 h post transfection, follow-up analysis of the cells was performed.

3.7 CCK8 assay

Cell viability or the effects of MAP17 on proliferation were determined using CCK8 assay. CAL62 and TPC-1 cells at a density of about 4×10^3 cells/well were seeded on the 96-well plates, and cultured overnight at 37 °C with 5% CO₂. Whereafter, 10 μ L CCK8 was added, and the cells were incubated for another 24 h. Optical density (OD) for each well was determined at 450 nm using microplate reader (Infinite® 200 PRO, Tecan).

3.8 Clone formation assay

After transfection, CAL62 and TPC-1 cells (400 cells/well) in the logarithmic phase were seeded in 6-well plates and incubated at 37 °C with 5% CO₂ for 14 days. Medium in each plate was refreshed every 3 days. After that, the cells were fixed with 4% paraformaldehyde (1 mL), and stained with 0.1% crystal violet. The colonies were photographed under a microscope and counted.

3.9 Terminal deoxynucleotidyl transferase deoxyuridine triphosphate (dUTP) nick end labeling (TUNEL) staining

Apoptosis was determined by TUNEL staining using *In Situ* Cell Death Detection Kit (Roche). CAL62 and TPC-1 cells were inoculated into 24-well plates at a density of about 2×10^4 cells/well, and then incubated. After infection or transfection, the cells were fixed with 4% paraformaldehyde solution (100 μ L), and then exposed to TUNEL reaction mixture (50 μ L) in dark, followed by incubation of 1 h. After that, fluorescence labeled DNA were visualized using fluorescence microscopy (Leica, Wetzlar, Germany).

3.10 Wound healing assay

CAL62 and TPC-1 cells were plated in 24-well plates and cultured until monolayers formed. The monolayers were subjected to mitomycin C at 75 μ g/mL to suppress cell self-replication. The monolayer was scratched using a 200 μ L sterile pipette tip, and subsequently washed twice with warm PBS. After the cells were cultured at 37 °C for 48 h and 0 h post injury, the scratches were visualized using Nikon ECLIPSE TE2000-U inverted microscope.

3.11 Transwell assay

Invasion was determined by transwell assay. CAL62 and TPC-1 cells were serum-starved for 24 h, and then inoculated into upper chambers coated with Matrigel

(BD Biosciences, Bedford, MA, USA) at a density of 2×10^4 . The lower chambers were filled with 500 μ L of 1640 RPMI containing 10% FBS, followed by incubation at 37 °C for 24 h. Then, the cells on the dorsal side of the lower chamber were fixed with 100% methanol and stained with 0.1% crystal violet. The crystal was dissolved in DMSO, and the OD value at 570 nm was measured with a microplate analyzer.

3.12 Immunofluorescence staining

The levels of E-cadherin and N-cadherin were detected by immunofluorescence staining. CAL62 and TPC-1 cells grown in 96 well plates were washed with PBS, fixed with 4% paraformaldehyde, and sealed with 5% BSA/PBS for 30 min. The cells were then incubated with primary antibodies of E-cadherin (1:200, #3195, Cell Signaling Technology, Danvers, MA, USA) and N-cadherin (1:200, #13116, Cell Signaling Technology, Danvers, MA, USA). Thereafter, the cells were stained with fluorescein isothiocyanate or Cy3 linked secondary antibody, and imaged by cellomics ArrayScan HCS reader (Thermo Science, Waltham, MA, USA).

3.13 Animal modeling and randomization

Female BALB/c nude SPF mice (6 weeks of age, 18–22 g in weight) were provided by Vital River (Beijing, China). Animals were kept at room temperature of 18–25 °C, with a 12-hour light/dark cycle and free access to food and water. TPC-1 cells were cultured *in vitro* and infected with adenovirus overexpressing MAP17 or transfected with shNC or shRNA#2, respectively. Then, infected or transfected TPC-1 cells (3.5×10^6 cells per mouse) were subcutaneously injected into female BALB/c nude mice. Tumor bearing mice were randomly divided into four groups (10 mice per group): NC group, MAP17 group, shNC group and shRNA#2 group. Tumor size was measured once every 3 days. Thirty days after injection, the mice were executed by cervical dislocation. The tumor was removed, and the weight was measured, and immunohistochemical analysis was performed.

3.14 Immunohistochemistry

The expression of Ki67, MAP17 and p53 in tumor tissues was determined using immunohistochemical staining as previously described [20]. In brief, tissue sections were dewaxed with xylene and hydrated with different concentration gradients of ethanol. The sections were then immersed in a pH 6.0 sodium citrate buffer and boiled for 30 min. Sections were incubated at 37 °C with anti-Ki67 (#9449, 1:200, Cell Signaling Technology, Danvers, MA, USA), anti-MAP17 (ab156014, 1:200, Abcam, Cambridge, UK) and anti-p53 (ab32389, 1:100, Abcam, Cambridge, UK), followed by observation under light microscope (Olympus Corporation, Tokyo, Japan).

3.15 Statistical analysis

Data are presented as means \pm SD and analyzed with SPSS (version 19.0, Chicago, IL, USA). Comparisons were carried out using Student's *t* test (two groups) and analysis of variance (three groups or above). $p < 0.05$ was deemed as statistically significant difference.

4. Results

4.1 The expression of MAP17 was increased in patients with PTC

We first analyzed the expression of MAP17 in PTC using GEPIA database and UALCAN database. The expression of MAP17 was increased in PTC (Fig. 1A,B), and the level of MAP17 increased over with lymph node metastasis (Fig. 1C). Besides, the expression of MAP17 in tall papillary thyroid carcinoma was the highest (Fig. 1D). In addition, tumor tissue samples and para-cancerous tissue samples from patients with PTC were collected during the surgery, and MAP17 expression was detected by PCR, immunohistochemistry and Western blotting. MAP17 expression was increased at both mRNA (Fig. 1E) and protein levels in tumor tissues compared with adjacent tissues (Fig. 1F–G). Overall, these evidences consistently indicated that the expression of MAP17 was increased in patients with PTC.

4.2 MAP17 promoted the growth of PTC cells and inhibited cell apoptosis *in vitro*

We investigated the expression and role of MAP17 in PTC *in vitro*. The expression of MAP17 in Htori-3 cells and PTC cell lines (HTH-7, CAL62, KHM-5M, TPC-1 and KTC-1) was determined by Western blotting. As shown in Fig. 2A, MAP17 expression in PTC cell lines was significantly increased compared with that in Htori-3 cells, and CAL62 and TPC-1 cells were selected for further study. To further explore the effect of MAP17 on PTC cells, MAP17 was overexpressed or knocked down in CAL62 and TPC-1 cells, respectively. As shown in Fig. 2B, the levels of MAP17 in CAL62 and TPC-1 cells were significantly increased after the cells were infected with adenovirus overexpressing MAP17, while decreased after the cells were transfected with shRNA (shRNA#1 and shRNA#2), indicating that MAP17 had been successfully overexpressed or knocked down in this study. In addition, the effects of abnormal MAP17 expression on the growth and apoptosis of CAL62 and TPC-1 cells were determined by clone formation assay and TUNEL staining, respectively. As shown in Fig. 2, the overexpression of MAP17 significantly increased cell viability of PTC cells (Fig. 2C), and promoted the growth of PTC cells (Fig. 2D), as well as inhibited the apoptosis of PTC cells (Fig. 2E), while the low expression of MAP17 showed opposite effects. These results indicated that MAP17 promoted the growth of PTC cells and inhibit cell apoptosis *in vitro*.

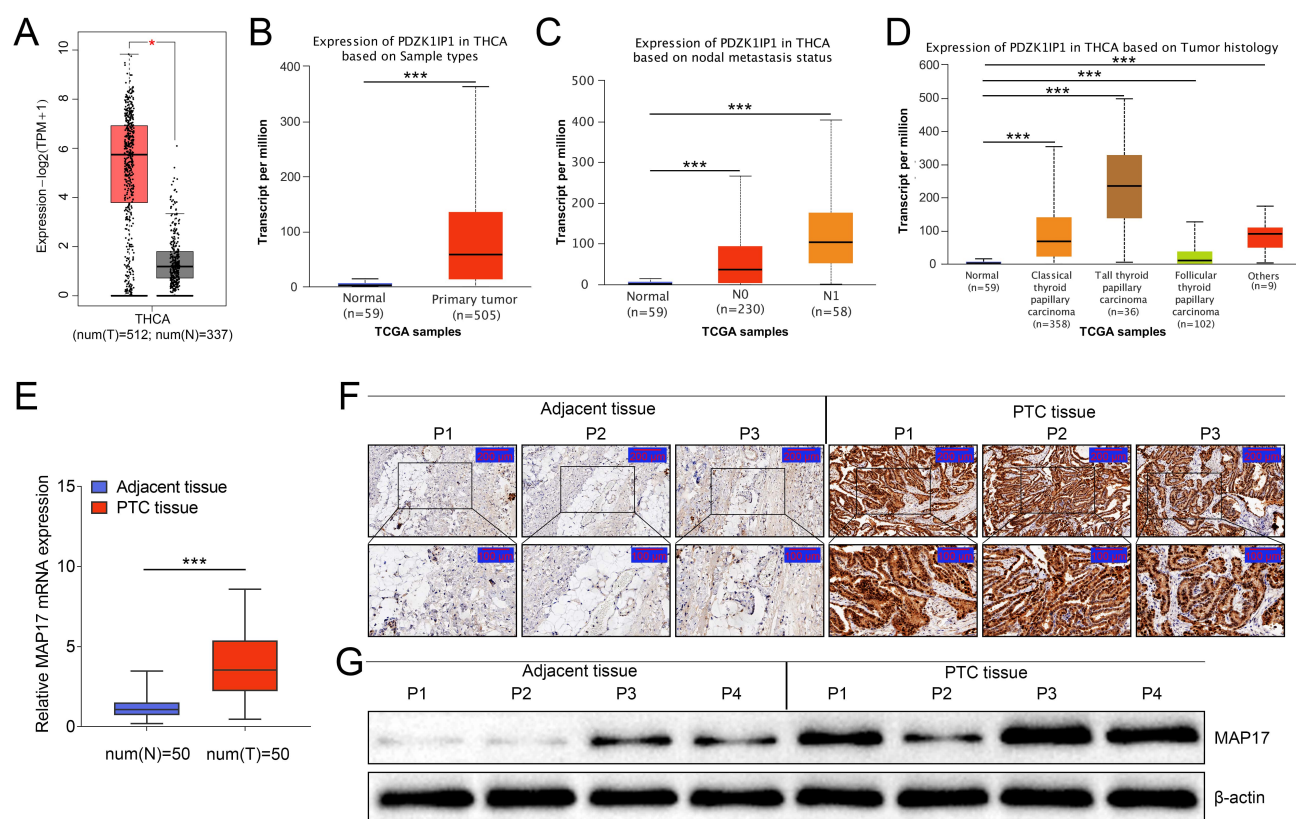


Fig. 1. The expression of MAP17 was increased in patients with PTC. (A–D) MAP17 expression was analyzed in GEPIA database and UALCAN database (* $p < 0.05$, *** $p < 0.001$ vs. Normal group). (E) The expression of MAP17 in PTC tissues and adjacent tissues was detected by RT-qPCR. (F) The expression of MAP17 in PTC tissues and adjacent tissues was detected by immunohistochemistry. (G) The expression of MAP17 in PTC tissues and adjacent tissues was detected by western blotting. (*** $p < 0.001$ vs. Normal group).

4.3 MAP17 promoted the migration, invasion and epithelial-mesenchymal metastasis (EMT) of PTC cells

In addition, we further investigated the effect of abnormal MAP17 expression on the motility of CAL62 and TPC-1 cells. As shown in Fig. 3A,B, the overexpression of MAP17 significantly promoted the invasion and migration of PTC cells, while the low expression of MAP17 inhibited the invasion and migration of PTC cells. Furthermore, immunofluorescence analysis showed that MAP17 overexpression significantly increased the level of N-cadherin, but decreased the level of E-cadherin. The low expression of MAP17 showed an opposite effect, indicating that MAP17 promoted the process of EMT (Fig. 3C). In summary, these results showed that MAP17 promoted the motility of PTC cells.

4.4 MAP17 reversed NUMB-mediated p53 stabilization

Studies have shown that NUMB could inhibit p53 ubiquitination, and enhance the stability of p53, and MAP17 could physically interact with NUMB. Therefore, this study investigated the effect of MAP17 on NUMB mediated p53 stabilization. As shown in the Fig. 4A–B, RT-qPCR showed that the abnormal expression of MAP17 had

no significant effect on the transcription of p53, but significantly decreased the p53 expression in CAL62 and TPC-1 cells, while MAP17 knockdown significantly increased the protein level of p53 (Fig. 4A,B). After MAP17 was overexpressed or knocked down in TPC-1 cells, TPC-1 cells were subjected to the protein synthesis inhibitor cycloheximide (CHX) for 0, 0.5, 1 and 2 h, respectively. Western blotting showed that in the presence of CHX, the overexpression of MAP17 accelerated the degradation of p53, while the knockdown of MAP17 slowed the degradation of p53 (Fig. 4C). Furthermore, CAL62 or TPC-1 cells were infected with adenovirus overexpressing MAP17 or NUMB, either alone or in combination. The results showed that MAP17 overexpression significantly promoted p53 degradation in CAL62 or TPC-1 cells, while NUMB overexpression significantly inhibited p53 degradation in CAL62 or TPC-1 cells (Fig. 4D). Further study showed that p53 degradation was significantly slowed in CAL62 or TPC-1 cells after co-infection with adenovirus overexpressing MAP17 and NUMB. Notably, NUMB did not affect the expression of MAP17 (Fig. 4E). Taken together, these results suggested that MAP17 reversed NUMB-mediated p53 stabilization.

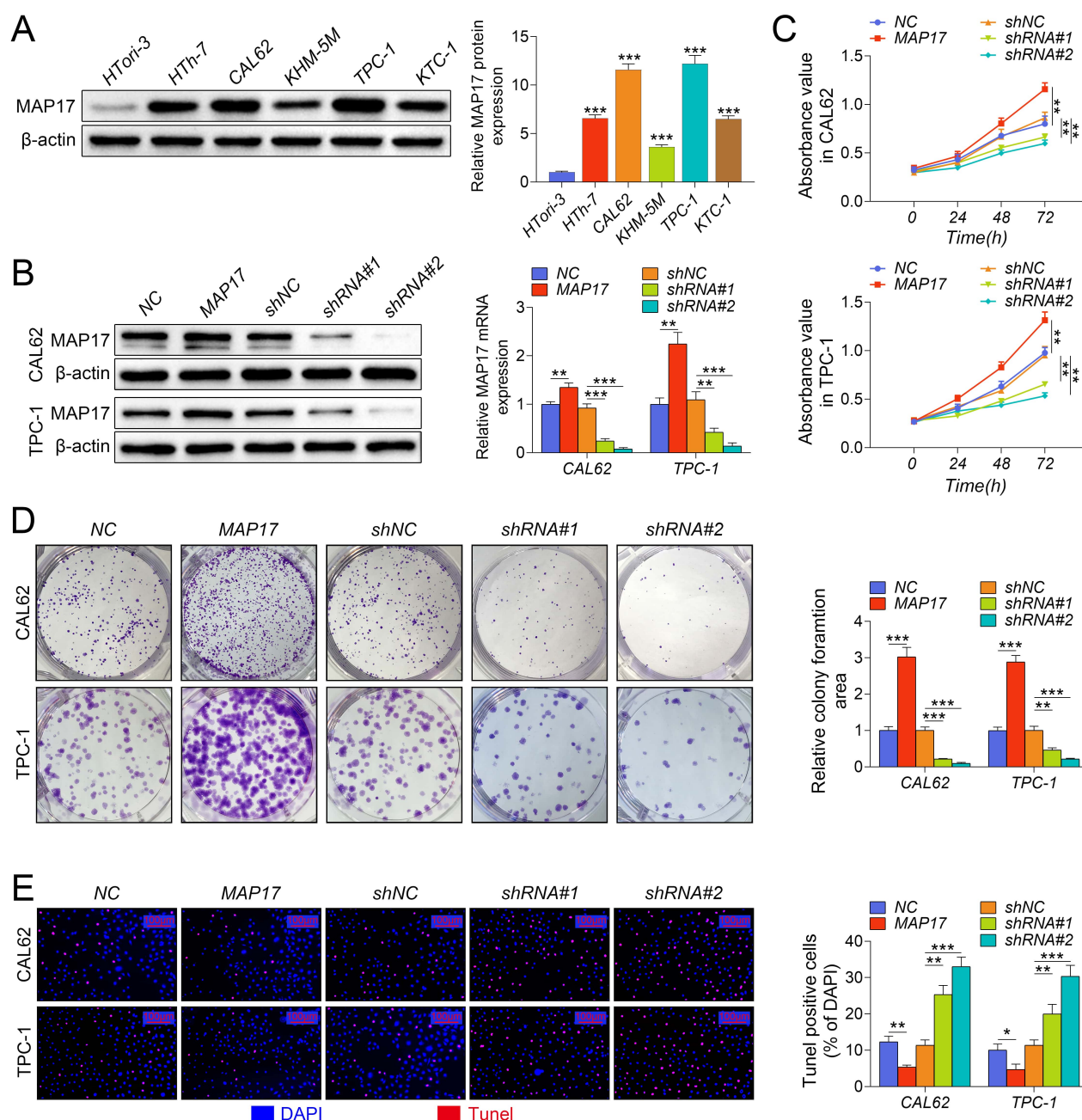


Fig. 2. MAP17 promoted the growth of PTC cells and inhibited cell apoptosis *in vitro*. (A) The expression of MAP17 in Htori-3 cells and PTC cell lines (HTH-7, CAL62, KHM-5M, TPC-1 and KTC-1) was detected by western blotting (** $p < 0.001$ vs. Htori-3 cells). (B) CAL62 and TPC-1 cells were infected with adenovirus containing MAP17, or transfected with shRNAs (shRNA#1 or shRNA#2). (C) Cell viability was measured by CCK 8 assay. (D) Cell growth was measured by clone formation assay. (E) Cell apoptosis was measured by TUNEL staining (** $p < 0.01$, *** $p < 0.001$ vs. NC group or shNC group) (* indicates $p < 0.05$, i.e., there was statistically significant difference).

4.5 MAP17 promoted the growth and metastasis of PTC cells by p53

Previous results in this study showed that MAP17 accelerated p53 degradation, thereby reducing p53 levels in cells. Therefore, to investigate whether p53 is involved in the progression of the tumor in this study, TPC-1 cells were infected with adenovirus containing MAP17 or p53 to overexpress MAP17 or p53, respectively. Cell viabil-

ity was examined by CCK8 assay, while cell growth was measured by clone formation assay. As shown in Fig. 5A and B, MAP17-induced low expression of p53, which significantly improved the viability of TPC-1 cells, and promoted cell growth, but p53 overexpression exhibited the opposite effect. However, the co-transfection of MAP17 and p53 significantly reversed the promotional effect of MAP17 overexpression on the growth of TPC-1 cells. Similarly,

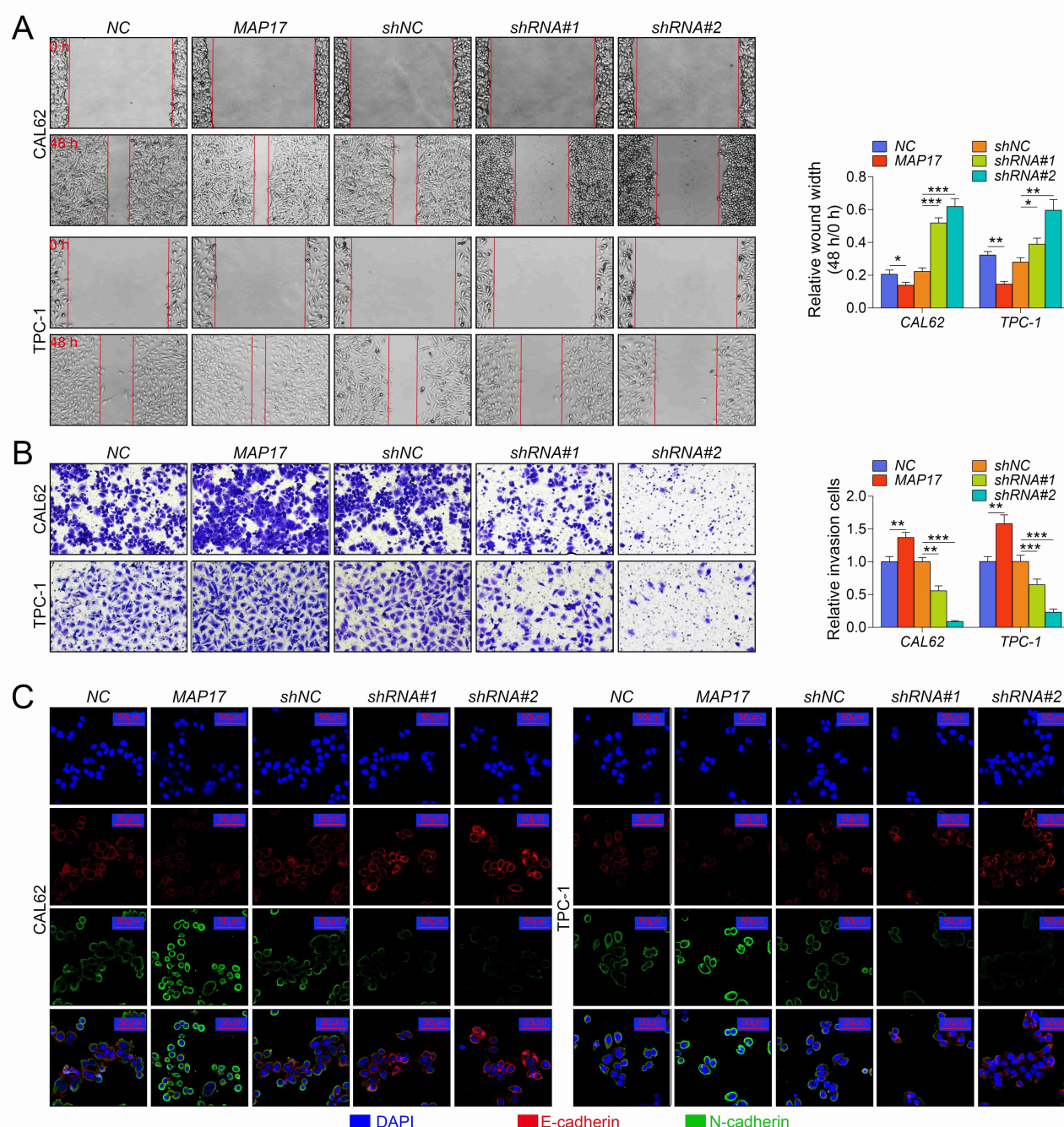


Fig. 3. MAP17 promoted the migration, invasion and epithelial-mesenchymal metastasis of PTC cells. CAL62 and TPC-1 cells were infected with adenovirus containing MAP17, or transfected with shRNAs (shRNA#1 or shRNA#2). (A) Cell migration was measured by wound healing assay. (* $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ vs. NC group or shNC group). (B) Cell invasion was measured by transwell assay. (C) The levels of E-cadherin and N-cadherin were measured by immunofluorescence assay. (** $p < 0.01$, *** $p < 0.001$ vs. NC group or shNC group).

MAP17-induced low expression of p53 significantly inhibited apoptosis in TPC-1 cells, while p53 overexpression showed the opposite effect. However, the co-transfection of MAP17 and p53 significantly reversed the inhibitory effect of MAP17 overexpression on the apoptosis of TPC-1 cells (Fig. 5C). Furthermore, MAP17-induced low expression of p53 significantly promoted the mobility of TPC-1 cells, while p53 overexpression showed the opposite effect.

However, the co-transfection of MAP17 and p53 significantly reversed the promotional effect of MAP17 overexpression on the mobility of TPC-1 cells (Fig. 5D–E). Taken together, these results suggested that MAP17 promoted the growth and metastasis of PTC cells through degrading p53.

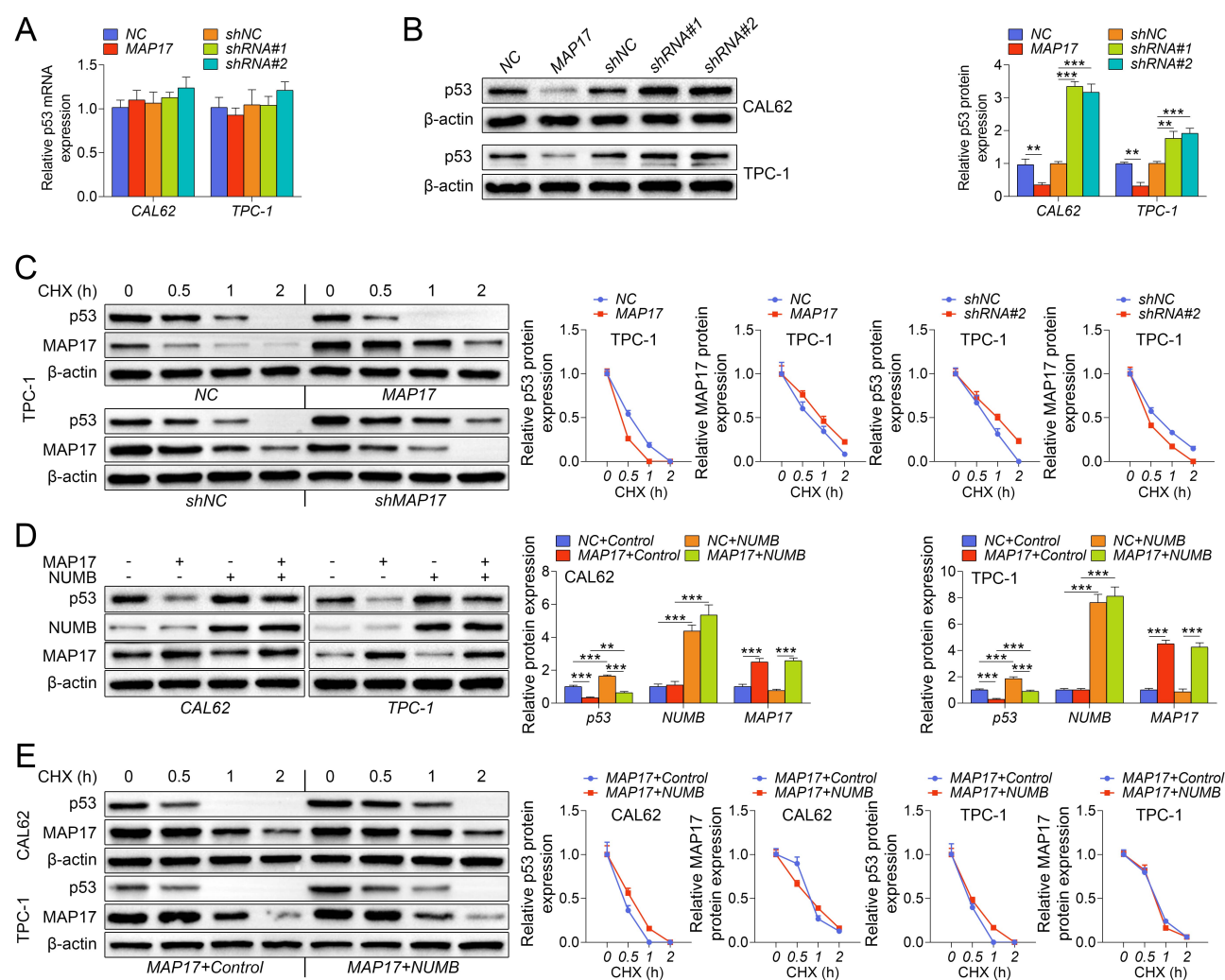


Fig. 4. MAP17 reversed NUMB-mediated p53 stabilization. CAL62 and TPC-1 cells were infected with adenovirus containing MAP17, or transfected with shRNAs (shRNA#1 or shRNA#2). (A) The expression of p53 in cells was detected by RT-qPCR. (B) The expression of p53 in cells was detected by western blotting. (C) TPC-1 cells were infected with adenovirus containing MAP17, or transfected with shRNA#2, and then subjected to the protein synthesis inhibitor cycloheximide (CHX) for 0, 0.5, 1 and 2 h, respectively. The expression of p53 and MAP17 in cells was detected by western blotting. (D) CAL62 or TPC-1 cells were infected with adenovirus overexpressing MAP17 or NUMB, either alone or in combination. The expression of p53, NUMB and MAP17 in cells was detected by western blotting (** $p < 0.01$, *** $p < 0.001$ vs. NC + Control group or NC + NUMB group). (E) CAL62 or TPC-1 cells were infected with adenovirus overexpressing alone or in combination with NUMB, and then subjected to CHX for 0, 0.5, 1 and 2 h, respectively. The expression of p53 and MAP17 in cells was detected by western blotting.

4.6 Knockdown of MAP17 inhibited tumor growth in vivo

TPC-1 cells over-expressing and low-expressing MAP17 were injected subcutaneously into BALB/c nude mice, and tumor volume and weight were measured. As shown in Fig. 6A–C, MAP17 overexpression significantly promoted tumor growth, while MAP17 knockdown significantly inhibited tumor growth. Besides, immunohistochemistry showed that MAP17 overexpression significantly promoted the expression of Ki67, but inhibited the expression of p53 in tumor tissues. On the contrary, MAP17 knockdown got an opposite result (Fig. 6D). Taken together, these results suggested that knockdown of MAP17 inhibited tumor growth *in vivo*.

5. Discussion

In recent years, the incidence of PTC has increased globally, but the mortality rate is decreasing [21, 22]. Therefore, novel molecular markers are essential for risk stratification and personalized management of PTC [23]. Previous studies have revealed multiple potential PTC markers, such as miRNAs (miR-136, miR-21, and miR-127) [19], lncRNA SLC26A4-AS1 [24], and Lactate Dehydrogenase A [25]. Fortunately, in this study, we identified a novel potential molecular marker for PTC treatment, i.e., MAP17. Bioinformatics analysis demonstrated that MAP17 was highly expressed in PTC, which was further confirmed by *in vitro* and *in vivo* studies. Specifically, the

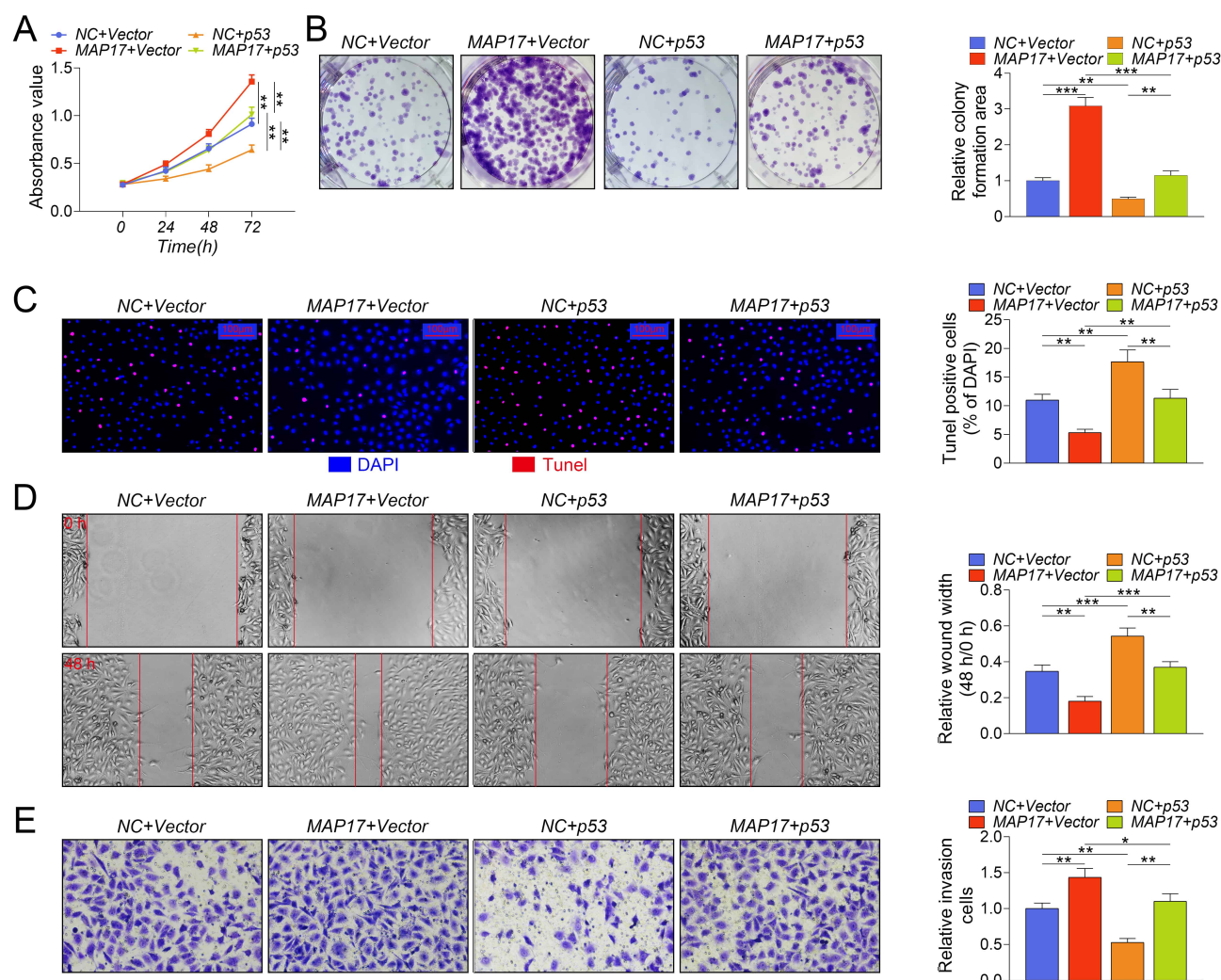


Fig. 5. MAP17 promoted the growth and metastasis of PTC cells by p53. TPC-1 cells were infected with adenovirus containing MAP17 or p53 to overexpress MAP17 or p53, respectively. (A) Cell viability was examined by CCK8 assay. (B) Cell growth was measured by clone formation assay. (C) Cell apoptosis was measured by TUNEL staining. (D) Cell migration was measured by wound healing assay. (E) Cell invasion was measured by transwell assay (* $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ vs. NC + Vector group or MAP17 + Vector or NC + p53 group).

high expression of MAP17 was associated with excessive proliferation and reduced apoptosis in PTC cells, as well as tumor growth. Consistently, the carcinogenic effects of MAP17 have also been demonstrated in non-small cell lung cancer and breast cancer [26, 27]. Collectively, it suggested that MAP17 may serve as a potential prognostic marker for PTC.

MAP17 is a small protein related to tumor progression that regulates inflammation and tumor progression [28]. Studies have shown that MAP17 has potential prognostic roles in tumors, such as lung cancer, especially in lung adenocarcinoma [28]. Besides, MAP17 overexpression enhanced stem cell characterization of cells, and promoted microsphere formation and stem cell-related gene transcription by interacting with NUNB [16]. However, MAP17 is still poorly studied in PTC. Previously, Sandra Rodríguez-Rodero *et al.* [29] found that abnormal overex-

pression of MAP17 promoter associated with hypomethylation may promote tumor growth in thyroid cancer. As supplementary, in this study, it was found that the level of MAP17 was significantly up-regulated in PTC, which was further verified by the analysis of the GEPIA database and the UALCAN database. Besides, we also found that overexpression of MAP17 significantly promoted the progression of PTC. Indeed, it was considered that the carcinogenic effect of MAP17 was achieved mainly by activating the Notch pathway [30], which changed the phenotype of cancer stem cells [16]. Inconsistently, in our study, it was found that MAP17 exhibited carcinogenic effects through interacting with NUMB to reduce the stability of p53.

MAP17 was found to be physically combined with NUMB, leading to a mislocalization of NUMB, thereby activating the Notch pathway [16]. Interestingly, NUMB was found to bind to and stabilize p53, thereby exhibiting a sup-

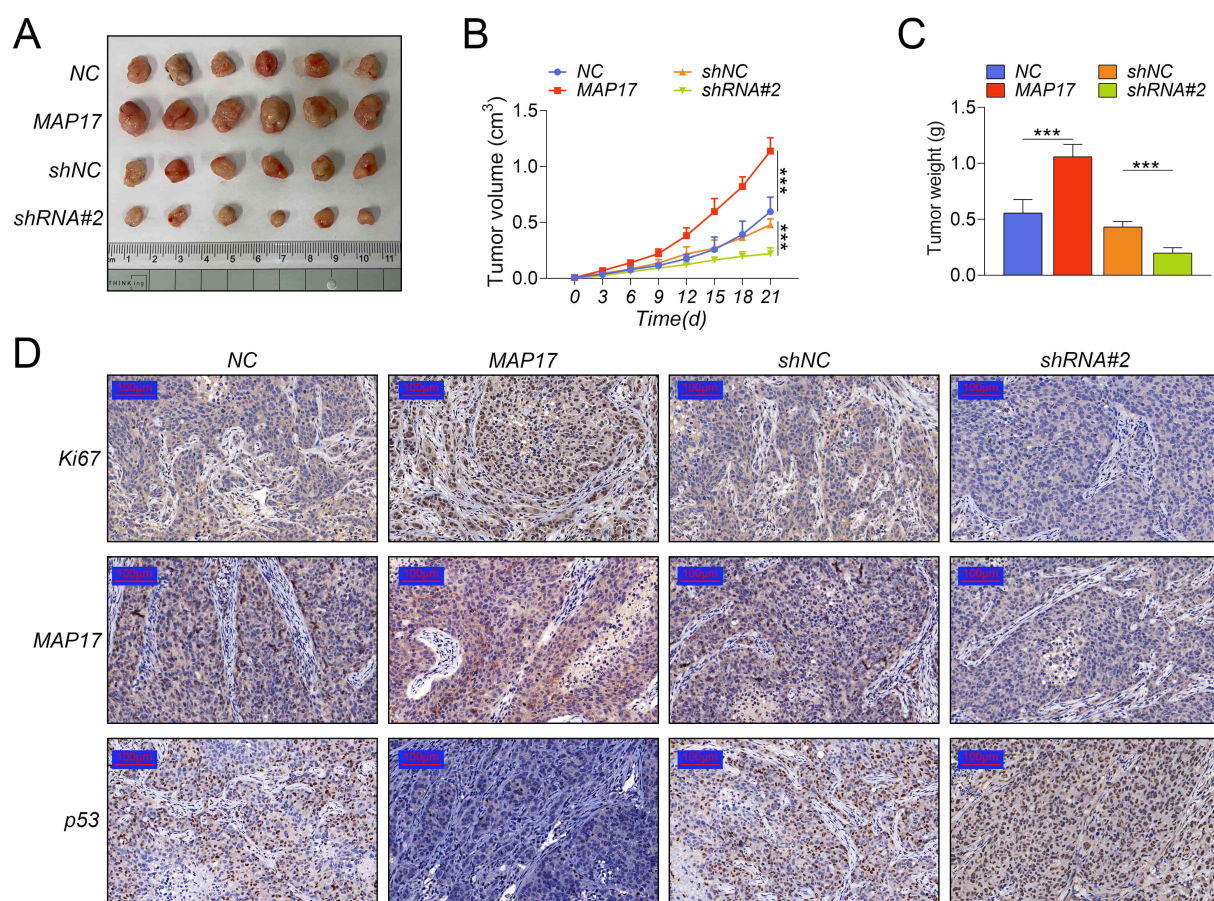


Fig. 6. Knockdown of MAP17 inhibited tumor growth *in vivo*. TPC-1 cells over-expressing and low-expressing MAP17 were injected subcutaneously into BALB/c nude mice. Tumor bearing mice were randomly divided into four groups (10 mice per group): NC group, MAP17 group, shNC group and shRNA#2 group. (A) Tumors. (B) Tumor volume. (C) Tumor weight. (D) The expression of Ki67, MAP17 and p53 were measured by immunohistochemistry (***) $p < 0.001$ vs. NC group or shNC group).

pressive effect on tumor [13]. However, methylation can uncouple NUMB from p53, thereby increasing p53 ubiquitination and degradation. In this study, we found that abnormal expression of MAP17 significantly reduced the protein level of p53, but has no effect on the transcription of *p53 in vitro*. To further clarify the mechanism of action between MAP17 and p53, cells were subjected to synthesis inhibitor CHX for various hours. After CHX treatment, the protein levels of p53 and MAP17 in TPC-1 cells were significantly reduced, indicating that CHX successfully inhibited the protein synthesis of p53 and MAP17 in TPC-1 cells. Interestingly, overexpression of MAP17 enhanced the inhibitory effect of CHX on p53 protein synthesis, while MAP17 knockdown counteracted the inhibitory effect of CHX on p53 protein synthesis. Furthermore, we found that overexpression of NUMB reversed the reduction of p53 protein levels mediated by MAP17. As expected, we found that compared with overexpression of MAP17 alone, simultaneous overexpression of MAP17 and NUMB not only maintained the stability of p53, but NUMB also bound to a part of MAP17, thereby reversed the inhibitory

effect of MAP17 on p53 synthesis. In addition, our study further confirmed that p53 is involved in the carcinogenic effects of MAP17. It was found that the reduction of p53 induced by MAP17 significantly promoted cell growth and motility, and inhibited cell apoptosis, which was significantly reversed by forced overexpression of p53, indicating that p53 is involved in the carcinogenic effects of MAP17.

6. Conclusions

In this study, the expression of MAP17 in PTC was measured *in vitro* and *in vivo*, and the effects of MAP17 on PTC were measured from *in vitro* studies and animal experiments. The results showed that MAP17 expression was increased in PTC, which is associated with poor prognosis of patients with PTC, cell growth and motility, as well as tumor growth. MAP17 knockdown inhibited the proliferation and motility of PTC cells and promoted apoptosis. Taken together, our results suggested that MAP17 exhibited carcinogenic effects by interacting with NUMB to reduce the stability of p53.

7. Author contributions

KY and HL designed the study, supervised the data collection, YC and YX analyzed the data, interpreted the data, ZT and QY prepare the manuscript for publication and reviewed the draft of the manuscript. All authors have read and approved the manuscript.

8. Ethics approval and consent to participate

The study was approved by the Medical Ethics Committee of Zhejiang Provincial People's Hospital (Approval No. 2016QT160). Animal experiment was approved by the Laboratory Animal Welfare and Ethics Committee of Zhejiang University (Approval No. ZJU20190092).

9. Acknowledgment

Not applicable.

10. Funding

This work was supported by The General Project of the Medical and Health of Zhejiang Province (Grant Number: 2021KY493).

11. Conflict of interest

The authors declare no conflict of interest.

12. References

- [1] Kwon H, Cheon Y, Shin S, Sung E, Lee J, Kim JJ, *et al.* Risk factors of suprasternal lymph node metastasis in papillary thyroid carcinoma with clinical lateral cervical lymph node metastasis. *Gland Surgery*. 2021; 10: 512–520.
- [2] Ho AS, Davies L, Nixon IJ, Palmer FL, Wang LY, Patel SG, *et al.* Increasing diagnosis of subclinical thyroid cancers leads to spurious improvements in survival rates. *Cancer*. 2015; 121: 1793–1799.
- [3] Ito Y, Kudo T, Kobayashi K, Miya A, Ichihara K, Miyauchi A. Prognostic Factors for Recurrence of Papillary Thyroid Carcinoma in the Lymph Nodes, Lung, and Bone: Analysis of 5,768 Patients with Average 10-year Follow-up. *World Journal of Surgery*. 2012; 36: 1274–1278.
- [4] Moo T, McGill J, Allendorf J, Lee J, Fahey T, Zarnegar R. Impact of prophylactic central neck lymph node dissection on early recurrence in papillary thyroid carcinoma. *World Journal of Surgery*. 2010; 34: 1187–1191.
- [5] Vriens MR, Weng J, Suh I, Huynh N, Guerrero MA, Shen WT, *et al.* MicroRNA expression profiling is a potential diagnostic tool for thyroid cancer. *Cancer*. 2012; 118: 3426–3432.
- [6] Guijarro MV, Link W, Rosado A, Leal JFM, Carnero A. MAP17 inhibits Myc-induced apoptosis through PI3K/AKT pathway activation. *Carcinogenesis*. 2007; 28: 2443–2450.
- [7] Pribanic S, Gisler SM, Bacic D, Madjdpour C, Hernando N, Sorribas V, *et al.* Interactions of MAP17 with the NaPi-IIa/PDZK1 protein complex in renal proximal tubular cells. *American Journal of Physiology. Renal Physiology*. 2003; 285: F784–F791.
- [8] Carnero A. MAP17 and the double-edged sword of ROS. *Biochimica Et Biophysica Acta*. 2012; 1826: 44–52.
- [9] Perez M, Praena-Fernandez JM, Felipe-Abrio B, Lopez-Garcia MA, Lucena-Cacace A, Garcia A, *et al.* MAP17 and SGLT1 protein expression levels as prognostic markers for cervical tumor patient survival. *PLoS ONE*. 2013; 8: e56169.
- [10] Guijarro MV, Vergel M, Marin JJ, Muñoz-Galván S, Ferrer I, Ramon y Cajal S, *et al.* P38 α limits the contribution of MAP17 to cancer progression in breast tumors. *Oncogene*. 2012; 31: 4447–4459.
- [11] Guijarro MV, Leal JFM, Fominaya J, Blanco-Aparicio C, Alonso S, Leonart M, *et al.* MAP17 overexpression is a common characteristic of carcinomas. *Carcinogenesis*. 2007; 28: 1646–1652.
- [12] Ragazzi M, Torricelli F, Donati B, Ciarrocchi A, de Biase D, Tallini G, *et al.* Coexisting well-differentiated and anaplastic thyroid carcinoma in the same primary resection specimen: immunophenotypic and genetic comparison of the two components in a consecutive series of 13 cases and a review of the literature. *Virchows Archiv*. 2020; 478: 265–281.
- [13] Dhami GK, Liu H, Galka M, Voss C, Wei R, Muranko K, *et al.* Dynamic methylation of Numb by Set8 regulates its binding to p53 and apoptosis. *Molecular Cell*. 2013; 50: 565–576.
- [14] Siddique HR, Feldman DE, Chen C, Punj V, Tokumitsu H, Machida K. NUMB phosphorylation destabilizes p53 and promotes self-renewal of tumor-initiating cells by a NANOG-dependent mechanism in liver cancer. *Hepatology*. 2015; 62: 1466–1479.
- [15] Liu L, Lanner F, Lendahl U, Das D. Numblike and Numb differentially affect p53 and Sonic Hedgehog signaling. *Biochemical and Biophysical Research Communications*. 2011; 413: 426–431.
- [16] Garcia-Heredia JM, Lucena-Cacace A, Verdugo-Sivianes EM, Pérez M, Carnero A. The Cargo Protein MAP17 (PDZK1IP1) Regulates the Cancer Stem Cell Pool Activating the Notch Pathway by Abducting NUMB. *Clinical Cancer Research*. 2017; 23: 3871–3883.
- [17] Chandrashekar DS, Bashel B, Balasubramanya SAH, Creighton CJ, Ponce-Rodriguez I, Chakravarthi BVSK, *et al.* UALCAN: a Portal for Facilitating Tumor Subgroup Gene Expression and Survival Analyses. *Neoplasia*. 2017; 19: 649–658.
- [18] Tang Z, Li C, Kang B, Gao G, Li C, Zhang Z. GEPIA: a web server for cancer and normal gene expression profiling and interactive analyses. *Nucleic Acids Research*. 2017; 45: W98–W102.
- [19] Park JL, Kim SK. MicroRNA Profile for Diagnostic and Prognostic Biomarkers in Thyroid Cancer. *Cancers*. 2021; 13: 632.
- [20] Wang W, Wen Q, Luo J, Chu S, Chen L, Xu L, *et al.* Suppression of β -catenin Nuclear Translocation by CGP57380 Decelerates Poor Progression and Potentiates Radiation-Induced Apoptosis in Nasopharyngeal Carcinoma. *Theranostics*. 2017; 7: 2134–2149.
- [21] Kim WB, Jeon MJ, Kim WG, Kim TY, Shong YK. Unmet Clinical Needs in the Treatment of Patients with Thyroid Cancer. *Endocrinology and Metabolism*. 2020; 35: 14–25.
- [22] Lim H, Devesa SS, Sosa JA, Check D, Kitahara CM. Trends in Thyroid Cancer Incidence and Mortality in the United States, 1974–2013. *Journal of the American Medical Association*. 2017; 317: 1338–1348.
- [23] Iñiguez-Ariza NM, Brito JP. Management of Low-Risk Papillary Thyroid Cancer. *Endocrinology and Metabolism*. 2018; 33: 185–194.
- [24] Li Z, Lin W, Zheng J, Hong W, Zou J, Zhang T, *et al.* Identification of immune-related lncRNAs to improve the prognosis prediction for patients with papillary thyroid cancer. *Bioscience Reports*. 2021; 41: BSR20204086.
- [25] Ban EJ, Kim D, Kim JK, Kang S, Lee J, Jeong JJ, *et al.* Lactate Dehydrogenase a as a Potential New Biomarker for Thyroid Cancer. *Endocrinology and Metabolism*. 2021; 36: 96–105.

- [26] Liang Q, Zhang H. MAP17 contributes to non-small cell lung cancer progression via suppressing miR-27a-3p expression and p38 signaling pathway. *Cancer Biology & Therapy*. 2021; 22: 19–29.
- [27] Tampaki EC, Tampakis A, Nonni A, von Flüe M, Patsouris E, Kontzoglou K, *et al*. Combined Fascin-1 and MAP17 Expression in Breast Cancer Identifies Patients with High Risk for Disease Recurrence. *Molecular Diagnosis & Therapy*. 2019; 23: 635–644.
- [28] Ferrer I, Quintanal-Villalonga Á, Molina-Pinelo S, Garcia-Heredia JM, Perez M, Suárez R, *et al*. MAP17 predicts sensitivity to platinum-based therapy, EGFR inhibitors and the proteasome inhibitor bortezomib in lung adenocarcinoma. *Journal of Experimental & Clinical Cancer Research*. 2018; 37: 195.
- [29] Rodríguez-Rodero S, Fernández AF, Fernández-Morera JL, Castro-Santos P, Bayon GF, Ferrero C, *et al*. DNA methylation signatures identify biologically distinct thyroid cancer subtypes. *The Journal of Clinical Endocrinology and Metabolism*. 2013; 98: 2811–2821.
- [30] Borggreffe T, Oswald F. The Notch signaling pathway: Transcriptional regulation at Notch target genes. *Cellular and Molecular Life Sciences*. 2009; 66: 1631–1646.

Keywords: Papillary thyroid cancer; MAP17; NUMB; p53; Tumorigenesis

Send correspondence to: Qiong Yang, Department of Breasts Surgery, Center of Otolaryngology-Head and Neck Surgery, Zhejiang Provincial People's Hospital, People's Hospital of Hangzhou Medical College, Key Laboratory of Endocrine Gland Diseases of Zhejiang Province, 310014 Hangzhou, Zhejiang, China, E-mail: tidiq@163.com

† These authors contributed equally.